1. **Hitomi Kyogoku, Yasuaki Yamaguchi, Momoyo Miyano-Ono, Ayano Yamaguchi *and* Takanori Satoh :** Protein Expression Analysis in Temparature-adaptated Mutant Strains of Moderately thermophile Geobacillus stearothermophilus, *Natural Science Research,* **Vol.23,** *No.3,* 87-95, 2009.
2. **柏木 桂子, 藤原 好恒, 坂尾 智美, 古野 伸明, 柳澤 誠, 花田 秀樹, 谷本 能文, 山下 雅道, 渡部 稔, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 内藤 富夫, 鈴木 賢一, 柏木 昭彦 :** 両生類の生活環における強磁場の影響ー初期発生の形態学的および分子生物学的解析ー, *Space Utilization Research,* **Vol.26,** 232-235, 2010年.
3. **山本 真由美, 長積 仁, 大橋 眞, 金丸 芳, 寺嶋 吉保, 長宗 雅美 :** 特別支援教育における学生ボランティアの活用の試み, *大学教育研究ジャーナル,* **Vol.0,** *No.6,* 102-107, 2010年.
4. **大橋 眞, 齊藤 隆仁 :** 北欧とドイツの生涯学習と，徳島大学の「地域社会人を活用した教養教育」から見えてくる視点とは, *大学教育研究ジャーナル, No.7,* 66-71, 2010年.
5. **大橋 眞, 光永 雅子, 中恵 真理子, 齊藤 隆仁 :** 地域社会に学ぶ体験型学習∼第7回国際珍島学会と韓国の地域から学ぶことの意義∼, *大学教育研究ジャーナル,* **Vol.7,** *No.7,* 72-77, 2010年.
6. **大橋 眞, 光永 雅子, 中恵 真理子, 福田 利久 スティーブ, 齊藤 隆仁 :** 高等教育と生涯教育を考えるInternational Conference-地域社会人を活用した教養教育の一環としての日韓中交流-, *大学教育研究ジャーナル,* **Vol.7,** *No.7,* 78-84, 2010年.
7. **光永 雅子, 中恵 真理子, 齊藤 隆仁, 的場 一将, 大橋 眞 :** 自主的な学びを目指す「学びのコミュニティー」活動-学生・社会人・教員で創る生涯学習の形-, *大学教育研究ジャーナル, No.7,* 102-109, 2010年.
8. **大橋 眞, 岸江 信介, 光永 雅子, 中恵 真理子, 富永 朝己, 宮武 秀考, 齊藤 隆仁 :** 学際的分野における学士課程構築を視野に入れた授業評価アンケートとその活用, *大学教育研究ジャーナル, No.7,* 120-130, 2010年.
9. **佐藤 高則, 佐山 哲雄, 谷 啓二, 中原 由加里, 藤本 順子, 渡部 稔 :** 高大連携による高校生への生命科学体験型講座の実施報告, --- ∼SPPおよびSSHを例として∼ ---, *大学教育研究ジャーナル, No.7,* 173-186, 2010年.
10. **渡部 稔, 山城 考, 佐藤 高則, 大橋 眞 :** 「高校生の大学研究室への体験入学形学習プログラム」実施報告(第2報), *平成21年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 28-29, 2010年.
11. **齊藤 隆仁, 佐藤 高則, 大橋 眞, 嵯峨山 和美, 中恵 真理子, 光永 雅子 :** 社会人ボランティアを活用した教養教育, *平成21年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 18-19, 2010年.
12. **Yuta Tagami, Yuki Okamoto, Takeshi Nishiyama *and* Minoru Watanabe :** Application of RNAi technology to Xenopus beta-catenin gene, *日本発生生物学会第42回大会 発表要旨集,* 125, May 2009.
13. **Taku Aoki, Tomomi Maemura, Yuiko Hanno, Hayata Sakai, Hiroto Itoh, Haruka Shibagaki *and* Takanori Satoh :** Contribution of Trp residues to lipocalin function of E.coli yodA protein, *The 9th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 71, May 2009.
14. **Yuiko Hanno, Tomomi Maemura, Hiroto Itoh, Hayata Sakai, Haruka Shibagaki *and* Takanori Satoh :** Effect of substitutions of tetrameric type amino acids on the conformational stability of E.coli nucleoside diphosphate kinase., *The 9th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 104, May 2009.
15. **大橋 眞, 中恵 真理子, 光永 雅子, 齊藤 隆仁, 香川 順子, 廣渡 修一 :** 地域社会人を活用した教養教育―地域に広がる知の循環型社会実現を目指して―, *日本高等教育学会第12回要旨集禄,* **Vol.12,** 218-221, 2009年5月.
16. **渡部 稔, 西山 剛司 :** RNAi法によりbeta-catenin遺伝子の機能を抑制されたアフリカツメガエル初期胚の解析, *第3回日本ツメガエル研究集会 要旨集,* 25, 2009年10月.
17. **Hiroto Itoh, Hayata Sakai, Yousuke Endoh, Takamasa Kageji, Ayaka Matsunaga, Haruka Shibagaki *and* Takanori Satoh :** Effect of substitutions of hexameric type amino acids on the conformational stability of Geobacillus stearothermophilus nucleoside diphosphate kinase., *The 82nd Annual Meeting of the Japanese Biochemical Society,Abstracts,* **Vol.81,** *No.9,* 3P137, Oct. 2009.
18. **Haruka Shibagaki, Hiroto Itoh, Hayata Sakai, Yousuke Endoh, Takamasa Kageji, Ayaka Matsunaga *and* Takanori Satoh :** Contribution of C-terminal amino acids to the conformational stability of Geobacillus stearothermophilus nucleoside diphosphate kinase(2)., *The 82nd Annual Meeting of the Japanese Biochemical Society,Abstracts,* **Vol.81,** *No.9,* 3P138, Oct. 2009.
19. **佐藤 高則, 山城 考, 西山 賢一, 石田 啓祐, Haruka Shibagaki :** Oxidase activity of Cu,Fe-resistant bacteria isolated from Shirataki copper mine., *第82回日本生化学会年会プログラム·要旨集,* **Vol.81,** *No.9,* 3P233, 2009年10月.
20. **西山 剛司, 渡部 稔 :** RNAi法によりbeta-catenin遺伝子の機能を抑制されたアフリカツメガエル初期胚の解析, *第32回日本分子生物学会年会 プログラム,* 182, 2009年12月.
21. **柏木 啓子, 藤原 好恒, 板尾 智美, 古野 伸明, 柳澤 誠, 花田 秀樹, 谷本 能文, 山下 雅道, 渡部 稔, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 内藤 富夫, 鈴木 賢一, 柏木 昭彦 :** 両生類の生活環に対する強磁場の影響ー初期発生の形態学的および分子生物学的解析ー, *第26回宇宙利用シンポジウム,* 2010年1月.
22. **渡部 稔, 山城 考, 佐藤 高則, 大橋 眞 :** 「高校生の大学研究室への体験入学形学習プログラム」実施報告(第2報), *平成21年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 28-29, 2010年3月.
23. **齊藤 隆仁, 佐藤 高則, 大橋 眞, 嵯峨山 和美, 中恵 真理子, 光永 雅子 :** 社会人ボランティアを活用した教養教育, *平成21年度全学FD 徳島大学教育カンファレンス in 徳島,* 18-19, 2010年3月.
24. **佐藤 高則 :** 特殊環境微生物のタンパク質・酵素・ペプチドの構造―機能相関と有効利用に関する研究, *徳島大学知財本部講演会,* 2010年1月.
25. **渡部 稔, 山城 考, 佐藤 高則, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第2報), *平成21年度全学FD大学教育カンファレンスin徳島，発表抄録集,* 28-29, 2010年3月.
26. **Olivier Tassy, Delphine Dauga, Fabrice Daian, Daniel Sobral, François Robin, Pierre Khoueiry, David Salgado, Vanessa Fox, Danièle Caillol, Renaud Schiappa, Baptiste Laporte, Anne Rios, Guillaume Luxardi, Takehiro Kusakabe, Jean-Stéphane Joly, Sébastien Darras, Lionel Christiaen, Magali Contensin, Hélène Auger, Clément Lamy, Clare Hudson, Ute Rothbächer, Michael J. Gilchrist, Kazuhiro W. Makabe, Kohji Hotta, Shigeki Fujiwara, Nori Satoh, Yutaka Satou *and* Patrick Lemaire :** The ANISEED database: digital representation, formalization, and elucidation of a chordate developmental program., *Genome Research,* **Vol.20,** *No.10,* 1459-1468, 2010.
27. **大橋 眞, 中恵 真理子, 光永 雅子, 齊藤 隆仁, 廣渡 修一 :** 大学教育ボランティアを活用した教養教育―地域に知の循環型社会の構築を目指す新しい形の生涯学習―, *日本生涯教育学会年報, No.31,* 83-96, 2010年.
28. **柳澤 誠, 古野 伸明, 渡部 稔, 柏木 啓子, 花田 秀樹, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 鈴木 賢一, 山下 雅道, 柏木 昭彦 :** 過重力によって生じたアフリカツメガエル初期胚の頭部形成異常の解析 II, *Space Utilization Research,* **Vol.27,** 198-200, 2011年.
29. **大橋 眞, 光永 雅子, 齊藤 隆仁 :** 徳島大学「地域社会人を活用した教養教育の視点からの開発途上国をモデルとした環境教育プログラムの可能性, *大学教育研究ジャーナル,* **Vol.8,** 76-81, 2011年.
30. **大橋 眞, 光永 雅子, 佐藤 高則, 齊藤 隆仁 :** 日本とモンゴルの大学教育改革を考える国際会議「International Conference on Global Trends in Educational Culture」の成果と課題, *大学教育研究ジャーナル, No.8,* 82-90, 2011年.
31. **光永 雅子, 大橋 眞, 佐藤 高則, 齊藤 隆仁 :** グローバル化時代に即した環境教育プログラム開発を目的とした体験型異文化交流, *大学教育研究ジャーナル, No.8,* 91-100, 2011年.
32. **豊田 哲也, 石田 基広, 小野 公輔, 掛井 秀一, 佐藤 高則, 佐藤 充宏, 内藤 徹, 西山 賢一, 村上 公一, 真岸 孝一, 矢部 拓也, 行實 鉄平, 川野 卓二, 吉田 博, 田中 さやか :** 学部共通科目「情報処理の基礎」の授業開発と実施に関する研究, *平成22年度徳島大学教育カンファレンスin徳島 発表抄録集,* 16-17, 2011年.
33. **Takanori Satoh, Yousuke Endoh, Takamasa Kageji, Ayaka Matsunaga, Masami Toyoshima *and* Haruka Shibagaki :** Utilization of Extremophilic Bacteria and Their Enzymes as Bioresource., *HSUM and Tokushima University Joint Research Conference,and 1st International Conference on Mongoloid Cultures.,* 13-29, Ulaanbaatar, Mongolia, Sep. 2010.
34. **Takanori Satoh :** *35th anniversary International Conference "Chemistry in Medicine" at the chemistry and Biochemistry department, HSUM,* 20-21, Ulaanbaatar, Mongolia, Dec. 2010.
35. **Takanori Satoh :** Prospects for Regional Sciences by Learning Cultures and Histories of Ainu, *Proceedings of 3rd International Conference in Higher Education and Lifelong Learning"Educational program renewals by employment of ethnic biography of pacific region",* 10-11, Tokushima, Jan. 2011.
36. **柴垣 明佳, 伊藤 寛人, 酒井 勇太, 遠藤 陽介, 松永 亜也佳, 影治 孝政, 佐藤 高則 :** Geobacillus stearothermophilus nucleoside diphosphate kinaseの高次構造安定性に寄与する部位の探索, *第51回日本生化学会中国・四国支部例会要旨集,* 50, 2010年5月.
37. **Haruka Shibagaki, Hayata Sakai, Yousuke Endoh, Takamasa Kageji, Ayaka Matsunaga, Hiroto Itoh *and* Takanori Satoh :** Contribution of C-terminal three amino acids to the conformational stability of Geobacillus stearothermophilus nucleoside diphosphate kinase., *The 10th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 123, Jun. 2010.
38. **Hayata Sakai, Hiroto Itoh, Yousuke Endoh, Ayaka Matsunaga, Takamasa Kageji, Haruka Shibagaki *and* Takanori Satoh :** Contribution of hexamer-conserved residues to the conformational stability of Geobacillus stearothermophilus nucleoside diphosphate kinase., *The 10th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 124, Jun. 2010.
39. **佐藤 高則, 西山 賢一, 石田 啓祐, 遠藤 陽介, 柴垣 明佳, 山城 考 :** 吉野川流域の鉱山跡周辺における重金属耐性微生物の探索, *第11回極限環境生物学会年会,* **Vol.9,** 91-92, 2010年11月.
40. **佐藤 高則, 西山 賢一, 石田 啓祐, 遠藤 陽介, 柴垣 明佳, 山城 考 :** 吉野川流域鉱山跡採掘滓に存在する重金属耐性細菌の探索, *第125回徳島生物学会会報,* **Vol.63,** *No.2,* 9, 2010年12月.
41. **Akihiro Marumo, Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Yousuke Endoh, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of Thymidine kinase gene from Geobacillus genus bacteria., *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,Abstracts,* 404, Dec. 2010.
42. **Masami Toyoshima, Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Thymidylate kinase gene from Geobacillus genus bacteria., *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,Abstracts,* 404, Dec. 2010.
43. **Yousuke Endoh, Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of nucleoside diphosphate kinase gene from Geobacillus genus bacteria, *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,Abstracts,* 404, Dec. 2010.
44. **Takamasa Kageji, Yousuke Endoh, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of Thymidylate synthase gane from Geobacillus genus bacteria., *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,Abstracts,* 404, Dec. 2010.
45. **Ayaka Matsunaga, Yousuke Endoh, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of peroxiredoxin gene from Geobacillus genus bacteria, *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,Abstracts,* 405, Dec. 2010.
46. **Akihiro Marumo, Yousuke Endoh, Masami Toyoshima *and* Takanori Satoh :** 16SrRNA gene analysis and enzymatic activities of bacteria isolated from cattle compost., *The 83rd Annual Meeting of the Japanese Biochemical Society and The 33rd Annual Meeting of the Molecular Biology Society of Japan,LBA Abstracts,* 81, Dec. 2010.
47. **西山 剛司, 有家 翔子, 前高 明典 *and* Minoru Watanabe :** cDNA microarray analysis of Xenopus laevis beta-catenin knock-down embryos by RNAi, *第33回日本分子生物学会年会プログラム,* 218, Dec. 2010.
48. **柳澤 誠, 古野 伸明, 渡部 稔, 柏木 啓子, 花田 秀樹, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 鈴木 賢一, 山下 雅道, 柏木 昭彦 :** 過重力によって生じたアフリカツメガエル初期胚の頭部形成異常の解析II, *第27回宇宙利用シンポジウム,* 2011年1月.
49. **豊田 哲也, 石田 基広, 小野 公輔, 掛井 秀一, 佐藤 高則, 佐藤 充宏, 内藤 徹, 西山 賢一, 村上 公一, 真岸 孝一, 矢部 拓也, 行實 鉄平, 川野 卓二, 吉田 博, 田中 さやか :** 学部共通科目「情報処理の基礎」の授業開発と実施に関する研究, *平成22年度徳島大学教育カンファレンスin徳島発表抄録集,* 16-17, 2011年1月.
50. **大橋 眞, 齊藤 隆仁, 中恵 真理子, 光永 雅子 :** 地域社会人を活用した教養教育―地域に広がる地の循環型社会を目指して―, *平成22年度全学FD 徳島大学教育カンファレンス in 徳島,* 26-27, 2011年1月.
51. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 271B, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
52. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 271D, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
53. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 272-2, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
54. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 272-3, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
55. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2121A, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
56. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2121B, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
57. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2122A, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
58. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2122B, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
59. **Takanori Satoh, KY. Satoh, KI. Satoh *and* H. Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2123C1, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
60. **Takanori Satoh, KY. Satoh, KI. Satoh *and* Haruka Shibagaki :** Uncultured compost bacterium gene for 16S ribosomal RNA, partial sequence, clone: 2123C2, --- Thermostable bacteria isolated from cattle compost, Minami-soma,Fukushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2010.
61. **佐藤 高則, 石田 啓祐, 西山 賢一, 山城 考 :** 吉野川流域の鉱山跡周辺における重金属耐性微生物と酵素機能の調査・研究, *平成21年度学部長裁量経費プロジェクト成果発表会,* 2010年9月.
62. **佐藤 高則 :** バイオリソースを利用した耐熱性酵素遺伝子ライブラリーの構築と展開, *第3回徳島大学研究者との集い~生き物に優しい化学に取り組む研究者の素顔~,* 2010年10月.
63. **佐藤 高則, 石田 啓祐, 西山 賢一, 山城 考 :** 吉野川流域の鉱山跡周辺における重金属耐性微生物と 酵素機能の調査・研究, *平成21年度 学部長裁量経費 報告書,* 91-110, 徳島, 2010年10月.
64. **佐藤 高則 :** ホタルはなぜ光るのか? ∼生物発光の不思議にせまる∼, *平成21年度サイエンス・パートナーシップ・プロジェクト 講座型学習活動プランA 実施報告書,* 1-6, 東京, 2010年10月.
65. **大橋 眞, 中恵 真理子, 光永 雅子, 齊藤 隆仁 :** 地域社会人，学生，教員でつくる学びのコミュニティから創出される新たな視点, *日本生涯教育学会論集,* **Vol.32,** 3-11, 2011年.
66. **柳澤 誠, 古野 伸明, 渡部 稔, 柏木 啓子, 花田 秀樹, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 鈴木 賢一, 山下 雅道, 柏木 明彦 :** 過重力はアフリカツメガエルの頭部形成(特に眼と脳)に特異的に影響を与える, *Space Utilization Research,* **Vol.28,** 172-175, 2012年.
67. **大橋 眞, 佐藤 高則, 齊藤 隆仁 :** 国際交流による生涯学習と大学教育の融合の意義―韓国珍島の伝統文化を生かしたソウル大学の取り組みから何を学ぶか―, *大学教育研究ジャーナル, No.9,* 59-65, 2012年.
68. **Yousuke Endoh, Hiroto Itoh, Hayata Sakai, Masami Toyoshima, Akihiro Marumo, Haruka Shibagaki *and* Takanori Satoh :** PCR cloning and Mutagenesis Studies of Nucleoside Diphosphate Kinase From Thermophilic Geobacillus Bacteria, *Proceedings of the 3rd Asia Pacific Protein Association Conference in conjunction with the 3rd Symposium of the Chinese Protein Society (APPA-2011)-Protein and Beyond-,* 129, 2011.
69. **Takanori Satoh, Takamasa Kageji, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Akihiro Marumo, Masami Toyoshima *and* Yousuke Endoh :** Cloning and comparison of genes for dTTP synthesizing enzymes from Geobacillus bacteria, *Proceedings of the 8th Intenational Congress of Comparative Physiology and Biochemistry (ICCPB2011),* 161, 2011.
70. **大橋 眞, 佐藤 高則, 齊藤 隆仁 :** モンゴル国との学生交流から，何を学ぶのか-International Student Conference と医学系実習体験を通して-, *大学教育研究ジャーナル, No.9,* 66-73, 2012年.
71. **三隅 友子, 大橋 眞 :** 交流と対話を通した学内の連携を考える-「異文化交流の体験から何を学ぶのか」と「日本事情Ⅳ」の連携, *徳島大学国際センター紀要,* 31-43, 2012年.
72. **渡部 稔, 佐藤 高則, 山城 考, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第3報), *平成23年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 66-67, 2012年.
73. **原田 愛弓, 永廣 卓哉, 高岡 祐里, 佐藤 高則, 大橋 眞 :** モンゴル国における1st International Student Conference開催の意義, *平成23年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 32-33, 2012年.
74. **Yousuke Endoh, Hiroto Itoh, Hayata Sakai, Masami Toyoshima, Akihiro Marumo, Haruka Shibagaki *and* Takanori Satoh :** PCR cloning and Mutagenesis Studies of Nucleoside Diphosphate Kinase From Thermophilic Geobacillus Bacteria, *the 3rd Asia Pacific Protein Association Conference in conjunction with the 3rd Symposium of the Chinese Protein Society (APPA-2011)-Protein and Beyond-The Program and Abstracts,* 129, Shanghai,China, May 2011.
75. **Takanori Satoh, Takamasa Kageji, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Akihiro Marumo, Masami Toyoshima *and* Yousuke Endoh :** Cloning and comparison of genes for dTTP synthesizing enzymes from Geobacillus bacteria, *Proceedings of the 8th Intenational Congress of Comparative Physiology and Biochemistry (ICCPB2011),* 161, Nagoya, May 2011.
76. **Takanori Satoh :** *3rd HSUM and Tokushima University Joint Research Conference,* 5, Ulaanbaatar, Mongolia, Sep. 2011.
77. **Minoru Watanabe *and* Takeshi Nishiyama :** Involvement of the Nk3 class homeobox gene in the regulation of mesoderm gene expression during eraly Xenopus development, *第44回日本発生生物学会年会,* May 2011.
78. **丸茂 晃裕, 松永 亜也佳, 影治 孝政, 江原 和也, 栗原 健, 松本 千奈美, 豊嶋 政美, 遠藤 陽介, 佐藤 高則 :** 中等度好熱性細菌Thymidine kinase遺伝子のPCRクローニングと遺伝子解析, *日本分子生物学会第11 回春季シンポジウム「分子生物学の明日 – 金沢シンフォニー」要旨集,* 75, 2011年5月.
79. **Takamasa Kageji, Kazuya Ebara, Ayaka Matsunaga, Yousuke Endoh, Ken Kurihara, Chinami Matsumoto, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes, *The 11th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 133, Jun. 2011.
80. **Yousuke Endoh, Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of Geobacillus Nucleoside diphosphat e kinase genes, *The 11th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 133, Jun. 2011.
81. **Ayaka Matsunaga, Yousuke Endoh, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Masahiro Sakata, Jun Kurochi, Masami Toyoshima *and* Takanori Satoh :** PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes, *The 11th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 134, Jun. 2011.
82. **Kazuya Ebara, Ken Kurihara, Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Uridine kinase gene from Geobacillus genus bacteria, *The 84th Annual Meeting of the Japanese Biochemical Society,Abstracts,* Sep. 2011.
83. **Jun Kurochi, Masahiro Sakata, Masami Toyoshima, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Uridylate kinase gene from Geobacillus genus bacteria, *The 84th Annual Meeting of the Japanese Biochemical Society,Abstracts,* Sep. 2011.
84. **Masahiro Sakata, Jun Kurochi, Masami Toyoshima, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Cytidylate kinase gene from Geobacillus genus bacteria, *The 84th Annual Meeting of the Japanese Biochemical Society,Abstracts,* Sep. 2011.
85. **Yousuke Endoh, Masahiro Sakata, Masami Toyoshima, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Jun Kurochi *and* Takanori Satoh :** PCR cloning and genetic analysis of nucleoside diphosphate kinase gene from Geobacillus genus bacteria(2), *The 84th Annual Meeting of the Japanese Biochemical Society,Abstracts,* Sep. 2011.
86. **西山 剛司, 麻植 汐那, 水本 菜摘, 山根 真由香, 渡部 稔 :** KN3クラスホメオボックス遺伝子のアフリカツメガエル初期発生における機能解析, *第5回日本ツメガエル研究集会要旨集,* 20, 2011年10月.
87. **大橋 眞, 中恵 真理子, 光永 雅子, 齊藤 隆仁 :** 世代間交流による生涯学習―大学教養教育における対話型学習―, *日本生涯教育学会,* 2011年11月.
88. **西山 賢一, 石田 啓祐, 東明 省三, 森江 孝志, 橋本 寿夫, 阿部 肇, 佐藤 高則 :** 山川町の地質:段丘堆積物と鉱山の分布, *平成23年度 総合学術調査発表会 -吉野川市「山川町」-について(,* 2011年12月.
89. **Takanori Satoh, Takamasa Kageji, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Matsumoto, Akihiro Marumo, Masami Toyoshima, Yousuke Endoh, Jun Kurochi *and* Masahiro Sakata :** Comparison of genes for enzymes synthesizing pyrimidine nucleotides from four Geobacillus bacteria, *Program Booklet of The 34th Annual Meeting of the Molecular Biology Society of Japan(LBA),* Dec. 2011.
90. **柳澤 誠, 古野 伸明, 渡部 稔, 柏木 啓子, 花田 秀樹, 新海 正, 吉留 賢, 久保 英夫, 坂井 雅夫, 藤井 博匡, 鈴木 賢一, 山下 雅道, 柏木 明彦 :** 過重力はアフリカツメガエルの頭部形成(特に眼と脳)に特異的に影響を与える, *第28回宇宙利用シンポジウム,* 2012年1月.
91. **渡部 稔, 佐藤 高則, 山城 考, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第3報), *平成23年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 2012年1月.
92. **原田 愛弓, 永廣 卓哉, 高岡 祐里, 佐藤 高則, 大橋 眞 :** モンゴル国における1st International Student Conference開催の意義, *平成23年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 2012年1月.
93. **H Nakagawa, Jun Kurochi, Kozue Edo, Yuika Watanabe, H Sakai, Ayaka Miki, Shun Morikawa, Takanori Satoh, M Shinohara *and* K Ohura :** Partial purification and characterization of a glycoprotein from the coelomic fluid of the sea urchin, Toxopneustes pileolus, *JAPANESE PHARMACOLOGICAL SOC, EDITORIAL OFF, KANTOHYA BLDG GOKOMACHI-EBISUGAWA NAKAGYO-KU, KYOTO, 604, JAPAN,* **Vol.118,** *No.suppl.1,* 258P, Mar. 2012.
94. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst ThyA gene for thymidylate synthase, complete cds, --- Geobacillus stearothermophilus DNA for Thymidylate synthase ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
95. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct ThyA gene for thymidylate synthase, complete cds, strain: NBRC 15314, --- PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
96. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 ThyA gene for thymidylate synthase,complete cds, --- Geobacillus stearothermophilus DNA for Thymidylate synthase ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
97. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl ThyA gene for thymidylate synthase,complete cds, strain: NBRC 15313., --- PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
98. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv ThyA gene for thymidylate synthase,complete cds, strain: NBRC 15315, --- PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
99. **T. Kageji, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc ThyA gene for thymidylate synthase, complete cds, strain: NBRC 15316, --- PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
100. **A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl Prx gene for peroxiredoxin, complete cds, strain: NBRC 15313, --- PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
101. **Takamasa Kageji, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Akihiro Marumo, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza ThyA gene for thymidylate synthase,complete cds, strain: NBRC 101842, --- PCR cloning and genetic analysis of Geobacillus Thymidylate synthase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
102. **Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Akihiro Marumo, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct Prx gene for peroxiredoxin, complete cds,strain: NBRC 15314, --- PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
103. **Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Akihiro Marumo, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv Prx gene for peroxiredoxin, complete cds,strain: NBRC 15315, --- PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
104. **Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Akihiro Marumo, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc Prx gene for peroxiredoxin,complete cds, strain: NBRC 15316, --- PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
105. **Ayaka Matsunaga, Takamasa Kageji, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Akihiro Marumo, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza Prx gene for peroxiredoxin, complete cds,strain: NBRC 101842, --- PCR cloning and genetic analysis of Geobacillus Peroxiredoxin genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
106. **Akihiro Marumo, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Takamasa Kageji, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst tdk gene for thymidine kinase, complete cds,strain: ATCC7953, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
107. **Akihiro Marumo, Ayaka Matsunaga, Kazuya Ebara, Ken Kurihara, Chinami Mastumoto, Takamasa Kageji, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl tdk gene for thymidine kinase, complete cds, strain: NBRC 15313, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
108. **Akihiro Marumo, Ayaka Matsunaga, Kazuhiro Ebara, Ken Kurihara, Chinami Mastumoto, Takamasa Kageji, Masami Toyoshima, Yosuke Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct tdk gene for thymidine kinase, complete cds,strain: NBRC 15314, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
109. **A. Marumo, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, T. Kageji, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv tdk gene for thymidine kinase, complete cds,strain: NBRC 15315, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
110. **A. Marumo, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, T. Kageji, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc tdk gene for thymidine kinase,complete cds, strain: NBRC 15316, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
111. **A. Marumo, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, T. Kageji, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 tdk gene for thymidine kinase,complete cds,strain: NBRC102445, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
112. **A. Marumo, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, T. Kageji, M. Toyoshima, Y. Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza tdk gene for thymidine kinase,complete cds, strain: NBRC 101842, --- PCR cloning and genetic analysis of Geobacillus Thymidine kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
113. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst tmk gene for thymidylate kinase, complete cds,strain: ATCC7953, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
114. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl tmk gene for thymidylate kinase, complete cds, strain: NBRC 15313, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
115. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct tmk gene for thymidylate kinase, complete cds,strain: NBRC 15314, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
116. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv tmk gene for thymidylate kinase, complete cds,strain: NBRC 15315, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
117. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc tmk gene for thymidylate kinase,complete cds, strain: NBRC 15316, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
118. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 tmk gene for thymidylate kinase,complete cds,strain: NBRC102445, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
119. **M. Toyoshima, A. Matsunaga, K. Ebara, K. Kurihara, C. Mastumoto, A. Marumo, T. Kageji, Y. Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza tmk gene for thymidylate kinase,complete cds, strain: NBRC 101842, --- PCR cloning and genetic analysis of Geobacillus Thymidylate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
120. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Bacillus caldolyticus Bcl ndk gene for nucleoside diphosphate kinase, complete cds, strain: NBRC 15313, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
121. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Bacillus caldotenax Bct ndk gene for nucleoside diphosphate kinase, complete cds,strain: NBRC 15314, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
122. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Bacillus caldovelox Bcv ndk gene for nucleoside diphosphate kinase, complete cds,strain: NBRC 15315, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
123. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc ndk gene for nucleoside diphosphate kinase,complete cds, strain: NBRC 15316, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
124. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 ndk gene for nucleoside diphosphate kinase,complete cds,strain: NBRC102445, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
125. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Geobacillus zalihae Gza ndk gene for nucleoside diphosphate kinase,complete cds, strain: NBRC 101842, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
126. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Geobacillus thermoglucosidasius Gtg ndk gene for nucleoside diphosphate kinase,complete cds, strain: MAFF 118538, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
127. **Y. Endoh, A. Matsunaga, T. Kageji, K. Ebara, K. Kurihara, C. Mastumoto, M. Toyoshima *and* Takanori Satoh :** Geobacillus toebii Gto ndk gene for nucleoside diphosphate kinase,complete cds, strain: MAFF 118539, --- PCR cloning and genetic analysis of Geobacillus nucleoside diphosphate kinase genes ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Aug. 2011.
128. **佐藤 高則 :** コミュニケーションコラム「ヤマモモ」∼汚染された人々∼, *徳島生物学会会報,* **Vol.64,** *No.2,* 17-18, 2011年11月.
129. **渡部 稔, 佐藤 高則, 山城 考, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第3報), *平成23年度全学FD大学教育カンファレンス in 徳島 発表抄録集,* 66-67, 2012年1月.
130. **Kazuhiro W. Makabe *and* Hiroki Nishida :** Cytoplasmic localization and reorganization in ascidian eggs: role of postplasmic/PEM RNAs in axis formation and fate determination., *Wiley Interdisciplinary Reviews. Developmental Biology,* **Vol.1,** *No.4,* 501-518, 2012.
131. **阿部 肇, 西山 賢一, 石田 啓祐, 東明 省三, 森江 孝志, 橋本 寿夫, 阿部 肇, 佐藤 高則 :** 吉野川市山川町の地質:段丘堆積物と鉱山の分布と現況, *阿波学会紀要,* **Vol.58,** 1-12, 2012年.
132. **Makoto Yanagisawa, Keiko Kashiwagi, Hideki Hanada, Tadashi Shinkai, Satoshi Yoshitome, Hideo Kubo, Masao Sakai, Hirotada Fujii, Masamichi Yamashita, Akihiko Kashiwagi, Nobuaki Furuno *and* Minoru Watanabe :** Analysis of Head-Defects Caused by Hypergravity in Early Xenopus Embryos, *Biological Sciences in Space,* **Vol.26,** 1-6, 2012.
133. **Takanori Satoh, Yousuke Endoh, Haruka Shibagaki, Ken-ichi Nishiyama, Keisuke Ishida *and* Tadashi Yamashiro :** Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku., *Proceedings of Awagakkai,* **Vol.58,** 187-196, 2012.
134. **Makoto Yanagisawa, Keiko Kashiwagi, Hideki Hanada, Tadashi Shinkai, Satoshi Yoshitome, Hideo Kubo, Masao Sakai, Hirotada Fujii, Masamichi Yamashita, Akihiko Kashiwagi, Minoru Watanabe *and* Nobuaki Furuno :** Hypergravity Specifically Affects Anterior Head Formation in Early Xenopus Embryos, *Biological Sciences in Space,* **Vol.26,** 47-52, 2013.
135. **大橋 眞, 齊藤 隆仁 :** Peer learningを主体としたサマースクールプログラム, *大学教育研究ジャーナル,* **Vol.10,** *No.10,* 31-38, 2013年.
136. **Yousuke Endoh, Hiroto Itoh, Hayata Sakai, Tomomi Maemura, Yuiko Hanno, Haruka Shibagaki-Hirose *and* Takanori Satoh :** Contribution of amino acids to the conformational thermostabilities of nucleoside diphosphate kinases from Geobacillus stearothermophilus and E.coli, *Proceedings of 13th FAOBMB International Congress of Biochemistry and Molecular Biology,* 100, 2012.
137. **劉 潔, Makoto Ohashi *and* Shinsuke Kishie :** Study on linguistic landscapes of the public transportation facilities of Qingdao city in China, *Regional Science Research, the University of Tokushima,* **Vol.2,** *No.2,* 67-74, 2012.
138. **劉 潔, Makoto Ohashi *and* Shinsuke Kishie :** Linguistic landscapes of the famous tourist spots of Qingdao city in China, *Regional Science Research, the University of Tokushima,* **Vol.2,** *No.2,* 75-83, 2012.
139. **渡部 稔, 佐藤 高則, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第4報), *平成24年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 54-55, 2012年.
140. **Momoko Abe, Yuto Oe, Yousuke Endoh, Masahiro Sakata, Ichinkhorloo Zesem, Munkhtsetseg Janlav, Makoto Ohashi *and* Takanori Satoh :** 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products, *4th Taiwan-Korea-Japan International Symposium on Microbial Ecology and the 28th Annual Meeting of the Japanese Society of Microbial Ecology (JSME2012),* 165, Toyohashi, Sep. 2012.
141. **Yousuke Endoh, Hiroto Itoh, Hayata Sakai, Tomomi Maemura, Yuiko Hanno, Haruka Shibagaki-Hirose *and* Takanori Satoh :** Contribution of amino acids to the conformational thermostabilities of nucleoside diphosphate kinases from Geobacillus stearothermophilus and E.coli, *13th FAOBMB International Congress of Biochemistry and Molecular Biology,* 100, Bangkok, Nov. 2012.
142. **Jun Kurochi, Masahiro Sakata, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Uridylate kinase gene from Geobacillus genus bacteria(2), *The 12th Annual Meeting of The Protein Science Society of Japan,Abstracts,* Jun. 2012.
143. **Masahiro Sakata, Jun Kurochi, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and genetic analysis of Cytidylate kinase gene from Geobacillus genus bacteria(2), *The 12th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 2P-074, Jun. 2012.
144. **Kazuya Ebara, Yousuke Endoh, Chinami Matsumoto, Ken Kurihara, Jun Kurochi, Masahiro Sakata *and* Takanori Satoh :** PCR cloning and Protein expression of Uridine kinase gene from Geobacillus genus Bacteria, *The 12th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 2P-075, Jun. 2012.
145. **Chinami Matsumoto, Kazuya Ebara, Ken Kurihara, Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** Expression of four Geobacillus Peroxiredoxins in E.coli, *The 12th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 2P077, Jun. 2012.
146. **渡部 稔 :** ツメガエルを用いた高大連携事業, *第6回XCIJ首都圏支部会(XCIJ-MA)兼第6回日本ツメガエル研究集会,* 2012年7月.
147. **三隅 友子, 大橋 眞 :** 交流と対話を通した学内の協同・連携を考える, *日本協同教育学会第9回全国大会予稿集,* 2012年9月.
148. **大橋 眞, 齊藤 隆仁 :** グローバル化社会に向けた生涯教育, --- 大学教育の国際化との連携 ---, *日本生涯教育学会第33回発表要旨集録,* 24, 2012年11月.
149. **Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and Protein Expression of eight Geobacillus Uridylate kinase genes, *Journal of The Japanese Society for Extremophiles,* **Vol.11,** 144, Dec. 2012.
150. **前高 明典, 西山 剛司, 渡部 稔 :** アフリカツメガエル初期胚におけるβ-catenin遺伝子の下流因子の解析, *第128回徳島生物学会 会報,* 13, 2012年12月.
151. **Masahiro Sakata, Jun Kurochi, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Yousuke Endoh *and* Takanori Satoh :** PCR cloning and Protein Expression of Geobacillus Cytidylate kinase gene, *The 85th Annual Meeting of The Japanese Biochemical Society,Abstracts,* 3P-333, Dec. 2012.
152. **Yousuke Endoh, Masahiro Sakata, Asuka Maegawa, Nana Kawano, Satomi Takeuchi, Jun Kurochi *and* Takanori Satoh :** Protein expression and purification of Geobacillus Nucleoside diphosphate kinases, *The 85th Annual Meeting of The Japanese Biochemical Society,Abstracts,* 3P-335, Dec. 2012.
153. **Asuka Maegawa, Kazuya Ebara, Nana Kawano, Satomi Takeuchi, Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** Protein expression and Purification of Geobacillus Uridine kinases, *The 85th Annual Meeting of The Japanese Biochemical Society,Abstracts,* 3P-331, Dec. 2012.
154. **Yuto Oe, Momoko Abe, Yousuke Endoh, Masahiro Sakata, Ichinkhorloo Zesem, Munkhtsetseg Janlav, Makoto Ohashi *and* Takanori Satoh :** Comparison of bacteria in the Mongolian animal milks and their dairy products by 16S rRNA gene analysis, *The 85th Annual Meeting of The Japanese Biochemical Society,Abstracts,* 2P721, Dec. 2012.
155. **Satomi Takeuchi, Ken Kurihara, Asuka Maegawa, Nana Kawano, Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** Protein expression and Purification of Geobacillus Thymidylate synthases, *The 85th Annual Meeting of The Japanese Biochemical Society,Abstracts,* 3P-337, Dec. 2012.
156. **渡部 稔, 佐藤 高則, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第4報), *平成24年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 2012年12月.
157. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Pseudomonas sp. HYC24 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
158. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Staphylococcus sp. K2F10 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
159. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Bacillus sp. K2F13gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
160. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Staphylococcaceae bacterium K2F20gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
161. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. C2-1 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
162. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. C2-5 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
163. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. C2-7 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
164. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F47-3 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
165. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F47-6 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
166. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F47-7 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
167. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F56-1 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
168. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F56-2 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
169. **Takanori Satoh, Y. Endoh, H. Shibagaki, K. Nishiyama, K. Ishida *and* T. Yamashiro :** Streptomyces sp. F56-3 gene for 16S rRNA, partial sequence, --- Exploration for Cu,Fe-resistant bacteria from copper mine spoils in the Yoshino River basin, Shikoku ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
170. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst UK gene for uridine kinase, complete cds,strain:ATCC 7953, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
171. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl UK gene for uridine kinase, complete cds,strain:NBRC15313, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
172. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct UK gene for uridine kinase, complete cds,strain:NBRC15314, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
173. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv UK gene for uridine kinase, complete cds,strain:NBRC15315, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
174. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc UK gene for uridine kinase, complete cds,strain:NBRC15316, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
175. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 UK gene for uridine kinase, complete cds,strain: NBRC102445, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
176. **K. Ebara, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, J. Kurochi, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza UK gene for uridine kinase, complete cds, strain: NBRC 101842, --- PCR cloning of uridine kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
177. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst CMPK gene for cytidylate kinase, complete cds, strain: ATCC 7953, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
178. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 CMPK gene for cytidylate kinase,complete cds, strain: NBRC 102445, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
179. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl CMPK gene for cytidylate kinase,complete cds, strain: NBRC 15313, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
180. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct CMPK gene for cytidylate kinase, complete cds, strain: NBRC 15314, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
181. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv CMPK gene for cytidylate kinase, complete cds, strain: NBRC 15315, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
182. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc CMPK gene for cytidylate kinase, complete cds, strain: NBRC 15316, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
183. **M. Sakata, K. Kurihara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Ebara, J. Kurochi, Y. Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza CMPK gene for cytidylate kinase, complete cds, strain: NBRC 101842, --- PCR cloning of cytidylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
184. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus stearothermophilus Bst UMPK gene for uridylate kinase, complete cds,strain:ATCC 7953, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
185. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldolyticus Bcl UMPK gene for uridylate kinase, complete cds,strain:NBRC15313, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
186. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldotenax Bct UMPK gene for uridylate kinase, complete cds,strain:NBRC15314, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
187. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Bacillus caldovelox Bcv UMPK gene for uridylate kinase, complete cds,strain:NBRC15315, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
188. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus thermocatenulatus Gtc UMPK gene for uridylate kinase, complete cds,strain:NBRC15316, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
189. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus kaustophilus Gka2 UMPK gene for uridylate kinase, complete cds,strain: NBRC102445, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
190. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus zalihae Gza UMPK gene for uridylate kinase, complete cds, strain: NBRC 101842, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
191. **J. Kurochi, K. Ebara, C. Mastumoto, N. Kawano, S. Takeuchi, A. Maegawa, K. Kurihara, M. Sakata, Y. Endoh *and* Takanori Satoh :** Geobacillus caldoxylosilyticus Gcx UMPK gene for uridylate kinase, complete cds,strain:MAFF 118540, --- PCR cloning of uridylate kinase gene from Geobacillus genus bacteria ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* Apr. 2012.
192. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A1-14 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750687, Dec. 2012.
193. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A1-17 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750688, Dec. 2012.
194. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A1-18 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750689, Dec. 2012.
195. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A1-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750690, Dec. 2012.
196. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-5 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750691, Dec. 2012.
197. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-11 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750692, Dec. 2012.
198. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-12 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750693, Dec. 2012.
199. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-13 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750694, Dec. 2012.
200. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-18 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750695, Dec. 2012.
201. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750696, Dec. 2012.
202. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-25 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750697, Dec. 2012.
203. **Yousuke Endo, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactobacillus sp. A2-27 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750698, Dec. 2012.
204. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. H1-20 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750699, Dec. 2012.
205. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. H1-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750700, Dec. 2012.
206. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. H1-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750701, Dec. 2012.
207. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Streptococcus sp. H1-29 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750702, Dec. 2012.
208. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. H1-30 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750703, Dec. 2012.
209. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. H2-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750704, Dec. 2012.
210. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. H2-3 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750705, Dec. 2012.
211. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Staphylococcus sp. H2-5 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750706, Dec. 2012.
212. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. H2-8 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750707, Dec. 2012.
213. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. H2-10 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750708, Dec. 2012.
214. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. H2-12 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750709, Dec. 2012.
215. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Streptococcus sp. H2-18 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750710, Dec. 2012.
216. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. H2-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750711, Dec. 2012.
217. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. G1-1 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750712, Dec. 2012.
218. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G1-6 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750713, Dec. 2012.
219. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G1-20 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750714, Dec. 2012.
220. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G1-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750715, Dec. 2012.
221. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. G1-28 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750716, Dec. 2012.
222. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. G2-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750717, Dec. 2012.
223. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. G2-3 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750718, Dec. 2012.
224. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G2-12 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750719, Dec. 2012.
225. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G2-14 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750720, Dec. 2012.
226. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G2-15 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750721, Dec. 2012.
227. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured bacterium G2-18 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750722, Dec. 2012.
228. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. G2-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750723, Dec. 2012.
229. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. G2-32 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB750724, Dec. 2012.
230. **渡部 稔, 佐藤 高則, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第4報), *平成24年度全学FD大学教育カンファレンス in 徳島 発表抄録集,* 54-55, 2012年12月.
231. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Trichococcus sp. Ca1-11 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749360, Jan. 2013.
232. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca1-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749361, Jan. 2013.
233. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Janthinobacterium sp. Ca1-23 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749362, Jan. 2013.
234. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-4 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749363, Jan. 2013.
235. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Serratia sp. Ca2-5 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749364, Jan. 2013.
236. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Trichococcus sp. Ca2-8 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749365, Jan. 2013.
237. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Serratia sp. Ca2-14 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749366, Jan. 2013.
238. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Serratia sp. Ca2-17 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749367, Jan. 2013.
239. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749368, Jan. 2013.
240. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-11 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749369, Jan. 2013.
241. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-12 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749370, Jan. 2013.
242. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-16 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749371, Jan. 2013.
243. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-17 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749372, Jan. 2013.
244. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-19 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749373, Jan. 2013.
245. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Munkhtsetseg Janlav, Ichinkhorloo Zesem *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749374, Jan. 2013.
246. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749375, Jan. 2013.
247. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. Co1-26 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749376, Jan. 2013.
248. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp.. Co2-6 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749377, Jan. 2013.
249. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. Co2-9 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749378, Jan. 2013.
250. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C1-57 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749379, Jan. 2013.
251. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Vogesella sp. C2-9 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749380, Jan. 2013.
252. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C2-13 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749381, Jan. 2013.
253. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp.C2-15 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749382, Jan. 2013.
254. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C2-18 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749383, Jan. 2013.
255. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C2-24 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749384, Jan. 2013.
256. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Acetobacter sp. C2-26 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749385, Jan. 2013.
257. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C3-8 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749386, Jan. 2013.
258. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C3-20 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749387, Jan. 2013.
259. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. C3-35 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB749388, Jan. 2013.
260. **矢野 千晴, 奇 旭瓊, VICTORIA MARIE BLOYER, CHIMEDDORJ TSOLMON, 佐藤 高則, 齊藤 隆仁, 石田 三千雄, 松尾 義則, 大橋 眞 :** INTERNATIONAL STUDENT CONFERENCE OF 2012, *平成24年度総合科学教育部プロジェクト研究I 発表会,* 2013年2月.
261. **佐藤 高則 :** バイオリソースを利用した耐熱性酵素遺伝子ライブラリーの構築, *平成23年度研究成果展開事業 研究成果最適展開支援プログラム フィージビリティスタディ【FS】ステージ 完了報告書,* 1-10, 東京, 2012年4月.
262. **矢野 千晴, 奇 旭瓊, VICTORIA MARIE BLOYER, CHIMEDDORJ TSOLMON, 佐藤 高則, 齊藤 隆仁, 石田 三千雄, 松尾 義則, 大橋 眞 :** プロジェクト研究Ⅰ(International Student Conference 開催プロジェクト)報告書, *平成24年度総合科学教育部プロジェクト研究I 報告書,* 徳島, 2013年1月.
263. **Noriko Sugiyama, Ayumi Maeda, Hiroshi Tanaka, Eiichiro Hanafusa, Taku Kanai *and* Takanori Satoh :** Thermus thermophilus Inorganic Pyrophosphatase Variants by Error-prone PCR Suppressed Thermal Aggregation, *Natural Science Research,* **Vol.27,** *No.4,* 73-81, 2013.
264. **Yasuaki Yamaguchi, Momoyo Miyano-Ono, Hitomi Kyogoku-Tsujinaka, Ayano Yamaguchi, Taku Aoki *and* Takanori Satoh :** Protein Expression Analysis in Amitrole-Resistant E.coli Strains, *Natural Science Research,* **Vol.27,** *No.4,* 83-90, 2013.
265. **k Tanaka, A Kato, C Angelocci, Minoru Watanabe *and* Y Kato :** A potential molecular pathogenesis of cardiac/laterality defects in Oculo-Facio-Cardio-Dental syndrome., *Developmental Biology,* **Vol.387,** *No.1,* 28-36, 2014.
266. **大橋 眞, 齊藤 隆仁, 胡 萌萌, 入口 幸子, 間賀田 悠吾 :** グローバル化社会に向けた大学教養教育とは, *大学教育研究ジャーナル,* **Vol.11,** *No.11,* 117-124, 2014年.
267. **Takanori Satoh, Masahiro Sakata, Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi *and* Yousuke Endoh :** Comparison of Thermostabilities among Nucleoside Diphosphate Kinases from Five Geobacillus Bacteria, *Proceedings of 8th Asian Biophysics Association (ABA) Symposium &2013 Annual Meeting of Korean Biophysical Society,* 131-132, 2013.
268. **Takanori Satoh, Asuka Maegawa, Kazuya Ebara, Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Jun Kurochi, Masahiro Sakata *and* Yousuke Endoh :** Characteristics and Thermostabilities of Four Geobacillus Uridine kinases, *Proceedings of 5th Taiwan-Korea-Japan International Symposium on Microbial Ecology,* 48, 2013.
269. **Takanori Satoh, Masahiro Sakata, Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi *and* Yousuke Endoh :** Comparison of Thermostabilities among Nucleoside Diphosphate Kinases from Five Geobacillus Bacteria, *8th Asian Biophysics Association (ABA) Symposium &2013 Annual Meeting of Korean Biophysical Society,* 131-132, Jeju,Korea, May 2013.
270. **Takanori Satoh, Asuka Maegawa, Kazuya Ebara, Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Jun Kurochi, Masahiro Sakata *and* Yousuke Endoh :** Characteristics and Thermostabilities of Four Geobacillus Uridine kinases, *5th Taiwan-Korea-Japan International Symposium on Microbial Ecology,* 48, Jhongli,Taiwan, Oct. 2013.
271. **Mari Yamada, Takeshi Nishiyama, Yuta Tagami *and* Minoru Watanabe :** Suppression of gene function in Xenopus laevis, *International Symposium on RNAi and Genome Editing Methods,* Mar. 2014.
272. **Minoru Watanabe :** Application of CRISPR/Cas system to Xenopus laevis, *International Symposium Frontiers in Amphibian Biology: Endangered Specices Consevation and Genome Editing,* 59-60, Mar. 2014.
273. **黒地 潤, 前川 明日香, 竹内 里美, 河野 奈菜, 坂田 真宏, 遠藤 陽介, 佐藤 高則 :** Geobacillus属細菌由来Uridylate kinaseのクローニング,発現と耐熱性の検討, *第54回日本生化学会中国・四国支部例会要旨集,* 2013年5月.
274. **Masahiro Sakata, Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Yousuke Endoh *and* Takanori Satoh :** Thermostabilities of Cytidylate kinases from three thermophilic Geobacillus bacteria, *The 13th Annual Meeting of The Protein Science Society of Japan,Abstracts,* 2P-080, Jun. 2013.
275. **Nana Kawano, Yuto Oe, Momoko Abe, Satomi Takeuchi, Asuka Maegawa, Chinami Matsumoto, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** High conformational thermostability of four Geobacillus Peroxiredoxins, *The 86th Annual Meeting of The Japanese Biochemical Society,Abstracts,,* 2LBA-012, Sep. 2013.
276. **Satomi Takeuchi, Asuka Maegawa, Nana Kawano, Momoko Abe, Yuto Oe, Jun Kurochi, Masahiro Sakata, Yousuke Endoh *and* Takanori Satoh :** Expression, Purification and Thermostability of Geobacillus thymidylate synthases, *The 86th Annual Meeting of The Japanese Biochemical Society,Abstracts,,* 3P-160, Sep. 2013.
277. **Yuto Oe, Momoko Abe, Yousuke Endoh, Masahiro Sakata, Ichinkhorloo Zesem, Munkhtsetseg Janlav, Makoto Ohashi *and* Takanori Satoh :** 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products,Biosorbent, *The 29th Annual Meeting of the Japanese Society of Microbial Ecology (JSME2013),Program and abstracts,* 130, Nov. 2013.
278. **齊藤 隆仁, 大橋 眞 :** 生涯学習の導入教育としての 大学教養教育, --- 徳島大学における事例から ---, *日本生涯教育学会第34回発表要旨集録,* 5, 2013年11月.
279. **大橋 眞, 齊藤 隆仁 :** グローバル化社会に向けた大学教養教育, *平成25年度全学FD 徳島大学教育カンファレンスin徳島,* 26-27, 2013年12月.
280. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BC1-31 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797136, Apr. 2013.
281. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BC1-32 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797137, Apr. 2013.
282. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Acetobacter sp. BC1-33 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797138, Apr. 2013.
283. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BC1-36 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797139, Apr. 2013.
284. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BC1-38 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797140, Apr. 2013.
285. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BCC1-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797141, Apr. 2013.
286. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Streptococcus sp. BCC1-8 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797142, Apr. 2013.
287. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Acetobacter sp. BCC1-10 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797143, Apr. 2013.
288. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Streptococcus sp. BH1-30 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797144, Apr. 2013.
289. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Streptococcus sp. BH1-31 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797145, Apr. 2013.
290. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. H12-9 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB797146, Apr. 2013.
291. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH1-29 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828401, Aug. 2013.
292. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH1-34 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828402, Aug. 2013.
293. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Streptococcus sp. BH2-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828403, Aug. 2013.
294. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH2-3 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828404, Aug. 2013.
295. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH2-12 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828405, Aug. 2013.
296. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH2-14 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828406, Aug. 2013.
297. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Streptococcus sp. BH2-16 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828407, Aug. 2013.
298. **Momoko Abe, Yuto Oe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BH2-19 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and biosorbent ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828408, Aug. 2013.
299. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BC1-34 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828409, Aug. 2013.
300. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BC1-37 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828410, Aug. 2013.
301. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BC2-4 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828411, Aug. 2013.
302. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BC2-13 gene for 19S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828412, Aug. 2013.
303. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC1-1 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828413, Aug. 2013.
304. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC1-3 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828414, Aug. 2013.
305. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC1-6 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828415, Aug. 2013.
306. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC2-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828416, Aug. 2013.
307. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BCC2-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828417, Aug. 2013.
308. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC3-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828418, Aug. 2013.
309. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BCC3-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828419, Aug. 2013.
310. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BCC3-24 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828420, Aug. 2013.
311. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. BCC3-26 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828421, Aug. 2013.
312. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactobacillus sp. BCC3-28 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828422, Aug. 2013.
313. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. H12-2 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828423, Aug. 2013.
314. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. H12-3 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828424, Aug. 2013.
315. **Yuto Oe, Momoko Abe, Nana Kawano, Satomi Takeuchi, Asuka Maegawa, Jun Kurochi, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Lactococcus sp. H12-6 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB828425, Aug. 2013.
316. **前高 明典, 西山 剛司, 渡部 稔 :** C/EBPα遺伝子による中胚葉遺伝子の発現制御, *第7回 日本ツメガエル研究集会要旨集,* 20, 2013年9月.
317. **渡部 稔, 佐藤 高則, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第5報), *平成25年度全学FD大学教育カンファレンス in 徳島 発表抄録集,* 46-47, 2013年12月.
318. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca1-16 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858501, Jan. 2014.
319. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca1-28 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858502, Jan. 2014.
320. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-16 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858505, Jan. 2014.
321. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-10 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858504, Jan. 2014.
322. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-21 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858506, Jan. 2014.
323. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-22 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858507, Jan. 2014.
324. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-6 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858503, Jan. 2014.
325. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Masahiro Sakata, Yousuke Endo, Makoto Ohashi, Janlav Munkhtsetseg, Zesem Ichinkhorloo *and* Takanori Satoh :** Uncultured Leuconostoc sp. Ca2-24 gene for 16S rRNA, partial sequence, --- 16S rRNA gene analysis of bacteria from Mongolian animal milks and their dairy products ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB858508, Jan. 2014.
326. **馬 根兄, MOLOMJAMTS ENKHJARGAL, 李 蓉蓉, 孫 元至, 山口 祐佳, 石田 三千雄, 佐藤 高則, 中山 信太郎 :** プロジェクト研究Ⅰ(中山・石田・佐藤グループ)発表会, --- 東日本大震災と原発事故に対する徳島県民調査 ---, *平成25年度総合科学教育部プロジェクト研究I 発表会,* 2014年2月.
327. **佐藤 高則, 大橋 眞, 渡部 稔, Munkhtsetseg Janlav, Ichinkhorloo Zesem :** モンゴル国家畜乳および乳製品の生物資源としての利用を指向した免疫作用 および細菌・機能性生体分子の調査研究, *平成24年度 学部長裁量経費・創生研究プロジェクト 報告書,* 20-22, 徳島, 2013年9月.
328. **馬 根兄, MOLOMJAMTS ENKHJARGAL, 李 蓉蓉, 孫 元至, 山口 祐佳, 石田 三千雄, 佐藤 高則, 中山 信太郎 :** プロジェクト研究Ⅰ(中山・石田・佐藤グループ)報告書, --- 東日本大震災と原発事故に対する徳島県民調査 ---, *平成25年度総合科学教育部プロジェクト研究I 報告書,* 1-9, 徳島, 2014年1月.
329. **阿波学会60周年記念誌編集委員会 編, 佐藤 高則 :** 発見! 徳島の自然と文化, --- トクシマシタ! 阿波学会 ---, 阿波学会, 徳島, 2014年12月.
330. **Yuto Oe, Momoko Abe, Yousuke Endoh, Masahiro Sakata, Zesem Ichinkhorloo, Janlav Munkhtsetseg, Makoto Ohashi *and* Takanori Satoh :** Phylogenetic analysis of bacteria from Mongolian animal milks and their dairy products by clone library method, *Natural Science Research,* **Vol.28,** *No.4,* 31-40, 2014.
331. **佐藤 高則, 原 将希, 濱地 由衣, 井上 翔太, 高橋 昇汰, 中山 美咲, 瀬部 直之, 牧野 祐未子, 坂口 雅俊, 大岩 由利恵, 石田 啓祐 :** 阿南市石灰鉱山由来の炭酸カルシウム耐性微生物の探索, *阿波学会紀要, No.60,* 201-210, 2015年.
332. **大橋 眞, 齊藤 隆仁, 胡 萌萌, 入口 幸子, 間賀田 悠吾 :** グローバル教育の課題-「持続可能な開発のための教育」の視点から, *大学教育研究ジャーナル,* **Vol.12,** *No.12,* 54-61, 2015年.
333. **Takanori Satoh, Momoko Abe, Shota Takahashi, Shota Inoue, Misaki Nakayama, Yuto Oe *and* Masahiro Sakata :** Contribution of Different Amino Acids to the Conformational Thermostabilities of Cytidylate kinases from Thermophilic Geobacillus bacteria, *Proceedings of 4th Asia Pacific Protein Association Conference (APPA2014,* 169-170, 2014.
334. **Takanori Satoh, Yuto Oe, Momoko Abe, Yousuke Endoh, Masahiro Sakata, Ichinkhorloo Zesem, Munkhtsetseg Janlav *and* Makoto Ohashi :** Phylogenetic analysis of bacteria from Mongolian animal milks and their dairy products by clone library method, *Proceedings of 15th International Symposium on Microbial Ecology(ISME15),* 55-56, 2014.
335. **Yurie Ohiwa, Yuto Oe, Hiroto Itoh, Hayata Sakai, Haruka Shibagaki-Hirose, Yousuke Endoh *and* Takanori Satoh :** Thermostabilizing Factors For Nucleoside Diphosphate Kinase From Moderately Thermophile Geobacillus stearothermophilus, *Proceedings of 15th IUBMB-24th FAOBMB-TSBMB conference,* 211, 2014.
336. **三隅 友子, 大橋 眞 :** 共生社会を支える教養教育の可能性Ⅰ-パフォーマティブ・ラーニングを導入する-, *徳島大学国際センター紀要,* **Vol.2014,** 1-12, 2015年.
337. **Takanori Satoh, Momoko Abe, Shota Takahashi, Shota Inoue, Misaki Nakayama, Yuto Oe *and* Masahiro Sakata :** Contribution of Different Amino Acids to the Conformational Thermostabilities of Cytidylate kinases from Thermophilic Geobacillus bacteria, *The 4th Asia Pacific Protein Association Conference (APPA2014),* 169-170, Seogwipo,Jeju,Korea, May 2014.
338. **Takanori Satoh, Yuto Oe, Momoko Abe, Yousuke Endoh, Masahiro Sakata, Ichinkhorloo Zesem, Munkhtsetseg Janlav *and* Makoto Ohashi :** Phylogenetic analysis of bacteria from Mongolian animal milks and their dairy products by clone library method, *The 15th International Symposium on Microbial Ecology(ISME15),* 55-56, Seoul, Aug. 2014.
339. **Yurie Ohiwa, Yuto Oe, Hiroto Itoh, Hayata Sakai, Haruka Shibagaki-Hirose, Yousuke Endoh *and* Takanori Satoh :** Thermostabilizing Factors For Nucleoside Diphosphate Kinase From Moderately Thermophile Geobacillus stearothermophilus, *15th IUBMB-24th FAOBMB-TSBMB conference,* 211, Taipei, Oct. 2014.
340. **北村 友哉, 渡部 稔, 吉留 賢, 古野 伸明 :** ネッタイツメガエルMyt-1遺伝子のクローニングと初期発生における機能解析, *日本動物学会中国四国支部会報第66号,* 14, 2014年5月.
341. **山田 真里, 渡部 稔 :** 人工ヌクレアーゼのアフリカツメガエル初期胚への応用, *日本動物学会中国四国支部会報第66号,* 15, 2014年5月.
342. **Mari Yamada *and* Minoru Watanabe :** Taegeted mutagenesis of Xenopus laevis gene by CRISPR/Cas system, *第47回日本発生生物学会年会,* 76, May 2014.
343. **Asuka Maegawa, Shota Inoue, Momoko Abe, Yuto Oe, Masahiro Sakata, Jun Kurochi, Yousuke Endoh *and* Takanori Satoh :** Thermostabilities of recombinant uridine kinases from four thermophilic Geobacillus bacteria, *The 14th Annual Meeting of The Protein Science Society of Japan,Abstracts,* Jun. 2014.
344. **Yuto Oe, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Yurie Ohiwa *and* Takanori Satoh :** Fluorescence spectroscopic studies on conformational changes and Nucleotide binding of thermophilic nucleotide kinases and their variants, *The 52nd Annual Meeting of the Biophysical Society of Japan,The Program and Abstracts,* **Vol.54,** *No.Suppl.1-2,* S214, Sep. 2014.
345. **Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa, Keisuke Ishida *and* Takanori Satoh :** Exploration for calcium carbonate-resistant bacteria from Limestone mine, Tokushima, *The 87th Annual Meeting of the Japanese Biochemical Society,The Program and Abstracts,* Oct. 2014.
346. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library method, *The 87th Annual Meeting of the Japanese Biochemical Society,The Program and Abstracts,* Oct. 2014.
347. **Momoko Abe, Shota Takahashi, Shota Inoue, Misaki Nakayama, Yurie Ohiwa, Yuto Oe, Masahiro Sakata *and* Takanori Satoh :** Contribution of Arg23 and Glu26 to the conformational Thermostabilities of Geobacillus Cytidylate kinases, *The 87th Annual Meeting of the Japanese Biochemical Society,The Program and Abstracts,* Oct. 2014.
348. **Yuto Oe, Yurie Ohiwa, Momoko Abe, Shota Inoue, Shota Takahashi, Misaki Nakayama, Yousuke Endoh *and* Takanori Satoh :** Amino acids contributing to the hexamer formation and conformational thermostabilities of Geobacillus stearothermophilus Nucleoside diphosphate kinase, *The 87th Annual Meeting of the Japanese Biochemical Society,The Program and Abstracts,* Oct. 2014.
349. **Yui Hamaji, Masaki Hara, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa, Ken-ichi Nishiyama, Keisuke Ishida *and* Takanori Satoh :** Exploration for environmental adaptive bacteria from mine spoils from Tokushima and Wakayama prefectures, *The Joint Meeting of Japanese Environmental Microbiology-related Associations 2014,The Program and abstracts,* Oct. 2014.
350. **大橋 眞, 中恵 真理子, 光永 雅子, 齊藤 隆仁 :** グローバル化社会に向けた生涯教育, --- 留学生と学び合う大学教育との融合 ---, *日本生涯教育学会,* 2014年11月.
351. **原田 新, 小野 覚久, 高橋 直樹, 伊槻 悟, 大岩 由利恵, 坂本 和歌子, 山田 真里, 渡邉 凌, 佐藤 高則, 中山 信太郎 :** ふくしま，とくしま，ともに輝こうプロジェクトの活動報告, *平成26年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 52-53, 2014年12月.
352. **大橋 眞, 齊藤 隆仁, モロムジャム ツエンヘジャルガルガル, 孫 元至, ウガンバートル ガートルガー, 齊藤 隆仁 :** グローバル化社会に向けた教養教育の課題, *平成26年度全学FD 徳島大学教育カンファレンスin徳島,* 2014年12月.
353. **渡部 稔, 山城 考, 小川 誠 :** タンポポを題材とした遺伝子解析の実験, *日本生物教育学会第98回研究発表予稿集,* 30, 2015年1月.
354. **佐藤 高則 :** 生物と教養と雑学, *先生のコバナシ,* 2014年6月.
355. **渡部 稔 :** CRISPR/Cas9法によるアフリカツメガエル遺伝子の改変, *第9回XCIJ-MA・第8回ツメガエル研究会ジョイント研究集会,* 2014年11月.
356. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Pseudomonas sp. Asec003 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987914, Dec. 2014.
357. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. Asec004 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987915, Dec. 2014.
358. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Pseudomonas sp. Asec005 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987916, Dec. 2014.
359. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Pseudomonas sp. ACT001 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987917, Dec. 2014.
360. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT0051 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987918, Dec. 2014.
361. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT0052 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987919, Dec. 2014.
362. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT008 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987920, Dec. 2014.
363. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT0091 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987921, Dec. 2014.
364. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Lysinibacillus sp. ACT0092 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987922, Dec. 2014.
365. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT010 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987923, Dec. 2014.
366. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT013 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987924, Dec. 2014.
367. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT016 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987925, Dec. 2014.
368. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT031 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987926, Dec. 2014.
369. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT035 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987927, Dec. 2014.
370. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Staphylococcus sp. ACT039 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987928, Dec. 2014.
371. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Stenotrophomonas sp. ACT041 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987929, Dec. 2014.
372. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT045 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987930, Dec. 2014.
373. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT048 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987931, Dec. 2014.
374. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT058 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987932, Dec. 2014.
375. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT059 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987933, Dec. 2014.
376. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT060 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987934, Dec. 2014.
377. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT061 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987935, Dec. 2014.
378. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT064 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987936, Dec. 2014.
379. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT068 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987937, Dec. 2014.
380. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Arthrobacter sp. ACT071 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987938, Dec. 2014.
381. **Takanori Satoh, Masaki Hara, Yui Hamaji, Shota Inoue, Shota Takahashi, Misaki Nakayama, Naoyuki Sebe, Yumiko Makino, Masatoshi Sakaguchi, Yurie Ohiwa *and* Keisuke Ishida :** Bacillus sp. ACT104 gene for 16S ribosomal RNA, partial sequence, --- Exploration for calcium carbonate-resistant bacteria from Asebi Limestone mine, Tokushima ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* AB987939, Dec. 2014.
382. **渡部 稔, 大橋 眞 :** 高大連携事業「高校生の大学研究室への体験入学型学習プログラム」実施報告(第6報), *平成26年度全学FD大学教育カンファレンス in 徳島 発表抄録集,* 48-49, 2014年12月.
383. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG1, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001846, Feb. 2015.
384. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG2, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001847, Feb. 2015.
385. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG3, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001848, Feb. 2015.
386. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Geobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG4, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001849, Feb. 2015.
387. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG5, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001850, Feb. 2015.
388. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Psychrobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG6, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001851, Feb. 2015.
389. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Shewanella sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG7, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001852, Feb. 2015.
390. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG9, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001853, Feb. 2015.
391. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG10, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001854, Feb. 2015.
392. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG13, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001855, Feb. 2015.
393. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG15, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001856, Feb. 2015.
394. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG16, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001857, Feb. 2015.
395. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG18, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001858, Feb. 2015.
396. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG19, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001859, Feb. 2015.
397. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: FSSG20, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001860, Feb. 2015.
398. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Bacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ1, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001831, Feb. 2015.
399. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ2, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001832, Feb. 2015.
400. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Bacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ3, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001833, Feb. 2015.
401. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ4, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001834, Feb. 2015.
402. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ5, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001835, Feb. 2015.
403. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ6, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001836, Feb. 2015.
404. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ7, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001837, Feb. 2015.
405. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ8, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001838, Feb. 2015.
406. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Bacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ9, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001839, Feb. 2015.
407. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ10, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001840, Feb. 2015.
408. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ11, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001841, Feb. 2015.
409. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ12, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001842, Feb. 2015.
410. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ13, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001843, Feb. 2015.
411. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ14, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001844, Feb. 2015.
412. **Yui Hamaji, Masaki Hara, Natsumi Yoshioka, Mayu Kanamoto, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: AJKZ15, --- 16S rRNA gene analysis of bacteria from samma narezushi (fermented sushi of Saury) and leaf of Alpinia japonica by clone library ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC001845, Feb. 2015.
413. **小野 覚久, 高橋 直樹, 伊槻 悟, 大岩 由利恵, 坂本 和歌子, 山田 真里, 渡邉 凌, 原田 新, 佐藤 高則, 中山 信太郎 :** ふくしま，とくしま，ともに輝こうプロジェクトの活動報告, *平成26年度総合科学教育部プロジェクト研究I 発表会,* 2015年2月.
414. **佐藤 高則 :** 15th International Symposium on Microbial Ecology(ISME)での国際学会発表, *平成26年度総合科学部創生研究プロジェクト 研究成果発表旅費支援 報告書,* 1-2, 徳島, 2014年10月.
415. **佐藤 高則 :** 15th International Union of Biochemistry and Molecular Biology(IUBMB)-24th The Federation of Asian and Oceanian Biochemists and Molecular Biologists (FAOBMB)-The Taiwan society for Biochemistry and Molecular Biology(TSBMB)国際会議での研究成果発表, *平成26年度総合科学部創生研究プロジェクト 研究成果発表旅費支援 報告書,* 1-2, 徳島, 2014年10月.
416. **佐藤 高則, 宮田 政徳, 堤 和博, 福田 利久 スティーブ, 齊藤 隆仁, 古屋 S. 玲, 吉本 勝彦, 三好 德和, 荒木 秀夫 :** 全学共通教育センター FDの歴史, *徳島大学FDの歴史,* **Vol.2,** 91-111, 徳島, 2014年12月.
417. **伊槻 悟, 大岩 由利恵, 小野 覚久, 坂本 和歌子, 高橋 直樹, 山田 真里, 渡邉 凌, 原田 新, 佐藤 高則, 中山 信太郎 :** プロジェクト研究Ⅰ(中山・佐藤・原田グループ)報告書, --- ふくしま，とくしま，ともに輝こうプロジェクトの活動報告 ---, *平成26年度総合科学教育部プロジェクト研究I 報告書,* 1-13, 徳島, 2015年1月.
418. **佐藤 高則 :** 教育改革推進のための全学FD支援事業(共通教育センター), *平成26年度学長裁量経費による成果報告書,* 1-2, 徳島, 2015年3月.
419. **Taku Aoki, Kayo Takechi, Tomomi Maemura, Yuiko Hanno, Hayata Sakai, Hiroto Itoh, Haruka Shibagaki-Hirose *and* Takanori Satoh :** Lipocalin function of E.coli zinT (yodA) protein, *Natural Science Research,* **Vol.29,** *No.4,* 35-47, 2015.
420. **大橋 眞, 齊藤 隆仁 :** 教養教育における持続可能な社会を目指す体験型学習, *大学教育研究ジャーナル,* **Vol.13,** *No.13,* 56-62, 2016年.
421. **Shota Inoue, Asuka Maegawa, Kazuya Ebara, Yuto Oe, Momoko Abe, Mayu Kanamoto, Natsumi Yoshioka, Yui Hamaji, Shota Takahashi, Yurie Ohiwa *and* Takanori Satoh :** Contribution of different amino acids among Geobacillus uridine kinases to the conformational thermostabilities, *Proceedings of 9th Asian Biophysics Association (ABA) Symposium,* 237, 2015.
422. **小野 覚久, 村中 貴恵, 伊槻 悟, 井上 翔太, 小林 大起, 都築 弘充, 森 健太朗, 大岩 由利恵, 坂本 和歌子, 山田 真里, 高橋 直樹, 渡邉 凌, 佐藤 高則, 山本 真由美, 中山 信太郎 :** 東日本大震災から学ぶ，防災教育の実態とあり方 ∼福島と徳島における放射線教育についての調査報告を含めて∼, *平成27年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 60-61, 2016年.
423. **Shota Inoue, Asuka Maegawa, Kazuya Ebara, Yuto Oe, Momoko Abe, Mayu Kanamoto, Natsumi Yoshioka, Yui Hamaji, Shota Takahashi, Yurie Ohiwa *and* Takanori Satoh :** Contribution of different amino acids among Geobacillus uridine kinases to the conformational thermostabilities, *9th Asian Biophysics Association (ABA) Symposium,* 237, Hangzhou,China, May 2015.
424. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Phylogenetic analysis of Cu,Fe-tolelant bacteria from Myoho copper mine spoils in Wakayama prefecture, *The 7th Japan-Taiwan-Korea International Symposium on Microbial Ecology and the 30th Annual Meeting of the Japanese Society of Microbial Ecology (JSME2015),* Tsuchiura, Ibaraki, Oct. 2015.
425. **山田 真里, 渡部 稔 :** CRISPR/Cas9 法によるアフリカツメガエル初期胚のゲノム編集, *日本動物学会中国四国支部会報第67号,* 13, 2015年5月.
426. **Shota Takahashi, Momoko Abe, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Contribution of different amino acids to thermostabilities of Geobacillus Cytidylate kinase, *The 15th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* Jun. 2015.
427. **Shota Inoue, Shota Takahashi, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Yurie Ohiwa *and* Takanori Satoh :** Exploration for amino acids contributing to the thermostabilities of Geobacillus Uridine kinase, *The 15th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* Jun. 2015.
428. **Yurie Ohiwa, Yuto Oe, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Shota Takahashi, Shota Inoue *and* Takanori Satoh :** Exploration for amino acids contributing to thermostabilities of Geobacillus stearothermophilus Nucleoside diphosphate kinase, *The 15th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* Jun. 2015.
429. **大橋 眞, 齊藤 隆仁 :** グローバル社会に向けた生涯教育, --- 大学教育との融合による医療制度の検証 ---, *日本生涯教育学会,* 2015年11月.
430. **Yurie Ohiwa, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue *and* Takanori Satoh :** Exploration for amino acids contributing to the conformational thermostabilities of Geobacillus stearothermophilus Nucleoside diphosphate kinase, *Annual Meeting of The 88th Japanese Biochemical Society and the 38th Molecular Biology Society of Japan,The Program and Abstract,* Dec. 2015.
431. **Shota Inoue, Shota Takahashi, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Yurie Ohiwa *and* Takanori Satoh :** Amino acids contributing to the conformational thermostabilities of Geobacillus Uridine kinase, *Annual Meeting of The 88th Japanese Biochemical Society and the 38th Molecular Biology Society of Japan,The Program and Abstract,* Dec. 2015.
432. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Exploration for Cu,Fe-tolelant bacteria from Myoho copper mine spoils in Wakayama prefecture, *Annual Meeting of The 88th Japanese Biochemical Society and the 38th Molecular Biology Society of Japan,The Program and Abstract,* Dec. 2015.
433. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** 16S rRNA gene analysis of bacteria from narezushi (fermented sushi) produced at Wakayama prefecture by clone library method, *Annual Meeting of The 88th Japanese Biochemical Society and the 38th Molecular Biology Society of Japan,The Program and Abstract,* Dec. 2015.
434. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus, *Annual Meeting of The 88th Japanese Biochemical Society and the 38th Molecular Biology Society of Japan,The Program and Abstract(LBA),* Dec. 2015.
435. **佐藤 高則, 濵地 由衣, 井上 翔太, 高橋 昇汰, 瀬部 直之, 牧野 祐未子, 坂口 雅俊, 大岩 由利恵, 石田 啓祐 :** 阿南市石灰鉱山由来の炭酸カルシウム耐性微生物の探索, *徳島生物学会会報,* **Vol.68,** *No.2,* 11, 2015年12月.
436. **小野 覚久, 村中 貴恵, 伊槻 悟, 井上 翔太, 小林 大起, 都築 弘充, 森 健太朗, 大岩 由利恵, 坂本 和歌子, 山田 真里, 高橋 直樹, 渡邉 凌, 佐藤 高則, 山本 真由美, 中山 信太郎 :** 東日本大震災から学ぶ，防災教育の実態とあり方 ∼福島と徳島における放射線教育についての調査報告を含めて∼, *平成27年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 60-61, 2016年1月.
437. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaH1, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075522, Sep. 2015.
438. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaH2, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075523, Sep. 2015.
439. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaH3, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075524, Sep. 2015.
440. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaH4, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075525, Sep. 2015.
441. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaH5, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075526, Sep. 2015.
442. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaR1, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075527, Sep. 2015.
443. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaR2, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075528, Sep. 2015.
444. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaR3, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075529, Sep. 2015.
445. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaR4, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075530, Sep. 2015.
446. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaR5, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075531, Sep. 2015.
447. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaS1, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075532, Sep. 2015.
448. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaS2, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075533, Sep. 2015.
449. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaS3, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075534, Sep. 2015.
450. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaS4, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075535, Sep. 2015.
451. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaS5, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075536, Sep. 2015.
452. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaU1, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075537, Sep. 2015.
453. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaU2, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075538, Sep. 2015.
454. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaU3, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075539, Sep. 2015.
455. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: KozaU4, --- 16S rRNA gene analysis of bacteria from narezushi (fermented sushi of Saury) produced at Kozagawa,Wakayama prefecture by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC075540, Sep. 2015.
456. **佐藤 高則, 渡部 稔 :** 「高校生物の復習テスト」結果について, *FD企画 平成27年度大学入門講座・オリエンテーション反省会,* 2015年9月.
457. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM11, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076843, Oct. 2015.
458. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM12, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076844, Oct. 2015.
459. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM13, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076845, Oct. 2015.
460. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM14, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076846, Oct. 2015.
461. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM15, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076847, Oct. 2015.
462. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM16, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076848, Oct. 2015.
463. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM17, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076849, Oct. 2015.
464. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM18, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076850, Oct. 2015.
465. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM19, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076851, Oct. 2015.
466. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM20, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076852, Oct. 2015.
467. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM22, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076853, Oct. 2015.
468. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM23, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076854, Oct. 2015.
469. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM29, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076855, Oct. 2015.
470. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM31, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076856, Oct. 2015.
471. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM33, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076857, Oct. 2015.
472. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM35, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076858, Oct. 2015.
473. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM36, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076859, Oct. 2015.
474. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM37, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076860, Oct. 2015.
475. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM38, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076861, Oct. 2015.
476. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM39, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076862, Oct. 2015.
477. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM41, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076863, Oct. 2015.
478. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM42, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076864, Oct. 2015.
479. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM44, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076865, Oct. 2015.
480. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM48, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076866, Oct. 2015.
481. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM49, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076867, Oct. 2015.
482. **Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Natsumi Yoshioka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: WYM50, --- 16S rRNA gene analysis of bacteria from saba narezushi (fermented sushi of Mackerel) produced at Wakayama by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database.,* LC076868, Oct. 2015.
483. **井上 翔太, 都築 弘充, 小林 大起, 村中 貴恵, 森 健太朗, 佐藤 高則, 山本 真由美, 中山 信太郎 :** 東日本大震災から学ぶ，防災教育の実態とあり方 ∼福島と徳島における放射線教育についての調査報告を含めて∼, *平成27年度総合科学教育部プロジェクト研究I 発表会,* 2016年2月.
484. **井上 翔太, 都築 弘充, 小林 大起, 村中 貴恵, 森 健太朗, 佐藤 高則, 山本 真由美, 中山 信太郎 :** プロジェクト研究Ⅰ(中山・佐藤・山本グループ)報告書, --- 東日本大震災から学ぶ，防災教育の実態とあり方 ∼福島と徳島における放射線教育についての調査報告を含めて∼ ---, *平成27年度総合科学教育部プロジェクト研究I 報告書,* 1-13, 徳島, 2016年1月.
485. **中山 信太郎, 佐藤 高則, 山本 真由美, 都築 弘充, 井上 翔太, 小林 大起, 森 健太朗, 村中 貴恵 :** 東日本大震災から学ぶ防災 ∼当事者は君だ!∼, *平成27年度総合科学部部局長裁量経費「地域交流プロジェクト」成果報告書,* 1, 徳島, 2016年2月.
486. **渡部 稔, 佐藤 高則, 大橋 眞 :** 平成27年度全学共通教育授業方法等改善経費使用実績報告書, --- 基礎生物学実験 ---, 1, 徳島, 2016年3月.
487. **佐藤 高則 :** The 9th Symposium of Asian Biophysics Association (ABA2015)国際会議での研究成果発表, *平成27年度総合科学部創生研究プロジェクト 研究成果発表旅費支援 報告書,* 1, 徳島, 2016年3月.
488. **佐藤 高則 :** 教育改革推進のための全学FD支援事業(共通教育センター), *平成27年度学長裁量経費による成果報告書,* 1-2, 徳島, 2016年3月.
489. **齊藤 隆仁, 小山 晋之 :** 基礎物理学実験テキスト (2016年度後期改訂版), 徳島大学生協, 2016年10月.
490. **西山 賢一, 日本地形学連合編 :** 地形の辞典, 朝倉書店, 東京都, 2017年1月.
491. **徳島県教育委員会(編), 石田 啓祐 :** 南海地震阿波地震津波碑及び津波標石碑調査報告書, --- 地質学から診た地震津波碑 ー立地・石材・保存の意義ー ---, 徳島県教育委員会, 徳島, 2017年3月.
492. **徳島県教育委員会(編), 石田 啓祐 :** 赤色顔料生産遺跡及び関連遺跡の調査 採掘遺跡 石器編, --- 赤色顔料採掘遺跡出土の石杵・石臼石材の産地と石材の特性 ---, 徳島県教育委員会, 徳島, 2017年3月.
493. **徳島県教育委員会(編), 石田 啓祐 :** 赤色顔料生産遺跡及び関連遺跡の調査 採掘遺跡 土器編, --- 徳島県丹生谷-津乃峰山塊の赤色顔料鉱物はどのように産するのか –地質学の調査研究から- ---, 徳島県教育委員会, 徳島, 2017年3月.
494. **西本 和哉, 石田 啓祐, 植地 岳彦 :** 赤色顔料生産遺跡及び関連遺跡の調査 採掘遺跡 土器編, 徳島県教育委員会, 徳島, 2017年3月.
495. **今井 昭二, 山本 祐平, 上村 健 :** 本州日本海沿いの遠隔地における降雪中の無機小球体粒子の組成と長距離輸送, *分析化学,* **Vol.65,** *No.4,* 211-219, 2016年.
496. **L. M. Ahnen, Reiko Orito *and* et al (143 coauthors) :** MAGIC observations of the February 2014 flare of 1ES 1011+496 and ensuing constraint of the EBL density, *Astronomy & Astrophysics,* **Vol.590,** id.A24, 2016.
497. **L. M. Ahnen, Reiko Orito *and* et al (154 coauthors) :** Deep observation of the NGC 1275 region with MAGIC: search of diffuse -ray emission from cosmic rays in the Perseus cluster, *Astronomy & Astrophysics,* **Vol.589,** id.A33, 2016.
498. **K. Abe, Reiko Orito *and* et al (37 coauthors) :** Measurements of Cosmic-Ray Proton and Helium Spectra from the BESS-Polar Long-duration Balloon Flights over Antarctica, *The Astrophysical Journal,* **Vol.822,** *No.2,* id.65, 2016.
499. **Ritesh Haldar, Munehiro Inukai, Satoshi Horike, Kazuhiro Uemura, Susumu Kitagawa *and* Kumar Tapas Maji :** 113Cd Nuclear Magnetic Resonance as a Probe of Structural Dynamics in a Flexible Porous Framework Showing Selective O2/N2 and CO2/N2 Adsorption, *Inorganic Chemistry,* **Vol.55,** *No.9,* 4166-4172, 2016.
500. **L. M. Ahnen, Reiko Orito *and* et al (153 coauthors) :** Super-orbital variability of LS I +61°303 at TeV energies, *Astronomy & Astrophysics,* **Vol.591,** id.A76, 2016.
501. **J. Aleksić, Reiko Orito *and* et al (164 coauthors) :** Insights into the emission of the blazar 1ES 1011+496 through unprecedented broadband observations during 2011 and 2012, *Astronomy & Astrophysics,* **Vol.591,** id.A10, 2016.
502. **Ken-Ichi Fushimi, Hiroyasu Ejiri, R. Hazama, H. Ikeda, K. Imagawa, K. Inoue, G. Kanzaki, A. Kozlov, Reiko Orito, T. Shima, Y. Takemoto, Y. Teraoka, S. Umehara, K. Yasuda *and* S. Yoshida :** Dark matter search project PICO-LON, *Journal of Physics: Conference Series,* **Vol.718,** 042022, 2016.
503. **T Iida, K Nakajima, S Ajimura, T Batpurev, W M Chan, Ken-Ichi Fushimi, R Hazama, H Kakubata, B T Khai, T Kishimoto, X Li, T Maeda, A Masuda, K Matsuoka, N Nakatani, M Nomachi, S Noshiro, I Ogawa, T Ohata, H Ohsumi, K Suzuki, Y Tamagawa, K Tetsuno, V T T Trang, T Uehara, S Umehara *and* S Yoshida :** Status and future prospect of 48Ca double beta decay search in CANDLES, *Journal of Physics: Conference Series,* **Vol.718,** 062026, 2016.
504. **Queaño L. Karlo, Dimalanta B. Carla, Jr. P. Yumul Graciano, Marquez J. Edanjarlo, Faustino-Eslava V. Decibel, Suzuki Shigeyuki *and* Keisuke Ishida :** Stratigraphic units overlying the Zambales Ophiolite Complex in Luzon, Philippines: Tectonostratigraphic significance and regional implications, *Journal of Asian Earth Sciences,* **Vol.Special issue "Crust Mantle Evolution",** *No.published online,* in-press, 2016.
505. **Naoki Ogiwara, Munehiro Inukai, Tomoya Itakura, Satoshi Horike *and* Susumu Kitagawa :** Fast Conduction of Organic Cations in Metal Sulfate Frameworks, *Chemistry of Materials,* **Vol.28,** *No.11,* 3968-3975, 2016.
506. **L. M. Ahnen, Reiko Orito *and* et al (161 coauthors) :** Investigating the peculiar emission from the new VHE gamma-ray source H1722+1, *Monthly Notices of the Royal Astronomical Society,* **Vol.459,** *No.3,* 3271-3281, 2016.
507. **L. M. Ahnen, Reiko Orito *and* et al (160 coauthors) :** Multiwavelength observations of the blazar 1ES 1011+496 in Spring 2008, *Monthly Notices of the Royal Astronomical Society,* **Vol.459,** *No.3,* 2286-2298, 2016.
508. **Munehiro Inukai, Satoshi Horike, Tomoya Itakura, Ryota Shinozaki, Naoki Ogiwara, Daiki Umeyama, Sanjog Nagarkar, Yusuke Nishiyama, Michal Malon, Akari Hayashi, Takashi Ohhara, Ryoji Kiyanagi *and* Susumu Kitagawa :** Encapsulating Mobile Proton Carriers into Structural Defects in Coordination Polymer Crystals: High Anhydrous Proton Conduction and Fuel Cell Application, *Journal of the American Chemical Society,* **Vol.138,** *No.27,* 8505-8511, 2016.
509. **H. Takiya, K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, T. Norita, H. Ogawa, H. Sekiya, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y.I tow, R. Kegasa, K. Kobayashi, K. Masuda, Ken-Ichi Fushimi, K. Martens, Y. Suzuki, R. Fujita, K. Hosokawa, K. Miuchi, N. Oka, Y. Onishi, Y. Takeuchi, Y.H. Kim, J.S. Lee, K.B. Lee, M.K. Lee, Y. Fukuda, K. Nishijima *and* S. Nakamura :** A measurement of the time profile of scintillation induced by low energy gamma-rays in liquid xenon with the XMASS-I detector, *Nuclear Instruments and Methods in Physics Research Section A: Accelerators, Spectrometers, Detectors and Associated Equipment,* **Vol.834,** 192-196, 2016.
510. **Satoshi D. Ohmura, Choki MUROI, Hajime SAKATA, Makoto Wada *and* Norikazu Miyoshi :** Depolymerization of Waste PET with Phosphoric Acid-Modified Silica Gel Under Microwave Irradiation, *Journal of Polymers and the Environment,* **Vol.25,** *No.2,* 250-257, 2016.
511. **Takashi Yamamoto, Aoi Teramachi, Akihiro Orita, Akihito Kurimoto, Takashi Motoi *and* Tsunehiro Tanaka :** Generation of Strong Acid Sites on Yttrium-Doped Tetragonal ZrO2-Supported Tungsten Oxides: Effects of Dopant Amounts on Acidity, Crystalline Phase, Kinds of Tungsten Species, and Their Dispersion, *The Journal of Physical Chemistry C,* **Vol.120,** *No.35,* 19705-19713, 2016.
512. **Suzuki Shigeyuki, Peña E. Rolando, Tam A. Tomas, Jr. P. Yumul Graciano, Dimalanta B. Carla, Usui Mayumi *and* Keisuke Ishida :** Development of the Philippine Mobile Belt in northern Luzon from Eocene to Pliocene, *Journal of Asian Earth Sciences,* **Vol.Special issue "Crust Mantle Evolution",** *No.(published online),* in-press, 2016.
513. **Shinobu Takao, Oki Sekizawa, Gabor Samjeské, Shin-ichi Nagamatsu, Takuma Kaneko, Kotaro Higashi, Takashi Yamamoto, Kensaku Nagasawa, Xiao Zhao, Tomoya Uruga *and* Yasuhiro Iwasawa :** Spatially Non-Uniform Degradation of Pt/C Cathode Catalysts in Polymer Electrolyte Fuel Cells Imaged by Combination of Nano XAFS and STEM-EDS Techniques, *Topics in Catalysis,* **Vol.59,** *No.19,* 1722-1731, 2016.
514. **L. M. Ahnen, Reiko Orito *and* et al (150 coauthors) :** Long-term multi-wavelength variability and correlation study of Markarian 421 from 2007 to 2009, *Astronomy & Astrophysics,* **Vol.593,** id.A91, 2016.
515. **Suzuki Shigeyuki, Keisuke Ishida, Yamashita Shinji *and* Takesue Norito :** Deformation in shear zone of the Butsuzo Tectonic Line under syn-sedimentary to diagenetic conditions, *Abstracts Volume of GSA 2016 Denber, Colorado, USA,* **Vol.Online,** *No.#281331,* Session265-Booth438, 2016.
516. **Takuma Kaneko, Gabor Samjeské, Shin-ichi Nagamatsu, Kotaro Higashi, Sekizawa Oki, Takao Shinobu, Takashi Yamamoto, Xiao Zhao, Tomoya Uruga *and* Yasuhiro Iwasawa :** Key Structural Kinetics for Carbon Effects on the Performance and Durability of Pt/Carbon Cathode Catalysts in Polymer Electrolyte Fuel Cells Characterized by in Situ Time-Resolved XAFS, *The Journal of Physical Chemistry C,* **Vol.120,** *No.42,* 24250-24264, 2016.
517. **Y. Toyomori, S. Tsuji, S. Mitsuda, Y. Okayama, A. Mori, T. Kobayashi, Y. Miyazaki, T. Yaita, S. Arae, T. Takahashi *and* Masamichi Ogasawara :** Bithiophene with Winding Vine-Shaped Molecular Asymmetry: Preparation, Structural Characterization, and Enantioselective Synthesis, *Bulletin of the Chemical Society of Japan,* **Vol.89,** *No.12,* 1480-1486, 2016.
518. **西山 賢一 :** 粗粒砂岩の風化による岩石物性の変化, *応用地質,* **Vol.57,** *No.5,* 213-218, 2016年.
519. **Takehito Watanabe, Masafumi Uematsu, Yasser Hehamed, Hiroshi Eguchi, Shoji Imai *and* Takashi Kitaoka :** Corneal Erosion With Pigments Derived From a Cosmetic Contact Lens: A Case Report, *Eye and Contact Lens,* 2017.
520. **Hao Hu, Yi Wei Wu, Tamotsu Takahashi, Kazuhiro Yoshida *and* Masamichi Ogasawara :** Enantioselective Synthesis of Ferrocene- or Cymantrene-Fused Planar-Chiral Phospholes, *European Journal of Inorganic Chemistry,* **Vol.2017,** *No.2,* 325-329, 2017.
521. **山本 祐平, 喜多 郁弥, 磯野 成美, 今井 昭二 :** 鉄(III)イオン競合によるフミン酸と有害微量金属イオンの錯生成反応への影響, *分析化学,* **Vol.66,** *No.12,* 875-883, 2017年.
522. **Ken Kamikawa, Ya-Yi Tseng, Jia-Hong Jian, Tamotsu Takahashi *and* Masamichi Ogasawara :** Planar-Chiral Phosphine-Olefin Ligands Exploiting a (Cyclopentadienyl)manganese(I) Scaffold to Achieve High Robustness and High Enantioselectivity, *Journal of the American Chemical Society,* **Vol.139,** 1545-1553, 2017.
523. **Yuki Saito, Haruro Ishitani, Masaharu Ueno *and* Shu Kobayashi :** Selective Hydrogenation of Nitriles to Primary Amines Catalyzed by a Polysilane/SiO2-Supported Palladium atalyst unde Continuous-Flow Conditions, *ChemistryOpen,* **Vol.6,** *No.2,* 211-215, 2017.
524. **今井 昭二, 山本 祐平, 佐名川 洋右, 耒見 祐哉, 黒谷 功, 西本 潤, 菊地 洋一 :** 2008年から2014年における高知県梶ヶ森山頂の樹氷と降雪中のカドミウム，鉛，非海塩性硫酸イオンの長距離輸送機構, *分析化学,* **Vol.66,** *No.2,* 95-113, 2017年.
525. **Masayuki Torii, Mitsuru Okuno, Ken-ichi Nishiyama, Yoshito Kitazono, Toshiaki Hasenaka *and* Shuichiro Yokota :** Three types of sediment movement in the Sannoudani River, Minamiaso Village, Kumamoto Prefecture, southwestern Japan, *Journal of the Sedimentological Society of Japan,* **Vol.75,** *No.2,* 64, 2017.
526. **Masamichi Ogasawara, Ya-Yi Tseng, Qiang Liu, Ninghui Chang, Xicheng Yang, Tamotsu Takahashi *and* Ken Kamikawa :** Kinetic Resolution of Planar-Chiral (5-Bromocyclopentadienyl)manganese(I) Complexes by Molybdenum-Catalyzed Asymmetric Ring-Closing Metathesis, *Organometallics,* **Vol.36,** *No.7,* 1430-1435, 2017.
527. **Yu Kawasaki, Ryo MORIOKA, Yutaka Kishimoto, Koichi Nakamura, Kohei NISHIYAMA, Takehide KOYAMA, Takeshi MITO, Takuyuki BABA, Toru YAMAUCHI, Masahiko ISOBE *and* Yutaka UEDA :** 51V-NMR study of charge order induced by cation order in -Ag2/3V2O5, *Journal of Physics: Conference Series,* **Vol.807,** *No.062001-1 062001-6,* 2017.
528. **西山 賢一, 瀬部 直之, 石田 啓祐, 中尾 賢一, 辻野 泰之 :** 地形・地質から見た鳴門海峡の成立, *阿波学会紀要,* **Vol.61,** 1-10, 2017年.
529. **西山 賢一, 石田 啓祐, 中尾 賢一, 辻野 泰之 :** 「ドイツ橋」および「ドイツ兵の慰霊碑」に使用された撫養石石材の産地と経年風化, *阿波学会紀要,* **Vol.61,** 177-184, 2017年.
530. **山本 孝, 近藤 真季, 入江 智章, 谷間 直人 :** タングステンジルコニウムオキシ水酸化物結晶脱水過程のXRD/XAFS観察およびその酸触媒特性, *X線分析の進歩,* **Vol.48,** 137-148, 2017年.
531. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Isolation and Identification of Cu, Fe-tolerant Bacteria from Myoho Copper Mine Spoils, Wakayama, *Proceedings of 11th International Congress on Extremophiles,* 155, 2016.
532. **Takanori Satoh, Yui Hamaji, Daiki Fujimoto, Mitsuki Fujii, Kouta Nagai, Narumi Maegawa, Natsumi Yoshioka, Mayu Kanamoto, Kenji Akiyoshi-Hiraoka, Yurie Ohiwa *and* Shota Inoue :** Exploration for bacteria from Japanese fermented foods, Narezushi and Awa Bancha tea by 16S rRNA gene analysis, *Proceedings of 8th Asian Symposium on Microbial Ecology(ASME2016),* S7-17, 2016.
533. **S. Umehara, T. Kishimoto, M. Nomachi, S. Ajimura, N. Nakatani, K. Mtasuoka, K. Ichimura, M. Saka, T. Ishikawa, D. Tanaka, M. Tanaka, S. Yoshida, K. Suzuki, G. Ito, H. Kakubata, W. Wang, J. Takemoto, W.M. Chan, M. Doihara, Y. Tamagawa, I. Ogawa, T. Ueno, S. Maeda, A. Yamamoto, S. Tomita, G. Fujita, A. Kawamura, T. Harada, Ken-Ichi Fushimi, Ryuta Hazama, H. Ohsumi *and* K. Okada :** Search for neutrino-less double beta decay of 48 Ca CANDLES, *Proceedings of Physics(INPC2916) The 26th International Nuclear Physics Conference,* 246, 2016.
534. **K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, T. Norita, H. Ogawa, H. Sekiya, O. Takachio, A. Takeda, M. Yamashita, B.S. Yanga, N.Y. Kim, Y.D. Kim, S. Tasaka, Ken-Ichi Fushimi, J. Liue, K. Martense, Y. Suzuki, B.D. Xue, R. Fujita, K. Hosokawa, K. Miuchi, Y. Onishi, N. Oka, Y. Takeuchi, Y.H. Kim, J.S. Lee, K.B. Lee, M.K. Lee, Y. Fukuda, Y. Itow, R. Kegasa, K. Kobayashi, K. Masuda, H. Takiya, K. Nishijima *and* S. Nakamura :** Direct dark matter search by annual modulation in XMASS-I, *Physics Letters B,* **Vol.759,** 272-276, 2016.
535. **A. Gando, Y. Gando, T. Hachiya, A. Hayashi, S. Hayashida, H. Ikeda, K. Inoue, K. Ishidoshiro, Y. Karino, M. Koga, S. Matsuda, T. Mitsui, K. Nakamura, S. Obara, T. Oura, H. Ozaki, I. Shimizu, Y. Shirahata, J. Shirai, A. Suzuki, T. Takai, K. Tamae, Y. Teraoka, K. Ueshima, H. Watanabe, A. Kozlov, Y. Takemoto, S. Yoshida, Ken-Ichi Fushimi, T.I. Banks, B.E. Berger, B.K. Fujimura, T. O'Donnell, L.A. Winslow, Y. Efremenko, H.J. Kanwowski, D.M. Markoff, W. Tornow, J.A. Detwiler, S. Enomoto *and* M.P. Decowski :** Search for Majorana Neutrinos Near the Inverted Mass Hierarchy Region with KamLAND-Zen, *Physical Review Letters,* **Vol.117,** 082503, 2016.
536. **A. Gando, Y. Gando, T. Hachiya, A. Hayashi, S. Hayashida, H. Ikeda, K. Inoue, K. Ishidoshiro, Y. Karino, M. Koga, S. Matsuda, T. Mitsui, K. Nakamura, S. Obara, T. Oura, H. Ozaki, I. Shimizu, Y. Shirahata, J. Shirai, A. Suzuki, T. Takai, K. Tamae, Y. Teraoka, K. Ueshima, H. Watanabe, A. Kozlov, Y. Takemoto, S. Yoshida, Ken-Ichi Fushimi, T.I. Banks, B.E. Berger, B.K. Fujimura, T. O'Donnell, L.A. Winslow, Y. Efremenko, H.J. Kanwowski, D.M. Markoff, W. Tornow, J.A. Detwiler, S. Enomoto *and* M.P. Decowski :** A SEARCH FOR ELECTRON ANTINEUTRINOS ASSOCIATED WITH GRAVITATIONAL-WAVE EVENTS GW150914 AND GW151226 USING KAMLAND, *The Astrophysical Journal Letters,* **Vol.829,** *No.2,* L34, 2016.
537. **Zenro Hioki, Kazumasa Ohkuma *and* Akira Uejima :** Refined analysis and updated constraints on general non-standard tbW couplings, *Physics Letters B,* **Vol.761,** 219-222, 2016.
538. **佐藤 高則, 秋吉 研二, Ahmed Ibrahim Elhossany Elmarhomy, 武岡 大輔, 時久 栞, 井上 翔太, 小林 大起, 都築 弘充, 村中 貴恵, 森 健太朗, 小野 覚久, 山本 真由美, 中山 信太郎 :** 東日本大震災から学ぶ防災教育の実態と課題, *平成28年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 38-39, 2016年.
539. **佐藤 高則, 石田 啓祐, 小山 晋之, 久田 旭彦, 渡部 稔, 秋山 治彦, 三好 徳和 :** 高校生向け課題研究研修会によるアクティブ・ラーニング型高大連携とFDへの展開, *平成28年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 28-29, 2016年.
540. **伏見 賢一, 齊藤 隆仁, 小山 晋之, 中山 信太郎, 折戸 玲子, 久田 旭彦, 真岸 孝一 :** 基礎物理学実験テキスト 総合科学部・医学部医学科・歯学部歯学科, *基礎物理学実験テキスト,* 2016年4月.
541. **伏見 賢一 :** 観測指針4月, *天文回報,* **Vol.885,** 2-4, 2016年4月.
542. **伏見 賢一 :** 観測指針5月, *天文回報,* **Vol.886,** 2-4, 2016年5月.
543. **伏見 賢一 :** 観測指針6月, *天文回報,* **Vol.887,** 2-4, 2016年6月.
544. **伏見 賢一 :** 観測指針7月, *天文回報,* **Vol.888,** 2-3, 2016年7月.
545. **日置 善郎 :** ディラック方程式入門, --- 量子論と相対論を結びつける ---, *数理科学,* **Vol.54,** *No.7,* 7-13, 2016年7月.
546. **伏見 賢一 :** 観測指針8月, *天文回報,* **Vol.889,** 2-3, 2016年8月.
547. **伏見 賢一 :** 観測指針9月, *天文回報,* **Vol.890,** 2-4, 2016年9月.
548. **村田 明広 :** 布田川・日奈久断層帯という活断層が引き起こした熊本地震, *いのち輝く, No.83,* 7-11, 2016年9月.
549. **伏見 賢一 :** 観測指針10月, *天文回報,* **Vol.891,** 2-5, 2016年10月.
550. **伏見 賢一 :** 観測指針11月, *天文回報,* **Vol.891,** 2-5, 2016年11月.
551. **伏見 賢一 :** 観測指針12月, *天文回報,* **Vol.891,** 2-7, 2016年12月.
552. **Masamichi Ogasawara :** Cross-Coupling Reactions Other Than Suzuki-Miyaura Coupling, *Science of Synthesis, Reference Library,* **Vol.2016/5a,** 141-159, Jan. 2017.
553. **Yu Kawasaki, Ryo MORIOKA, Yutaka Kishimoto, Koichi Nakamura, Kohei NISHIYAMA, Takehide KOYAMA, Takeshi MITO, Takuyuki BABA, Toru YAMAUCHI, Masahiko ISOBE *and* Yutaka UEDA :** 51V-NMR study of charge order induced by cation order in -Ag2/3V2O5, *The International Conference on Strongly Correlated Electron Systems,* Hangzhou,China, May 2016.
554. **Yuhei Yamamoto, Yosuke Sanagawa, Yuya Kurumi, Takeshi Kamimura, Kenji Kodama *and* Shoji Imai :** Chemical properties of wet depositions at a remote site in Shikoku highland, Japan, *Goldschmidt Conference 2016,* Yokohama, Jun. 2016.
555. **Zenro Hioki, Kazumasa Ohkuma *and* Akira Uejima :** Constraints on generalized nonstandard tbW couplings, *38th International Conference on High Energy Physics,* Chicago, Aug. 2016.
556. **Masamichi Ogasawara :** Enantioselective Synthesis of Planar-Chiral Zirconocenes by Asymmetric Ring-Closing Metathesis and Their Application in ZACA Reaction, *International Symposium on Pure & Applied Chemistry 2016 (ISPAC 2016),* Kuching, Aug. 2016.
557. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Isolation and Identification of Cu, Fe-tolerant Bacteria from Myoho Copper Mine Spoils, Wakayama, *11th International Congress on Extremophiles,* 155, Kyoto, Sep. 2016.
558. **Takanori Satoh, Yui Hamaji, Daiki Fujimoto, Mitsuki Fujii, Kouta Nagai, Narumi Maegawa, Natsumi Yoshioka, Mayu Kanamoto, Kenji Akiyoshi-Hiraoka, Yurie Ohiwa *and* Shota Inoue :** Exploration for bacteria from Japanese fermented foods, Narezushi and Awa Bancha tea by 16S rRNA gene analysis, *The 8th Asian Symposium on Microbial Ecology(ASME2016),* S7-17, Taipei, Oct. 2016.
559. **Mitsuru OKUNO, Masayuki TORII, Ken-ichi Nishiyama *and* Shuichiro YOKOTA :** Landslides and earth-flow deposits triggered by the 2016 Kumamoto Earthquake in Aso caldera, SW Japan., *Geological Society of America Annual Meeting 2016,* Nov. 2016.
560. **Masamichi Ogasawara :** Enantioselective Synthesis of P-Stereogenic Phosphine Derivatives by Molybdenum-Catalyzed "Formal" Cross-Metathesis: Use of Chromium-Template for High Enantio-, Chemo-, and E/Z-Selectivities, *International Symposium on Catalysis and Fine Chemicals 2016 (C&FC 2016),* Taipei, Nov. 2016.
561. **Ken-Ichi Fushimi, Hiroyasu Ejiri, Ryuta Hazama, H. Ikeda, K. Inoue, K. Imagawa, G. Kanzaki, A. Kozlov, Reiko Orito, Tatsushi Shima, Y. Takemoto, Y. Teraoka, S. Umehara *and* S. Yoshida :** High purity NaI(Tl) scintillator to search for dark matter, *JPS Conference Proceedings: Proceedings of International Symposium on Radiation Detectors and Their Uses (ISRD2016),* **Vol.11,** 020003(1)-020003(6), つくば市, Nov. 2016.
562. **Akihiko Hisada, Kie Muranaka, Kuniyuki Koyama, Ko-ichi Magishi, Takahito Saito *and* Makoto Haguwara :** Oxidation and reduction effects of successive superconducting transitions in ultra-fineYBa2Cu4O8 ceramics, 29th International Symposium on Superconductivity (ISS2016), *ISS2016,* Tokyo, Dec. 2016.
563. **S. Umehara, T. Kishimoto, M. Nomachi, S. Ajimura, T. Iida, Y. Takemoto, K. Matsuoka, V.T.T. Trang, S. Yoshida, W. Wang, W.M. Chang, T. Maeda, T. Ohata, K. Tetsuno, T. Uehara, X. Lee, B. Temuge, K. Akutagawa, K. Kanagawa, S. Katagiri, M. Tsuzuki, N. Yotsunaga, Y. Tamagawa, I. Ogawa, K. Nakajima, S. Noshiro, A. Masuda, K. Morishita, M. Washino, N. Takahashi, K. Teranishi, F. Tokaku, T. Hiyama, Ken-Ichi Fushimi, K. Mori, Ryuta Hazama, N. Nakatani, K. Suzuki *and* H. Ohsumi :** Search For Neutrino-less Double Beta Decay Of 48Ca- Candles -, *Proceedings of Science,* **Vol.281,** 2017.
564. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Applications, *TMU and PetroMat Joint Mini Symposium,* Bangkok, Feb. 2017.
565. **Masamichi Ogasawara :** Total Synthesis of Axially Chiral Naturally Occurring Allenes Utilizing Palladium-Catalyzed Allene Synthesis Reaction, *Pure and Applied Chemistry International Conference 2017 (PACCON 2017),* Bangkok, Feb. 2017.
566. **犬飼 宗弘, 田村 優実, 中村 浩一 :** 規則性ナノ空間に閉じ込められた二酸化炭素の特異的なダイナミクス, *第59回 固体NMR・材料フォーラム,* 2016年5月.
567. **山本 祐平, 佐名川 洋右, 耒見 祐哉, 上村 健, 児玉 憲治, 今井 昭二 :** 四国の山岳地帯における湿性沈着の化学的特徴, *第25回環境化学討論会,* 2016年6月.
568. **Natsumi Yoshioka, Mayu Kanamoto, Yui Hamaji, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa, Zesem Ichinkhorloo, Janlav Munkhtsetseg *and* Takanori Satoh :** Functional analysis for whey protein and peptides in Mongolian animal milks, *The 16th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* 2P084, Jun. 2016.
569. **Shota Inoue, Kouta Nagai, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Daiki Fujimoto, Narumi Maegawa, Yurie Ohiwa *and* Takanori Satoh :** Contribution of Thr91 to the conformational stabilities in Geobacillus thermocatenulatus Uridine kinase, *The 16th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* Jun. 2016.
570. **Yurie Ohiwa, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Mayu Kanamoto, Yui Hamaji, Natsumi Yoshioka, Shota Inoue *and* Takanori Satoh :** Contribution of amino acids to the thermostability of Geobacillus stearothermophilus Nucleoside diphosphate kinase, *The 16th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* Jun. 2016.
571. **中村 浩一, 犬飼 宗弘, 森賀 俊広, 岩井 良樹, 桑田 直明, 河村 純一 :** 局所構造の擾乱にともなうニオブ酸ナトリウムのイオン伝導挙動の変化, *第19回超イオン導電体物性研究会講演要旨集,* 79-82, 2016年6月.
572. **上野 雅晴 :** 実践的なグリーンケミストリーへの挑戦, *日本プロセス学会第20回プロセス化学東四国フォーラムセミナー,* 2016年6月.
573. **小笠原 正道 :** 均一系遷移金属触媒による遷移金属錯体の立体選択的分子変換とその応用, *第49回有機金属若手の会 夏の学校,* 2016年7月.
574. **村中 貴恵, 浜崎 寛子, 中島 弘貴, 小山 晋之, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 萩原 亮 :** 酸素量の変化しない超微細セラミックスYBa2Cu4O8単相試料の二段階転移, *2016年度 応用物理・物理系学会中国四国支部 合同学術講演会,* 2016年7月.
575. **齋藤 あゆみ, 塩見 太嘉良, 日達 涼太, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 小山 晋之 :** Fe-Se二元系の合成及びその輸送特性, *2016年度 応用物理・物理系学会中国四国支部 合同学術講演会,* 2016年7月.
576. **久田 旭彦 :** 高ドープ領域の梯子格子銅酸化物の圧力誘起超伝導, *2016年度 応用物理・物理系学会中国四国支部 合同学術講演会,* 9, 2016年7月.
577. **西山 賢一, 鳥井 真之, 奥野 充, 井口 隆 :** 2016年熊本地震で発生した地すべり・斜面崩壊の地形・地質的特徴, *日本地質学会第123年学術大会講演要旨集,* 149, 2016年9月.
578. **三好 德和, 北方 咲, 立田 彩和, 西澤 高宏, 大村 D. 聡, 上野 雅晴 :** ストロンチウム-メタラサイクル中間体の合成とその反応性に関する研究, *第27回基礎有機化学討論会, 広島国際会議場(広島), 講演番号 1C05,* 2016年9月.
579. **大村 D. 聡, 森田 健司, 三好 德和, 平尾 俊一, 森内 敏之 :** アミノ酸部位の導入により誘起されたレドックス活性フェニレンジアミンの不斉構造, *第27回基礎有機化学討論会, 広島国際会議場(広島), 講演番号 2P067,* 2016年9月.
580. **山名 啓介, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 有機機能性材料を指向したトリアリールアミン誘導体の合成と電子的特性, *第27回基礎有機化学討論会, 広島国際会議場(広島), 講演番号 2P084,* 2016年9月.
581. **山本 孝, 寺町 葵 :** イットリア-ジルコニア担持酸化タングステン触媒表面上での複合酸化物の生成, *第19回XAFS討論会,* 2016年9月.
582. **北尾 泰広, 青矢 睦月 :** 後期白亜紀ユーラシア東縁沈み込み帯の地質を巡る徳島周辺のジオ鉄コース, *日本地質学会第123年学術大会講演要旨集,* 225, 2016年9月.
583. **野田 篤, 植木 岳雪, 川畑 博, 松浦 浩久, 青矢 睦月 :** 5 万分の1 地質図幅「観音寺」地域の地質, *日本地質学会第123年学術大会講演要旨集,* 2016年9月.
584. **石田 啓祐, Sizuki Shigeyuki, Dimalanta B. Carla, Jr. Graciano P. Yumul, R. Lawrence Zamoras, Michel Faure, Francis Hirsch, Murat Ali Kilic, Pablo Placencia :** Litho- and microbiostratigraphy of Triassic carbonates in the Buruanga Peninsula, Panay Island, Philippines, *Abstract Volume of the GSJ 123th Annual Meeting,* **Vol.123,** R9O6, 2016年9月.
585. **中村 浩一, 竹内 智史, 犬飼 宗弘, 森賀 俊広 :** NaCoO2のNa組成とNaサイト周辺の局所構造, *日本物理学会2016年秋季大会 講演概要集,* 2016年9月.
586. **今井 昭二, 山本 祐平, 耒見 祐哉, 佐名川 洋右, 齋藤 あゆみ :** 日本海沿岸および四国の遠隔地における降雪中のPb/Cd比を用いた長距離輸送機構解明のためのバックグラウンド, *日本分析化学会第65年会 北海道大学・札幌,* 2016年9月.
587. **神川 憲, 曾 雅怡, 小笠原 正道 :** 面不斉シクロペンタジエニルマンガンカルボニル錯体を基盤とした第二世代型P-オレフィン配位子の開発, *第63回有機金属化学討論会,* 2016年9月.
588. **小笠原 正道, 胡 淏, 呉 威毅, 吉田 和弘, 高橋 保 :** メタロセン縮環した面不斉ホスホールの不斉合成, *第63回有機金属化学討論会,* 2016年9月.
589. **村中 貴恵, 小山 晋之, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 萩原 亮 :** 酸素量の変化しない超微細セラミックスYBa2Cu4O8単相試料の二段階転移の酸化・還元効果, *日本物理学会講演概要集,* 2016年9月.
590. **蕨野 龍世, Ka S., 出口 博之, 美藤 正樹, 萩原 亮, 小山 晋之 :** YBCO超伝導セラミクスにおけるグレイン間相転移の磁場-温度相図, *日本物理学会講演概要集,* 2016年9月.
591. **久田 旭彦, 小山 晋之, 齊藤 隆仁, 真岸 孝一, 藤原 直樹, 上床 美也, G Deng, E Pomjakushina, K Conder, D Mohan Radheep, R Thiyagarajan, S Esakkimuthu, S Arumugam :** 高圧下NMRによる 梯子格子銅酸化物Sr1.4Ca12.6Cu24O41の圧力誘起超伝導の研究, *日本物理学会講演概要集,* **Vol.71,** *No.1,* 2016年9月.
592. **真岸 孝一, 久田 旭彦, 齊藤 隆仁, 小山 晋之, 野坂 洸太, 雷前 伸, 並木 孝洋, 西村 克彦 :** Nd(V,Ti)2Al20のNMR, *日本物理学会講演概要集,* 2016年9月.
593. **山本 祐平, 今井 昭二, 白崎 俊浩, 米谷 明, 田上 梓 :** 黒鉛炉型原子吸光光度計による高温原子化元素の定量における化学修飾剤の化学状態変化が吸光度に及ぼす影響, *日本分析化学会 第65回年会,* 2016年9月.
594. **瀧本 涼介, 川崎 祐, 岸本 豊, 中村 浩一, 上田 寛 :** Bi4V2O11-δ のNMRによる研究, *日本物理学会2016年秋季大会 講演概要集,* 2016年9月.
595. **山本 孝, 栗本 彰人 :** 13 族イオン添加酸化ジルコニウムによるエタノール転換反応, *第118回触媒討論会,* 2016年9月.
596. **伏見 賢一 :** PICO-LON報告:高感度NaI(Tl)検出器による宇宙暗黒物質探索の現状, *日本物理学会2016年秋季大会(宮崎大学木花キャンパス),* 2016年9月.
597. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Genetic analysis and expression of nucleoside diphosphate kinase gene from Anoxybacillus flavithermus, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 2P-183, Sep. 2016.
598. **Daiki Fujimoto, Kouta Nagai, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Characterization of genes for pyrimidine-nucleoside monophosphate kinases from Anoxybacillus flavithermus, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 2P-189, Sep. 2016.
599. **Shota Inoue, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Yurie Ohiwa *and* Takanori Satoh :** Genetic analysis and expression of uridine kinase gene from Anoxybacillus flavithermus, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 2P-185, Sep. 2016.
600. **Kenji Akiyoshi-Hiraoka, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Genetic analysis and expression of genes for dTMP-forming enzymes from Anoxybacillus flavithermus, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 2P188, Sep. 2016.
601. **Natsumi Yoshioka, Narumi Maegawa, Mayu Kanamoto, Yui Hamaji, Kohta Nagai, Daiki Fujimoto, Kenji Akiyoshi-Hiraoka, Shota Inoue, Yurie Ohiwa, Zesem Ichinkhorloo, Janlav Munkhtsetseg *and* Takanori Satoh :** Lipocalin function of whey proteins from Mongolian animal milks, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 2P-144-2P144, Sep. 2016.
602. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method, *Annual Meeting of The 89th Japanese Biochemical Society, Abstracts,* 3P-277, Sep. 2016.
603. **西山 賢一, 鳥井 真之, 奥野 充, 井口 隆, 加藤 靖郎 :** 2016年熊本地震により熊本県阿蘇地方で発生した斜面崩壊・土石流の地形・地質的特徴, *日本応用地質学会平成28年度研究発表会講演論文集,* 51-52, 2016年10月.
604. **田近 淳, 薄井 敏彦, 矢野 寛幸, 西山 賢一 :** 2016年熊本地震で発生した崩壊ー土石流:とくに垂玉川の例, *日本応用地質学会平成28年度研究発表会講演論文集,* 243-244, 2016年10月.
605. **井口 隆, 高見 智之, 加藤 靖郎, 西山 賢一, 山本 茂雄 :** 熊本地震による京都大学阿蘇火山センター周辺の斜面災害の特徴, *日本応用地質学会平成28年度研究発表会講演論文集,* 145-146, 2016年10月.
606. **日本応用地質学会2016年熊本・大分地震災害調査団陥没性亀裂調査班, 西山 賢一 :** 阿蘇カルデラ内に出現した陥没性亀裂と周辺の地形変位, *日本応用地質学会平成28年度研究発表会講演論文集,* 147-148, 2016年10月.
607. **高見 智之, 井口 隆, 加藤 靖郎, 西山 賢一, 日本応用地質学会九州支部WG斜面災害班 :** 2016年熊本地震で生じた斜面災害の特徴と類型化, *日本応用地質学会平成28年度研究発表会講演論文集,* 149-150, 2016年10月.
608. **朽津 信明, 森井 順之, 西山 賢一 :** 風化形態の違いによる砂岩の侵食速度の違い, *日本応用地質学会平成28年度研究発表会講演論文集,* 201-202, 2016年10月.
609. **西山 賢一, 横田 修一郎 :** 2016年熊本地震で発生した立野地区の斜面崩壊と宇土・御船地区の盛土地すべり, *日本応用地質学会中国四国支部平成28年度研究発表会発表論文集,* 33-38, 2016年10月.
610. **犬飼 宗弘, 田村 優実, 中村 浩一 :** 多孔性配位高分子内部の架橋配位子と二酸化炭素のダイナミクス, *第60回 固体NMR・材料フォーラム,* 2016年10月.
611. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Bacterial analysis of Awa Bancha tea leaves (fermented tea) by clone library method, *Program and Abstracts of the 31st Annual Meeting of the Japanese Society of Microbial Ecology (JSME2016),* 119, Oct. 2016.
612. **山本 孝, 近藤 真季, 入江 智章 :** タングステンジルコニウムオキシ水酸化物の脱水結晶化過程および酸触媒特性, *第52回X線分析討論会,* 2016年10月.
613. **奥野 充, 鳥井 真之, 西山 賢一, 中西 利典, 横田 修一郎 :** 平成28年熊本地震による阿蘇カルデラ内の斜面崩壊堆積物の地形, *地形,* **Vol.38,** *No.1,* 56-57, 2016年11月.
614. **立田 彩和, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウムアミドを利用した合成手法の開発, *2016年日本化学会中国四国支部大会,香川大学(香川),講演番号 1J12,* 2016年11月.
615. **西澤 高宏, 北方 咲, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるメタラサイクル中間体の合成とそれを用いた反応の開発, *2016年日本化学会中国四国支部大会,香川大学(香川),講演番号 1J13,* 2016年11月.
616. **劉 強, 木村 亮介, 和田 志郎, 高橋 保, 吉田 和弘, 小笠原 正道 :** 面不斉4-ジアルキルアミノピリジン誘導体のエナンチオ選択的合成法の開発と応用, *2016年日本化学会中国四国支部大会,* 2016年11月.
617. **瓜生 瑞穂, 曾 雅怡, 石本 寛人, 高橋 保, 神川 憲, 小笠原 正道 :** 不斉閉環メタセシス反応を用いた面不斉マンガン錯体の触媒的不斉合成, *2016年日本化学会中国四国支部大会,* 2016年11月.
618. **日下 亮, 大村 D. 聡, 三好 德和, 上野 雅晴 :** ビスマス塩を触媒として用いるRitter反応の開発, *2016年日本化学会中国四国支部大会,香川大学(香川),講演番号 2P84,* 2016年11月.
619. **上野 雅晴, 喜多 亜希子, 室井 超帰, 大村 D. 聡, 和田 眞, 三好 德和 :** ビスマス塩を触媒として用いるアルコール類の空気酸化反応およびアルデヒドの酸化的エステル化反応の開発, *第49回酸化反応討論会, 講演番号 1O-05,* 2016年11月.
620. **田村 優実, 犬飼 宗弘, 中村 浩一 :** 多孔性配位高分子の空隙に物理吸着させた気体分子のダイナミクス, *第55回 NMR討論会,* 2016年11月.
621. **今井 昭二 :** PM2.5粒子の東アジアからの冬季長距離輸送器項と起源評価, *山口機器分析研究会第85回学術講演会,* 2016年12月.
622. **西山 賢一, 鳥井 真之, 奥野 充, 横田 修一郎 :** 2016年熊本地震により阿蘇崋山地域で発生した地すべり・斜面崩壊の地形・地質的特徴, *第16回日本地質学会四国支部講演会要旨,* 8, 2016年12月.
623. **犬飼 宗弘 :** 多孔性配位高分子の多核NMR, *16-2 NMR 研究会 最新の多核NMR,* 2016年12月.
624. **北尾 泰広, 青矢 睦月 :** 後期白亜紀ユーラシア東縁沈み込み帯の地質を巡る徳島周辺のジオ鉄コース, *第16回日本地質学会四国支部講演要旨集,* 2, 2016年12月.
625. **近藤 真季, 入江 智章, 谷間 直人, 山本 孝 :** タングステンジルコニウムオキシ水酸化物結晶の脱水過程のXRD/XAFS観察およびその酸触媒特性, *触媒学会第26回キャラクタリゼーション講習会,* 2016年12月.
626. **井藤 弘章, 犬飼 宗弘, 森賀 俊広, 岩井 良樹, 桑田 直明, 河村 純一, 中村 浩一 :** LiMPO4(M=Fe, Mn, Co)における高温でのイオン運動と局所構造の変化, *第42回固体イオニクス討論会,* 100-101, 2016年12月.
627. **水田 悠介, 梅岡 優, 中村 浩一, 村井 啓一郎, 森賀 俊広 :** Li量を変化させたLiFe0.2Co0.8O2の合成および電気化学的特性評価, *第23回ヤングセラミストミーティング in 中四国,* 68-69, 2016年12月.
628. **石田 啓祐 :** 古代人が求めた朱の正体, --- 地質学の調査研究から ---, *講演会要旨集「朱を考古学する part II」,* 20-26, 2016年12月.
629. **佐藤 高則, 秋吉 研二, Ahmed Ibrahim Elhossany Elmarhomy, 武岡 大輔, 時久 栞, 井上 翔太, 小林 大起, 都築 弘充, 村中 貴恵, 森 健太朗, 小野 覚久, 山本 真由美, 中山 信太郎 :** 東日本大震災から学ぶ防災教育の実態と課題, *平成28年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 38-39, 2016年12月.
630. **佐藤 高則, 石田 啓祐, 小山 晋之, 久田 旭彦, 渡部 稔, 秋山 治彦, 三好 徳和 :** 高校生向け課題研究研修会によるアクティブ・ラーニング型高大連携とFDへの展開, *平成28年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 28-29, 2016年12月.
631. **金井 純子, 森本 恵美, 井上 貴文, 佐々木 千鶴, 北岡 和義, 日下 一也, 浮田 浩行, 岡本 敏弘, 岸本 豊, 出口 祥啓, 久保 智裕, 安澤 幹人, 寺田 賢治, 藤澤 正一郎 :** 学生の自己能力評価アンケート調査からみたイノベーション教育の課題, *大学教育カンファレンスin徳島,* 2016年12月.
632. **小笠原 正道 :** 均一系遷移金属触媒による遷移金属錯体の立体選択的分子変換, *有機金属部会平成28年度第4回例会,* 2017年1月.
633. **犬飼 宗弘 :** 固体NMRによる多孔性配位高分子のダイナミクス解析と機能発現, *1. 環境調和セラミック材料研究会 名工大-FS調査共同研究支援事業 共同研究 講演会,* 2017年1月.
634. **伏見 賢一 :** PICO-LONの現状報告, *LBGT2017(飛騨市),* 2017年2月.
635. **木村 将大, 久保 誠輝, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるアミド化合物のモノアルキル化反応, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 1PB-110,* 2017年3月.
636. **大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるアミド化合物の還元反応, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 1PB-111,* 2017年3月.
637. **陳 彦洲, 小笠原 正道 :** 3-ブロモ-2,4-ペンタジエニル・エステル類を基質とするパラジウム触媒二重求核置換反応によるC2-対称軸不斉アレンの触媒的不斉合成, *日本化学会第97春季年会,* 2017年3月.
638. **植田 潤一, 小笠原 正道, 吉田 和弘 :** ドミノ酸化/分子内環化反応を用いる含窒素縮合環の形成, *日本化学会第97春季年会,* 2017年3月.
639. **瓜生 瑞穂, 曾 雅怡, 石本 寛人, 大矢 直輝, 高橋 保, 神川 憲, 小笠原 正道 :** モリブデン触媒閉環メタセシス反応による面不斉ホスファシクロペンタジエニル-マンガン錯体の触媒的不斉合成, *日本化学会第97春季年会,* 2017年3月.
640. **劉 強, 木村 亮介, 和田 志郎, 吉田 和弘, 小笠原 正道 :** メタロセン縮環面不斉4-ジアルキルアミノピリジン誘導体のエナンチオ選択的合成法の開発と応用(2), *日本化学会第97春季年会,* 2017年3月.
641. **木村 亮介, 劉 強, 和田 志郎, 小笠原 正道, 吉田 和弘 :** メタロセン縮環面不斉4-ジアルキルアミノピリジン誘導体のエナンチオ選択的合成法の開発と応用(1), *日本化学会第97春季年会,* 2017年3月.
642. **胡 淏, 関 直樹, 小笠原 正道 :** ジフェロセニルホスフィノ基を有する新規不斉配位子の合成及び応用, *日本化学会第97春季年会,* 2017年3月.
643. **長岡 玄, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 発光特性を有するトリアリールアミン誘導体の合成とストロンチウム試剤を用いた置換基効果の検討, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 2PC-060,* 2017年3月.
644. **上野 雅晴, 喜多 亜希子, 室井 超帰, 大村 D. 聡, 和田 眞, 三好 德和 :** ビスマス塩を触媒として用いた簡便かつ多様な酸化反応の開発, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 2E5-38,* 2017年3月.
645. **日下 亮, 大村 D. 聡, 三好 德和, 上野 雅晴 :** ビスマス塩を環境調和型触媒として用いるRitter反応の開発, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 2E5-39,* 2017年3月.
646. **大熊 一正, 上嶋 明, 日置 善郎 :** 有効ラグランジアンに基づくtbW結合の拡張可能性, *日本物理学会年次大会,* 2017年3月.
647. **鷹尾 忍, 関澤 央輝, Samjeske Gabor, 金子 拓真, 東 晃太朗, 東 晃太朗, 趙 暁, 吉田 祐介, 坂田 智裕, 山本 孝, 宇留賀 朋哉, 岩澤 康裕 :** ナノXAFS-TEM/STEM-EDS同視野測定法による燃料電池MEA内の合金カソード触媒層の劣化分布・因子の解明に関する研究, *日本化学会第97春季年会,* 2017年3月.
648. **村中 貴恵, 小山 晋之, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 萩原 亮 :** サイトレイト・パイロリシス法を用いて作成された超微細セラミックス高温超伝導体の酸化・還元効果, *日本物理学会講演概要集,* **Vol.72,** *No.1,* 2017年3月.
649. **伊東 拓, 六人部 陽介, 萩原 亮, 島 龍夫, 小山 晋之 :** サイトレイト・パイロリシス前駆体の焙焼処理によってPr247単一相を生成させる手法, *日本物理学会講演概要集,* **Vol.72,** *No.1,* 2121, 2017年3月.
650. **折口 直輝, 竹内 正俊, 藤原 直樹, 久田 旭彦, 上床 美也 :** 一軸圧用加圧セル装置の開発, *日本物理学会講演概要集,* **Vol.72,** *No.1,* 2126, 2017年3月.
651. **西澤 高宏, 北方 咲, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるメタラサイクル中間体の調製とそれを用いる新たな反応の開発, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 4D4-01,* 2017年3月.
652. **下田 亮介, 阿比留 保奈美, 小松 翔, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウムアルコキシドを用いる新規合成手法の開発, *日本化学会第97春季年会, 慶應義塾大学日吉キャンパス(神奈川),講演番号 4D4-02,* 2017年3月.
653. **安藝 将也, 川崎 祐, 岸本 豊, 中村 浩一, 小山 岳秀, 水戸 毅, 八島 光晴, 椋田 秀和, 小手川 恒, 菅原 仁 :** A15型超伝導体V3SiのSi-NMRによる研究, *日本物理学会第72回年次大会 講演概要集,* 2017年3月.
654. **伏見 賢一 :** NaI(Tl)による宇宙暗黒物質探索(PICO-LON)報告, *日本物理学会第72回年次大会(大阪大学待兼山キャンパス),* 2017年3月.
655. **竹内 智史, 犬飼 宗弘, 川崎 祐, 森賀 俊広, 中村 浩一 :** NaxCoO2の電気伝導と局所構造変化, *日本物理学会第72回年次大会 講演概要集,* 2017年3月.
656. **犬飼 宗弘 :** 配位高分子のダイナミクスとプロトン伝導能, *第1 回有機・バイオイオニクス研究会・第67 回固体イオニクス研究会,* 2017年3月.
657. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus ndk gene for nucleoside diphosphate kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128785, May 2016.
658. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus uk gene for uridine kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128786, May 2016.
659. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus cmpk gene for cytidylate kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128787, May 2016.
660. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus thyA gene for thymidylate synthase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128788, May 2016.
661. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus tdk gene for thymidine kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128789, May 2016.
662. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus tmk gene for thymidylate kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128790, May 2016.
663. **Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Shota Inoue, Yurie Ohiwa *and* Takanori Satoh :** Anoxybacillus flavithermus umpk gene for uridylate kinase, complete cds, strain: NBRC 15317, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Anoxybacillus flavithermus ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC128791, May 2016.
664. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Geobacillus sp. S5-1-11 gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129500, May 2016.
665. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Paenibacillus sp. S5-1Fe gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129501, May 2016.
666. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Streptomyces sp.S10G5B gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129502, May 2016.
667. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Streptomyces sp.S96-001 gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129503, May 2016.
668. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Rahnella sp. S15-1 gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129504, May 2016.
669. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Streptomyces sp.S10G5R gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129505, May 2016.
670. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Streptomyces sp.S10Cu gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129506, May 2016.
671. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Staphylococcus sp. S5-1-2Y gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129507, May 2016.
672. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Staphylococcus sp. S5-1-2WY gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129508, May 2016.
673. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Staphylococcus sp. S5-1-2W gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129509, May 2016.
674. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Streptomyces sp.S3Cu gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129510, May 2016.
675. **Yui Hamaji, Yurie Ohiwa, Natsumi Yoshioka, Mayu Kanamoto, Shota Inoue, Ken-ichi Nishiyama *and* Takanori Satoh :** Paenibacillus sp. S5-1-1 gene for 16S ribosomal RNA, partial sequence, --- Exploration for Cu,Fe-tolerant bacteria from Myoho copper mine spoils in Wakayama prefecture ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC129511, May 2016.
676. **佐藤 高則 :** まなびの森講演会 「藍染めの世界へようこそ!」, *放送大学徳島学習センター公開講演会「まなびの森講演会」,* 2016年5月.
677. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2016年10月.
678. **佐藤 高則 :** FD企画「教育の質保証に向けたTAの活用」, --- 実験・実習授業におけるTAの活用 ---, *大学院総合科学教育部FD企画「教育の質保証に向けたTAの活用」,* 2016年10月.
679. **山本 孝 :** XAFS, *近畿化学協会触媒・表面部会 ナノ材料の表面分析講習,* 2016年11月.
680. **秋吉 研二, Ahmed Ibrahim Elhossany Elmarhomy, 武岡 大輔, 時久 栞, 山本 真由美, 中山 信太郎, 佐藤 高則 :** 社会的弱者，特に外国人と女性の視点から学ぶ減災, *平成28年度総合科学教育部プロジェクト研究I 発表会,* 2017年2月.
681. **川崎 祐 :** Aサイト秩序型マンガン酸化物における電子相分離, *核磁気共鳴を主とした物性研究の新展開,* 2017年3月.
682. **秋吉 研二, Ahmed Ibrahim Elhossany Elmarhomy, 武岡 大輔, 時久 栞, 山本 真由美, 中山 信太郎, 佐藤 高則 :** 平成28年度プロジェクト研究I(佐藤グループ)報告書, --- 社会的弱者，特に外国人と女性の視点から学ぶ減災 ---, *平成28年度総合科学教育部プロジェクト研究I 報告書,* 1-11, 徳島, 2017年1月.
683. **西山 賢一 :** 越高遺跡B地点 地形・地質調査結果, *熊本大学 考古学研究室報告,* **Vol.52,** 36-40, 2017年3月.
684. **佐藤 高則 :** 平成28年度 教養教育授業方法等改善経費使用実績報告書, --- イノベーション科目「つたえることとものづくり∼藍染めの科学∼」における授業内容充実のための経費申請 ---, *平成28年度 教養教育授業方法等改善経費使用実績報告書,* 1-2, 徳島, 2017年3月.
685. **山本 真由美, 中山 信太郎, 佐藤 高則, 荒木 秀夫, 小野 覚久 :** 東北大震災から5年，福島は今 ―福島から学ぶ― シンポジウム, *平成28年度 総合科学部部局長裁量経費 「地域交流プロジェクト」成果報告書,* 徳島, 2017年3月.
686. **野田 篤, 植木 岳雪, 川畑 博, 松浦 浩久, 青矢 睦月 :** 観音寺地域の地質(5 万分の1 地質図幅), 産総研地質調査総合センター, つくば, 2017年7月.
687. **山本 孝 :** 硬XANES 「XAFSの基礎と応用 第3章4節2項」, 株式会社 講談社, 2017年7月.
688. **西山 賢一, 日本地質学会編 :** はじめての地質学, ペレ出版, 2017年9月.
689. **齊藤 隆仁, 小山 晋之 :** 基礎物理学実験テキスト (2017年度後期改訂版), 徳島大学生協, 2017年10月.
690. **中村 浩一 :** 次世代電池用電極材料の高エネルギー密度，高出力化, --- 第2章 第5節 リチウム過剰系LixV2O5とLi2MnO3における局所構造とリチウムイオン拡散挙動 ---, 株式会社 技術情報協会, 東京, 2017年11月.
691. **Toshiyuki Moriuchi, Satoshi D. Ohmura *and* Toshikazu Hirao :** Chapter-4: Chirality Organization of Peptide and pi-Conjugated Polyanilines in "Asymmetric Synthesis of Drugs and Natural Products", Taylor & Francis, Feb. 2018.
692. **F. Acero, R. Aloisio, J. Amans, Reiko Orito *and* et al (380 coauthors) :** Prospects for Cherenkov Telescope Array Observations of the Young Supernova Remnant RX J1713.7-3946, *The Astrophysical Journal,* **Vol.840,** *No.2,* id.74, 2017.
693. **Yue Wang, Takunori Harada, Yoshihito Shiota, Kazunari Yoshizawa, Heng Wang, Sheng Wang, Xichong Ye, Masamichi Ogasawara *and* Tamaki Nakano :** Isolation and Photo Transformation of Enatiomerically Pure Iridium(III) Bis[(4,6-difuorophenyl)pyridinato-N,C2]picolinate, *RSC Advances,* **Vol.7,** 29550-29553, 2017.
694. **Satoshi D. Ohmura, Yasuaki Miyazaki, Daisuke Kanehiro, Yuka Yamaguchi, Saki Kitakata, Sawa Tateda, Takahiro Nishizawa, Ryosuke Shimoda, Gen Nagaoka, Masaharu Ueno *and* Norikazu Miyoshi :** Superior Alkylating Agents for Bulky Esters/Ketones via Strontium-Mediated Barbier-Type Reaction, *Asian Journal of Organic Chemistry,* **Vol.6,** *No.7,* 821-824, 2017.
695. **Hiroaki Ichio, Hidetoshi Murakami, Yen-Chou Chen, Tamotsu Takahashi *and* Masamichi Ogasawara :** Palladium-Catalyzed SN2'-Cyclization of Ambivalent (Bromoalkadienyl)malonates: Preparation of Medium- to Large-Membered Endocyclic Allenes, *The Journal of Organic Chemistry,* **Vol.82,** 7503-7511, 2017.
696. **Masamichi Ogasawara, Hirotaka Sasa, Hao Hu, Yuta Amano, Hikaru Nakajima, Naoko Takenaga, Kiyohiko Nakajima, Yasuyuki Kita, Tamotsu Takahashi *and* Toshifumi Dohi :** Atropisomeric Chiral Diiododienes (Z,Z)-2,3-Di(1-iodoalkylidene)tetralins: Synthesis, Enantiomeric Resolution, and Application in Asymmetric Catalysis, *Organic Letters,* **Vol.19,** 4102-4105, 2017.
697. **Akihiko Hisada, Kie Muranaka, Kuniyuki Koyama, Ko-ichi Magishi, Takahito Saito *and* Makoto Haguwara :** Oxidation and reduction effects of successive superconducting transitions in ultra-fineYBa2Cu4O8 ceramics, *Journal of Physics: Conference Series,* **Vol.871,** 012010-1-012010-4, 2017.
698. **Hiroyuki Deguti, R. Warabino, S. Ka, M. Mito, Haguwara Makoto *and* Kuniyuki Koyama :** Field-Temperature Phase Diagram of Intergrain Ordering in Superconducting Ceramic YBCO, *Journal of Physics: Conference Series,* **Vol.871,** 012011-1-012011-5, 2017.
699. **Keisuke Oyama, Norikazu Miyoshi *and* Yasuo Oyama :** Cytometric analysis on cytotoxicity of 4,4 -methylenediphenyl diisocyanate, a chemical allergen, in rat thymocytes, *Fundamental Toxicological Sciences,* **Vol.4,** *No.4,* 173-178, 2017.
700. **H. Abe, H. Fuke, Reiko Orito *and* 20 coauthors :** The results from BESS-Polar experiment, *Advances in Space Research,* **Vol.60,** *No.4,* 806-814, 2017.
701. **村田 明広, 高橋 司 :** 黒瀬川構造帯の模式地と四国西部の秩父帯を巡る:四国西予ジオパーク, *地質学雑誌,* **Vol.123,** *No.8,* 585-597, 2017年.
702. **西山 賢一, 横田 修一郎, 長谷川 修一, 菅原 大助 :** 火山角礫岩からなる急崖に発達するノッチ・タフォニの形成プロセスと岩盤崩落との関係:香川県高松市五剣山の例．, *地学雑誌,* **Vol.126,** *No.4,* 425-437, 2017年.
703. **山本 祐平, 田上 梓, 白崎 俊浩, 米谷 明, 山本 孝, 今井 昭二 :** 鉄マトリクス修飾剤を用いた黒鉛炉原子吸光法によるホウ素の原子化機構, *分析化学,* **Vol.66,** *No.9,* 629-637, 2017年.
704. **Masatsugu Oishi, Shohei Shiomi, Takashi Yamamoto, Tomoyuki Ueki, Yoichiro Kai, Shigefusa F. Chichibu, Aiko Takatori *and* Kazunobu Kojima :** High temperature degradation mechanism of a red phosphor, CaAlSiN3:Eu for solid-state lighting, *Journal of Applied Physics,* **Vol.122,** *No.11,* 113104-113111, 2017.
705. **青矢 睦月, 遠藤 俊祐 :** 初期三波川変成作用の認識，及び後期白亜紀三波川沈み込み帯の描像, *地質学雑誌,* **Vol.123,** *No.9,* 677-698, 2017年.
706. **Masamichi Ogasawara, Ya-Yi Tseng, Mizuho Uryu, Naoki Ohya, Ninghui Chang, Hiroto Ishimoto, Sachie Arae, Tamotsu Takahashi *and* Ken Kamikawa :** Molybdenum-Catalyzed Enantioselective Synthesis of Planar-Chiral (5-Phosphacyclopentadienyl)manganese(I) Complexes and Application in Asymmetric Catalysis, *Organometallics,* **Vol.36,** *No.20,* 4061-4069, 2017.
707. **井澤 健一, 谷口 智紀 :** 量子論におけるレトロディクション I, *徳島大学総合科学部自然科学研究,* **Vol.30,** 1-4, 2017年.
708. **N. Oka, K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, T. Norita, H. Ogawa, K. Sato, H. Sekiya, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, R. Kegasa, K. Masuda, H. Takiya, Ken-Ichi Fushimi, G. Kanzaki, K. Martens, Y. Suzuki, B.D. Xu, R. Fujita, K. Hosokawa, K. Miuchi, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fukuda, M. Miyasaka, K. Nishijima *and* S. Nakamura :** Search for solar KaluzaKlein axions by annual modulation with the XMASS-I detector, *Progress of Theoretical and Experimental Physics,* 103C01, 2017.
709. **K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, T. Norita, H. Ogawa, K. Sato, H. Sekiya, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, R. Kegasa, K. Masuda, H. Takiya, Ken-Ichi Fushimi, G. Kanzaki, K. Martens, Y. Suzuki, B.D. Xu, R. Fujita, K. Hosokawa, K. Miuchi, N. Oka, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fukuda, M. Miyasaka, K. Nishijima *and* S. Nakamura :** Identification of 210Pb and 210Po in the bulk od copper samples with a low-background alpha particle counter, *Nuclear Instruments and Methods in Physics Research Section A: Accelerators, Spectrometers, Detectors and Associated Equipment,* **Vol.884,** 157-161, 2017.
710. **W.M. Chan, T. Kishimoto, S. Umehara, K. Matsuoka, K. Suzuki, S. Yoshida, K. Nakajima, T. Iida, Ken-Ichi Fushimi, M. Nomachi, I. Ogawa, Y. Tamagawa, Ryuta Hazama, Y. Takemoto, N. Nakatani, Y. Takihira, M. Tozawa, H. Kakubata, V.T.T. Trang, T. Ohata, K. Tetsuno, T. Maeda, B.T. Khai, X. Li *and* T. Batprev :** Development of CANDLES low background HPGe detector and half-life measurement of 180Tam, *AIP Conference Proceedings,* **Vol.1921,** 030004, 2018.
711. **今井 昭二, 山本 祐平, 清水 魁人, 兼清 恵理, 西本 潤, 菊地 洋一 :** 本州日本海沿いの遠隔地における北東中国北部・ロシア沿海地方からの大気塊に由来する降雪中のPb/Cd比および冬季の長距離輸送機構, *分析化学,* **Vol.67,** *No.2,* 95-101, 2018年.
712. **K. Nakajima, T. Iida, K. Akutagawa, T. Batprev, W.M. Chan, F. Dotaku, Ken-Ichi Fushimi, H. Kakubata, K. Kanagawa, S. Katagiri, K. Kawasaki, B.T. Khai, H. Kino, E. Kinoshita, T. Kishimoto, R. Hazama, H. Hiraoka, T. Hiyama, M. Ishikawa, X. Li, T. Maeda, K. Matsuoka, M. Moser, M. Nomachi, I. Ogawa, T. Ohata, H. Sato, K. Shamoto, M. Shimada, M. Shokati, N. Takahashi, Y. Takemoto, Y. Takihira, Y. Tamagawa, M. Tozawa, K. Teranishi, K. Tetsuno, V.T.T. Trang, M. Tsuzuki, S. Umehara, W. Wang, S. Yoshida *and* N. Yotsunaga :** Background studies of high energy γrays from ( n , γ) reactions in the CANDLES experiment, *Astroparticle Physics,* **Vol.100,** 54-60, 2018.
713. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Microbial diversity on the leaves of Japanese and Taiwan teas, *Proceedings of 9th Asian Symposium on Microbial Ecology(ASME2017),* 204, 2017.
714. **Kouta Nagai, Yousuke Endoh, Yurie Ohiwa, Yuto Oe, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Comparison of Thermostabilities among Nucleoside Diphosphate Kinases from Geobacillus and Anoxybacillus bacteria, *Proceedings of 5th Asia Pacific Protein Association Conference (APPA2017),* 185, 2017.
715. **Ken-Ichi Fushimi, D. Chenyak, Hiroyasu Ejiri, R. Hazama, Shoko Hirata, H. Ikeda, K. Inoue, K. Imagawa, G. Kanzaki, A. Kozlov, Reiko Orito, T. Shima, Y. Takemoto, Y. Teraoka, S. Umehara *and* S. Yoshida :** Search for cosmic dark matter by means of ultra high purity NaI(Tl) scintillator, *Proceedings of the 31 st Workshop on Radiation Detectors and Their Uses,* 42-50, 2017.
716. **佐藤 高則, Gantulga Uuganbaatar, Erdenebat Erdenedalai, 岡 里美, 谷口 智紀, 平田 晶子, 藤巻 晃, 山本 真由美, 中山 信太郎 :** 東日本大震災被災地支援と連携したPBL型大学院教育, *平成29年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 46-47, 2018年.
717. **佐藤 高則 :** 教養教育におけるシチュエーション・ラーニング(Situated learning)の実践, *平成29年度全学FD 徳島大学教育カンファレンス in 徳島 発表抄録集,* 26-27, 2018年.
718. **小笠原 正道 :** モリブデン触媒不斉オレフィン・メタセシスによる面不斉遷移金属錯体の触媒的不斉合成, *Organometallic News,* 52-57, 2017年4月.
719. **山本 孝 :** 低出力X線管を利用したX線吸収分光, *ぶんせき,* 150-151, 2017年4月.
720. **青矢 睦月, 水上 知行, 遠藤 俊祐 :** 巡検案内書 三波川帯トラバース:最浅部変成岩から最深部超苦鉄質岩まで, *地質学雑誌,* **Vol.123,** *No.7,* 491-514, 2017年7月.
721. **小笠原 正道 :** 均一系オレフィン・メタセシス反応による面不斉遷移金属錯体のエナンチオ選択的合成法の開発と不斉触媒反応への応用, *Bulletin of Japan Society of Coordination Chemistry,* **Vol.70,** 14-21, 2017年12月.
722. **山本 孝 :** X線を利用したタングステンジルコニウム複合酸化物系固体強酸触媒の構造解析, *触媒,* **Vol.60,** *No.1,* 8-13, 2018年1月.
723. **石田 啓祐 :** 新たな国指定天然記念物及び名勝「大歩危小歩危」, *日本地質学会NEWS, No.21,* 2-3, 2018年1月.
724. **Ken-Ichi Fushimi :** Low Background Measurement by Means of NaI(Tl) Scintillator ~ Improvement of Sensitivity for Cosmic Dark Matter ~, *Radioisotopes,* **Vol.67,** *No.3,* 101-111, Mar. 2018.
725. **今井 昭二, 上村 健, 児玉 憲治, 山本 祐平 :** WDXRF による樹氷と雪の中の非水溶性イオウ化合物の化学形態別分析と東アジアの石炭燃焼排出物の冬期モンスーン下での長距離輸送機構, *X線分析の進歩,* **Vol.49,** 125-142, 2018年3月.
726. **Masamichi Ogasawara :** Design and Synthesis of Phosphine-Olefin Ligands with Planar-Chiral Metal Complex Scaffold and Application in Asymmetric Catalysis, *The 253rd ACS National Meeting,* San Francisco, Apr. 2017.
727. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Microbial diversity on the leaves of Japanese and Taiwan teas, *The 9th Asian Symposium on Microbial Ecology(ASME2017),* 204, Busan, Apr. 2017.
728. **Mutsuki Aoya, Shunsuke Endo, Tomoyuki Mizukami *and* Simon Wallis :** The Early-Cretaceous Sambagawa metamorphism and a cross-sectional view of the Late-Cretaceous Sambagawa subduction zone, *JpGU-AGU Joint Meeting 2017, No.SGL33-08,* May 2017.
729. **Masamichi Ogasawara :** Mo-Catalyzed Asymmetric Synthesis of P-Stereogenic Phosphine Derivatives by "Formal" Cross-Metathesis on Arene-Cr Complexes, *International Symposium on Pure & Applied Chemistry 2017 (ISPAC 2017),* Ho Chi Minh, Jun. 2017.
730. **Kouta Nagai, Yousuke Endoh, Yurie Ohiwa, Yuto Oe, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Comparison of Thermostabilities among Nucleoside Diphosphate Kinases from Geobacillus and Anoxybacillus bacteria, *The 5th Asia Pacific Protein Association Conference (APPA2017),* 185, Bangsaen,Thailand, Jul. 2017.
731. **Masamichi Ogasawara :** Enantioselective Synthesis of P-Stereogenic Phosphine Derivatives by Molybdenum-Catalyzed "Formal" Cross-Metathesis: Use of Chromium-Template for High Enantio-, Chemo-, and E/Z-Selectivities, *The 22nd International Symposium on Olefin Metathesis and Related Chemistry (ISOM 22),* Zuerich, Jul. 2017.
732. **Takashi Yamamoto :** Relation between XANES Pre-edge and XRF Kb5 as the Changes of Chemical State, Local Symmetry and Coordination Number, *The 66th annual Denver X-ray Conference,* Big Sky, USA, Aug. 2017.
733. **Ko-ichi Magishi, Ryo Watanabe, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama, S. Okada, Y. Kamihara, N. Ohkubo, S. Ban *and* M. Matoba :** NMR study of the layered cobalt oxyphosphide Sr2Sc(Co1-xFex)PO3, 28th International Conference on Low Temperature Physics (LT28), *LT28,* Gothenburg, Aug. 2017.
734. **Mitsuru OKUNO, Masayuki TORII, Ken-ichi Nishiyama *and* Shuichiro YOKOTA :** Failure Process of the Western Slope of Takano-obane Volcano in Aso Caldera Triggered by the 2016 Kumamoto Earthquake, SW Japan, Based on Geomorphological Classification, *The 3rd ASQUA Conference 2017,* Jeju Island, Korea, Oct. 2017.
735. **Takanori Satoh, Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Bacterial diversity on the Japanese and Taiwanese tea leaves, *2017 International Seminar on Sustainable Global Society,* 10, Tokushima, Oct. 2017.
736. **Takashi Yamamoto *and* Kazunori Miyamoto :** Decomposition Behavior of Platinum Precursor on Silica and Alumina Studied by in-situ XAFS, *Int. Symp. Novel Energy Nanomater., Catal. Surf. Future Earth,* Chofu, Oct. 2017.
737. **Shinobu Takao, Oki Sekizawa, Takashi Yamamoto, Gabor Samjeské, Takuma Kaneko, Kotaro Higashi, Yuske Yoshida, Xiao Zhao, Tomohiro Sakata, Tomoya Uruga *and* Yasuhiro Iwasawa :** In situ Visualization of Platinum Chemical species in a Nano Fuel-cell, *Int. Symp. Novel Energy Nanomater., Catal. Surf. Future Earth,* Chofu, Oct. 2017.
738. **Masamichi Ogasawara :** Atropisomeric Chiral Dienes in Asymmetric Catalysis, *2017 International Symposium on Chemistry Frontiers,* Shanghai, Dec. 2017.
739. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Applications in Asymmetric Catalysis, *SIOC Special Seminar,* Shanghai, Dec. 2017.
740. **Masamichi Ogasawara :** Pd-Catalyzed Asymmetric Synthesis of Axially Chiral Allenes and Their Application in Organic Transformations, *Fudan University Special Seminar,* Shanghai, Dec. 2017.
741. **Masamichi Ogasawara :** Synthesis and Catalytic Applications of Novel Atropisomeric Chiral Dienes, *Pure and Applied Chemistry International Conference 2018 (PACCON 2018),* Hat Yai (Thailand), Feb. 2018.
742. **Masamichi Ogasawara :** Atropisomeric Chiral Dienes in Asymmetric Catalysis, *International Congress on Pure & Applied Chemistry 2018 (ICPAC 2018),* Siem Reap, Mar. 2018.
743. **Takashi Yamamoto, Kazunori Miyamoto *and* Masaya Kondo :** Thermal Decomposition Behavior of Supported Platinum Salt for Catalyst Preparation, *International Congress on Pure & Applied Chemistry (ICPAC) 2018,* Siem Reap, Cambodia, Mar. 2018.
744. **奧野 充, 鳥井 真之, 西山 賢一, 横田 修一郎 :** 詳細地形区分から見た阿蘇カルデラ・高野尾羽根火山西側斜面の崩壊プロセス, *平成28年度自然災害総合研究班西部地区部会・研究発表会講演論文集,* 2017年4月.
745. **西山 賢一, 鳥井 真之, 奥野 充 :** 2016年熊本地震で流下・堆積した崩壊堆積物, *日本地球惑星科学連合2017年大会講演要旨集,* 2017年5月.
746. **今井 昭二, 山本 祐平 :** 黒鉛炉原子吸光法における吸光度プロファイルの減衰過程-1:標準消失過程-, *第77回分析化学討論会,* **Vol.G2007,** 2017年5月.
747. **山本 祐平, 喜多 郁弥, 磯野 成美, 今井 昭二 :** 腐植物質と金属イオンの錯生成における競合の影響, *第77回分析化学討論会,* 2017年5月.
748. **奧野 充, 鳥井 真之, 西山 賢一, 横田 修一郎 :** 阿蘇・高野尾羽根火山西斜面の崩壊過程, *日本地質学会西日本支部第168回例会,* 2017年6月.
749. **山本 祐平, 清水 魁人, 今井 昭二 :** 遠隔山岳地域で採取されたエアロゾルに対する人口密集地域からの寄与, *第26回環境化学討論会,* 2017年6月.
750. **Narumi Maegawa, Natsumi Yoshioka, Mayu Kanamoto, Kouta Nagai, Daiki Fujimoto, Yui Hamaji, Kenji Akiyoshi-Hiraoka, Shota Inoue, Yurie Ohiwa, Ichinkhorloo Zesem, Munkhtsetseg Janlav *and* Takanori Satoh :** Binding properties for Lipocalin function of whey proteins in Mongolian animal milks, *The 17th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,,* 65, Jun. 2017.
751. **中村 浩一, 井藤 弘章, 犬飼 宗弘, 森賀 俊広, 岩井 良樹, 桑田 直明, 河村 純一 :** LiMPO4(M=Fe,Mn)におけるLi+イオン運動と局所構造の変化, *第20回超イオン導電体物性研究会講演要旨集,* 71-74, 2017年7月.
752. **山本 孝 :** XANESの見かけ上の吸収端シフトおよび価数評価, *関西分析研究会平成29年度第一回例会,* 2017年7月.
753. **小笠原 正道 :** 不斉遷移金属錯体の触媒的不斉合成と有機合成への応用, *関西大学 化学生命工学部 特別講演会,* 2017年7月.
754. **西山 賢一, 鳥井 真之, 横田 修一郎, 若月 強, 井上 弦, 中尾 賢一, 星出 和裕 :** 阿蘇カルデラ斜面における斜面崩壊・土石流の発生頻度の推定, *日本第四紀学会2017年大会講演要旨集,* **Vol.47,** 129-130, 2017年8月.
755. **鳥井 真之, 長谷中 利昭, 北園 芳人, 西山 賢一, 横田 修一郎, 奥野 充 :** 2016年熊本地震における災害遺産の第四紀地質学, *日本第四紀学会2017年大会講演要旨集,* **Vol.47,** 124-125, 2017年8月.
756. **西山 賢一, 柳田 裕三, 辻本 裕也 :** 長崎県佐世保市に位置する史跡・福井洞窟の形成過程, *日本第四紀学会2017年大会講演要旨集,* **Vol.47,** 24, 2017年8月.
757. **佐藤 高則, 中山 信太郎, 山本 真由美 :** 東日本大震災被災地支援と連携したAL型大学院教育, *SPODフォーラム2017 プログラム,* P-23, 2017年8月.
758. **Mitsuki Fujii, Kouta Nagai, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Shota Inoue, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method, *2017 Joint Conference of the Societies for Environmental Microbiology (JCSEM2017) program,* 56, Aug. 2017.
759. **西山 賢一, 丹野 祥一, 岡林 眞姫, 山上 陽平, 中尾 賢一, 川村 教一 :** ボーリング資料に基づく徳島平野の地下地質, *日本地質学会第124年学術大会講演要旨集,* 125, 2017年9月.
760. **安藝 将也, 川崎 祐, 岸本 豊, 中村 浩一, 小山 岳秀, 水戸 毅, 八島 光晴, 椋田 秀和, 小手川 恒, 菅原 仁 :** A15型超伝導体V3SiのSi-NMRによる研究II, *日本物理学会秋季大会,* 2017年9月.
761. **小笠原 正道 :** 均一系遷移金属触媒による遷移金属錯体の不斉合成, *第33回若手化学者のための化学道場,* 2017年9月.
762. **胡 淏, 一柳 浩輝, 関 直樹, 中島 清彦, 小笠原 正道 :** ジフェロセニルホスフィノ基を有する新規不斉ホスフィン配位子の合成と応用, *第64回有機金属化学討論会,* 2017年9月.
763. **陳 彦洲, 小笠原 正道 :** 3-ブロモ-2,4-ペンタジエニル・エステル類へのパラジウム触媒二重求核置換反応によるアレン合成, *第64回有機金属化学討論会,* 2017年9月.
764. **小笠原 正道, 劉 強, 和田 志郎, 木村 亮介, 安江 里紗, 吉田 和弘 :** メタロセン縮環面不斉4-ピリドンおよび4-ジアルキルアミノピリジン類のエナンチオ選択的合成法の開発と不斉反応への応用, *第64回有機金属化学討論会,* 2017年9月.
765. **大村 D. 聡, 長岡 玄, 山名 啓介, 上野 雅晴, 三好 德和 :** アルキル置換基導入に基づくトリアリールアミン誘導体の発光特性制御, *第28回基礎有機化学討論会, 九州大学伊都キャンパス(福岡), 講演番号 2P085,* 2017年9月.
766. **下田 亮介, 宮崎 恭彰, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウムアルコキシドを用いる新規合成手法の開発, *第28回基礎有機化学討論会, 九州大学伊都キャンパス(福岡), 講演番号 2P107,* 2017年9月.
767. **今井 昭二, 山本 祐平, 宮本 征弥, 峯崎 正樹, 小堺 久美子 :** 黒鉛炉原子吸光法における吸光度プロファイルの減衰過程-2:Tmax>1000K金属・酸化物原子化中間体-, *日本分析化学会第66年会,* 2017年9月.
768. **今井 昭二, 清水 魁人, 兼清 恵理, 山本 祐平, 西本 潤, 菊地 洋一 :** 本州の日本海沿いの山岳における降雪中のPb/Cd濃度比と石炭起源の影響, *日本分析化学会第66年会,* 2017年9月.
769. **西澤 高宏, 北方 咲, 立田 彩和, 大村 D. 聡, 上野 雅晴, 三好 德和 :** Preparation of Metallacycle Intermediates Using Metallic Strontium and Its Application for New Reaction, *第64回有機金属化学討論会, 東北大学川内キャンパス(宮城), 講演番号 P3-01,* 2017年9月.
770. **木村 将大, 久保 誠輝, 大村 D. 聡, 上野 雅晴, 三好 德和 :** Metallic Strontium-Mediated Monoalkylation of N,N-Dimethylamides, *第64回有機金属化学討論会, 東北大学川内キャンパス(宮城), 講演番号P3-02,* 2017年9月.
771. **今井 昭二, 山本 祐平, 山本 孝, 児玉 憲治, 西本 潤, 菊地 洋一 :** 蛍光X線分析法を用いた四国の樹氷と日本海沿いの遠隔地における降雪中の不溶性イオウのスペシエーション, *日本分析化学会第66年会,* 2017年9月.
772. **山本 祐平, 山本 孝, 児玉 憲治, 早川 慎二郎, 今井 昭二 :** 汎用型蛍光X線分析装置を用いた大気エアロゾル中の微量硫黄の化学形態分析, *日本地球化学会第64回年会,* 2017年9月.
773. **山本 孝, 寺町 葵, 原 領汰 :** 金属イオン添加ジルコニア固溶体担持酸化タングステン触媒の固体強酸特性とタングステン種との相関, *第120回触媒討論会,* 2017年9月.
774. **Sato Takehiro, Kamikawa Ken, 小笠原 正道, Mori Seiji :** Theoretical Investigation of Enantioselective 1,4-Addition using Rhodium Catalysts with Planar-Chiral Phosphine-Olefin Ligands, *錯体化学会第67回討論会,* 2017年9月.
775. **上野 雅晴 :** ビスマス塩を用いた環境にやさしい分子変換反応の開発, *第17回社会産業理工学研究交流会2017,* 2017年9月.
776. **梅岡 優, 仡川 昂平, 胡 魁, 水田 悠介, 中村 浩一, 村井 啓一郎, 森賀 俊広 :** Li-Ti系酸窒化物の合成および充放電特性の評価, *日本セラミックス協会第30回秋季シンポジウム,* 2017年9月.
777. **中村 浩一, 竹内 智史, 犬飼 宗弘, 川崎 祐, 森賀 俊広 :** Na系遷移金属酸化物における局所構造とNa+イオンの運動状態, *日本物理学会2017年秋季大会 講演概要集,* 2017年9月.
778. **小笠原 正道 :** 均一系遷移金属触媒による立体選択的分子変換反応, *第11回触媒道場,* 2017年9月.
779. **西山 賢一, 柳田 裕三, 辻本 裕也 :** 長崎県佐世保市に位置する史跡・福井洞窟遺跡の地形・地質的特徴, *日本応用地質学会平成29年度研究発表会講演論文集,* 165-166, 2017年10月.
780. **西山 賢一 :** 徳島市の地盤, *地盤工学会・表層地盤の特性抽出と解析手法についてのシンポジウム資料,* 2017年10月.
781. **犬飼 宗弘 :** 配位高分子の固体NMR, *第62回固体NMR・材料フォーラム,* 2017年10月.
782. **今井 昭二, 上村 健, 児玉 憲治, 山本 祐平 :** WDXRF による樹氷と雪の中の非水溶性イオウ化合物の化学形態別分析と東アジアの石炭燃焼排出物の冬期モンスーン下での長距離輸送機構, *第53回X線分析討論会,* 2017年10月.
783. **山本 祐平, 山本 孝, 早川 慎二郎, 今井 昭二 :** X線吸収分光法を用いた大気エアロゾル中の非水溶性硫黄の化学形態分析, *第53回X線分析討論会,* 2017年10月.
784. **谷 和紀, 大村 D. 聡, 三好 德和, 上野 雅晴 :** 薗頭-鈴木タンデムカップリング反応による水溶媒中でのワンポット多成分連結法の開発, *2017年日本化学会中国四国支部大会,鳥取産業体育館(鳥取),講演番号 OR44P,* 2017年11月.
785. **藤田 真美, 宮崎 泰彰, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いたアリルシランの新規合成法, *2017年日本化学会中国四国支部大会,鳥取産業体育館(鳥取),講演番号 OR54P,* 2017年11月.
786. **中村 光裕, 中原 舞, 松田 和生 :** セルロースを用いたチオール化合物の精製法の開発, *2017年日本化学会中国四国支部大会 鳥取大会,* 2017年11月.
787. **西山 賢一, 丹野 祥一, 岡林 眞姫, 山上 陽平, 中尾 賢一, 川村 教一, 田村 俊之 :** ボーリング資料に基づく徳島平野の地下地質, *日本地質学会四国支部講演会資料,* 6, 2017年12月.
788. **佐藤 丈弘, 神川 憲, 小笠原 正道, 森 聖治 :** 面不斉ホスフィンーオレフィン配位子を用いたRh触媒不斉1,4-付加反応の理論的研究, *第27回日本化学会関東支部茨城地区研究交流会,* 2017年12月.
789. **竹内 智史, 中村 浩一, 犬飼 宗弘, 川崎 祐, 森賀 俊広, 桑田 直明, 河村 純一 :** NaxMO2(M=Co,Cr)の電気伝導と局所構造変化, *第43回固体イオニクス討論会,* 4-5, 2017年12月.
790. **犬飼 宗弘, 高木 翼, 中村 浩一 :** 構造欠陥を利用したプロトン伝導性配位高分子, *第43回固体イオニクス討論会,* 2017年12月.
791. **Haruka Iwata, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Hajime Nonoichi, Ai Hasegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Genetic analysis and thermostabilities of nucleoside diphosphate kinases from two moderately thermophiles, *ConBio2017 Abstracts,* Dec. 2017.
792. **Ai Hasegawa, Daiki Fujimoto, Kouta Nagai, Narumi Maegawa, Haruka Iwata, Hajime Nonoichi, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Genetic analysis and thermostabilities of cytidylate kinases from two moderately thermophiles, *ConBio2017 Abstracts,* Dec. 2017.
793. **Hajime Nonoichi, Shota Inoue, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Haruka Iwata, Ai Hasegawa, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Genetic analysis and expression of genes for uridine nucleotide synthesizing enzymes from two moderately thermophiles, *ConBio2017 Abstracts,* Dec. 2017.
794. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Phylogenetic analysis of bacteria from Awa Bancha and non-fermented tea leaves produced at Miyoshi City, Tokushima, *ConBio2017 Abstracts,* Dec. 2017.
795. **Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Mitsuki Fujii, Shota Inoue, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Microbial phylogenetic analysis from the leaves of fermented and semi-fermented teas produced at Tokushima, and Taiwan, *ConBio2017 Abstracts,* Dec. 2017.
796. **Masatsugu Oishi, Takuya Doi, Takashi Yamamoto, Takashi Nakamura, Koji Amezawa *and* Yuji Okuyama :** Evaluation of the electronic and local structure of proton-conducting oxide, (CaZr1-x,Mnx)O3-d, to elucidate a novel protonation mechanism, *43rd symposium on solid state ionics of Japan,* Dec. 2017.
797. **蕨野 龍世, 長渕 康之, 出口 博之, 美藤 正樹, 萩原 亮, 小山 晋之 :** YBCO超伝導セラミクスにおけるグレイン間相転移の磁場依存性, *日本物理学会 九州支部例会 講演概要集,* 21, 2017年12月.
798. **小笠原 正道 :** 均一系遷移金属触媒による遷移金属錯体の不斉合成法の開発と応用, *2017年日本化学会中国四国支部徳島地区化学講演会,* 2017年12月.
799. **佐藤 高則, Gantulga Uuganbaatar, Erdenebat Erdenedalai, 岡 里美, 谷口 智紀, 平田 晶子, 藤巻 晃, 山本 真由美, 中山 信太郎 :** 東日本大震災被災地支援と連携したPBL型大学院教育, *平成29年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 46-47, 2018年1月.
800. **佐藤 高則 :** 教養教育におけるシチュエーション・ラーニング(Situated learning)の実践, *平成29年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 26-27, 2018年1月.
801. **梅岡 優, 仡川 昂平, 胡 魁, 水田 悠介, 中村 浩一, 村井 啓一郎, 森賀 俊広 :** 岩塩型Li-Ti系酸窒化物の合成及び充放電過程における相変化の検討, *第56回セラミックス基礎科学討論会,* 2018年1月.
802. **鳥井 真之, 北園 芳人, 長谷中 利昭, 奥野 充, 遠田 晋次, 西山 賢一, 横田 修一郎 :** 南阿蘇村河陽地区における2016年熊本地震とテフラ層との関係, *国際火山噴火史情報研究集会講演要旨集2017-2,* 107-109, 2018年2月.
803. **伏見 賢一 :** PICO-LONの現状報告, *LBGT2018 (Workshop on Low Background Technique) (天童市),* 2018年3月.
804. **大石 昌嗣, 塩見 昌平, 山本 孝, 植木 智之, 改井 陽一郎, 秩父 重英, 高取 愛子, 小島 一信 :** 赤色蛍光体(CaAlSiN3:Eu)の高温劣化機構の評価, *第65回応用物理学会 春季学術講演会,東京 (2018.3.17-20).,* 2018年3月.
805. **Munehiro Inukai, 田村 優実, 堀毛 悟史, 樋口 雅一, 北川 進 *and* Koichi Nakamura :** Storage and dynamics of CO2 into porous coordination polymer controlled by ligand dynamics, *日本化学会第98春季年会,* Mar. 2018.
806. **日下 亮, 大村 D. 聡, 三好 德和, 上野 雅晴 :** 協奏機能型ビスマス触媒を用いた新規カップリング反応の研究, *日本化学会第98春季年会, 日本大学理工学部(千葉), 講演番号2H2-08,* 2018年3月.
807. **石井 悠佑, 立田 彩和, 西澤 高宏, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いる含有窒素化合物に対するアルキル化 反応とその応用, *日本化学会第98春季年会, 日本大学理工学部(千葉), 講演番号2PC-097,* 2018年3月.
808. **藤田 真美, 宮崎 泰彰, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いる Peterson 反応を利用したオレフィン 類の新しい合成手法の開発, *日本化学会第98春季年会, 日本大学理工学部(千葉), 講演番号2PC-098,* 2018年3月.
809. **谷 和紀, 林 俊輝, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 水中でのワンポット薗頭-鈴木タンデムカップリング反応の開発, *日本化学会第98春季年会, 日本大学理工学部(千葉), 講演番号2PC-122,* 2018年3月.
810. **中村 浩一, 竹内 智史, 犬飼 宗弘, 川崎 祐, 森賀 俊広, 桑田 直明, 河村 純一 :** Na系遷移金属酸化物における電気伝導のNa組成依存性, *日本物理学会第73回年次大会 講演概要集,* 2018年3月.
811. **陳 彦洲, Wang Yue, 中野 環, 小笠原 正道 :** Palladium-Catalyzed Synthesis of Allenic Oligomers/Polymers, *日本化学会第98春季年会,* 2018年3月.
812. **瓜生 瑞穂, 中島 清彦, 小笠原 正道 :** 面不斉フェロセンを基盤とするP-オレフィン配位子の開発と応用, *日本化学会第98春季年会,* 2018年3月.
813. **木村 将大, 久保 誠輝, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるアミド化合物へのアルキル化反応の 応用と発展, *日本化学会第98春季年会, 日本大学理工学部(千葉), 講演番号4H2-03,* 2018年3月.
814. **久田 旭彦, 真岸 孝一, 齊藤 隆仁, 小山 晋之, 出口 博之, 萩原 亮 :** 超微細セラミックスYBa2Cu4O8のグレイン間超伝導転移の酸化・還元効果, *日本物理学会講演概要集,* **Vol.73,** *No.1,* 2107, 2018年3月.
815. **Kenji Akiyoshi-Hiraoka, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Narumi Maegawa, Daiki Fujimoto, Kouta Nagai, Shota Inoue *and* Takanori Satoh :** Alicyclobacillus acidocaldarius subsp. acidocaldarius ndk gene for nucleoside diphosphate kinase, complete cds., --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Alicyclobacillus acidocaldarius subsp. acidocaldarius ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC218426, Apr. 2017.
816. **Kenji Akiyoshi-Hiraoka, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Narumi Maegawa, Daiki Fujimoto, Kouta Nagai, Shota Inoue *and* Takanori Satoh :** Alicyclobacillus acidocaldarius subsp. acidocaldarius umpk gene for uridylate kinase, complete cds., --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Alicyclobacillus acidocaldarius subsp. acidocaldarius ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC218427, Apr. 2017.
817. **Kenji Akiyoshi-Hiraoka, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Narumi Maegawa, Daiki Fujimoto, Kouta Nagai, Shota Inoue *and* Takanori Satoh :** Alicyclobacillus acidocaldarius subsp. acidocaldarius cmpk gene for cytidylate kinase, complete cds, --- Genetic analysis of genes for pyrimidine nucleotide synthesizing enzymes from Alicyclobacillus acidocaldarius subsp. acidocaldarius ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC218428, Apr. 2017.
818. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2017年10月.
819. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw13, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317317, Nov. 2017.
820. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw105, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317318, Nov. 2017.
821. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Burkholderia sp.gene for 16S ribosomal RNA, partial sequence, clone: Tw106, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317319, Nov. 2017.
822. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw111, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317320, Nov. 2017.
823. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw112, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317321, Nov. 2017.
824. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Finegoldia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw102, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317322, Nov. 2017.
825. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Paenibacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw11, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317323, Nov. 2017.
826. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Propionibacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw108, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317324, Nov. 2017.
827. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw12, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317325, Nov. 2017.
828. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw14, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317326, Nov. 2017.
829. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw103, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317327, Nov. 2017.
830. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw104, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317328, Nov. 2017.
831. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw107, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317329, Nov. 2017.
832. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Weissella sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw15, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317330, Nov. 2017.
833. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Weissella sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw110, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317331, Nov. 2017.
834. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Acidovorax sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw215, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317332, Nov. 2017.
835. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw27, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317333, Nov. 2017.
836. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw28, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317334, Nov. 2017.
837. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw216, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317335, Nov. 2017.
838. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw258, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317336, Nov. 2017.
839. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Devosia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw24, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317337, Nov. 2017.
840. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Devosia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw29, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317338, Nov. 2017.
841. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Devosia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw260, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317339, Nov. 2017.
842. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Methylobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw259, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317340, Nov. 2017.
843. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Variovorax sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw213, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317341, Nov. 2017.
844. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Agrobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3024, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317342, Nov. 2017.
845. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Agrobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3025, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317343, Nov. 2017.
846. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Aurantimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw319, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317344, Nov. 2017.
847. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Aurantimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw323, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317345, Nov. 2017.
848. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Methylobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3019, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317346, Nov. 2017.
849. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw321, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317347, Nov. 2017.
850. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw322, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317348, Nov. 2017.
851. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Microbacterium sp.gene for 16S ribosomal RNA, partial sequence, clone: Tw3023, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317349, Nov. 2017.
852. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pedobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw31, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317350, Nov. 2017.
853. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pedobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw35, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317351, Nov. 2017.
854. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pedobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw36, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317352, Nov. 2017.
855. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pedobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw320, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317353, Nov. 2017.
856. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pedobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3020, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317354, Nov. 2017.
857. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Rhizobium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3021, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317355, Nov. 2017.
858. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Rhizobium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw3022, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317356, Nov. 2017.
859. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Brevundimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw409, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317357, Nov. 2017.
860. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw46, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317358, Nov. 2017.
861. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw401, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317359, Nov. 2017.
862. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Edaphobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw424, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317360, Nov. 2017.
863. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Finegoldia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw428, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317361, Nov. 2017.
864. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Propionibacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw412, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317362, Nov. 2017.
865. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pseudonocardia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw47, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317363, Nov. 2017.
866. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Pseudonocardia sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw427, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317364, Nov. 2017.
867. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw406, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317365, Nov. 2017.
868. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw408, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317366, Nov. 2017.
869. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw416, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317367, Nov. 2017.
870. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw417, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317368, Nov. 2017.
871. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptomyces sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw414, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317369, Nov. 2017.
872. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Streptomyces sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw429, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317370, Nov. 2017.
873. **Takanori Satoh, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Mitsuki Fujii, Kouta Nagai, Shota Inoue *and* Kenji Akiyoshi-Hiraoka :** Uncultured Coleodesmium sp. gene for 16S ribosomal RNA, partial sequence, clone: Tw422, --- Microbial diversity on the leaves of Japanese and Taiwan teas ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317371, Nov. 2017.
874. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A1, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317909, Nov. 2017.
875. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A2, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317910, Nov. 2017.
876. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A4, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317911, Nov. 2017.
877. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A5, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317912, Nov. 2017.
878. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A8, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317913, Nov. 2017.
879. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A9, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317914, Nov. 2017.
880. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A10, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317915, Nov. 2017.
881. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Paenibacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: A11, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317916, Nov. 2017.
882. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: A12, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317917, Nov. 2017.
883. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone: A13, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317918, Nov. 2017.
884. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A14, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317919, Nov. 2017.
885. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: A19, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317920, Nov. 2017.
886. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Geobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: A21, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317921, Nov. 2017.
887. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A23, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317922, Nov. 2017.
888. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: A25, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317923, Nov. 2017.
889. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Paenibacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: A28, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317924, Nov. 2017.
890. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Propionibacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: 2B28, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317925, Nov. 2017.
891. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: 2B31, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317926, Nov. 2017.
892. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Veillonella sp.gene for 16S ribosomal RNA, partial sequence, clone: B1, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317927, Nov. 2017.
893. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Bosea sp.gene for 16S ribosomal RNA, partial sequence, clone: B6, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317928, Nov. 2017.
894. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp..gene for 16S ribosomal RNA, partial sequence, clone: B7, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317929, Nov. 2017.
895. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Bosea sp.gene for 16S ribosomal RNA, partial sequence, clone:B10, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317930, Nov. 2017.
896. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: B12, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317931, Nov. 2017.
897. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B13, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317932, Nov. 2017.
898. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Anaerococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: B14, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317933, Nov. 2017.
899. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B15, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317934, Nov. 2017.
900. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B16, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317935, Nov. 2017.
901. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Streptomyces sp. gene for 16S ribosomal RNA, partial sequence, clone: B17, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317936, Nov. 2017.
902. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B19, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317937, Nov. 2017.
903. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B20, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317938, Nov. 2017.
904. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: B24, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317939, Nov. 2017.
905. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Brevundimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: B29, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317940, Nov. 2017.
906. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Brevundimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: B30, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317941, Nov. 2017.
907. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Rahnella sp.gene for 16S ribosomal RNA, partial sequence, clone: C1, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317942, Nov. 2017.
908. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: C33, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317943, Nov. 2017.
909. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Rahnella sp.gene for 16S ribosomal RNA, partial sequence, clone: C39, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317944, Nov. 2017.
910. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone:C42, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317945, Nov. 2017.
911. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone:C47, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317946, Nov. 2017.
912. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Rahnella sp. gene for 16S ribosomal RNA, partial sequence, clone: C50, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317947, Nov. 2017.
913. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Streptomyces sp. gene for 16S ribosomal RNA, partial sequence, clone: C51, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317948, Nov. 2017.
914. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Leuconostoc sp. gene for 16S ribosomal RNA, partial sequence, clone:C52, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317949, Nov. 2017.
915. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: C56, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317950, Nov. 2017.
916. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: C58, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317951, Nov. 2017.
917. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: C59, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317952, Nov. 2017.
918. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone: C64, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317953, Nov. 2017.
919. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Paenibacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: C66, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317954, Nov. 2017.
920. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Rahnella sp. gene for 16S ribosomal RNA, partial sequence, clone: C67, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317955, Nov. 2017.
921. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: C69, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317956, Nov. 2017.
922. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Streptomyces sp. gene for 16S ribosomal RNA, partial sequence, clone: C70, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317957, Nov. 2017.
923. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: C72, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317958, Nov. 2017.
924. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: C73, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317959, Nov. 2017.
925. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D1, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317960, Nov. 2017.
926. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D2, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317961, Nov. 2017.
927. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D3, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317962, Nov. 2017.
928. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Burkholderia sp. gene for 16S ribosomal RNA, partial sequence, clone: D4, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317963, Nov. 2017.
929. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Enterobacteriaceae bacterium gene for 16S ribosomal RNA, partial sequence, clone: D5, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317964, Nov. 2017.
930. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D6, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317965, Nov. 2017.
931. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: D8, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317966, Nov. 2017.
932. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D10, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317967, Nov. 2017.
933. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D13, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317968, Nov. 2017.
934. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D14, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317969, Nov. 2017.
935. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D15, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317970, Nov. 2017.
936. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D16, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317971, Nov. 2017.
937. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: D17, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317972, Nov. 2017.
938. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D18, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317973, Nov. 2017.
939. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D19, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317974, Nov. 2017.
940. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D20, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317975, Nov. 2017.
941. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: D21, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317976, Nov. 2017.
942. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D22, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317977, Nov. 2017.
943. **Mitsuki Fujii, Kouta Nagai, Daiki Fujimoto, Narumi Maegawa, Kenji Akiyoshi-Hiraoka, Shota Inoue *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: D23, --- 16S rRNA gene analysis of bacteria from Awa Bancha tea (fermented tea) produced at Tokushima by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC317978, Nov. 2017.
944. **伏見 賢一 :** 宇宙暗黒物質レビュー, *SMART2017(Scintillator for Medical, Astroparticle and environmental Radiation Technologies),* 2017年11月.
945. **山本 孝 :** XAFS, *近畿化学協会触媒・表面部会 ナノ材料の表面分析講習,* 2017年12月.
946. **Gantulga Uuganbaatar, Erdenebat Erdenedalai, 岡 里美, 谷口 智紀, 平田 晶子, 藤巻 晃, 山本 真由美, 中山 信太郎, 佐藤 高則 :** 大震災に備える放射線・防災教育 ∼留学生および日本人学生に向けて∼, *平成29年度総合科学教育部プロジェクト研究I 成果発表会兼FDプログラム,* 2018年2月.
947. **犬飼 宗弘 :** 多孔性配位高分子に吸着させた気体分子の固体NMR解析, *NIMS微細構造解析プラットフォーム 2017年度 地域セミナー,* 2018年2月.
948. **西山 賢一, 鳥井 真之, 田近 淳, 奧野 充, 井口 隆 :** 地震で流下・堆積した阿蘇カルデラ西部，山王谷川・垂玉川流域の崩壊堆積物, *日本応用地質学会熊本・大分地震災害調査団調査報告書,* 128-138, 2017年4月.
949. **西山 賢一, 鳥井 真之, 井口 隆, 釜井 俊孝, 田近 淳 :** 阿蘇火山研究所周辺で発生した地すべり, *日本応用地質学会熊本・大分地震災害調査団調査報告書,* 120-127, 2017年4月.
950. **高見 智之, 井口 隆, 加藤 靖郎, 西山 賢一, 井口 隆 :** 地震で生じた斜面災害の特徴と地形地質条件からの類型化, *日本応用地質学会熊本・大分地震災害調査団調査報告書,* 85-95, 2017年4月.
951. **日本応用地質学会 2016年熊本・大分地震災害調査団 陥没性亀裂調査班, 西山 賢一, 井口 隆 :** 阿蘇カルデラ内に出現した陥没性亀裂と側方移動, *日本応用地質学会熊本・大分地震災害調査団調査報告書,* 64-72, 2017年4月.
952. **奧野 充, 鳥井 真之, 西山 賢一, 中西 利典, 横田 修一郎 :** 地震で誘発された阿蘇カルデラ・高野尾羽根火山の西側斜面の崩壊プロセスについて, *日本応用地質学会熊本・大分地震災害調査団調査報告書,* 114-119, 2017年4月.
953. **西山 賢一 :** 2016年熊本地震の災害調査報告, *めらんじゅ(宮崎応用地質研究会誌), No.28,* 52-59, 2017年5月.
954. **西山 賢一 :** 徳島市の地盤, *全国電子地盤図の拡張と運用に関する研究報告書,* 97-100, 2017年10月.
955. **Gantulga Uuganbaatar, Erdenebat Erdenedalai, 岡 里美, 谷口 智紀, 平田 晶子, 藤巻 晃, 山本 真由美, 中山 信太郎, 佐藤 高則 :** 平成29年度プロジェクト研究I(佐藤グループ)報告書, --- 大震災に備える放射線・防災教育 ∼留学生および日本人学生に向けて∼ ---, *平成29年度総合科学教育部プロジェクト研究I 報告書,* 1-12, 徳島, 2018年1月.
956. **佐藤 高則, 山本 真由美, 中山 信太郎 :** 大震災に備える放射線・防災教育 ∼留学生および日本人学生に向けて∼, *平成29年度プロジェクト研究 授業実施報告書,* 1, 徳島, 2018年2月.
957. **中村 浩一, 河村 純一 :** 多核NMR による酸化物の局所構造変化とイオン拡散挙動に関する研究, *物質・デバイス領域共同研究拠点研究成果報告書(平成29年度),* 大阪, 2018年3月.
958. **西山 賢一 :** 越高遺跡B地点 地形・地質調査結果, *熊本大学文学部考古学研究室考古学研究室報告,* **Vol.53,** 43-48, 2018年3月.
959. **西山 賢一, 撰田 克哉, 井口 隆 :** 朝倉市妙見川流域で発生した地すべり・斜面崩壊の地形・地質的特徴, *日本応用地質学会九州北部豪雨災害調査団報告書,* 70-77, 2018年3月.
960. **西山 賢一, 田近 淳, 太田 岳洋, 千田 敬二, 長谷川 怜思, 碓井 敏彦, 田村 浩行, 西村 智博, 松澤 真 :** 火山岩地域でおきた特徴的な斜面変動-日田市小野地すべりを中心に, *日本応用地質学会九州北部豪雨災害調査団報告書,* 34-42, 2018年3月.
961. **山本 孝 :** 第66回デンバーX線会議報告, *X線分析の進歩,* **Vol.49,** 143-148, 2018年3月.
962. **山本 孝 :** 第53回X線分析討論会報告, *X線分析の進歩,* **Vol.49,** 95-99, 2018年3月.
963. **原 英俊, 青矢 睦月, 野田 篤, 田辺 晋, 山崎 徹, 大野 哲二, 駒澤 正夫 :** 20万分の1地質図幅「高知」(第2版), 産総研地質調査総合センター, つくば, 2018年7月.
964. **山本 孝 :** 広域X線吸収微細構造(EXAFS) 「X線分光法 第5章4節」, 株式会社 講談社, 2018年9月.
965. **山本 孝 :** 触媒「X線分光法 第6章5節」, 株式会社 講談社, 2018年9月.
966. **村松 康司, 山本 孝 :** X線吸収端近傍構造(XANES)「X線分光法 第5章3節」, 株式会社 講談社, 2018年9月.
967. **齊藤 隆仁, 小山 晋之 :** 基礎物理学実験テキスト (2018年度後期改訂版), 徳島大学生協, 2018年10月.
968. **中村 浩一 :** リチウムイオン電池における高容量化・高電圧化技術と安全対策, --- 第11章 第5節 リチウム遷移金属酸化物における組成変化と構造擾乱による伝導・拡散挙動の変化 ---, 株式会社 技術情報協会, 東京, 2018年10月.
969. **徳島県教育委員会(編), 石田 啓祐, 安間 了 :** 赤色顔料生産遺跡及び関連遺跡の調査 若杉山遺跡 発掘調査編, --- 若杉山遺跡「石灰岩露頭部」におけるHg値のpXRF計測 -辰砂採掘候補地の地球科学的解析(その1)- ---, 徳島県教育委員会, 徳島, 2019年3月.
970. **徳島県教育委員会(編), 石田 啓祐, 安間 了 :** 赤色顔料生産遺跡及び関連遺跡の調査 若杉山遺跡 発掘調査編, --- 若杉山遺跡「石灰岩露頭部」発掘土壌からの赤色重鉱物(辰砂)の検出と同定 -辰砂採掘候補地の地球科学的解析(その2)- ---, 徳島県教育委員会, 徳島, 2019年3月.
971. **阿南市教育委員会(編), 石田 啓祐, 安間 了 :** 若杉山遺跡発掘調査報告書, --- 第5章 自然科学分析 1 若杉山遺跡「採掘坑跡1」におけるHG値のpXRF計測 -辰砂採掘候補地の地球科学的解析(その1)- ---, 阿南市教育委員会, 阿南市, 2019年3月.
972. **阿南市教育委員会(編), 石田 啓祐, 安間 了 :** 若杉山遺跡発掘調査報告書, --- 第5章 自然科学分析 2 若杉山遺跡「採掘坑跡1」発掘土壌からの赤色重鉱物(辰砂)の検出と同定 -辰砂採掘候補地の地球科学的解析(その2)- ---, 阿南市教育委員会, 阿南市, 2019年3月.
973. **Koichi Nakamura, Kosuke Shimokita, Yoichiro Sakamoto, Kuniyuki Koyama, Toshihiro Moriga, Naoaki Kuwata *and* Juinichi Kawamura :** Milling effect on the local structure, site occupation, and site migration in aluminum substituted lithium manganese oxides, *Solid State Ionics,* **Vol.317,** 214-220, 2018.
974. **Yuhei Yamamoto, Azusa Tagami, Toshihiro Shiarasaki, Akira Yonetani, Takashi Yamamoto *and* Shoji Imai :** Role of iron modifier on boron atomization process using graphite furnace-atomic absorption spectrometry based on speciation of iron using X-ray absorption fine structure, *Spectrochimica Acta. Part B: Atomic Spectroscopy,* **Vol.142,** 55-61, 2018.
975. **Takashi Yamamoto *and* Akira Yukumoto :** Discrepancy in the quantitative oxidation-state analysis of Eu species in sulfide phosphors by K-, L1-, and L3-edge XANES spectrometry: Choice of absorption edge and measurement mode, *Journal of Analytical Atomic Spectrometry,* **Vol.33,** *No.4,* 585-592, 2018.
976. **Ko-ichi Magishi, Ryo Watanabe, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama, S. Okada, Y. Kamihara, N. Ohkubo, S. Ban *and* M. Matoba :** NMR study of the layered cobalt oxyphosphide Sr2Sc(Co1-xFex)PO3, *Journal of Physics: Conference Series,* **Vol.969,** *No.1,* 012068-1-012068-5, 2018.
977. **Shoji Imai, Yuhei Yamamoto, Takashi Yamamoto, Kenji KODAMA, Jun Nishimoto *and* Yoichi KIKUCHI :** Sulfur Chemical State and Chemical Composition of Insoluble Substance in Soft Rime, Hard Rime and Snow collected on Remote and Rural Areas in Japan using WDXRF, *Analytical Sciences,* **Vol.34,** *No.5,* 589-598, 2018.
978. **Munehiro Inukai, Masanori Tamura, Satoshi Horike, Masakazu Higuchi, Susumu Kitagawa *and* Koichi Nakamura :** Storage of CO2 into Porous Coordination Polymer Controlled by Molecular Rotor Dynamics, *Angewandte Chemie International Edition,* **Vol.57,** *No.28,* 8687-8690, 2018.
979. **M Takahashi, Y. Inome, S. Yoshii, Reiko Orito *and* 20 coauthors :** A technique for estimating the absolute gain of a photomultiplier tube, *Nuclear Instruments and Methods in Physics Research Section A: Accelerators, Spectrometers, Detectors and Associated Equipment,* **Vol.894,** 1-7, 2018.
980. **K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, H. Ogawa, K. Sato, H. Sekiya, T. Suzuki, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, K. Masuda, K. Martens, Y. Suzuki, B.D. Xu, K. Miuchi, N. Oka, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fukuda, M. Miyasaka, K. Nishijima, Ken-Ichi Fushimi, G. Kanzaki *and* S. Nakamura :** Direct dark matter search by annual modulation with 2.7 years of XMASS-I data, *Physical Review D,* **Vol.97,** 102006, 2018.
981. **K. Abe, H. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, T. Norita, H. Ogawa, K. Sato, H. Sekiya, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, R. Kegasa, K. Masuda, H. Takiya, Ken-Ichi Fushimi, G. Kanzaki, K. Martens, Y. Suzuki, B.D. Xu, R. Fujita, K. Hosokawa, K. Miuchi, N. Oka, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fujita, M. Miyasaka *and* K. Nishijima :** Improved search for two-neutrino double electron capture on 124 Xe and 126 Xe using particle identification in XMASS-I, *Progress of Theoretical and Experimental Physics,* **Vol.2018,** *No.5,* 053D03, 2018.
982. **山本 祐平, 佐名川 洋右, 兼清 恵理, 清水 魁人, 山本 孝, 今井 昭二 :** 実験室型X線吸収分光装置を用いた遠隔山岳地域で採取された湿性沈着中の長距離輸送エアロゾル中に含まれる鉄の化学状態分析, *分析化学,* **Vol.67,** *No.6,* 349-354, 2018年.
983. **Takumi Kishida, Kenji Akiyoshi, Erdenedalai Erdenebat, Enhetomuru Anu, Shoji Imai *and* Yasuo Oyama :** Cellular adverse actions of dibromoacetonitrile, a by-product in water bacterial control, at sublethal levels in rat thymocytes, *Toxicology In Vitro,* **Vol.51,** 145-149, 2018.
984. **Satoshi D. Ohmura, Masaharu Ueno *and* Norikazu Miyoshi :** Strontium-mediated selective protonation of unsaturated linkage of aromatic hydrocarbons and these derivatives, *Tetrahedron Letters,* **Vol.59,** *No.23,* 2268-2271, 2018.
985. **Takashi Yamamoto :** Relation between intensity of XANES pre-edge and XES Kb5 of vanadium and manganese compounds, *Advances in X-ray Analysis,* **Vol.61,** 211-221, 2018.
986. **Shinobu Takao, Oki Sekizawa, Gabor Samjeské, Takuma Kaneko, Kotaro Higashi, Yuske Yoshida, Xiao Zhao, Tomohiro Sakata, Takashi Yamamoto, Tomoya Uruga *and* Yasuhiro Iwasawa :** Observation of Degradation of Pt and Carbon Support in Polymer Electrolyte Fuel Cell Using Combined Nano-X-ray Absorption Fine Structure and Transmission Electron Microscopy Techniques, *ACS Applied Materials & Interfaces,* **Vol.10,** *No.33,* 27734-27744, 2018.
987. **Hitoshi Mizuguchi, Ryota Ishida, Yasushi Kouno, Tadahiko Tachibana, Tomomi Honda, Tatsuro Kijima, Yuhei Yamamoto *and* Toshio Takayanagi :** A Rapid Enrichment Technique for the Ultratrace Determination of Nickel in Water Samples Using a Nanofiber-composite Membrane Filter, *Analytical Sciences,* **Vol.34,** *No.8,* 907-912, 2018.
988. **西山 賢一, 鳥井 真之, 横田 修一郎, 奥野 充 :** 阿蘇火山周辺における斜面崩壊・土石流の発生頻度, *第9回土砂災害に関するシンポジウム論文集,* 157-160, 2018年.
989. **Takanori Satoh :** Supporting Activities and Preservation of the Traditional Food Culture for Sustainable Development in Fukushima, Japan, *Proceedings of International Conference on Sustainable Agriculture,* 144-152, 2018.
990. **S. Kotani, Y. Yoshiwara, Masamichi Ogasawara, M. Sugiura *and* M. Nakajima :** Catalytic Enantioselective Aldol Reaction of Unprotected Carboxylic Acids under Phosphine Oxide Catalysis, *Angewandte Chemie International Edition,* **Vol.57,** 15877-15881, 2018.
991. **山本 祐平, 兼清 恵理, 清水 魁人, 髙橋 玄樹, 橋本 元輝, 今井 昭二 :** 2015年から2018年における国内遠隔山岳地域で採取した降雪中のカドミウムと鉛の組成及びその起源, *分析化学,* **Vol.67,** *No.21,* 733-741, 2018年.
992. **西山 賢一 :** 日本における岩石の風化研究の進展と課題, *地質学雑誌,* **Vol.124,** *No.11,* 877-888, 2018年.
993. **山本 祐平, 佐名川 洋右, 耒見 祐哉, 齋藤 あゆみ, 西本 潤, 菊地 洋一, 今井 昭二 :** 日本の遠隔地における清浄な降雨・雪中のカドミウム, 鉛及び非海塩性硫酸イオンの人為活動に依存しない自然レベルの濃度, *分析化学,* **Vol.68,** *No.1,* 51-64, 2019年.
994. **Kyohei Kawashima, Takehiro Sato, Masamichi Ogasawara, Ken Kamikawa *and* Seiji Mori :** Theoretical Investigations of Rh-Catalyzed Asymmetric 1,4-Addition to Enones using Planar-Chiral Phosphine-Olefin Ligands, *Journal of Computational Chemistry,* **Vol.40,** *No.1,* 113-118, 2019.
995. **Masamichi Ogasawara, Yuuki Enomoto, Mizuho Uryu, Xicheng Yang, Ayami Kataoka *and* Atsushi Ohnishi :** Application of Polysaccharide-Based Chiral HPLC Columns for Separation of Non-Enantiomeric Isomeric Mixtures of Organometallic Compounds, *Organometallics,* **Vol.38,** *No.2,* 512-518, 2019.
996. **Kyohei Nakajima, Temuge Batpurev, Wei M Chan, Fumiya Dokaku, Ken-Ichi Fushimi, Kazuki Kanagawa, Seiya Katagiri, Kohei Kawasaki, Bui T Khai, Hidetoshi Kino, Enki Kinoshita, Tadafumi Kishimoto, Ryuta Hazama, Hirokazu Hiraoka, Daiki Hiyama, Takashi Iida, Masahiro Ishikawa, Xiaolong Li, Tsuyoshi Maeda, Kenji Matsuoka, Michael Moser, Masaharu Nomachi, Izumi Ogawa, Takaki Ohata, Hiroyoshi Sato, Kazuki Shamoto, Makiko Shimada, Masoumeh Shokati, Naruki Takahashi, Yasuhiro Takemoto, Yukichi Takihira, Yoichi Tamagawa, Masashi Tozawa, Kanau Teranishi, Konosuke Tetsuno, Van T T Trang, Masahito Tsuzuki, Saori Umehara, Wei Wang, Sei Yoshida *and* Naoki Yotsunaga :** Upgrading of shielding for rare decay search in CANDLES, *Progress in Nuclear Science and Technology,* **Vol.6,** 144-147, 2019.
997. **Masaharu Ueno, Kusaka Ryo, Satoshi D. Ohmura *and* Norikazu Miyoshi :** Environmentally Benign Ritter Reaction Using Bismuth Salt as a Catalyst, *European Journal of Organic Chemistry,* **Vol.2019,** *No.8,* 1796-1800, 2019.
998. **川村 教一, 西山 賢一 :** 四国地方の主要臨海平野における上部更新統および完新統の対比:現状と課題, *地質学雑誌,* **Vol.125,** *No.1,* 87-105, 2019年.
999. **Masaharu Ueno, Satoshi D. Ohmura, Makoto Wada *and* Norikazu Miyoshi :** Aerobic oxidation of alcohols using bismuth bromide as a catalyst, *Tetrahedron Letters,* **Vol.60,** *No.7,* 570-573, 2019.
1000. **Naoki Nakao, Masaharu Ueno, Shota Sakai, Daich Egawa, Hiroyuki Hanzawa, Shohei Kawasaki, Keigo Kumagai, Makoto Suzuki, Shu Kobayashi *and* Kentaro Hanada :** Natural ligand-nonmimetic inhibitors of the lipid-transfer protein CERT. (N.N., M.U., and S.S. equally contributed to this study and are the co-first authors), *Communications Chemistry,* **Vol.2,** Articlenumber:20, 2019.
1001. **Jr. P. Yumul Graciano, Dimalanta B. Carla, Salapare C. Ricky, Queano L. Karlo, Faustino-Eslava V. Decibel, Marquez J. Edanjarlo, Ramos T. Noelynna, Payot D. Betchaida, Guotana Miguel R. Juan, Gabo-Ratio Aira S. Jillian, Armada T. Leo, Padrones T. Jenielyn, Keisuke Ishida *and* Suzuki Shigeyuki :** Slab rollback and microcontinent subduction in the evolution of the Zambales Ophiolite Complex (Philippines): A review, *Geoscience Frontiers,* **Vol.11,** 23-36, 2019.
1002. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Miyoshi city and Tokushima prefecture, *Proceedings of Awagakkai,* **Vol.62,** 179-188, 2019.
1003. **西山 賢一, 石田 啓祐, 橋本 寿夫, 辻野 泰之, 中尾 賢一, 殿谷 梓 :** 三好市の地質と地形:その生い立ちと特質, *阿波学会紀要,* **Vol.62,** 1-11, 2019年.
1004. **石田 啓祐 :** 新たな国指定天然記念物及び名勝「大歩危小歩危」, *阿波学会紀要, No.62,* 153-160, 2019年.
1005. **A Kozlov, D Chernyal, Y Takemoto, Ken-Ichi Fushimi, K Imagawa, K Yasuda, H Ejiri, Ryuta Hazama, H Ikeda, K Inoue, S Yoshida, R A Etezov, Yu M Gavrilyuk, V V Kazalov *and* V V Kuzminov :** The Dark Matter Search at KamLAND, *Journal of Physics: Conference Series,* **Vol.1390,** 2019.
1006. **井澤 健一, 谷口 智紀 :** 量子論におけるレトロディクション II, *徳島大学総合科学部自然科学研究,* **Vol.31,** 1-4, 2019年.
1007. **井澤 健一, 谷口 智紀 :** 量子レトロディクションと履歴, *徳島大学総合科学部自然科学研究,* **Vol.31,** 5-8, 2019年.
1008. **山本 孝, 宮園 拓自, 栗本 彰人 :** 形状の異なる焦電結晶積層体によるX線発生挙動および放電特性, *X線分析の進歩,* **Vol.50,** 291-298, 2019年.
1009. **Haruka Iwata, Kouta Nagai, Yousuke Endoh, Yurie Ohiwa, Shota Inoue, Kenji Akiyoshi *and* Takanori Satoh :** Different Conformational Thermostabilities among Hexameric Nucleoside Diphosphate Kinases from Moderately thermophilic bacteria, *Proceedings of 24th IUBMB Congress and 15th FAOBMB Congress,* 758, 2018.
1010. **Takanori Satoh, Mayumi YAMAMOTO *and* Shintaro Nakayama :** Activities Supporting Fukushima in Japan for Seven Years, Towards the Sustainable Development, *Proceedings of SUSTAINABILITY SEMINAR 2018: TOWARDS SUSTAINABLE SOCIETY,* 1-9, 2018.
1011. **K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, H. Ogawa, K. Sato, H. Sekiya, T. Suzuki, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, K. Masuda, K. Martens, Y. Suzuki, B.D. Xu, K. Miuchi, N. Oka, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fukuda, M. Miyasaka, K. Nishijima, Ken-Ichi Fushimi, G. Kanzaki *and* S. Nakamura :** Search for dark matter in the form of hidden photons and axion-like particles in the XMASS detector, *Physics Letters B,* **Vol.787,** 153-158, 2018.
1012. **佐藤 高則 :** 大人数教養科目でのSituated learning(SL)によるAL 実践報告, *平成30年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 50-51, 2018年.
1013. **佐藤 高則 :** 教養教育における体験型学習の実践と効果∼「藍染めの科学」の事例∼, *平成30年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 14-15, 2018年.
1014. **K. Abe, K. Hiraide, K. Ichimura, Y. Kishimoto, K. Kobayashi, M. Kobayashi, S. Moriyama, M. Nakahata, H. Ogawa, K. Sato, H. Sekiya, T. Suzuki, O. Takachio, A. Takeda, S. Tasaka, M. Yamashita, B.S. Yang, N.Y. Kim, Y.D. Kim, Y. Itow, K. Kanzawa, K. Masuda, K. Martens, Y. Suzuki, B.D. Xu, K. Miuchi, N. Oka, Y. Takeuchi, Y.H. Kim, K.B. Lee, M.K. Lee, Y. Fukuda, M. Miyasaka, K. Nishijima, Ken-Ichi Fushimi, G. Kanzaki *and* S. Nakamura :** A direct dark matter search in XMASS-I, *Physics Letters B,* **Vol.789,** 45-53, 2019.
1015. **青矢 睦月 :** 1.10 岩石の埋没と上昇, *地球科学の事典,* 20-21, 2018年4月.
1016. **ウォリス サイモン, 青矢 睦月 :** 1.7 大陸衝突, *地球科学の事典,* 14-15, 2018年4月.
1017. **Toshiyuki Moriuchi, Satoshi D. Ohmura *and* Takayo Moriuchi-Kawakami :** Chirality Induction in Bioorganometallic Conjugates, *Inorganics,* **Vol.6,** *No.4,* 110, Oct. 2018.
1018. **青矢 睦月, 西山 賢一 :** 眉山の青色片岩と岩盤クリープ, *第18回日本地質学会四国支部総会巡検案内書,* 1-15, 2018年12月.
1019. **山本 祐平 :** 中国の国内におけるPM2.5中の微量元素分析, *ぶんせき,* 2019年.
1020. **Takashi Yamamoto :** Pre-edge peak in XANES spectra of transition metals for empirical chemical state analysis, *National Conference on Optics, Photonics and Synchrotron Radiation for Technological Application (OPSR-2018),* Indore, India, May 2018.
1021. **Haruka Iwata, Kouta Nagai, Yousuke Endoh, Yurie Ohiwa, Shota Inoue, Kenji Akiyoshi *and* Takanori Satoh :** Different Conformational Thermostabilities among Hexameric Nucleoside Diphosphate Kinases from Moderately thermophilic bacteria, *24th IUBMB Congress and 15th FAOBMB Congress,* 758, Seoul, Jun. 2018.
1022. **Ken-Ichi Fushimi :** Dark matter search project PICOLON ~Present status and future~, *6th Symposium on Neutrinos and Dark Matter in Nuclear Physics 2018, from 29 June 2018 to 4 July 2018,* Jul. 2018.
1023. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Application in Organic Transformations, *Université de Reims Special Seminar,* Reims, Jul. 2018.
1024. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan., *the 32nd annual meeting of Japanese Society for Microbial Ecology (JSME) and 10th Asian symposium on Microbial Ecology,* 248, Ginowan,Okinawa, Jul. 2018.
1025. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Application in Organic Transformations, *Ecole Nationale Supérieure de Chimie de Montpellier Special Seminar,* Montpellier, Jul. 2018.
1026. **Takashi Yamamoto *and* Kazunori Miyamoto :** Differences in Thermal Decomposition Behavior of Platinum Salt on Silica and Alumina Characterized by X-ray Absorption Spectroscopy, *The 8th Tokyo Conference on Advanced Catalytic Science and Technology (TOCAT8),* Yokohama, Aug. 2018.
1027. **Shinobu Takao, Oki Sekizawa, Gabor Samjeské, Takuma Kaneko, Tomohiro Sakata, Kotaro Higashi, Takashi Yamamoto, Tomoya Uruga *and* Yasuhiro Iwasawa :** First direct observation on the degradation of both Pt catalyst and carbon in MEA Pt/C cathode by the same-view nano-XAFS/STEM-EDS imaging technique, *The 8th Tokyo Conference on Advanced Catalytic Science and Technology (TOCAT8),* Yokohama, Aug. 2018.
1028. **Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Axially Chiral Allenic Polymers, *2018 Zhengzhou Chemistry Workshop (XII),* Siem Reap, Sep. 2018.
1029. **Takanori Satoh :** Activities Supporting Fukushima in Japan for Seven Years,Towards the Sustainable Development, *Sustainability Seminar 2018: Towards Sustainable Society,* 20, Kota Kinabalu,Malaysia, Sep. 2018.
1030. **Takanori Satoh :** Supporting Activities and Preservation of the Traditional Food Culture for Sustainable Development in Fukushima, Japan, *The International Conference on Sustainable Agriculture (ICSA),* 43, Sandakan,Malaysia, Sep. 2018.
1031. **Masamichi Ogasawara :** Pd-Catalyzed Synthesis of Optically Active Axially Chiral Allenic Polymers, *International Congress on Pure & Applied Chemistry Langkawi 2018 (ICPAC Langkawi),* Langkawi, Oct. 2018.
1032. **K. Yoshida, Q. Liu, R. Yasue, S. Wada, R. Kimura *and* Masamichi Ogasawara :** Versatile and Enantioselective Preparation of Planar-Chiral Metallocene-Fused 4- Dialkylaminopyridines and Their Application in Asymmetric Organocatalysis, *The 14th International Kyoto Conference on New Aspects of Organic Chemistry (IKCOC-14),* Kyoto, Nov. 2018.
1033. **Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Optically Active Axially Chiral Allenic Polymers, *International Symposium on Catalysis and Fine Chemicals 2018 (C&FC 2018),* Bangkok, Dec. 2018.
1034. **Ken-Ichi Fushimi, Chernyak Dmitry, Hiroyasu Ejiri, Hata Kazumi, Ryuta Hazama, Hirata Shoko, Ikeda Haruo, Imagawa Kyoshiro, Inoue Kunio, Kozlov Alexandre, Reiko Orito, Tatsushi Shima, Takemoto Yasuhiro, Umehara Saori, Yasuda Kensuke *and* Yoshida Sei :** Dark Matter Search by Means of Highly Radiopure NaI(Tl) Scintillator, *JPS conference Proceedings of the Second International Symposium on Radiation Detectors and Their Uses (ISRD2018),* **Vol.24,** 011011-1-011011-6, Jan. 2019.
1035. **Takanori Satoh :** Microbial diversity of Mongolian animal milks, dairy products, and Argal, *2019 INTERNATIONAL STUDENTS CONFERENCE:A collaborative project for university education between Mongolia and Japan,* Ulaanbaatar, Mongolia, Feb. 2019.
1036. **Mitsuki Fujii *and* Takanori Satoh :** Japanese traditional toy, Kendama, and jiggling performance, *2019 INTERNATIONAL STUDENTS CONFERENCE:A collaborative project for university education between Mongolia and Japan,* Ulaanbaatar, Mongolia, Feb. 2019.
1037. **Masamichi Ogasawara :** Synthesis and Applications of Chiral Conjugated- and Cumulated-Dienes, *NTUT Special Seminar,* Reims, Mar. 2019.
1038. **Masamichi Ogasawara, Y.-C. Chen, Y. Wang *and* T. Nakano :** Pd-Catalyzed Synthesis of Axially Chiral Allenic Polymers, *The Fifth International Forum on Advanced Technologies (IFAT 2019),* Taipei, Mar. 2019.
1039. **A. Okazaki, Q. Liu, R. Yasue, K. Yoshida *and* Masamichi Ogasawara :** Enantioselective Synthesis of Planar-Chiral Ferrocene-Fused Cyclic Phosphonic Acids, *The Fifth International Forum on Advanced Technologies (IFAT 2019),* Taipei, Mar. 2019.
1040. **T. Konishi, H. Ichio *and* Masamichi Ogasawara :** Unusual Regioselectivity in Palladium-Catalyzed Cyclization of Nucleophile-Tethered 2-Bromo-1,3-dienes, *The Fifth International Forum on Advanced Technologies (IFAT 2019),* Taipei, Mar. 2019.
1041. **H. Ichiryu, H. Hu, Y. Ura *and* Masamichi Ogasawara :** Synthesis and Application of C1-Symmetric Bisphosphine Ligand Having Diferrocenylphosphino-Donor Moiety, *The Fifth International Forum on Advanced Technologies (IFAT 2019),* Taipei, Mar. 2019.
1042. **Ken-Ichi Fushimi :** PICOLONの報告, *LBGT2019,* Mar. 2019.
1043. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Application in Organic Transformations, *Nanyang Technological University Special Seminar,* Singapore, Mar. 2019.
1044. **小笠原 正道 :** 不斉炭素を持たないキラル化合物の立体制御, *京都大学大学院特別講演,* 2018年4月.
1045. **伏見 賢一 :** *高純度結晶の開発と応用,* 2018年5月.
1046. **安間 了, 申 基澈, 横尾 頼子, 中野 孝教, 黒澤 正紀 :** メソポタミア氾濫源堆積物と粘土板の化学組成・同位体組成にみる都市鉱山のあけぼの, *地球惑星科学関連学会合同大会2018年大会講演要旨,* HTT18-13, 2018年5月.
1047. **山本 孝 :** XAFSによる固体触媒材料の 局所構造，電子状態解析, *第78回分析化学討論会,* 2018年5月.
1048. **山本 祐平, 兼清 恵理, 清水 魁斗, 高橋 玄樹, 橋本 元輝, 今井 昭二 :** 2015年から2018年における国内遠隔山岳地域での降雪降雨中の金属元素組成, *第78年分析化学討論会,* 2018年5月.
1049. **Ai Hasegawa, Haruka Iwata, Hajime Nonoichi, Seiji Arai, Daiki Komahara, Mitsuki Fujii, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis and thermostabilities of cytidylate kinase from Alicyclobacillus acidocaldarius, *The 18th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* 72, Jun. 2018.
1050. **Hajime Nonoichi, Haruka Iwata, Ai Hasegawa, Seiji Arai, Daiki Komahara, Mitsuki Fujii, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis and thermostabilities of uridine kinase from Anoxybacillus flavithermus, *The 18th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* 72, Jun. 2018.
1051. **小笠原 正道 :** キラルな遷移金属錯体の不斉合成と応用, *第30回記念万有札幌シンポジウム,* 2018年7月.
1052. **安間 了 :** 古代西アジアをめぐる水と土と都市の相生相克と都市鉱山の起源, *新学術領域研究キックオフミーティング(筑波大学東京キャンパス),* 2018年7月.
1053. **西山 賢一, 鳥井 真之, 横田 修一郎, 奧野 充, 田近 淳 :** 阿蘇カルデラ・高野尾羽根火山北斜面における地すべり移動体の内部構造, *日本地すべり学会第57回研究発表会,* 2018年8月.
1054. **西山 賢一, 後 誠介 :** 和歌山県大雲取山地に分布する崩壊堆積物の地形・地質的特徴, *日本地質学会第125年学術大会講演要旨集,* 148, 2018年9月.
1055. **森 寛央, 野尻野 旭, 川崎 祐, 岸本 豊, 中村 浩一, 小山 岳秀, 水戸 毅, 八島 光晴, 椋田 秀和, 小手川 恒, 菅原 仁 :** A15型超伝導体V3SiのSi-NMRによる研究, *日本物理学会秋季大会,* 2018年9月.
1056. **森 寛央, 川崎 祐, 岸本 豊, 中村 浩一, 何 長振, 伊藤 満 :** α-CoV2O6における磁気構造のNMRによる研究, *日本物理学会秋季大会,* 2018年9月.
1057. **山本 孝, 行本 晃 :** XANESスペクトルによる遷移金属の半経験的価数評価(2), *第21回XAFS討論会,* 2018年9月.
1058. **石田 啓祐 :** 「大歩危小歩危」新たな国指定天然記念物及び名勝, *日本地質学会125年学術大会講演要旨,* 14, 2018年9月.
1059. **大村 D. 聡, 木村 将大, 久保 誠輝, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを反応剤として用いる選択的官能基変換, *第29回基礎有機化学討論会，東京工業大学大岡山キャンパス(東京)，講演番号1P106,* 2018年9月.
1060. **三好 德和, 西澤 高宏, 北方 咲, 大村 D. 聡, 上野 雅晴 :** ストロンチウム-メタラサイクル中間体の構造推定とその反応性に関する研究, *第29回基礎有機化学討論会， 東京工業大学大岡山キャンパス(東京)，講演番号3C08,* 2018年9月.
1061. **土屋 真緒, 高野 祥太朗, 辻阪 誠, 今井 昭二, 山本 祐平, 申 基澈, 宗林 由樹 :** 雨水中溶存態Ni, Cu, Zn同位体比分析法の開発, *日本地球化学会第65回年会,* 2018年9月.
1062. **犬飼 宗弘, 堀毛 悟史, 西山 裕介 :** プロトン伝導経路とゲスト分子吸着サイトを両立する配位高分子の合成と構造解析, *第57回NMR討論会,* 2018年9月.
1063. **山本 孝, 馬木 良輔 :** 積層した焦電結晶によるX線発生挙動, *日本鉄鋼協会第176回秋季講演大会,* 2018年9月.
1064. **陳 彦洲, 王 ヤン, 中野 環, 小笠原 正道 :** パラジウム触媒による主鎖に軸不斉アレン構造を有するポリマーの合成, *第65回有機金属化学討論会,* 2018年9月.
1065. **瓜生 瑞穂, 片岡 綾美, 小笠原 正道 :** 面不斉フェロセンを基盤とするホスフィンーオレフィン配位子の開発と不斉触媒反応への応用, *第65回有機金属化学討論会,* 2018年9月.
1066. **劉 強, 安江 里紗, 岡崎 願之晋, 吉田 和弘, 小笠原 正道 :** フェロセン縮環した面不斉環状ホスホン酸類の不斉合成, *第65回有機金属化学討論会,* 2018年9月.
1067. **酒井 祥太, 中尾 直樹, 上野 雅晴, 江川 大地, 半沢 宏之, 川崎 祥平, 熊谷 圭吾, 鈴木 誠, 小林 修, 花田 賢太郎 :** 天然リガンドと構造類似性のない脂質輸送タンパク質CERTの新規阻害剤, *第91回日本生化学会大会，国立京都国際会館(京都)，講演番号1T14e(1P-369),* 2018年9月.
1068. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara *and* Kenji Akiyoshi :** Microbial phylogenetic analysis from Cucumber pickles produced at Soma area, Fukushima prefecture, *The 91st Annual Meeting of the Japanese Biochemical Society Abstracts,* 3P264, Sep. 2018.
1069. **Seiji Arai, Ai Hasegawa, Daiki Fujimoto, Haruka Iwata, Hajime Nonoichi, Daiki Komahara, Mitsuki Fujii, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis and thermostabilities of cytidylate kinases from Alicyclobacillus and Geobacillus bacteria, *The 91st Annual Meeting of the Japanese Biochemical Society Abstracts,* 3P153, Sep. 2018.
1070. **Mitsuki Fujii, Haruka Iwata, Kouta Nagai, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis and thermostabilities of nucleoside diphosphate kinases from Alicyclobacillus and Geobacillus bacteria, *The 91st Annual Meeting of the Japanese Biochemical Society Abstracts,* 3P151, Sep. 2018.
1071. **Daiki Komahara, Hajime Nonoichi, Shota Inoue, Haruka Iwata, Ai Hasegawa, Seiji Arai, Mitsuki Fujii, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis, expression, and enzymatic characteristics of uridine kinases from three genera of moderately thermophiles, *The 91st Annual Meeting of the Japanese Biochemical Society Abstracts,* 3P152, Sep. 2018.
1072. **Kenji Akiyoshi, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Mitsuki Fujii, Shota Inoue *and* Takanori Satoh :** Phylogenetic analysis of genes for TTP-synthesizing enzymes from three genera of moderately thermophiles, *The 91st Annual Meeting of the Japanese Biochemical Society Abstracts,* 3P155, Sep. 2018.
1073. **山本 孝, 行本 晃 :** 3d 遷移元素 XANES スペクトルの見かけ上の吸収端シフトおよび価数評価への留意事項, *第122回触媒討論会,* 2018年9月.
1074. **西山 賢一, 日本応用地質学会災害調査団四国班 :** 四国地区で発生した土砂災害, *日本応用地質学会平成30年度研究発表会,* 2018年10月.
1075. **西山 賢一 :** 福岡県朝倉・大分県日田地域における地すべり・土石流の発生頻度, *日本応用地質学会平成30年度研究発表会講演論文集,* 91-92, 2018年10月.
1076. **撰田 克哉, 西山 賢一, 井口 隆 :** 朝倉市妙見川流域で発生した初生岩盤地すべりの地質的特徴, *日本応用地質学会平成30年度研究発表会講演論文集,* 143-144, 2018年10月.
1077. **小笠原 正道 :** 不斉炭素を持たないキラル化合物の立体制御, *2018年度第2回プロセス化学会東四国地区フォーラム,* 2018年10月.
1078. **小笠原 正道 :** ねじれた分子のデザイン・合成・応用, *第8回CSJ化学フェスタ2018,* 2018年10月.
1079. **山本 孝, 宮園 拓自, 馬木 良輔, 栗本 彰人 :** 積層型焦電結晶デバイスによる放電特性およびX線発生挙動, *第54回X線分析討論会,* 2018年10月.
1080. **西山 賢一, 日本応用地質学会災害調査団四国班 :** 四国地区で発生した土砂災害, *平成30年7月豪雨災害調査速報会・日本応用地質学会中国四国支部設立25周年記念行事 記念シンポジウム 講演資料集,* I-12-I-15, 2018年11月.
1081. **西山 賢一, 日本応用地質学会災害調査団四国班 :** 平成30年7月豪雨により愛媛県宇和島市で発生した土砂災害, *防災科学技術研究所2018年度土砂災害に関する研究集会発表概要集,* 22-23, 2018年11月.
1082. **木村 将大, 久保 誠輝, 西澤 高宏, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるラジカル種の生成とその応用, *2018年日本化学会中国四国支部大会，愛媛大学城北キャンパス(愛媛)，講演番号1P-030,* 2018年11月.
1083. **石井 悠佑, 西澤 高宏, 立田 彩和, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いる嵩高い含窒素化合物の合成手法の開発, *2018年日本化学会中国四国支部大会，愛媛大学城北キャンパス(愛媛)，講演番号2G-07,* 2018年11月.
1084. **日下 亮, 大村 D. 聡, 三好 德和, 上野 雅晴 :** 有機ビスマスを含む協奏機能触媒を用いた新規カップリング反応の研究, *2018年日本化学会中国四国支部大会，愛媛大学城北キャンパス(愛媛)，講演番号2G-08,* 2018年11月.
1085. **山本 孝 :** X線吸収分光法によるタングステン-ジルコニウム系複合酸化物固体強酸触媒, *第六回 元素戦略に基づいた触媒設計シンポジウム,* 2018年11月.
1086. **塀内 聖奈, 近藤 真季, 入江 智章, 山本 孝 :** 水熱合成法により調製したタングステン酸ジルコニア(W/Zr=2)の構造と酸触媒活性, *触媒学会第28回キャラクタリゼーション講習会,* 2018年11月.
1087. **佐藤 里緒菜, 栗本 彰人, 谷間 直人, 鎌田 涼太, 山本 孝 :** 金属イオン添加酸化ジルコニウムによるエタノール転換反応, *触媒学会第28回キャラクタリゼーション講習会,* 2018年11月.
1088. **土屋 真緒, 高野 祥太朗, 辻阪 誠, 今井 昭二, 山本 祐平, 申 基澈, 宗林 由樹 :** 西日本における雨水中Ni, Cu, Zn安定同位体比, *第118回化学研究所研究発表会,* 2018年11月.
1089. **西山 賢一, 川村 教一 :** 徳島平野における電子地盤図の作成, *日本地質学会四国支部講演会資料,* 7, 2018年12月.
1090. **尻労安部洞窟遺跡発掘調査団, 西山 賢一 :** 安部遺跡, *平成30年度青森県埋蔵文化財発掘調査報告会,* 2018年12月.
1091. **尻労安部洞窟遺跡発掘調査団・慶應義塾大学, 西山 賢一 :** 青森県尻労安部洞窟, *第32回 東北日本の旧石器文化を語る会福島大会,* 2018年12月.
1092. **安間 了 :** pXRF(携帯型蛍光X線分析計)によるアッシリア浮彫石材の非破壊分析, *渡辺科研研究会,* 2018年12月.
1093. **犬飼 宗弘, 堀毛 悟史, 西山 裕介 :** プロトン伝導経路とゲスト分子吸着サイトを両立する配位高分子, *第44回固体イオニクス討論会,* 2018年12月.
1094. **土井 卓哉, 奥山 勇治, 山本 孝, 中村 崇司, 雨澤 浩史, 大石 昌嗣 :** Mn添加ペロブスカイト型酸化物における新規プロトン溶解反応と, *第44回固体イオニクス討論会,* 2018年12月.
1095. **中村 浩一, 犬飼 宗弘, 森賀 俊広 :** Li4Ti5O12の局所構造の擾乱と伝導挙動の変化, *第44回固体イオニクス討論会講演要旨集,* 100-101, 2018年12月.
1096. **Rie Takahara, Sun Jung-Ting, Kohei Shizukawa, Yusuke Mizuta, Koichi Nakamura, Kei-ichiro Murai *and* Toshihiro Moriga :** Effects of Different Titanium Lithium Ration in LiTin(0,N)z (1.25<n<2.0, Z~4.0) on Electrochemical Performance as an Anode Material, *第25回ヤングセラミストミーティングin 中四国,* 38-39, Dec. 2018.
1097. **西村 弘季, 青矢 睦月 :** 眉山地域三波川帯における変形構造方位と主変形の剪断センス, *第18回日本地質学会四国支部講演会要旨集,* O-2, 2018年12月.
1098. **長谷川 就人, 青矢 睦月 :** 泥質片岩を用いた眉山地域三波川帯の変成分帯とユニット区分, *第18回日本地質学会四国支部講演会要旨集,* P-1, 2018年12月.
1099. **西江 苑香, 青矢 睦月 :** 眉山地域三波川帯の変塩基性岩におけるガーネット仮像の認識, *第18回日本地質学会四国支部講演会要旨集,* P-2, 2018年12月.
1100. **安間 了 :** メソポタミア氾濫源堆積物・粘土製品にみる都市鉱山のあけぼの., *日本地質学会四国支部総会,* 2018年12月.
1101. **佐藤 高則 :** 大人数教養科目でのSituated learning(SL)によるAL 実践報告, *平成30年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 50-51, 2018年12月.
1102. **佐藤 高則 :** 教養教育における体験型学習の実践と効果∼「藍染めの科学」の事例∼, *平成30年度全学FD 徳島大学教育カンファレンス 発表抄録集,* 14-15, 2018年12月.
1103. **小笠原 正道 :** 共役ジエン/集積ジエンの不斉合成と応用, *首都大学東京 特別講演会,* 2019年1月.
1104. **小笠原 正道 :** キラルな遷移金属錯体の触媒的不斉合成, *理研セミナー,* 2019年2月.
1105. **安間 了 :** 古代西アジアをめぐる水と土と都市の相生相克と都市鉱山の起源, *新学術領域研究・領域全体研究会(筑波大学東京キャンパス),* 2019年2月.
1106. **野尻野 旭, 森 寛央, 川崎 祐, 岸本 豊, 中村 浩一, 小山 岳秀, 水戸 毅, 八島 光晴, 椋田 秀和, 小手川 恒, 菅原 仁 :** A15型超伝導体V3SiのSi-NMRによる研究, *日本物理学会秋季大会,* 2019年3月.
1107. **安間 了 :** 水路堆積物および粘土板胎土にみるメソポタミアの都市鉱山化., *文化地質研究会(大谷大学),* 2019年3月.
1108. **真岸 孝一, 久田 旭彦, 齊藤 隆仁, 小山 晋之, 石川 義和 :** PrCo2Sn2Zn18のNMR, *日本物理学会講演概要集,* **Vol.74,** *No.1,* 1983, 2019年3月.
1109. **加藤 優祈, 浦本 翔吾, 出口 博之, 美藤 正樹, 久田 旭彦, 小山 晋之, 萩原 亮 :** YBa2Cu3O7 セラミクスのグレイン間グラス転移および超伝導転移の磁場依存, *日本物理学会講演概要集,* **Vol.74,** *No.1,* 2151, 2019年3月.
1110. **中村 浩一, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一 :** ミリングによるLi4Ti5O12の結晶構造と電気伝導挙動の変化, *日本物理学会第74回年次大会 講演概要集,* 2019年3月.
1111. **小西 卓磨, 一尾 裕章, 小笠原 正道 :** 求核部位を有する2-ブロモ-1,3ジエン類のパラジウム触媒分子内環化における異常な選択性, *日本化学会第99春季年会,* 2019年3月.
1112. **一柳 浩輝, 胡 淏, 浦 康之, 小笠原 正道 :** ジフェロセニルホスフィノ基を有する非対称不斉二座ホスフィン配位子の開発と応用, *日本化学会第99春季年会,* 2019年3月.
1113. **山本 孝 :** XANESの使いどころ:酸化数とpre-edge peak, *日本化学会第99春季年会,* 2019年3月.
1114. **上野 雅晴, 奥村 眞優, 大村 D. 聡, 三好 德和 :** スチルバゾール類の一方的な光異性化反応の研究, *日本化学会第99春季年会, 甲南大学(兵庫), 講演番号1B4-42,* 2019年3月.
1115. **犬飼 宗弘, 栗原 拓也, 野田 泰斗, 吉川 輝, 中村 浩一 :** 高圧ガス雰囲気下のその場高分解能固体NMR, *日本化学会 第99春季年会,* 2019年3月.
1116. **鷹尾 忍, 関澤 央輝, 金子 拓真, 東 晃太朗, 坂田 智裕, 山本 孝, 宇留賀 朋哉, 岩澤 康裕 :** nano XAFS/STEM-EDS 同視野測定を用いた燃料電池Pt3Co/C カ ソード触媒の劣化機構解析, *日本化学会第99春季年会,* 2019年3月.
1117. **金子 拓真, 坂田 智裕, 東 晃太朗, 関澤 央輝, 鷹尾 忍, 吉田 祐介, 山本 孝, 宇留賀 朋哉, 岩澤 康裕 :** In-situ リアルタイムHR-XANES/XAFS/XRD 同時計測による実走行 モード下での固体高分子形燃料電池カソード触媒の吸着酸素種と 構造の動的挙動の研究, *日本化学会第99春季年会,* 2019年3月.
1118. **西江 苑香, 青矢 睦月 :** 眉山地域三波川帯の変塩基性岩におけるガーネット仮像の認識, *変成岩などシンポジウム2019蒲郡,* P-3, 2019年3月.
1119. **長谷川 就人, 青矢 睦月 :** 泥質片岩を用いた眉山地域三波川帯の変成分帯とユニット区分, *変成岩などシンポジウム2019蒲郡,* P-2, 2019年3月.
1120. **西村 弘季, 青矢 睦月 :** 眉山地域三波川帯における変形構造方位と主変形の剪断センス, *変成岩などシンポジウム蒲郡,* O-1, 2019年3月.
1121. **日下 亮, 大村 D. 聡, 三好 德和, 上野 雅晴 :** ビスマス化合物を触媒とする環境調和型有機合成反応の開発, *日本化学会第99春季年会, 甲南大学(兵庫), 講演番号3PC-027,* 2019年3月.
1122. **大村 D. 聡, 木村 将大, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを電子源とする芳香族カルボニル/アミド官能基変換反応の検討, *日本化学会第99春季年会, 甲南大学(兵庫), 講演番号3PC-056,* 2019年3月.
1123. **平田 和也, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いる含窒素ディスク状分子へのアルキル置換基導入の検討, *日本化学会第99春季年会, 甲南大学(兵庫), 講演番号3PC-057,* 2019年3月.
1124. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Propionibacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: E5, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318148, Apr. 2018.
1125. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone: E6, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318149, Apr. 2018.
1126. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: E11, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318150, Apr. 2018.
1127. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone: E16, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318151, Apr. 2018.
1128. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Brevundimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: E18, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318152, Apr. 2018.
1129. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Finegoldia sp. gene for 16S ribosomal RNA, partial sequence, clone: E7, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318153, Apr. 2018.
1130. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Finegoldia sp. gene for 16S ribosomal RNA, partial sequence, clone: E17, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318154, Apr. 2018.
1131. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Finegoldia sp. gene for 16S ribosomal RNA, partial sequence, clone: E32, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318155, Apr. 2018.
1132. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Geobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: E10, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318156, Apr. 2018.
1133. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: E8, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318157, Apr. 2018.
1134. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: E15, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318158, Apr. 2018.
1135. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: E26, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318159, Apr. 2018.
1136. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: E30, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318160, Apr. 2018.
1137. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: E1, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318161, Apr. 2018.
1138. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: E2, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318162, Apr. 2018.
1139. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: E13, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318163, Apr. 2018.
1140. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: E29, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318164, Apr. 2018.
1141. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Moraxella sp. gene for 16S ribosomal RNA, partial sequence, clone: E9, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318165, Apr. 2018.
1142. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: E12, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318166, Apr. 2018.
1143. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: E14, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318167, Apr. 2018.
1144. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: E24, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318168, Apr. 2018.
1145. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: E25, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318169, Apr. 2018.
1146. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: E28, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318170, Apr. 2018.
1147. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Brevundimonas sp. gene for 16S ribosomal RNA, partial sequence, clone: F50, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318171, Apr. 2018.
1148. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Corynebacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: F44, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318172, Apr. 2018.
1149. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Edaphobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: F6, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318173, Apr. 2018.
1150. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: F53, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318174, Apr. 2018.
1151. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Methylobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: F48, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318175, Apr. 2018.
1152. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Methyloversatilis sp. gene for 16S ribosomal RNA, partial sequence, clone: F131, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318176, Apr. 2018.
1153. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Peptoniphilus sp. gene for 16S ribosomal RNA, partial sequence, clone:F45, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318177, Apr. 2018.
1154. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Rahnella sp. gene for 16S ribosomal RNA, partial sequence, clone: F3, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318178, Apr. 2018.
1155. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: F47, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318179, Apr. 2018.
1156. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: F51, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318180, Apr. 2018.
1157. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured bacterium gene for 16S ribosomal RNA, partial sequence, clone: F137, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318181, Apr. 2018.
1158. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: F138, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318182, Apr. 2018.
1159. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: F140, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318183, Apr. 2018.
1160. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: F2, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318184, Apr. 2018.
1161. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: F4, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318185, Apr. 2018.
1162. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: F49, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318186, Apr. 2018.
1163. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone: F52, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318187, Apr. 2018.
1164. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone: F139, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318188, Apr. 2018.
1165. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Roseburia sp. gene for 16S ribosomal RNA, partial sequence, clone: G4, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318189, Apr. 2018.
1166. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G6, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318190, Apr. 2018.
1167. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G7, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318191, Apr. 2018.
1168. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G8, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318192, Apr. 2018.
1169. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Comamonas sp. gene for 16S ribosomal RNA, partial sequence, clone: G10, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318193, Apr. 2018.
1170. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G11, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318194, Apr. 2018.
1171. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G12, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318195, Apr. 2018.
1172. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G14, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318196, Apr. 2018.
1173. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G15, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318197, Apr. 2018.
1174. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: G19, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318198, Apr. 2018.
1175. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Comamonas sp. gene for 16S ribosomal RNA, partial sequence, clone: G21, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318199, Apr. 2018.
1176. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G22, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318200, Apr. 2018.
1177. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G23, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318201, Apr. 2018.
1178. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G24, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318202, Apr. 2018.
1179. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G25, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318203, Apr. 2018.
1180. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G30, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318204, Apr. 2018.
1181. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G32, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318205, Apr. 2018.
1182. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Pseudomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: G36, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318206, Apr. 2018.
1183. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Pseudomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: G37, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318207, Apr. 2018.
1184. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: G38, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318208, Apr. 2018.
1185. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: G43, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318209, Apr. 2018.
1186. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Pseudomonas sp. gene for 16S ribosomal RNA, partial sequence, clone: G45, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318210, Apr. 2018.
1187. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G47, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318211, Apr. 2018.
1188. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone: G53, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318212, Apr. 2018.
1189. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: H4, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318213, Apr. 2018.
1190. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingobium sp. gene for 16S ribosomal RNA, partial sequence, clone: H6, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318214, Apr. 2018.
1191. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Comamonas sp. gene for 16S ribosomal RNA, partial sequence, clone: H8, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318215, Apr. 2018.
1192. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H12, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318216, Apr. 2018.
1193. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H16, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318217, Apr. 2018.
1194. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H17, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318218, Apr. 2018.
1195. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H19, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318219, Apr. 2018.
1196. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H20, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318220, Apr. 2018.
1197. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Gluconacetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H22, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318221, Apr. 2018.
1198. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H23, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318222, Apr. 2018.
1199. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H26, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318223, Apr. 2018.
1200. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H27, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318224, Apr. 2018.
1201. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: H32, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318225, Apr. 2018.
1202. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Sphingobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone: H33, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318226, Apr. 2018.
1203. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H34, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318227, Apr. 2018.
1204. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H35, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318228, Apr. 2018.
1205. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H38, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318229, Apr. 2018.
1206. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone: H43, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318230, Apr. 2018.
1207. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H47, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318231, Apr. 2018.
1208. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H48, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318232, Apr. 2018.
1209. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H51, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318233, Apr. 2018.
1210. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Acetobacter sp. gene for 16S ribosomal RNA, partial sequence, clone: H53, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318234, Apr. 2018.
1211. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi-Hiraoka *and* Takanori Satoh :** Uncultured Comamonas sp. gene for 16S ribosomal RNA, partial sequence, clone: H55, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318235, Apr. 2018.
1212. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Methylobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:BB110, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318236, Apr. 2018.
1213. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Staphylococcus sp. gene for 16S ribosomal RNA, partial sequence, clone:BB111, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* Apr. 2018.
1214. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone:BB113, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318238, Apr. 2018.
1215. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Brachybacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:BB118, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318239, Apr. 2018.
1216. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Brachybacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:BB124, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318240, Apr. 2018.
1217. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone:BB133, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318241, Apr. 2018.
1218. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Brachybacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:BB138, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318242, Apr. 2018.
1219. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Brachybacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:BB210, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318243, Apr. 2018.
1220. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Acidovorax sp. gene for 16S ribosomal RNA, partial sequence, clone:IB206, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318244, Apr. 2018.
1221. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Bosea sp. gene for 16S ribosomal RNA, partial sequence, clone:IB211, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318245, Apr. 2018.
1222. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Chryseobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:IB213, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318246, Apr. 2018.
1223. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Escherichia sp. gene for 16S ribosomal RNA, partial sequence, clone:IB219, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318247, Apr. 2018.
1224. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Herbaspirillum sp. gene for 16S ribosomal RNA, partial sequence, clone:IB209, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318248, Apr. 2018.
1225. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Hymenobacter sp. gene for 16S ribosomal RNA, partial sequence, clone:IB210, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318249, Apr. 2018.
1226. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Methylobacterium sp. gene for 16S ribosomal RNA, partial sequence, clone:IB217, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318250, Apr. 2018.
1227. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Pseudomonas sp. gene for 16S ribosomal RNA, partial sequence, clone:IB203, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318251, Apr. 2018.
1228. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Sphingomonas sp. gene for 16S ribosomal RNA, partial sequence, clone:IB218, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318252, Apr. 2018.
1229. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured bacterium gene for 16S ribosomal RNA, partial sequence, clone:IB154, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318253, Apr. 2018.
1230. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone:SC8, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318254, Apr. 2018.
1231. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Enterobacteriaceae bacterium gene for 16S ribosomal RNA, partial sequence, clone:SC31, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318255, Apr. 2018.
1232. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone:SC209, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318256, Apr. 2018.
1233. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Enterobacter sp. gene for 16S ribosomal RNA, partial sequence, clone:SC212, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318257, Apr. 2018.
1234. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone:SC13, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318258, Apr. 2018.
1235. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone:SC39, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318259, Apr. 2018.
1236. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone:SC40, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318260, Apr. 2018.
1237. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Klebsiella sp. gene for 16S ribosomal RNA, partial sequence, clone:SC204, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318261, Apr. 2018.
1238. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC1, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318262, Apr. 2018.
1239. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC3, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318263, Apr. 2018.
1240. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC6, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318264, Apr. 2018.
1241. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC9, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318265, Apr. 2018.
1242. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC18, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318266, Apr. 2018.
1243. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC37, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318267, Apr. 2018.
1244. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC42, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318268, Apr. 2018.
1245. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC43, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318269, Apr. 2018.
1246. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC49, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318270, Apr. 2018.
1247. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC52, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318271, Apr. 2018.
1248. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC60, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318272, Apr. 2018.
1249. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC222, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318273, Apr. 2018.
1250. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactococcus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC35, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318274, Apr. 2018.
1251. **Mitsuki Fujii, Haruka Iwata, Hajime Nonoichi, Ai Hasegawa, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Uncultured Lactobacillus sp. gene for 16S ribosomal RNA, partial sequence, clone:SC10, --- Phylogenetic analysis for bacteria on fermented and non-fermented tea leaves by clone library method ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC318275, Apr. 2018.
1252. **Munehiro Inukai :** Storage and dynamics of gas molecules into porous coordination polymer, *1. 1st India-Japan NMR workshop,* Jun. 2018.
1253. **山本 孝 :** XANES pre-edge ピークによる遷移金属化合物の化学状態分析およびOPSR会議報告, *日本鉄鋼協会 誤差因子フォーラム 第1回フォーラム研究会,* 2018年7月.
1254. **久田 旭彦 :** 徳島大学社会産業理工学研究交流会2018, *超微細セラミックス高温超伝導体の酸化・還元効果,* 2018年9月.
1255. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB1 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379266, Oct. 2018.
1256. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB2 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379267, Oct. 2018.
1257. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB3 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379268, Oct. 2018.
1258. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB6 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379269, Oct. 2018.
1259. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB8 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379270, Oct. 2018.
1260. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB11 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379271, Oct. 2018.
1261. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB16 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379272, Oct. 2018.
1262. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB17 gene for 16S ribosomal RNA, partial sequence., --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379273, Oct. 2018.
1263. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB18 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379274, Oct. 2018.
1264. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB19 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379275, Oct. 2018.
1265. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB20 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379276, Oct. 2018.
1266. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB21 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379277, Oct. 2018.
1267. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. HIB23 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379278, Oct. 2018.
1268. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB24 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379279, Oct. 2018.
1269. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J2 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379379, Oct. 2018.
1270. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J7 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379380, Oct. 2018.
1271. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J8 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379381, Oct. 2018.
1272. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J10 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379382, Oct. 2018.
1273. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J11 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379383, Oct. 2018.
1274. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J16 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379384, Oct. 2018.
1275. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J17 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379385, Oct. 2018.
1276. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Methyloversatilis sp. J18 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379386, Oct. 2018.
1277. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Sphingobium sp. J19 gene for 16S ribosomal RNA, partial sequence., --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379387, Oct. 2018.
1278. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J20 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379388, Oct. 2018.
1279. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J22 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379389, Oct. 2018.
1280. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J109 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379390, Oct. 2018.
1281. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. J218 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379391, Oct. 2018.
1282. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L32 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379392, Oct. 2018.
1283. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L34 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379393, Oct. 2018.
1284. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Methyloversatilis sp. L35 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379394, Oct. 2018.
1285. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Sphingobacterium sp. L36 gene for 16S ribosomal RNA, partial sequence., --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379395, Oct. 2018.
1286. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L37 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379396, Oct. 2018.
1287. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L38 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379397, Oct. 2018.
1288. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L39 gene for 16S ribosomal RNA, partial sequence., --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379398, Oct. 2018.
1289. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L40 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379399, Oct. 2018.
1290. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Methyloversatilis sp. L41 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379400, Oct. 2018.
1291. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L42 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379401, Oct. 2018.
1292. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L43 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379402, Oct. 2018.
1293. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L45 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379403, Oct. 2018.
1294. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Paenibacillus sp. L48 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379404, Oct. 2018.
1295. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Acinetobacter sp. L53 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379405, Oct. 2018.
1296. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M4 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379406, Oct. 2018.
1297. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M5 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379407, Oct. 2018.
1298. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M9 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379408, Oct. 2018.
1299. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M15 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379409, Oct. 2018.
1300. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. M24 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379410, Oct. 2018.
1301. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M28 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379411, Oct. 2018.
1302. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M29 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379412, Oct. 2018.
1303. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Sphingomonas sp. M31 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379413, Oct. 2018.
1304. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Methyloversatilis sp. M36 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379414, Oct. 2018.
1305. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M37 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379415, Oct. 2018.
1306. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M39 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379416, Oct. 2018.
1307. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. M42 gene for 16S ribosomal RNA, partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379417, Oct. 2018.
1308. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M49 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379418, Oct. 2018.
1309. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M51 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC379419, Oct. 2018.
1310. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Klebsiella sp. HIB22 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC380664, Oct. 2018.
1311. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. J110 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC380665, Oct. 2018.
1312. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. L47 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC380666, Oct. 2018.
1313. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M44 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC380667, Oct. 2018.
1314. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Maki Takeda, Ai Hasegawa, Mariko Nakae *and* Kenji Akiyoshi :** Uncultured Lactobacillus sp. M53 gene for 16S ribosomal RNA,partial sequence, --- Comparison of bacterial diversity among the Awa Bancha tea leaves produced at Tokushima prefecture, Japan ---, *DDBJ/EMBL/GenBank International Nucleotide Sequence Database,* LC380668, Oct. 2018.
1315. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2018年10月.
1316. **久田 旭彦 :** 第11回理学系大学教育に関する研究フォーラム, *ICT機器を活用した物理実験授業の開発,* 2018年11月.
1317. **佐藤 高則 :** SUSTAINABILITY SEMINAR 2018およびICSA2018 (第140回徳島生物学会)での学会発表報告, *徳島生物学会会報,* **Vol.71,** *No.2,* 25-27, 2019年1月.
1318. **西山 賢一 :** 阿蘇の自然と歴史から斜面災害について考える, *第1回阿蘇サイエンスカフェ話題提供,熊本大学くまもと水循環・減災研究教育センター,* 2019年3月.
1319. **山本 孝 :** 応用理数コース自然科学系の学生実験におけるe-コンテンツの利用, *教育シンポジウム2019,* 2019年3月.
1320. **西山 賢一 :** 越高遺跡A地点 地形・地質調査結果, *熊本大学文学部考古学研究室報告,* **Vol.54,** 35-40, 2019年3月.
1321. **中村 浩一, 河村 純一 :** 多核NMR による酸化物の局所構造変化とイオン拡散挙動に関する研究, *物質・デバイス領域共同研究拠点研究成果報告書(平成30年度),* 大阪, 2019年3月.
1322. **安間 了, 前田 修, 常木 晃 :** イラク国北部Qalat Said Ahmadan出土の黒曜石製石器の化学組成と原産地推定, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2018年度,* 149-154, 2019年3月.
1323. **常木 晃, 渡部 展也, 安間 了, 辰巳 祐樹, ラワ カリム サリ :** 肥沃な三日月地帯東部の新石器化-イラク・クルディスタン，スレマニ地域チャルモ遺跡の調査(2018年)-, *第26回西アジア発掘調査報告会報告集,* 19-23, 2019年3月.
1324. **石田 啓祐, 安間 了 :** 若杉山遺跡「石灰岩露頭」におけるHg値のpXRF解析-辰砂採掘候補地の地球科学的解析(その1)-, *徳島県埋蔵文化財調査報告書第4集「赤色顔料生産遺跡及び関連遺跡の調査-若杉山遺跡 発掘調査編」,* **Vol.4,** 85-98, 2019年3月.
1325. **石田 啓祐, 安間 了 :** 若杉山遺跡「石灰岩露頭」発掘土壌からの赤色重鉱物(辰砂)の検出と同定-辰砂採掘候補地の地球科学的解析(その2)-, *徳島県埋蔵文化財調査報告書第4集「赤色顔料生産遺跡及び関連遺跡の調査-若杉山遺跡 発掘調査編」,* **Vol.4,** 99-111, 2019年3月.
1326. **石田 啓祐, 安間 了 :** 若杉山遺跡「採掘坑跡1」におけるHg値のpXRF計測-辰砂採掘候補地の地球科学的解析(その1)-, *阿南市埋蔵文化財発掘調査報告書第1冊「若杉山遺跡発掘調査報告書-平成30年度国庫補助事業報告書-」,* **Vol.1,** 40-48, 2019年3月.
1327. **石田 啓祐, 安間 了 :** 若杉山遺跡「採掘坑跡1」発掘土壌からの赤色重鉱物(辰砂)の検出と同定-辰砂採掘候補地の地球科学的解析(その2)-, *阿南市埋蔵文化財発掘調査報告書第1冊「若杉山遺跡発掘調査報告書-平成30年度国庫補助事業報告書-」,* **Vol.1,** 49-59, 2019年3月.
1328. **岩澤 康裕, 小林 修, 冨重 圭一, 関根 泰, 上野 雅晴, 唯 美津木 :** 化学の指針シリーズ 触媒化学, 株式会社 裳華房, 2019年5月.
1329. **杉山 茂, 森賀 俊広, 加藤 雅裕, 村井 啓一郎, 堀河 俊英, 霜田 直宏, 古部 昭広, 柳谷 伸一郎, 小笠原 正道, 山本 孝, 中村 嘉利, 浅田 元子, 佐々木 千鶴, 田中 秀治, 竹内 政樹, 竹谷 豊, 奥村 仙示, 増田 真志, 岡本 敏弘 :** 枯渇資源と技術開発, --- 徳島大学における分野融合型枯渇資源対応技術の開発 ---, 徳島大学産業院出版部, 徳島, 2020年3月.
1330. **西山 賢一, 鳥井 真之, 横田 修一郎, 若月 強, 井上 弦, 中尾 賢一, 星出 和裕, 奥野 充 :** 阿蘇カルデラ壁斜面における斜面崩壊の発生頻度, *第四紀研究,* **Vol.58,** *No.2,* 149-162, 2019年.
1331. **Satoshi D. Ohmura, Keisuke Yamana, Masaharu Ueno *and* Norikazu Miyoshi :** Methoxycarbonyl Group as a Conformational Regulator for the Benzene Ring of Triphenylamines, *ChemistrySelect,* **Vol.4,** *No.13,* 3799-3802, 2019.
1332. **A. A. Saddique, H. Masuda, Ryo Anma, P. Bhattacharya, Y. Yokoo *and* Y. Shimizu :** Hydrogeochemical and isotopic signatures for the identification of seawater intrusion in the paleobeach aquifer of Cox's Bazar city and its surrounding area, south-east Bangladesh., *Groundwater for Sustainable Development,* **Vol.9,** *No.100215,* 2019.
1333. **Hironao Shinjoe, Yuji Orihashi *and* Ryo Anma :** U-Pb ages of Miocene near-trench granitic rocks of the Southwest Japan arc: implications for magmatism related to hot subduction, *Geological Magazine,* 2019.
1334. **Masatsugu Oishi, Takashi Nakamura, Iwao Watanabe, Takashi Yamamoto, Takuya Doi, Toshiaki Ina, Kiyofumi Nitta, Koji Amezawa *and* Yuji Okuyama :** Evaluation of the Electronic and Local Structure of Mn in Proton-Conducting Oxide, Ca(Zr,Mn)O3-δ, to Elucidate a Direct Hydrogen-Dissolution Reaction, *The Journal of Physical Chemistry C,* **Vol.123,** 16034-16045, 2019.
1335. **Mitsuhiro Nakamura, Kazuo Matsuda, Misaki Nakamura, Kyohei Yamashita, Tomoko Suzuki *and* Satoshi Inouye :** Enzymatic conversion of Cypridina luciferyl sulfate to Cypridina luciferin with coenzyme A as a sulfate acceptor in Cypridina (Vargula) hilgendorfii, *Photochemistry and Photobiology,* **Vol.95,** *No.6,* 1376-1386, 2019.
1336. **Munehiro Inukai, Yusuke Nishiyama, Kayako Honjo, Chinmoy Das, Susumu Kitagawa *and* Satoshi Horike :** Glass-phase coordination polymer displaying proton conductivity and guest-accessible porosity, *Chemical Communications,* **Vol.55,** *No.59,* 8528-8531, 2019.
1337. **Takanori Satoh, Yousuke Endoh, Kayo Takechi, Mitsuki Fujii, Misato Tsuchie, Hiroto Itoh, Hayata Sakai, Yuiko Hanno, Tomomi Maemura, Yuto Oe, Kohta Nagai, Haruka Iwata, Yurie Ohiwa *and* Haruka Shibagaki :** Genetic analysis and thermostabilities of Nucleoside diphosphate kinases from moderately thermophilic Geobacillus bacteria, *Proceedings of 14th international symposium of the protein society of Thailand,* 200-206, 2019.
1338. **N. Hayman, M. Rioux, Ryo Anma, K. Tani, D. Dunkley, J. Crowley *and* M. Schmitz :** Accretion and oxidation of a superfast-spread axial melt lens: TIMS and SIMS zircon analysis of the IODP Hole 1256D gabbros, *Lithos,* **Vol.348-349,** *No.105184,* 2019.
1339. **A. Acharyya, I. Agudo, Reiko Orito *and* 468 coauthors :** Monte Carlo studies for the optimisation of the Cherenkov Telescope Array layout, *Astroparticle Physics,* **Vol.111,** 35-53, 2019.
1340. **A. Tsuneki, K. Rasheed, N. Watanabe, Ryo Anma, Y. Tatsumi *and* M. Minami :** Landscape and early farming at Neolithic sites in Slemani, Iraqi Kurdistan: A case study of Jarmo and Qalat Said Ahmadan., *Paleorient,* **Vol.45,** *No.2,* 33-51, 2019.
1341. **Yen-Chou Chen *and* Masamichi Ogasawara :** Palladium-Catalyzed Sequential Twofold Nucleophilic Substitution on 3-Bromopenta-2,4-dienyl Phosphate: Preparation of C1- and C2-Symmetric Doubly Functionalized Allenes, *The Journal of Organic Chemistry,* **Vol.84,** 12463-12470, 2019.
1342. **Sawano Eri *and* Masamichi Ogasawara :** Palladium-Catalyzed Three-Component Coupling of 1,1-Dibromoalkenes, Vinylzinc Chloride, and Soft Nucleophiles: One-Pot Synthesis of 1,3-Disubstituted Allenes, *ACS Omega,* **Vol.4,** 19499-19504, 2019.
1343. **Satoshi D. Ohmura, Toshikazu Hirao, Norikazu Miyoshi *and* Toshiyuki Moriuchi :** Cocrystal Structure of the Redox-active Phenylenediamine and Quinonediimine Derivatives, *X-ray Structure Analysis Online,* **Vol.35,** *No.11,* 63-65, 2019.
1344. **畑 和実, 崔 銀珠, 伏見 賢一, 折戸 玲子 :** Measurement of the scintillation response of neutron scattering in NaI(Tl) scintillator for dark matter search, *Proceedings of the 33rd Workshop on Radiation Detectors and Their Uses,* 69-76, 2019年.
1345. **Takashi Yamamoto *and* Akihito Kurimoto :** Ga ion-doped ZrO2 Catalyst Characterized by XRD, XAFS, and 2-Butanol Decomposition, *Analytical Sciences,* **Vol.36,** *No.1,* 41-46, 2020.
1346. **Yoshida Kazuhiro, Liu Qiang, Yasue Risa, Wada Shiro, Kimura Ryosuke, Konishi Takuma *and* Masamichi Ogasawara :** Versatile and Enantioselective Preparation of Planar-Chiral Metallocene-Fused 4-Dialkylaminopyridines and Their Application in Asymmetric Organocatalysis, *ACS Catalysis,* **Vol.10,** 292-301, 2020.
1347. **Masaharu Ueno, Norikazu Miyoshi, Hanada Kentaro *and* Kobayashi Shu :** Three-component, one-pot tandem Sonogashira/Suzuki-Miyaura coupling reactions for the synthesis of a library of ceramide-transport protein inhibitors that were designed in silico, *Asian Journal of Organic Chemistry,* **Vol.9,** *No.2,* 267-273, 2020.
1348. **A. A. Saddique, Y. Kon, Ryo Anma, H. Masuda, P. Bhattacharya, Y. Yokoo, S. Basak Bipulendu *and* K. Shinoda :** Source of U and Th in a paleobeach groundwater aquifer at Coxs Bazar, southeast Bangladesh, *Groundwater for Sustainable Development,* **Vol.10,** 100332, 2020.
1349. **Naoki Ogiwara, Hirokazu Kobayashi, Munehiro Inukai, Yusuke Nishiyama, Patricia Concepción, Fernando Rey *and* Hiroshi Kitagawa :** Ligand-Functionalization-Controlled Activity of MetalOrganic Framework-Encapsulated Pt Nanocatalyst toward Activation of Water, *Nano Letters,* **Vol.20,** *No.1,* 426-432, 2020.
1350. **Hu Hao, Ichiryu Hiroki, Naoki Sekitani, Nakajima Kiyohiko, Ohki Yasuhiro *and* Masamichi Ogasawara :** Synthesis, Characterization, and Application of Segphos Derivative Having Diferrocenylphosphino-Donor Moieties, *Organometallics,* **Vol.39,** 788-792, 2020.
1351. **Ko-ichi Magishi, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama, Namiki Takahiro *and* Nishimura Katsuhiko :** NMR Study of Caged Cubic Compound NdTr2Al20 (Tr = Ti and V), *JPS Conference Proceedings,* **Vol.29,** 015005-1-015005-6, 2020.
1352. **Satoshi Inouye, Yuiko Sahara-Miura, Mitsuhiro Nakamura *and* Takamitsu Hosoya :** Expression, purification, and characterization of recombinant apoPholasin, *Protein Expression and Purification,* 105615, 2020.
1353. **Shinobu Takao, Oki Sekizawa, Kotaro Higashi, Gabor Samjeské, Takuma Kaneko, Tomohiro Sakata, Takashi Yamamoto, Tomoya Uruga *and* Yasuhiro Iwasawa :** Visualization Analysis of Pt and Co Species in Degraded Pt3Co/C Electrocatalyst Layers of a Polymer Electrolyte Fuel Cell Using a Same-View Nano-XAFS/STEM-EDS Combination Technique, *ACS Applied Materials & Interfaces,* **Vol.12,** *No.2,* 2299-2312, 2020.
1354. **Kazuaki Kisu, Sangryun Kim, Munehiro Inukai, Hiroyuki Oguchi, Shigeyuki Takagi *and* Shin-ichi Orimo :** Magnesium Borohydride Ammonia Borane as a Magnesium Ionic Conductor, *ACS Applied Energy Materials,* **Vol.3,** *No.4,* 3174-3179, 2020.
1355. **Satoshi Inouye, Yuiko Miura-Sahara, Rie Iimori, Yuki Sakata, Yuki Hazama, Suguru Yoshida, Mitsuhiro Nakamura *and* Takamitsu Hosoya :** A novel yellow fluorescent protein of recombinant apoPholasin with dehydrocoelenterazine, *Biochemical and Biophysical Research Communications,* **Vol.526,** *No.2,* 404-409, 2020.
1356. **Ko-ichi Magishi, Haruki Yona, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama *and* Yosikazu Isikawa :** NMR Study of Caged Compound RCo2Sn2Zn18 (R = La and Pr), *JPS Conference Proceedings,* **Vol.30,** 011112-1-011112-6, 2020.
1357. **Akihiko Hisada, Kuniyuki Koyama, Ko-ichi Magishi, Takahito Saito, Deguchi Hiroyuki *and* Hagiwara Makoto :** Magnetic Field-Temperature Phase Diagram of Fine YBa2Cu3O7-δ Ceramics from Linear and Non-linear Resistivities, *JPS Conference Proceedings,* **Vol.30,** 011053-1-011053-5, 2020.
1358. **HIroyuki Deguchi, Takanori Fujii, Yuki Kato, Mito Masaki, Akihiko Hisada, Kuniyuki Koyama *and* Makoto Hagiwara :** Inter-grain Phase Transitions in Supervonducting Ceramic YBa2Cu3O7-δ under Low Magnetic Fields, International Conference on Strongly Correlated Electron Systems 2019 (SCES2019), *JPS Conference Proceedings,* **Vol.30,** 011029-1-011029-6, 2020.
1359. **Takanori Satoh, Kohei Okabe, Takuma Hayashi, Ryoma Minani, Yuzuka Kimura, Ganbat Orgilbayar, Logii Narantsetseg, Janlav Munkhtsetseg, Makoto Ohashi *and* Kenji Akiyoshi :** Bacterial analysis and exploration of thermophiles in Mongolian animal dungs, *1Proceedings of 1th Asian Symposium on Microbial Ecology(ASME2019),* 95, 2019.
1360. **中野 晋, 金井 純子, 田村 隆雄, 小川 宏樹, 蒋 景彩, 村田 明広, 三上 卓, 圓谷 政貴 :** 令和元年台風19号による千曲川と越辺川の氾濫被害(速報), *令和元年自然災害フォーラム論文集,* 61-66, 2019年.
1361. **Yuta Kanemitsu, D. Chernyak, Hiroyasu Ejiri, Ken-Ichi Fushimi, Kazumi Hata, Ryuta Hazama, H. Ikeda, K. Imagawa, K. Inoue, A. Kozlov, Reiko Orito, T. Shima, Y. Takemoto, S. Umehara, K. Yasuda *and* S. Yoshida :** Purification of the NaI(Tl) crystal for dark matter search project PICOLON, *Journal of Physics: Conference Series,* **Vol.1468,** 2020.
1362. **常木 晃, 渡部 展也, 安間 了, 辰巳 祐樹, サリ ジャンモ, ラワ カリム サリ :** 肥沃な三日月地帯東部の新石器化-イラク・クルディスタン，スレマニ地域チャルモ遺跡の調査(2019年)-, *第27回西アジア発掘調査報告会報告集,* **Vol.27,** 9-14, 2020年.
1363. **A. Gando, Y. Gando, T. Hachiya, A. Hayashi, S. Hayashida, H. Ikeda, K. Inoue, K. Ishidoshiro, Y. Karino, M. Koga, S. Matsuda, T. Mitsui, K. Nakamura, S. Obara, T. Oura, H. Ozaki, I. Shimizu, Y. Shirahata, J. Shirai, A. Suzuki, T. Takai, K. Tamae, Y. Teraoka, K. Ueshima, H. Watanabe, A. Kozlov, Y. Takemoto, S. Yoshida, Ken-Ichi Fushimi, T.I. Banks, B.E. Berger, B.K. Fujimura, T. O'Donnell, L.A. Winslow, Y. Efremenko, H.J. Kanwowski, D.M. Markoff, W. Tornow, J.A. Detwiler, S. Enomoto *and* M.P. Decowski :** Precision Analysis of the 136Xe Two-Neutrino ββ Spectrum in KamLAND-Zen and Its Impact on the Quenching of Nuclear Matrix Elements, *Physical Review Letters,* **Vol.122,** 2019.
1364. **西山 賢一 :** 広島県・愛媛県の山麓斜面における斜面崩壊・土石流の発生頻度の推定, *地学雑誌,* **Vol.129,** *No.1,* N1, 2020年.
1365. **Yuhei Yamamoto :** Atomization Mechanism of Boron in Graphite Furnace Atomic Absorption Spectrometry -Improvement of analytical performance by Fe modifier and its role-, *Scientific Instrument News,* **Vol.12,** 1-8, Apr. 2019.
1366. **山本 孝 :** 規則性メソポーラスシリカの固体酸性とそのキャラクタリゼーション, *化学工業,* **Vol.70,** *No.6,* 439-446, 2019年6月.
1367. **齊藤 隆仁, 伏見 賢一, 小山 晋之, 真岸 孝一, 折戸 玲子 :** 基礎物理学実験テキスト 理工学部, 医学部医学科, 歯学部歯学科, *基礎物理学実験テキスト,* 2019年9月.
1368. **犬飼 宗弘 :** 錯体結晶の固体NMR, *Bull. Nuc. Magn. Reson. Soc. Jpn.,* **Vol.10,** 108-112, 2019年11月.
1369. **Takanori Satoh, Kohei Okabe, Takuma Hayashi, Ryoma Minani, Yuzuka Kimura, Ganbat Orgilbayar, Logii Narantsetseg, Janlav Munkhtsetseg, Makoto Ohashi *and* Kenji Akiyoshi :** Bacterial analysis and exploration of thermophiles in Mongolian animal dungs, *11th Asian Symposium on Microbial Ecology(ASME2019),* 95, Taichung,Taiwan, May 2019.
1370. **Takashi Yamamoto, Seina Heiuchi, Maki Kondoh *and* Tomoaki Irie :** Hydrothermally Synthesized Zirconium-Tungsten Binary Oxides (W/Zr = 2) : Structure and Acidic Property, *The 17th Korea-Japan Symposium on Catalysis (KJCS17),* Jeju, Korea, May 2019.
1371. **Ryo Anma, K.-C. Shin, C. Watanabe *and* A. Tuji :** pXRF analyses on Assyrian sculptures., *Japan Geoscience Union Meeting 2019 Abstracts,* HTT21-05, Makuhari, Chiba, May 2019.
1372. **Y. Yokoo, K. Asai, S. Horii, Ryo Anma, M. Shiva *and* K-C. Shin :** Influences of soil dust, sea salt and anthropogenic activities on ionic and Sr isotopic compositions of wet deposition in Iran, *Japan Geoscience Union Meeting 2019 Abstracts,* HTT18-05, Makuhari, Chiba, May 2019.
1373. **Takanori Satoh, Mitsuki Fujii, Seiji Arai, Daiki Komahara, Yuzuka Kimura, Ryoma Minani, Takuma Hayashi *and* Kenji Akiyoshi :** Analysis and Comparison of bacteria on tea leaves produced at Japan, Taiwan, and Malaysia, *KSBMB International Conference 2019,* O-9, Seogwipo,Jeju,Korea, Jun. 2019.
1374. **Takanori Satoh, Narumi Maegawa, Natsumi Yoshioka, Mayu Kanamoto, Yui Hamaji, Janlav Munkhtsetseg *and* Makoto Ohashi :** Specific binding properties of Mongolian milk whey proteins to hydrophobic molecules as lipocalin, *3rd Global Summit on Nutritional Science & Food Chemistry(Nutritional Science-2019),* 57, Osaka, Jun. 2019.
1375. **Masamichi Ogasawara :** Synthesis and Applications of Chiral Conjugated- and Cumulated-Dienes, *IMU Special Seminar,* Hohhot, Jun. 2019.
1376. **Sakai Shota, Nakao Naoki, Masaharu Ueno, Egawa Daichi, Hanzawa Hiroyuki, Kawasaki Shohei, Kumagai Keigo, Suzuki Makoto, Kobayashi Shu *and* Hanada Kentaro :** Natural ligand-nonmimetic inhibitors of the lipid-transfer protein CERT, *The 60th International Conference on the Bioscience of Lipids (ICBL2019), Tokyo, Japan,* Tokyo, Jun. 2019.
1377. **Sakai Shota, Nakao Naoki, Masaharu Ueno, Egawa Daichi, Hanzawa Hiroyuki, Kawasaki Shohei, Kumagai Keigo, Suzuki Makoto, Kobayashi Shu *and* Hanada Kentaro :** Natural Ligand-nonmimetic inhibitors of the lipid-transfer protain CERT, *The 60th International Conference on the Bioscience of Lipids(ICBL2019). 17-21 June 2019, Tokyo, Japan,* Tokyo, Jun. 2019.
1378. **Masamichi Ogasawara :** RCM-Synthesis of Highly Enantioselective Phosphine-Olefin Ligands Based on Planar-Chiral Alkenylene-Bridged Transition-Metal Complexes, *The 23rd International Symposium on Olefin Metathesis and Related Chemistry (ISOM 23),* Barcelona, Jul. 2019.
1379. **Takanori Satoh, Yousuke Endoh, Kayo Takechi, Mitsuki Fujii, Misato Tsuchie, Hiroto Itoh, Hayata Sakai, Yuiko Hanno, Tomomi Maemura, Yuto Oe, Kohta Nagai, Haruka Iwata, Yurie Ohiwa *and* Haruka Shibagaki :** Genetic analysis and thermostabilities of Nucleoside diphosphate kinases from moderately thermophilic Geobacillus bacteria, *14th international symposium of the protein society of Thailand(PST2019),* 71, Bangkok, Jul. 2019.
1380. **Takashi Yamamoto, Akihito Kurimoto *and* Riona Sato :** Conversion of C2-C4 Alcohol over Ga-ion Doped ZrO2 Solid Solutions and the Structural Characterization, *The 8th Asia-Pacific Congress in Catalysis (APCAT),* Bangkok, Aug. 2019.
1381. **Takanori Satoh, Mitsuki Fujii, Ryoma Minani, Seiji Arai, Daiki Komahara, Yuzuka Kimura, Takuma Hayashi *and* Kenji Akiyoshi :** Thermostabilities and Nucleotide binding properties of Nucleoside Diphosphate Kinases from two moderately thermophilic bacteria, *27th FAOBMB & 44th MSBMB International Conference,* Kuala Lumpur, Malaysia, Aug. 2019.
1382. **Munehiro Inukai, Takuya Kurihara, Yasuto Noda *and* Koichi Nakamura :** Dynamics of carbon dioxides in a metal-organic framework under high pressure, *EUROISMAR 2019,* Aug. 2019.
1383. **Takanori Satoh, Kohei Okabe, Takuma Hayashi, Ryoma Minani, Yuzuka Kimura, Ganbat Orgilbayar, Logii Narantsetseg, Janlav Munkhtsetseg *and* Makoto Ohashi :** Exploration And Phylogenetic Analysis For Thermophiles From Mongolian Cattle Dung, Argal, *The 15th International Congress on Thermophiles(Thermophiles2019),* Fukuoka, Sep. 2019.
1384. **Ko-ichi Magishi, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama, Namiki Takahiro, Nishimura Katsuhiko *and* Kenya Tanaka :** NMR Study of Caged Cubic Compound NdTr2Al20 (Tr = Ti and V), J-Physics 2019 International Conference & KINKEN-WAKATE 2019 Multipole Physics (J-Physics 2019), *J-Physics 2019,* Kobe, Sep. 2019.
1385. **Akihiko Hisada, Kuniyuki Koyama, Ko-ichi Magishi, Takahito Saito, Deguchi Hiroyuki *and* Hagiwara Makoto :** Magnetic Field-Temperature Phase Diagram of Fine YBa2Cu3O7-δ Ceramics from Linear and Non-linear Resistivities, International Conference on Strongly Correlated Electron Systems 2019 (SCES2019), *SCES2019,* Okayama, Sep. 2019.
1386. **Hiroyuki Deguchi, Takanori Fujii, Yuki Kato, Masaki Mito, Akihiko Hisada, Kuniyuki Koyama *and* Makoto Hagiwara :** Inter-grain Phase Transitions in Superconducting Ceramic YBa2Cu3O7-δ under Low Magnetic Fields, International Conference on Strongly Correlated Electron Systems 2019 (SCES2019), *SCES2019,* Okayama, Sep. 2019.
1387. **Ko-ichi Magishi, Yona Haruki, Akihiko Hisada, Takahito Saito, Kuniyuki Koyama *and* Isikawa Yosikazu :** NMR Study of Caged Compound RCo2Sn2Zn18 (R = La and Pr), International Conference on Strongly Correlated Electron Systems 2019 (SCES2019), *SCES2019,* Okayama, Sep. 2019.
1388. **Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Optically Active Axially Chiral Allenic Polymers, *The 14th International Conference on Cutting-Edge Organic Chemistry in Asia (ICCEOCA-14),* Niseko Town, Sep. 2019.
1389. **Takanori Satoh, Mitsuki Fujii, Haruka Iwata, Ai Hasegawa, Hajime Nonoichi, Seiji Arai, Daiki Komahara, Yuzuka Kimura, Ryoma Minani, Takuma Hayashi, Mariko Nakae *and* Kenji Akiyoshi :** Comparison of microbial diversities in non-fermented, semi-fermented, fully-fermented, and post-fermented tea leaves, *4th World Summit & Expo on Food Technology and Probiotics,* Bangkok,Thailand, Oct. 2019.
1390. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes and Their Application in Organic Transformations, *Chiral India 2019: 8th International Conference & Exhibition,* Mumbai, Dec. 2019.
1391. **Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Optically Active Axially Chiral Allenic Polymers, *The 18th Asian Chemical Congress,* Taipei, Dec. 2019.
1392. **Munehiro Inukai, Yusuke Nishiyama, Susumu Kitagawa *and* Satoshi Horike :** Glass-phase coordination polymer displaying proton conductivity and guest-accessible porosity, *1st International symposium Hydrogenomics,* Jan. 2020.
1393. **Deguchi Tomoko, Yu Kawasaki, Yutaka Kishimoto, Nakamura Koichi, Nakai Yusuke, Mito Takeshi, Haque Zeba, Gupta Chand Laxmi *and* Ganguli Kumar Ashok :** NMR Study of Layered Eu-based Bismuth-Sulfide EuFBiS2, *JPS Conf.Proc.,* **Vol.30,** 011060-1-011060-6, Mar. 2020.
1394. **Yu Kawasaki, Mori Hiro, Kishimoto Yutaka, Nakamura Koichi, He Zhangzhen *and* Itoh Mitsuru :** NMR Study of Magnetic Structure in α-CoV2O6, *JPS Conf.Proc.,* **Vol.30,** 011088-1-011088-6, Mar. 2020.
1395. **Asahi Nojirino, Masaya Aki, Yu Kawasaki, Yutaka Kishimoto, Koichi Nakamura, Yusuke Nakai, Takeshi Mito, Mitsuharu Yashima, Hidekazu Mukuda, Hisashi Kotegawa *and* Hitoshi Sugawara :** Electronic State of V3Si Probed by 29Si NMR, *JPS Conf.Proc.,* **Vol.30,** 011050-1-011050-6, Mar. 2020.
1396. **小笠原 正道 :** キラルな遷移金属錯体の触媒的不斉合成, *大阪大学大学院 基礎工学研究科 特別講演会,* 2019年5月.
1397. **山本 祐平, 豊澤 大夢, 田中 稜真, 菊地 洋一, 西本 潤, 今井 昭二 :** 四国・本州の遠隔山岳地域における湿性沈着中の重金属のモニタリング, *第79回分析化学討論会,* 2019年5月.
1398. **中村 浩一, 富本 健介, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一 :** 遷移金属酸化物における局所構造の乱れとリチウムイオン拡散, *第22回超イオン導電体物性研究会講演要旨集,* 97-102, 2019年5月.
1399. **小笠原 正道 :** キラルな遷移金属錯体の触媒的不斉合成, *大阪市立大学大学院 理学研究科 談話会,* 2019年5月.
1400. **木下 正高, 岩森 光, 西川 友章, 安間 了, S. Lagarrigue, 阿部 なつ江, 横山 由香, A. Veloso-Espinosa, I. Cortes A. Rivero, P. G. Lazcano-Prado, A. I. Arriagada-Bascunan, 原口 悟, 太田 耕輔, 折橋 裕二, 中尾 魁史, 沼田 翔伍, 塩原 肇, 杉岡 裕子, 伊藤 亜紀, N. Perez-Estay :** チリ三重会合点におけるMR18-06 `EPIC'航海Leg2調査概要., *日本地球惑星科学連合2019年大会講演要旨,* SCG60-01, 2019年5月.
1401. **横尾 頼子, 阪本 千尋, 北村 篤志, 安間 了, M. Shiva :** イラン7都市の降水中に含まれる不溶性物質の鉱物組成と元素組成の地域的特徴., *日本地球惑星科学連合2019年大会講演要旨,* MIS09-02, 2019年5月.
1402. **朽津 信明, 栁沼 由可子, 後 誠介, 西山 賢一 :** 新宮市万歳の一遍上人名号碑と江戸時代に行われたその補修について, *日本文化財科学会,* 2019年6月.
1403. **Mitsuki Fujii, Ryoma Mitani, Takuma Hayashi, Yuzuka Kimura, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Expression and thermostabilities of nucleoside diphosphate kinases from two moderately thermophiles, *The 19th Annual Meeting of the Protein Science Society of Japan,The Program and Abstracts,* 29, Jun. 2019.
1404. **西山 賢一, 鳥井 真之 :** 阿蘇カルデラ・高野尾羽根火山北斜面における地すべり移動体の内部構造(その2), *日本地すべり学会第58回研究発表講演集(CD-ROM),* 2019年8月.
1405. **伏見 賢一 :** シンチレーター原料の高純度化, *新学術「地下宇宙」領域研究会,* 2019年8月.
1406. **野尻野 旭, 安藝 将也, 川崎 祐, 岸本 豊, 中村 浩一, 水戸 毅, 八島 光晴, 椋田 秀和, 小手川 恒, 菅原 仁 :** A15型超伝導体V3SiのSi-NMRによる研究, *日本物理学会 2019年秋季大会,* 2019年9月.
1407. **出口 智子, 川崎 祐, 岸本 豊, 中村 浩一, 水戸 毅, Z. Haque, L.C. Gupta, A.K. Ganguli :** BiS2系層状化合物EuFBiS2のNMRによる研究, *日本物理学会 2019年秋季大会,* 2019年9月.
1408. **西山 賢一, 後 誠介 :** 和歌山県南部，那智川流域に分布する中位段丘堆積物の風化, *日本地質学会第126年学術大会講演要旨,* 137, 2019年9月.
1409. **川崎 祐, 岸本 豊, 中村 浩一, 水戸 毅, 磯部 正彦, 馬場 拓行, 山内 徹, 上田 寛 :** バナジウムブロンズAg2/3V2O5における金属絶縁体転移のNMRによる研究, *日本物理学会 2019年秋季大会,* 2019年9月.
1410. **土井 卓哉, 山本 孝, 奥山 勇治, 中村 崇司, 雨澤 浩史, 大石 昌嗣 :** Mn添加ペロブスカイト型酸化物A(Zr,Mn)O3-dにおけるAサイト原子がプロトン伝導特性に与える影響, *第15回固体イオニクスセミナー(滋賀),* 2019年9月.
1411. **Takanori Satoh, Kohei Okabe, Yuzuka Kimura, Ryoma Minani, Takuma Hayashi, Ganbat Orgilbayar, Logii Narantsetseg, Janlav Munkhtsetseg *and* Kenji Akiyoshi :** Bacterial phylogenetic analysis and exploration for thermophiles in Mongolian animal dungs, *the 33rd annual meeting of Japanese Society for Microbial Ecology (JSME),* Sep. 2019.
1412. **豊澤 大夢, 村瀬 遼亮, 田中 稜真, 山本 祐平, 今井 昭二 :** 金属元素比を用いた山岳地域における湿性沈着中の微量亜鉛の発生源の推定, *日本分析化学会第68年会,* 2019年9月.
1413. **久田 旭彦, 真岸 孝一, 齊藤 隆仁, 小山 晋之, 出口 博之, 萩原 亮 :** YBa2Cu3O7-δ 微細セラミクスにおけるグレイン間転移の線形・非線形抵抗測定, *日本物理学会講演概要集,* **Vol.74,** *No.2,* 2019年9月.
1414. **浦本 翔吾, 出口 博之, 美藤 正樹, 堀出 朋哉, 松本 要, 久田 旭彦, 萩原 亮 :** YBCO薄膜およびセラミクスの低磁場磁束相における磁気グラス特性, *日本物理学会講演概要集,* **Vol.74,** *No.2,* 2019年9月.
1415. **中村 光裕, 松田 和生, 中村 美沙樹, 山下 喬平, 鈴木 智子, 井上 敏 :** アデニン誘導体を硫酸基受容体としたCypridina luciferyl sulfateからCypridina luciferinへの酵素的変換, *第61回 天然有機化合物討論会,* 2019年9月.
1416. **中村 浩一, 富本 健介, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一 :** チタン酸リチウムにおける局所的な構造擾乱とリチウムイオンの運動状態, *日本物理学会2019年秋季大会 講演概要集,* 2019年9月.
1417. **今井 昭二, 宮本 征弥, 山本 祐平 :** GFAAS における種々の原子蒸気とPG表面との物理化学的相互作用, *日本分析化学会第68年会,* 2019年9月.
1418. **山本 祐平, 豊澤 大夢, 村瀬 遼亮, 田中 稜真, 高橋 玄樹, 橋本 元輝, 今井 昭二 :** 四国の山岳地域における湿性沈着中のV，Ni，Cd，Pbの発生源と輸送ルート, *日本分析化学会第68年会,* 2019年9月.
1419. **陳 彦洲, 岩本 啓汰, 宋 志毅, 中野 環, 小笠原 正道 :** ビス(2ーブロモ1,3-ジエン)をモノマーとするパラジウム触媒による軸不斉アレン・ポリマーの合成, *第66回有機金属化学討論会,* 2019年9月.
1420. **榎本 裕貴, 大路 健仁, 瓜生 瑞穂, ヤン シーチェン, 片岡 綾美, 大西 敦, 小笠原 正道 :** 多糖類を固定相に有するキラルHPLCカラムによる非鏡像関係にある異性体の分離, *第66回有機金属化学討論会,* 2019年9月.
1421. **小西 卓磨, 一尾 裕章, 小笠原 正道 :** 求核部位を有する2-ブロモ-1,3-ジエン類のパラジウム触媒分子内環化における異常な選択性, *第66回有機金属化学討論会,* 2019年9月.
1422. **佐藤 亜樹, 折橋 裕二, 中井 俊一, 新正 裕尚, ナランホ ホセ, 安間 了 :** チリ南部火山地帯中央部における島弧火山岩中のホウ素およびその他微量元素の特徴, *Geochemical Journal,* 2019年9月.
1423. **Mitsuki Fujii, Ryoma Mitani, Takuma Hayashi, Yuzuka Kimura, Seiji Arai, Daiki Komahara, Kenji Akiyoshi *and* Takanori Satoh :** Expression and Thermostabilities of nucleoside diphosphate kinases from Geobacillus and Alicyclobacillus bacteria, *The 92nd Annual Meeting of the Japanese biochemical society,The Program and Abstracts,* Sep. 2019.
1424. **Takuma Hayashi, Ryoma Mitani, Yuzuka Kimura, Mitsuki Fujii, Kenji Akiyoshi *and* Takanori Satoh :** Genetic analysis and Expression of inorganic pyrophosphatases from two moderately thermophiles, *The 92nd Annual Meeting of the Japanese biochemical society,The Program and Abstracts,* Sep. 2019.
1425. **Kenji Akiyoshi, Yuzuka Kimura, Seiji Arai, Daiki Komahara, Mitsuki Fujii, Ryoma Mitani, Takuma Hayashi *and* Takanori Satoh :** Phylogenetic analysis and expression of Thymidylate synthase genes from three moderately thermophiles, *The 92nd Annual Meeting of the Japanese biochemical society,The Program and Abstracts,* Sep. 2019.
1426. **Takanori Satoh, Kohei Okabe, Yuzuka Kimura, Ryoma Minani, Takuma Hayashi, Ganbat Orgilbayar, Logii Narantsetseg, Janlav Munkhtsetseg *and* Kenji Akiyoshi :** Microbial phylogenetic analysis and exploration for thermophiles in Mongolian animal dungs, *The 92nd Annual Meeting of the Japanese biochemical society,The Program and Abstracts,* Sep. 2019.
1427. **伏見 賢一, D. Cernyak, 江尻 宏泰, 畑 和美, 硲 隆太, 池田 晴雄, 今川 恭四郎, 井上 邦雄, 金光 佑太, Kozlov A., 折戸 玲子, 嶋 達志, 梅原 さおり, 保田 賢輔, 吉田 斉 :** PICOLON報告︓超⾼感度NaI(Tl)検出器による宇宙暗⿊物質の探索・全体計画, *日本物理学会2019年秋季大会,* 2019年9月.
1428. **金光 佑太, D. Cernyak, 江尻 宏泰, 畑 和美, 硲 隆太, 池田 晴雄, 今川 恭四郎, 井上 邦雄, 伏見 賢一, Kozlov A., 折戸 玲子, 嶋 達志, 梅原 さおり, 保田 賢輔, 吉田 斉 :** PICOLON報告︓宇宙暗⿊物質探索のためのNaI(Tl)結晶の⾼純度化, *日本物理学会2019年秋季大会,* 2019年9月.
1429. **畑 和美, D. Cernyak, 江尻 宏泰, 金光 佑太, 硲 隆太, 池田 晴雄, 今川 恭四郎, 井上 邦雄, 伏見 賢一, Kozlov A., 折戸 玲子, 嶋 達志, 梅原 さおり, 保田 賢輔, 吉田 斉 :** PICOLON報告︓宇宙暗⿊物質探索のためのNaI(Tl)シンチレータの中性⼦応答評価, *日本物理学会2019年秋季大会,* 2019年9月.
1430. **小西 卓磨, 小笠原 正道 :** 求核性部位を有する2-ブロモ-1,3-ジエン類のパラジウム触媒分子内環化における異常な選択性, *社会産業理工学研究交流会2019,* 2019年9月.
1431. **青矢 睦月 :** 眉山地域三波川帯の変塩基性岩におけるガーネット仮像の認識, *第19回社会産業理工学研究交流会パンフレット,* 15, 2019年9月.
1432. **西山 賢一, 後 誠介 :** 和歌山県南部，那智川流域に分布する熊野層群の風化, *日本応用地質学会令和元年度研究発表会講演論文集,* 71-72, 2019年10月.
1433. **鳥井 真之, 古市 剛史, 長谷中 利昭, 北園 芳人, 遠田 晋次, 西山 賢一, 奥野 充 :** 地震に伴うテフラ層の変形パターンとその成因:熊本地震と北海道胆振東部地震との比較, *国際火山噴火史情報研究集会講演要旨集2019-1,* 1-2, 2019年10月.
1434. **小西 卓磨, 一尾 裕章, 小笠原 正道 :** 求核部位を有する2-ブロモ-1,3-ジエン類のパラジウム触媒分子内環化における異常な選択性, *第30回日本プロセス学会東四国地区フォーラムセミナー,* 2019年10月.
1435. **榎本 裕貴, 大路 健仁, 瓜生 瑞穂, ヤン シーチェン, 片岡 綾美, 大西 敦, 小笠原 正道 :** 多糖類を固定相に有するキラルHPLCカラムによる非鏡像関係にある異性体の分離, *第30回日本プロセス学会東四国地区フォーラムセミナー,* 2019年10月.
1436. **西山 賢一 :** 広島県東広島市・呉市に分布する崩壊堆積物の14C年代, *日本応用地質学会中国四国支部令和元年度研究発表会講演論文集,* 61-66, 2019年11月.
1437. **犬飼 宗弘 :** 錯体結晶の固体NMR, *第58回 NMR討論会,* 2019年11月.
1438. **犬飼 宗弘, 栗原 拓也, 野田 泰斗, 中村 浩一 :** 高圧環境下における配位高分子内部の二酸化炭素の運動, *第58回 NMR討論会,* 2019年11月.
1439. **大路 健仁, 榎本 裕貴, 大西 敦, 小笠原 正道 :** 多糖類ベースのキラルHPLCカラムによる非エナンチオマー異性体混合物の分離, *2019年日本化学会中国四国支部大会,* 2019年11月.
1440. **榎本 裕貴, 瓜生 瑞穂, 浦 康之, 小笠原 正道 :** 面不斉フェロセンを基盤とする新規ホスフィン-オレフィン配位子の合成と応用, *2019年日本化学会中国四国支部大会,* 2019年11月.
1441. **小西 卓磨, 一尾 裕章, 小笠原 正道 :** ω-位に求核部位を有する3-ブロモ-1,3-アルカジエン類のパラジウム触媒環化反応における異常な選択性, *2019年日本化学会中国四国支部大会,* 2019年11月.
1442. **田上 遥, 西本 昂司, 小笠原 正道 :** ビニルフェロセン類のホモメタセシスによる1,2-ジフェロセニルエチレン類の合成, *2019年日本化学会中国四国支部大会,* 2019年11月.
1443. **西本 昂司, 田上 遥, 小笠原 正道 :** 面不斉ビニルフェロセン類のホモメタセシス速度論分割, *2019年日本化学会中国四国支部大会,* 2019年11月.
1444. **岩本 啓汰, 陳 彦洲, 宋 志毅, 中野 環, 小笠原 正道 :** ジシロキサン架橋ビス(2-ブロモ-1,3-ジエン)のパラジウム触媒反応による軸不斉アレン・ポリマーの合成, *2019年日本化学会中国四国支部大会,* 2019年11月.
1445. **谷 有香里, 澤野 恵理, 小笠原 正道 :** 嵩高い置換基を有する新規Segphos誘導体の合成と遷移金属触媒不斉反応への応用, *2019年日本化学会中国四国支部大会,* 2019年11月.
1446. **中村 光裕, 松田 和生, 中村 美沙樹, 山下 喬平, 鈴木 智子, 小笠原 正道, 井上 敏 :** アデニン誘導体のCypridina luciferyl sulfateの硫酸基受容体としての機能性, *2019年日本化学会中国四国支部大会,* 2019年11月.
1447. **藤田 真美, 西澤 高宏, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウム-メタラサイクル中間体を用いた機能性ケイ素化合物を指向した簡便な合成手法の開発, *2019年日本化学会中国四国支部大会, 徳島大学(徳島), 講演番号1F-07,* 2019年11月.
1448. **中村 滉諒, 高橋 春香, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 多置換イソベンゾフラノン合成を指向した金属ストロンチウムを用いるアセチレン骨格を二つ持つ化合物の合成手法の検討, *2019年日本化学会中国四国支部大会, 徳島大学(徳島), 講演番号1P-026,* 2019年11月.
1449. **野田 大雅, 大村 D. 聡, 上野 雅晴, 三好 德和 :** N-アリールカルバゾール誘導体におけるアルキル基の導入による立体効果に基づく発光特性の検討, *2019年日本化学会中国四国支部大会, 徳島大学(徳島), 講演番号1P-029,* 2019年11月.
1450. **中村 浩一, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一 :** Li-Ti系酸化物の局所構造変化とLi+イオンの運動状態, *第45回固体イオニクス討論会講演要旨集,* 2-3, 2019年11月.
1451. **浦本 翔吾, 出口 博之, 美藤 正樹, 堀出 朋哉, 松本 要, 久田 旭彦, 萩原 亮 :** YBCO超伝導薄膜およびセラミクスの弱磁場下磁束グラス相, *第125回日本物理学会九州支部例会,* 2019年11月.
1452. **西山 賢一 :** 東広島市・呉市に分布する崩壊堆積物の14C年代, *日本地質学会四国支部講演会要旨集,* 13, 2019年12月.
1453. **犬飼 宗弘 :** ガラス状プロトン伝導性配位高分子の合成と NMR 解析, *第66回 固体NMR・材料フォーラム,* 2019年12月.
1454. **土井 卓哉, 奥山 勇治, 山本 孝, 大石 昌嗣 :** 新規プロトン溶解メカニズムを用いた低温作動型燃料電池の開発, *第26回ヤングセラミスト・ミーティング in 中四国(香川),* K16, 2019年12月.
1455. **西山 賢一 :** 愛媛県宇和島市に分布する崩壊堆積物の14C年代, *日本応用地質学会中国四国支部令和2年度研究発表会講演論文集,* 11-12, 2020年.
1456. **中村 滉諒, 高橋 春香, 大村 D. 聡, 上野 雅晴, 三好 德和 :** 多置換イソベンゾフラノン合成を指向した金属ストロンチウムを用いるアセチレン骨格を二つ以上持つ化合物の合成手法の検討, *2019年度第3回 日本プロセス化学会東四国地区フォーラムセミナー,* 2020年1月.
1457. **野田 大雅, 大村 D. 聡, 上野 雅晴, 三好 德和 :** N-アリールカルバゾール誘導体へのアルキル基の導入による立体効果に基づく発光特性変化の検討, *2019年度第3回 日本プロセス化学会東四国地区フォーラムセミナー,* 2020年1月.
1458. **川崎 祐, 岸本 豊, 中村 浩一, 真岸 孝一, 何 長振, 伊藤 満 :** α-CoV2O6における磁気構造とスピンダイナミクス, *日本物理学会年次大会,* 2020年3月.
1459. **羽板 進也, 戸波 勇人, 藤井 勇輔, 萩原 亮, 久田 旭彦, 小山 晋之 :** YBCO系におけるCuO複鎖領域超伝導の可能性の探索 -Pr置換系との対比で見た試料合成条件-, *日本物理学会講演概要集,* **Vol.75,** *No.1,* 2020年3月.
1460. **藤井 勇輔, 森下 真次, 戸波 勇人, 萩原 亮, 久田 旭彦, 小山 晋之 :** セラミック超伝導体の中間状態に現れる特異なI-V特性 -非線型・履歴応答を解析する新手法-, *日本物理学会講演概要集,* **Vol.75,** *No.1,* 2020年3月.
1461. **久田 旭彦, 真岸 孝一, 齊藤 隆仁, 小山 晋之, 出口 博之, 萩原 亮 :** YBa2Cu4O8超微細セラミクスにおけるグレイン間転移の磁場依存性, *日本物理学会講演概要集,* **Vol.75,** *No.1,* 2020年3月.
1462. **中村 浩一, 田中 康照, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一 :** チタン酸リチウムの局所構造とLi+イオン運動におけるミリング効果, *日本物理学会第75回年次大会 講演概要集,* 2020年3月.
1463. **山本 孝, 行本 晃 :** 蛍光XAFS法における測定吸収端により異なる低濃度Eu種の平均酸化数, *日本鉄鋼協会第179回春季講演大会,* 2020年3月.
1464. **大村 D. 聡, 野田 大雅, 上野 雅晴, 三好 德和 :** ベンゼン環上がアルキル置換されたN-アリールカルバゾール誘導体の発光特性, *日本化学会第100春季年会, 東京理科大学(千葉), 公演番号1C3-57,* 2020年3月.
1465. **上野 雅晴, 大村 D. 聡, 三好 德和 :** スチルバゾール類の光異性化/環化反応制御に関する研究, *日本化学会第100春季年会, 東京理科大学(千葉), 公演番号1C4-46,* 2020年3月.
1466. **上野 雅晴, 小畠 美穂, 大村 D. 聡, 三好 德和 :** ワンポットタンデムカップリング反応を用いた Riccardin Cの全合成研究, *日本化学会第100春季年会, 東京理科大学(千葉), 公演番号3B5-10,* 2020年3月.
1467. **藤田 真美, 西澤 高宏, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウムーメタラサイクル中間体を用いた機能性含ケイ素化合物の簡便な合成手法の開発及びその置換基効果による物性評価, *日本化学会第100春季年会, 東京理科大学(千葉), 公演番号3H8-49,* 2020年3月.
1468. **西山 賢一 :** 平成30年7月豪雨で発生した土砂災害, *日本技術士会四国本部第62回CPDセミナー・公開講座・防災セミナー講演,* 2019年6月.
1469. **西山 賢一 :** 徳島平野の地下地質, *産業技術総合研究所地質調査総合センター・沿岸域プロジェクト「紀伊水道」勉強会,* 2019年10月.
1470. **西山 賢一 :** 平成30年7月豪雨で発生した土砂災害から学ぶ, *令和元年度徳島県治山林道技術研修会,* 2019年10月.
1471. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2019年10月.
1472. **西山 賢一 :** 平成30年7月豪雨で発生した土砂災害(徳島県三好・高知県大豊・広島県), *令和元年度SSLアンカー・Q&Sフレーム工法技術講習会特別講義,* 2019年11月.
1473. **犬飼 宗弘 :** ゲスト分子吸着能を示すガラス状プロトン伝導性配位高分子, *金沢固体NMRセミナー2019,* 2020年1月.
1474. **西山 賢一 :** 越高遺跡の地形・地質とその特徴, *長崎県対馬市越高遺跡調査検討委員会,* 2020年2月.
1475. **中村 浩一, 小俣 孝久 :** 多核NMR による酸化物の局所構造変化とイオン拡散挙動に関する研究, *物質・デバイス領域共同研究拠点研究成果報告書(平成31年度),* 大阪, 2020年3月.
1476. **安間 了, 常木 晃, 三宅 裕 :** イラク国北部Jarmo遺跡およびトルコ国南東部Hasankeyf遺跡出土の石器材黒曜石の化学組成と原産地推定, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2019年度,* 197-204, 2020年3月.
1477. **堀川 恵司, 南 雅代, 安間 了 :** イラン北西部アリ・サドル洞窟のつらら石のU/Th年代, 14C年代, 炭素・酸素安定同位体比, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2019年度,* 187-195, 2020年3月.
1478. **横尾 頼子, 浅井 公輔, 堀井 彩衣, 濱口 弘平, 申 キチョル, 安間 了, メラバニ シバ :** イラン8都市の月別降水の化学組成, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2019年度,* 181-186, 2020年3月.
1479. **久田 旭彦 :** くわしい科学入門 大学の物理・力学 知っているようで知らないニュートンの世界, 吉岡書店, 京都, 2021年3月.
1480. **宗林 由樹, 辻 幸一, 藤原 学, 南 秀明, 今井 昭二 :** 機器分析ハンドブック3 固体・表面分析編, 株式会社 化学同人, 2021年3月.
1481. **朽津 信明, 栁沼 由可子, 後 誠介, 西山 賢一 :** 新宮市万歳の一遍上人名号碑とその補修の歴史, *保存科学,* **Vol.59,** 23-34, 2020年.
1482. **Munehiro Inukai, T. Kurihara, Y. Noda, W. Jiang, K. Takegoshi, N. Ogiwara, H. Kitagawa *and* Koichi Nakamura :** Probing dynamics of carbon dioxide in a metal-organic framework under high pressure by high-resolution solid-state NMR, *Physical Chemistry Chemical Physics,* **Vol.22,** *No.26,* 14465-14470, 2020.
1483. **K. Tesuno, S. Ajimura, K. Akutagawa, T. Batpurev, W.M. Chan, Ken-Ichi Fushimi, Ryuta Hazama, T. Iida, Y. Ikeyama, T. Kisimoto, K.K. Lee, X. Li, K. Matsuoka, K. Matsuoka, K. Mizukoshi, Y. Mori, K. Nakajima, P. Noithong, M. Nomachi, I. Ogawa, H. Ohsumi, K. Ozawa, K. Shimizu, M. Shokati, F. Soberi, K. Suzuki, Y. Takemoto, Y. Takihira, Y. tamagawa, M. Tozawa, V.T.T. Trang, S. Umehara, K. Yamamoto, S. Yoshida, D.H. Kwon, H.L. Kim, H.J. Lee *and* Y.H. Kim :** Status of 48Ca double beta decay search and its future prospect in CANDLES, *Journal of Physics: Conference Series,* **Vol.1468,** 2020.
1484. **Ueda Jun-ichi, Enomoto Yuuki, Seki Mizuki, Konishi Takuma, Masamichi Ogasawara *and* Yoshida Kazuhiro :** Oxidative Cyclization of o-(1-Hydroxy-2-alkynyl)-N-tosylanilides for Syn-thesis of 4-Quinolones, *The Journal of Organic Chemistry,* **Vol.85,** 6420-6428, 2020.
1485. **Yuhei Yamamoto, SANAGAWA Yosuke *and* Shoji Imai :** Determination of Trace Elements in Aerosols at a Rural Mountainous Area and a Local City of Eastern Shikoku Region, Japan, *Analytical Sciences,* **Vol.36,** *No.5,* 637-643, 2020.
1486. **井澤 健一 :** 超対称性の再考 I, *徳島大学総合科学部自然科学研究,* **Vol.32,** 1-4, 2020年.
1487. **Mitsuhiro Nakamura, Kazuo Matsuda, Takashi Morikawa, Kohsuke Ishizuka *and* Satoshi Inouye :** Efficient conversion to Cypridina luciferin from Cypridina luciferyl sulfate, coupled with enzymatic sulfation of acetic acid., *Biochemical and Biophysical Research Communications,* **Vol.529,** *No.3,* 678-684, 2020.
1488. **A. Kozlov, D. Chernyak, Y. Takemoto, Ken-Ichi Fushimi, K. Imagawa, K. Yasuda, Hiroyasu Ejiri, Ryuta Hazama, H. Ikeda, K. Inoue, S. Yoshida, R.A. Etezov, Yu.M. Gavrilyuk, V.V. Kazalov, V.V. Kuzminov *and* S.I. Panasenko :** Detectors for direct dark matter search at KamLAND, *Nuclear Instruments and Methods in Physics Research Section A: Accelerators, Spectrometers, Detectors and Associated Equipment,* **Vol.958,** 2020.
1489. **H. Azizi, Y. Asahara, M. Minami *and* Ryo Anma :** Sequential magma injection with a wide range of mixing and mingling in Late Jarassic plutons, southern Ghorveh, western Iran, *Journal of Asian Earth Sciences,* **Vol.200,** 104469, 2020.
1490. **K. Suzuki, T. Iizuka, K. Kitajima, K. Hattori, T. Hirata, Ryo Anma *and* Y. Sawaki :** Hf-O isotope systematics of zircons from the Taitao granitoids: Implications for slab-melting material, *Lithos,* **Vol.372-373,** 105665, 2020.
1491. **Yuji Orihashi, Ryo Anma, S. Machida, M. Sasaki, K. Nakao, Y. Takaku *and* N. Abe :** U-Pb dating of granitic cobble (dropstone) recovered from inner slope of the Chile Trench (48°S): Constraint for its provenance, *Geochemical Journal,* **Vol.54,** *No.4,* 195-201, 2020.
1492. **Inouye Satoshi, Mitsuhiro Nakamura, Taguchi Jumpei *and* Hosoya Takamitsu :** Identification of a novel oxidation product from yellow fluorophore in the complex of apoPholasin and dehydrocoelenterazine, *Bioorganic & Medicinal Chemistry Letters,* 127435, 2020.
1493. **Norikazu Miyoshi, Shohdai Kimura, Shigeki Kubo, Satoshi D. Ohmura *and* Masaharu Ueno :** Chemoselective Ketone Synthesis by the Strontium-mediated Alkylation or Arylation of N,N-Dimethylamides or Urea, *Asian Journal of Organic Chemistry,* **Vol.9,** *No.10,* 1660-1664, 2020.
1494. **A Acharyya, R Adam, Reiko Orito *and* 496 coauthors :** Sensitivity of the Cherenkov Telescope Array to a dark matter signal from the Galactic centre, *Journal of Cosmology and Astroparticle Physics,* **Vol.2021,** *No.1,* id.57, 2021.
1495. **H. Abdalla, H. Abe, Reiko Orito *and* 499 coauthors :** Sensitivity of the Cherenkov Telescope Array for probing cosmology and fundamental physics with gamma-ray propagation, *Journal of Cosmology and Astroparticle Physics,* **Vol.2,** 048, 2021.
1496. **Hao Hu, Hiroki Ichiryu, Kiyohiko Nakajima *and* Masamichi Ogasawara :** Estimating Effective Steric and Electronic Impacts of Ferrocenyl Group in Organophosphines, *ACS Omega,* **Vol.6,** 5981-5989, 2021.
1497. **Ken-Ichi Fushimi, Yuta Kanemitsu, Shoko Hirata, D. Chernyak, Ryuta Hazama, H. Ikeda, K. Imagawa, H. Ishiura, H. Ito, T. Kisimoto, A. Kozlov, Y. Takemoto, K. Yasuda, Hiroyasu Ejiri, Kazumi Hata, T. Iida, K. Inoue, M. Koga, K Nakamura, Reiko Orito, T. Shima, S. Umehara *and* S. Yoshida :** Development of highly radiopure NaI(Tl) scintillator for PICOLON dark matter search project, *Progress of Theoretical and Experimental Physics,* **Vol.2021,** *No.4,* 043F01, 2021.
1498. **西山 賢一, 中尾 賢一, 辻野 泰之, 石田 啓佑 :** 徳島県海陽町竹ヶ島の地質と地形, *阿波学会紀要, No.63,* 87-94, 2021年.
1499. **Yusuke Urano, Hata Kazumi, Reiko Orito, Kurosawa Shunsuke, Ken-Ichi Fushimi, Konishi Satoshi, Bakr Mahmoud *and* Mukai Keisuke :** Quenching factor measurement of NaI (Tl) using monochromatic neutrons for dark matter search, *Proceedings of the 35th Workshop on Radiation Detectors and Their Uses, No.2021-3,* 53-61, 2021.
1500. **Takashi Yamamoto, Akihito Kurimoto, Riona Sato, Shoki Katata, Hirotaka Mine, Naoto Tanima *and* Ryota Kamata :** Ethanol conversion over Ga2O3-ZrO2 solid solution: empirical evidence of the reaction pathway, the surface acid-base properties, and the role of gallium ions, *Catalysis Science & Technology,* **Vol.11,** *No.6,* 2047-2056, 2021.
1501. **西山 賢一 :** 洞窟・岩陰の形成と埋積:長崎県北西部に分布する福井洞窟とその周辺, *季刊考古学,* **Vol.151,** 47-49, 2020年.
1502. **中谷 是崇, 西山 賢一, 中尾 賢一, 佐藤 善輝, 羽田 裕貴, 鈴木 克明, 水野 清秀, 中島 礼 :** 徳島市中徳島町で掘削された第四系ボーリングの記載(速報), *地質調査総合センター速報( 令和 2 年度沿岸域の地質・活断層調査研究報告), No.82,* 7-20, 2021年.
1503. **西山 賢一 :** 熊本市西部，金峰火山で発生した土砂災害の伝承, *月刊地球号外「国際火山噴火史情報研究-III」,* 114-122, 2021年.
1504. **常木 晃, 渡部 展也, 安間 了, 辰巳 祐樹, サリ ジャンモ, サーベル アハマド サーベル, ラワ カリム サリ :** 肥沃な三日月地帯東部の新石器化-イラク・クルディスタン，スレマニ地域チャルモ遺跡の調査(2019年・2020年)-, *第28回西アジア発掘調査報告会報告集,* 9-13, 2021年.
1505. **K Abe, Y Chen, K Hiraide, K Ichimura, S Imaizumi, N Kato, K Kobayashi, M Kobayashi, S Moriyama, M Nakahata, K Sato, H Sekiya, T Suzuki, A Takeda, S Tasaka, M Yamashita, B.S Yang, N.Y Kim, Y.D Kim, T.H Kim, R Ishii, Y Itow, K Kanzawa, K Martens, A Mason, Y Suzuki, K Miuchi, Y Takeuchi, K.B Lee, M.K Lee, Y Fukuda, H Ogawa, Y Kishimoto, K Nishijima, Ken-Ichi Fushimi, B.D Xu *and* S Nakamura :** Search for exotic neutrino-electron interactions using solar neutrinos in XMASS-I, *Physics Letters B,* **Vol.809,** 135741, 2020.
1506. **Akira Hirata :** X-Ray Structure of the tRNA Methyltransferase Complex of Trm7 and Trm734 from Saccharomyces cerevisiae, *Photon Factory Highlights 2019,* 48-49, Oct. 2020.
1507. **Masataka Kinoshita, Ryo Anma, Yuka Yokoyama, K. Ohta, Yusuke Yokoyama, T. Nishikawa, N. Abe, H. Iwamori *and* L. Villar :** Thermal regime around the Chile Triple Junction based on JAMSTEC MR18-06 cruise EPIC, *European Union of Geoscience Annual Meeting 2020 Abstracts,* Online, May 2020.
1508. **B.T Khai, S Ajimura, W.M Chan, Ken-Ichi Fushimi, Ryuta Hazama, H Hiraoka, T Iida, K Kanagawa, H Kino, T Kishimoto, T Maeda, K Nakajima, M Nomachi, I Ogawa, T Ohata, K Suzuki, Y Takemoto, Y Takihira, M Tozawa, M Tsuzuki, S Umehara *and* S Yoshida :** A Study on Energy Resolution of CANDLES Detector, *IEEE Transactions on Nuclear Science,* **Vol.68,** *No.3,* 368-378, Jan. 2021.
1509. **山本 祐平, 檜垣 敦哉, 今井 昭二 :** 四国の山岳地域における冬季湿性沈着中の非水溶性CdおよびPb, *第80回分析化学討論会,* 2020年5月.
1510. **村瀬 遼亮, 田中 稜真, 中田 亮一, 永石 一弥, 山本 祐平, 今井 昭二 :** 四国の山岳地域で同日採取した雪および樹氷中の高分解能MC-ICP-MSを用いた鉛同位体比, *第80回分析化学討論会,* 2020年5月.
1511. **峰 広嵩, 片田 将生, 栗本 彰人, 佐藤 里緒菜, 山本 孝 :** ガリウム添加酸化ジルコニウム触媒によるC2-C4アルコール転換反応, *第9回 JACI/GSCシンポジウム,* 2020年6月.
1512. **森野 瑛介, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 小山 晋之 :** 鉄セレン系低次元化合物の合成および電気抵抗測定, *2020年度応用物理・物理系学会中国四国支部合同学術講演会予稿集,* 86, 2020年8月.
1513. **川崎 祐, 岸本 豊, 中村 浩一, 真岸 孝一, 何 長振, 伊藤 満 :** α-CoV2O6における磁気構造とスピンダイナミクス, *日本物理学会秋季大会,* 2020年9月.
1514. **吾郷 輝夕, 川崎 祐, 真岸 孝一, 岸本 豊, 中村 浩一, 何 長振, 伊藤 満 :** 擬一次元反強磁性体SrCo2V2O8のNMR, *日本物理学会秋季大会,* 2020年9月.
1515. **金山 真也, 出口 智子, 川崎 祐, 真岸 孝一, 岸本 豊, 中村 浩一, 中井 祐介, 水戸 毅, Z. Haque, L.C. Gupta, A.K. Ganguli :** BiS2系層状化合物EuFBiS2のNMR/NQRによる研究, *日本物理学会 2020年秋季大会,* 2020年9月.
1516. **久田 旭彦, 森野 瑛介, 真岸 孝一, 齊藤 隆仁, 小山 晋之, 出口 博之, 萩原 亮 :** 超微細セラミクス YBa2Cu3O7-d および YBa2Cu4O8 におけるグレイン間転移の磁場依存性, *日本物理学会講演概要集,* **Vol.75,** *No.2,* 1334, 2020年9月.
1517. **藤井 勇輔, 森下 真次, 戸波 勇人, 萩原 亮, 出口 博之, 久田 旭彦, 小山 晋之 :** Y系247構造相に内在する複数の超伝導機構 -多段階電気抵抗異常の精密観測, *日本物理学会講演概要集,* **Vol.75,** *No.2,* 2368, 2020年9月.
1518. **富本 健介, 犬飼 宗弘, 森賀 俊広, 桑田 直明, 河村 純一, 中村 浩一 :** チタン酸ナトリウムの結晶構造と伝導挙動におけるリチウム置換とミリングの効果, *日本物理学会2020年秋季大会 講演概要集,* 2020年9月.
1519. **山本 祐平, 村瀬 遼亮, 岡 健太郎, 今井 昭二 :** 四国山岳地域で採取した降雨中に含まれるNiおよびV, *日本分析化学会第69年会,* 2020年9月.
1520. **佐藤 里緒菜, 片田 将生, 峰 広嵩, 山本 孝 :** ガリウムイオン添加ジルコニア固溶体によるエタノール転換反応の反応経路, *第126回触媒討論会,* 2020年9月.
1521. **小笠原 正道 :** パラジウム触媒による主鎖に軸不斉アレン構造を有する光学活性ポリマーの合成, *第69回高分子討論会,* 2020年9月.
1522. **中村 光裕, 松田 和生, 森川 貴史, 石塚 幸扶, 井上 敏 :** 酢酸の硫酸化反応と共役したCypridina luciferyl sulfateからCypridina luciferinへの酵素的変換, *第62回 天然有機化合物討論会,* 2020年9月.
1523. **Munehiro Inukai :** Rotation of pillar ligands of MOFs and solid-state NMR analyses, *錯体化学会第70回討論会,* Sep. 2020.
1524. **西山 賢一 :** 広島県東広島市周辺における斜面崩壊・土石流の発生頻度, *日本応用地質学会令和2年度研究発表会講演論文集,* 111-112, 2020年10月.
1525. **小笠原 正道 :** 不斉炭素を持たないキラル化合物の立体制御 ~軸不斉アレンと面不斉メタロセンの触媒的不斉合成~, *奈良女子大学大学院 化学生物環境学専攻 特別講演会,* 2020年10月.
1526. **安間 了 :** pXRFの考古学利用の現状と課題-肥沃の三日月地帯出土の黒曜石原産地推定を例として, *日本地球化学会,* 2020年11月.
1527. **野利本 剛, 空 磨奈加, 平田 章, 堀 弘幸 :** 超好熱性アーキアThermococcus kodakarensis由来tRNA m2G6 メチル化酵素Trm14の 新たな部位特異性, *第43回日本分子生物学会年会,* 2020年12月.
1528. **堀 弘幸, 福本 修平, 長谷川 貴洋, 難波 実優, 平田 章, 河村 卓哉 :** SpoU-TrmD RNAメチル化酵素スーパーファミリーで，最も長いC末領域を持つTrm56の新しい精製法, *第43回日本分子生物学会年会,* 2020年12月.
1529. **青矢 睦月 :** 高越地域三波川帯における2段階の変形の伸長方向, *第20回日本地質学会四国支部講演会要旨集,* 12, 2020年12月.
1530. **富本 健介, 犬飼 宗弘, 森賀 俊広, 河村 純一, 中村 浩一 :** Na2Ti3O7の局所構造と伝導挙動におけるミリング効果と置換効果, *第46回固体イオニクス討論会講演要旨集,* 126-127, 2020年12月.
1531. **平田 章 :** 超好熱菌由来タンパク質の構造と機能に関する研究 ~遺伝情報を伝達する仕組みから生命進化の痕跡を探る~, *第144回徳島生物学会,* 2021年1月.
1532. **浦野 雄介, 畑 和美, 折戸 玲子, 伏見 賢一, 他 :** PICOLON宇宙暗黒物質探索:単色中性子を用いたNaI(Tl)の消光因子測定, *日本物理学会年次大会,* 2021年3月.
1533. **萩原 亮, 戸波 勇人, 藤井 勇輔, 出口 博之, 久田 旭彦, 小山 晋之 :** Y系247型化合物の超伝導転移過程と還元処理効果 ―Pr247の挙動との対比―, *日本物理学会講演概要集,* **Vol.76,** *No.1,* 2645, 2021年3月.
1534. **青矢 睦月 :** 高越地域三波川帯における2段階の変形の伸長方向, *変成岩などシンポジウム2021(オンライン),* 2021年3月.
1535. **中村 浩一, 富本 健介, 犬飼 宗弘, 森賀 俊広 :** チタン酸ナトリウムの局所構造と電気伝導におけるミリング効果, *日本物理学会第76回年次大会 講演概要集,* 2021年3月.
1536. **小畠 美穂, 三好 德和, 上野 雅晴 :** ワンポットタンデムカップリング反応を用いた環境調和型Riccardin Cの全合成研究, *日本化学会第101春季年会,オンライン開催,公演番号 A19-2pm-06,* 2021年3月.
1537. **野田 大雅, 木村 将大, 久保 誠輝, 上野 雅晴, 大村 D. 聡, 三好 德和 :** 金属ストロンチウムを用いる種々のアミド化合物への半アルキル化反応の応用, *日本化学会第101春季年会,オンライン開催,公演番号 A15-1vn-09,* 2021年3月.
1538. **中村 滉諒, 高橋 春香, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いたアセチレン骨格を複数持つ化合物を構築することによる短工程多置換イソベンゾフラノン合成手法の開発, *日本化学会第101春季年会,オンライン開催,公演番号 A15-1vn-10,* 2021年3月.
1539. **上野 雅晴, 田中 巧, 大村 D. 聡, 三好 德和 :** 照射波長によるスチルバゾール類光異性化/環化反応の厳密な制御, *日本化学会第101春季年会,オンライン開催,公演番号 A17-4am-08,* 2021年3月.
1540. **中尾 直樹, 上野 雅晴, 酒井 祥太, 江川 大地, 半沢 宏之, 川崎 祥平, 熊谷 圭悟, 鈴木 誠, 小林 修, 花田 賢太郎 :** 非天然骨格を有するCERT阻害剤のSBDD研究, *日本薬学会第141年会,広島国際会議場(広島),* 2021年3月.
1541. **西山 賢一 :** 2018年西日本豪雨災害被災地に残る災害伝承, *めらんじゅ(宮崎応用地質研究会誌),* **Vol.31,** 37-40, 2020年6月.
1542. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2020年10月.
1543. **西山 賢一 :** 徳島県三好市で発生した土砂災害の歴史, *京都大学防災研究所 一般研究集会「ジオパークを活用した地域住民ーレジデント型研究者ー行政連携の防災活動のあり方」,* 13, 2020年11月.
1544. **西山 賢一 :** 2018年西日本豪雨災害被災地に残る災害伝承, *めらんじゅ(宮崎応用地質研究会誌), No.31,* 37-40, 2020年6月.
1545. **西山 賢一, 佐世保市教育委員会 :** 史跡福井洞窟整備報告書, *史跡福井洞窟整備報告書,* 111-114, 2021年3月.
1546. **中村 浩一, 小俣 孝久 :** 多核NMR による酸化物の局所構造変化とイオン拡散挙動に関する研究, *物質・デバイス領域共同研究拠点研究成果報告書(令和2年度),* 大阪, 2021年3月.
1547. **安間 了, 西山 伸一, 三宅 裕, 常木 晃, 横尾 頼子 :** 肥沃な三日月地帯北縁部に分布する新石器時代~鉄器時代遺構の堆積物柱の元素濃度, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2020年度,* 195-204, 2021年3月.
1548. **黒澤 正紀, 池端 慶, 安間 了, 西山 伸一 :** ヤシン・テペ遺跡の石室内の金属濃集堆積物の分析, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2020年度,* 205-215, 2021年3月.
1549. **橋爪 正樹, 右手 浩一, 井﨑 ゆみ子, 大野 将樹, 中野 晋, 溝渕 啓, 玉谷 純二, 下村 直行, 上田 隆雄, 上野 勝利, 長尾 文明, 橋本 親典, 武藤 裕則, 渡邉 健, 木戸 崇博, 片山 哲郎, 山下 陽子, 田端 厚之, 佐々木 由香, 島木 美香, 音井 威重, 金丸 芳, 服部 武文, 濵野 龍夫, 宮脇 克行, 安間 了, 西山 賢一, 青矢 睦月 :** 安全マニュアル, 2021年4月.
1550. **西山 賢一, 佐世保市教育委員会編 :** 福井洞窟, 2022年2月.
1551. **山本 孝 :** 金属イオン添加酸化ジルコニウム「固体表面キャラクタリゼーション 機能性材料・ナノマテリアルのためのスペクトロスコピー 第16章13節」, 株式会社 講談社, 東京, 2022年3月.
1552. **西山 賢一, 熊本大学くまもと水循環・減災研究境域センター減災型社会システム部門編 :** 熊本地震の痕跡からの学び, 国立大学法人熊本大学, 2022年3月.
1553. **小川内 良人, 横山 修, 木下 篤彦, 山田 拓, 柴田 俊, 田中 健貴, 山越 隆雄, 西山 賢一 :** 14C法(AMS 法)による深層崩壊発生年代値と深層崩壊発生頻度の推定, *砂防学会誌,* **Vol.74,** *No.1,* 3-14, 2021年.
1554. **S. Abe, S. Asami, A. Gando, Y. Gando, T. Gima, A. Goto, T. Hachiya, K. Hata, S. Hayashida, K. Hosokawa, K. Ichimura, S. Ieki, H. Ikeda, K. Inoue, K. Ishidoshiro, Y. Kamei, N. Kawada, Y. Kishimoto, T. Kinoshita, M. Koga, N. Maemura, T. Mitsui, H. Miyake, K. Nakamura, R. Nakamura, H. Ozaki, T. Sakai, H. Sambonsugi, I. Shimizu, J. Shirai, K. Shiraishi, A. Suzuki, Y. Suzuki, A. Takeuchi, K. Tamae, K. Ueshima, Y. Wada, H. Watanabe, Y. Yoshida, S. Obara, A. Kozlov, D. Chernyak, Y. Takemoto, S. Yoshida, S. Umehara, Ken-Ichi Fushimi, A.K. Ichikawa, K.Z. Nakamura, M. Yoshida, B.E. Berger, B.K. Fujikawa, J.G. Learned, J. Maricic, S.N. Axani, L.A. Winslow, Z. Fu, J. Ouellet, Y. Efremenko, H.J. Karwowski, D.M. Markoff, W. Tornow, A. Li *and* J.A. Detwiler :** Search for Low-energy Electron Antineutrinos in KamLAND Associated with Gravitational Wave Events, *The Astrophysical Journal,* **Vol.909,** 2021.
1555. **Yuki Enomoto, Hiroki Ichiryu, Hao Hu, Yasuyuki Ura *and* Masamichi Ogasawara :** C1-Symmetric Binap Derivative Featuring Single Diferrocenylphosphino-Donor Moiety, *Organometallics,* **Vol.40,** 1020-1024, 2021.
1556. **S Ajimura, W.M Chain, K Ichimura, T Ishikawa, K Kanagawa, B.T Khai, T Kishimoto, H Kino, T Maeda, K Matsuoka, N Nakatani, M Nomachi, M Saka, K Seki, Y Takemoto, Y Takihira, D Tanaka, M Tanaka, K Tetsuno, V.T.T Trang, M Tsuzuki, S Umehara, K Akutagawa, T Batpurev, M Doihara, S Katagiri, E Kinoshita, Y Hirano, T Iga, M Ishikawa, G Ito, H Kakubata, K.K Lee, X Li, K Mizukoshi, M Moser, T Ohata, M Shokati, M.S Soberi, T Uehara, W Wang, K Yamamoto, K Yasuda, S Yoshida, N Yotsunaga, T Harada, H Hiraoka, T Hiyama, A Hirota, Y Ikeyama, A Kawamura, Y Kawashima, S Maeda, K Matsuoka, K Nakajima, I Ogawa, K Ozawa, K Shamoto, K Shimizu, Y Shinki, Y Tamagawa, M Tozawa, M Yoshizawa, Ken-Ichi Fushimi, Ryuta Hazama, P Noithong, A Rittrong, K Suzuki *and* T Iida :** Low background measurement in CANDLES-III for studying the neutrinoless double beta decay of 48Ca, *Physical Review D,* **Vol.103,** 092008, 2021.
1557. **Shotaro Takano, Mao Tsuchiya, Shoji Imai, Yuhei Yamamoto, Yusuke Fukami, Katsuhiko Suzuki *and* Yoshiki Sohrin :** Isotopic analysis of nickel, copper, and zinc in various freshwater samples for source identification, *Geochemical Journal,* **Vol.55,** *No.3,* 171-183, 2021.
1558. **Kazuyuki Matsubayashi, Hidekazu Okamura, Takashi Mizokawa, Naoyuki Katayama, Akitoshi Nakano, Hiroshi Sawa, Tatsuya Kaneko, Tatsuya Toriyama, Takehisa Konishi, Yukinori Ohta, Hiroto Arima, Rina Yamanaka, Akihiko Hisada, Taku Okada, Yuka Ikemoto, Taro Moriwaki, Koji Munakata, Akiko Nakano, Minoru Nohara, Yangfan Lu, Hidenori Takagi *and* Yoshiya Uwatoko :** Hybridization-Gap Formation and Superconductivity in the Pressure-Induced Semimetallic Phase of the Excitonic Insulator Ta2NiSe5, *Journal of the Physical Society of Japan,* **Vol.90,** 074706-1-074706-6, 2021.
1559. **Y. Gando, A. Gando, T. Hachiya, S. Hayashida, K. Hosokawa, H. Ikeda, A. Hyashi, Y. Honda, S. Ieki, K. Inoue, K. Ishidoshiro, S. Ishikawa, Y. Kamei, K. Kamizawa, Y. Karino, N. Kawada, T. Kinoshita, M. Koga, S. Matsuda, H. Miyake, K. Nakamura, K. Nemoto, A. Ono, N. Ota, S. Otsuka, Y. Shibukawa, I. Shimizu, Y. Shirahata, K. Soma, A. Suzuki, A.A. Suzuki, T. Takai *and* Ken-Ichi Fushimi :** The nylon balloon for xenon loaded liquid scintillator in KamLAND-Zen 800 neutrinoless double-beta decay search experiment, *Journal of Instrumentation,* **Vol.16,** P08023, 2021.
1560. **Fatemeh Nouri, Reza Ali Davoudian, B. Mark Allen, Hossein Azizi, Yoshihiro Asahara, Ryo Anma, N. Shabanian, Motohiro Tsuboi *and* Mahnaz Khodami :** Early Cambrian highly fractionated granite, Central Iran: evidence for drifting of Northern Gondwana and the evolution of the Proto-Tethys Ocean, *Precambrian Research,* **Vol.362,** *No.362,* 106291, 2021.
1561. **Fatemeh Nouri, Hossein Azizi, Yoshihiro Asahara, A. Scott Whattam, Motohiro Tsuboi, Osman Yousif Mohammad, Masayo Minami *and* Ryo Anma :** Coexistence of two types of Late Paleocene adakitic granitoids, Soursat complex, NW Iran, *Lithos,* **Vol.404-405,** *No.106438,* 2021.
1562. **Lucia Villar-Munoz, Masataka Kinoshita, P. J. Bento, I. Vargas-Cordero, E. Contreras-Reyes, U. Tinivella, M. Giustiniani, Natsue Abe, Ryo Anma, Yuji Orihashi, Hikaru Iwamori, Tomoaki Nishikawa, A. E. Veloso *and* Satoru Haraguchi :** A cold seep triggered by a hot ridge subduction, *Scientific Reports,* **Vol.11,** *No.1,* 20923, 2021.
1563. **Lucas Segura, Itai Massad, Masamichi Ogasawara *and* Ilan Marek :** Stereodivergent Access to Trisubstituted Alkenylboronate Esters through Alkene Isomerization, *Organic Letters,* **Vol.23,** 9194-9198, 2021.
1564. **Fatemeh Nouri, Reza Ali Davoudian, Nahid Shabanian, B. Mark Allen, Yoshihiro Asahara, Hossein Azizi, Ryo Anma, Mahnaz Khodami *and* Motohiro Tsuboi :** Tectonic transition from Ediacaran continental arc to early Cambrian rift in the NE Ardakan region, central Iran: constraints from geochronology and geochemistry of magmatic rocks, *Journal of Asian Earth Sciences,* **Vol.224,** *No.105011,* 2021.
1565. **Masamichi Ogasawara :** Enantioselective Preparation of Planar-Chiral Transition Metal Complexes by Asymmetric Olefin-Metathesis Reactions in Metal Coordination Spheres, *Chemical Record,* **Vol.21,** 3509-3519, 2021.
1566. **Satoshi Inouye, Mitsuhiro Nakamura *and* Takamitsu Hosoya :** Enzymatic conversion of dehydrocoelenterazine to coelenterazine using FMN-bound flavin reductase of NAD(P)H:FMN oxidoreductase, *Biochemical and Biophysical Research Communications,* **Vol.587,** 24-28, 2022.
1567. **Satoshi Inouye, Mitsuhiro Nakamura *and* Takamitsu Hosoya :** Formation of Coelenteramine from 2-Peroxycoelenterazine in the Ca2+-Binding Photoprotein Aequorin, *Photochemistry and Photobiology,* **Vol.98,** *No.5,* 1068-1076, 2022.
1568. **S. Abe, S. Asami, A. Gando, Y. Gando, T. Gima, A. Goto, T. Hachiya, K. Hata, S. Hayashida, K. Hosokawa, K. Ichimura, S. Ieki, H. Ikeda, K. Inoue, K. Ishidoshiro, Y. Kamei, N. Kawada, Y. Kishimoto, T. Kinoshita, M. Koga, N. Maemura, T. Mitsui, H. Miyake, K. Nakamura, R. Nakamura, H. Ozaki, T. Sakai, H. Sambonsugi, I. Shimizu, J. Shirai, K. Shiraishi, A. Suzuki, Y. Suzuki, A. Takeuchi, K. Tamae, K. Ueshima, Y. Wada, H. Watanabe, Y. Yoshida, S. Obara, A. Kozlov, D. Chernyak, Y. Takemoto, S. Yoshida, S. Umehara, Ken-Ichi Fushimi, A.K. Ichikawa, K.Z. Nakamura, M. Yoshida, B.E. Berger, B.K. Fujikawa, J.G. Learned, J. Maricic, S.N. Axani, L.A. Winslow, Z. Fu, J. Ouellet, Y. Efremenko, H.J. Karwowski, D.M. Markoff, W. Tornow, A. Li *and* J.A. Detwiler :** Search for Solar Flare Neutrinos with the KamLAND Detector, *The Astrophysical Journal,* **Vol.924,** 103-6, 2022.
1569. **Y. Gando, A. Gando, T. Hachiya, S. Hayashida, K. Hosokawa, H. Ikeda, A. Hyashi, Y. Honda, S. Ieki, K. Inoue, K. Ishidoshiro, S. Ishikawa, Y. Kamei, K. Kamizawa, Y. Karino, N. Kawada, T. Kinoshita, M. Koga, S. Matsuda, H. Miyake, K. Nakamura, K. Nemoto, A. Ono, N. Ota, S. Otsuka, Y. Shibukawa, I. Shimizu, Y. Shirahata, K. Soma, A. Suzuki, A.A. Suzuki, T. Takai *and* Ken-Ichi Fushimi :** Limits on Astrophysical Antineutrinos with the KamLAND Experiment, *The Astrophysical Journal,* **Vol.925,** *No.1,* 14, 2022.
1570. **Y. Minami, Ken-Ichi Fushimi, T. Harada, A. Hashimoto, Ryuta Hazama, T. Hiraiwa, T. Iga, T. Iida, Y. Kawashima, Yumiko Kishida, T. Kishimoto, K. Matsuoka, G. Miyoshi, Y. Muramatsu, K. Nakajima, I. Ogawa, A. Rittrong, T. Sakai, Akitoshi Sakaue, K. Suzuki, Y. Takihira, Y. Tamagawa, M. Tozawa, S. Umehara, A. Yamamoto, S. Yoshida, T. Yoshida *and* A. Yoshioka :** Status of 48Ca double beta decay search in CANDLES, *Journal of Physics: Conference Series,* **Vol.2156,** 012145, 2022.
1571. **Takashi Yamamoto, Seina Heiuchi, Maki Kondo, Kei Tabusadani *and* Atsushi Sakaki :** Hydrothermally Synthesized Poorly Crystalline Binary Oxides with ZrW2O8 composition: Preparation, Structural Analysis, and Catalytic Activity for the Alkylation of Anisole with Benzyl Alcohol, *RSC Advances,* **Vol.12,** *No.6,* 3774-3782, 2022.
1572. **Kakeru Masaoka, Manami Ohkubo, Haruka Taue, Masayuki Wakioka, Yasuhiro Ohki *and* Masamichi Ogasawara :** Synthesis of Monophosphaferrocenes Revisited, *ChemistrySelect,* **Vol.7,** e202104472-(1-6), 2022.
1573. **Ohji Takehito, Ohnishi Atsushi *and* Masamichi Ogasawara :** Application of Polysaccharide-Based Chiral HPLC Columns for Separation of Regio-, E/Z-, and Enantio-isomeric Mixtures of Allylic Compounds, *ACS Omega,* **Vol.7,** 5146-5153, 2022.
1574. **M. Eizuka, K. Ishidoshiro, S. Obara, KamLAND collaboration *and* Ken-Ichi Fushimi :** Supernova Neutrino Burst Search at KamLAND, *Journal of Physics: Conference Series,* **Vol.2156,** 012195, 2022.
1575. **Ken-Ichi Fushimi, D. Chernyak, Hiroyasu Ejiri, K. Hata, Ryuta Hazama, T. Iida, H. Ikeda, K. Imagawa, K. Inoue, H. Ishiura, H. Ito, T. Kishimoto, M. Koga, Kenta Kotera, A. Kozlov, K. Nakamura, Reiko Orito, Tatsushi Shima, Y. Takemoto, S. Umehara, Yusuke Urano, Yuhei Yamamoto, K. Yasuda *and* S. Yoshida :** PICOLON dark matter search project, *Journal of Physics: Conference Series,* **Vol.2156,** 012045, 2022.
1576. **S. Obara, S. Ieki, K. Ishidoshiro, T. Mitsui, H. Watanabe, M.P. Dekowski, KamLAND collaboration *and* Ken-Ichi Fushimi :** Search for Supernova Relic Neutrinos at KamLAND, *Journal of Physics: Conference Series,* **Vol.2156,** 012138, 2022.
1577. **Y. Gando, A. Gando, T. Hachiya, S. Hayashida, K. Hosokawa, H. Ikeda, A. Hyashi, Y. Honda, S. Ieki, K. Inoue, K. Ishidoshiro, S. Ishikawa, Y. Kamei, K. Kamizawa, Y. Karino, N. Kawada, T. Kinoshita, M. Koga, S. Matsuda, H. Miyake, K. Nakamura, K. Nemoto, A. Ono, N. Ota, S. Otsuka, Y. Shibukawa, I. Shimizu, Y. Shirahata, K. Soma, A. Suzuki, A.A. Suzuki, T. Takai *and* Ken-Ichi Fushimi :** A Search for Correlated Low-energy Electron Antineutrinos in KamLAND with Gamma-Ray Bursts, *The Astrophysical Journal,* **Vol.927,** *No.1,* 69-14, 2022.
1578. **Yancheng Liu, Yuko Takagi, Milyadi Sugijanto, Kieu My Duong Nguyen, Akira Hirata, Hiroyuki Hori *and* C. Kiong Ho :** Genetic and Functional Analyses of Archaeal ATP-Dependent RNA Ligase in C/D Box sRNA Circularization and Ribosomal RNA Processing, *Frontiers in Molecular Biosciences,* **Vol.9,** *No.8,* 811548, 2022.
1579. **常木 晃, 渡部 展也, 安間 了, 板橋 悠, 宮田 佳樹, 若狭 幸, サーリ ジャンモ, サーベル アハマド サーベル :** 肥沃な三日月地帯東部の新石器化-イラク·クルディスタン，スレマニ地域チャルモ遺跡の調査 (2021年)-, *第29回西アジア発掘調査報告会報告集,* 32-37, 2022年.
1580. **三上 文三, 安達 基泰, 水谷 公彦, 高橋 延行, 姜 有那, 平田 章, 山根 愛子, 坂 直樹, 河村 広和, 候 雪妮, 森 湖太郎 :** 非凍結結晶を用いたβ-アミラーゼと基質アナログ複合体構造のpH変化の解析, *応用糖質科学,* **Vol.11,** *No.2,* 79-86, 2021年5月.
1581. **犬飼 宗弘 :** 固体NMRによるCP/MOFの静的・動的構造解析, *NMRによる有機材料分析とその試料前処理,* 639-647, 2021年9月.
1582. **Hori Hiroyuki, Akira Hirata, Ueda Takuya, Watanabe Kimitsuna, Tomikawa Chie *and* Tomita Kozo :** Transfer RNA Synthesis and Regulation, *Encyclopedia of Life Sciences,* Nov. 2021.
1583. **中村 光裕, 井上 敏 :** コエンザイムA を硫酸基受容体とした硫酸化ウミホタル ルシフェリンからウミホタルルシフェリンへの変換, *硫酸と工業, No.12,* 153-160, 2021年12月.
1584. **西山 賢一 :** 地形・地質から見た鳴門海峡の成立, *瀬戸内海,* **Vol.83,** 11-14, 2022年3月.
1585. **Urano Yusuke, Hata Kazumi, Reiko Orito, Kurosawa S, Fushimi H, Konishi S, Bakr M *and* Mukai K :** Measurement of the quenching factor in NaI(Tl) scintillator for dark matter search, *17th International Conference on Topics in Astroparticle and Underground Physics,* Aug. 2021.
1586. **Masaharu Ueno, Kazuki Tani *and* Norikazu Miyoshi :** Development of Multicomponent, One-pot Sonogashira-Suzuki Tandem Coupling Reaction in Water, *The 2021 International Chemical Congress of Pacific Basin Societies (Pacifichem), Honolulu Hawaii, USA,* Honolulu Hawaii, USA, Dec. 2021.
1587. **Kobatake Miho, Norikazu Miyoshi *and* Masaharu Ueno :** Total synthesis of Riccardin C using one-pot tandem coupling reactions, *The 2021 International Chemical Congress of Pacific Basin Societies (Pacifichem), Honolulu Hawaii, USA (Online),* Honolulu Hawaii, USA, Dec. 2021.
1588. **Haruki Sato, Koichiro Miyanishi, Makoto Negoro, Akinori Kagawa, Koichi Nakamura *and* Munehiro Inukai :** Hyperpolarized NMR for cocrystals at room temperature using photoexcited triplet-states electron spins, *The 8th International Forum on Advanced Technologies 2022,* Mar. 2022.
1589. **豊澤 大夢, 岡 健太郎, 山本 祐平, 今井 昭二 :** 徳島大学で採取した大気エアロゾル中の微量金属元素, *第81回分析化学討論会,* 2021年5月.
1590. **小川内良人・横山 修・木下篤彦・山田 拓・柴田 俊・田中健貴・山越隆雄, 西山 賢一 :** 放射性炭素年代から推定される紀伊山地の深層崩壊発生頻度, *砂防学会研究発表会講演集,* 2021年6月.
1591. **朽津信明・白石明香・藤 隆宏・後 誠介・栁沼由可子, 西山 賢一 :** 新宮市・九重土砂災害慰霊碑の三次元印刷, *日本文化財科学会第38回大会研究発表会,* 2021年6月.
1592. **峰 広嵩, 片田 将生, 山本 孝 :** 種々の酸化ジルコニウム担持銅触媒によるエタノールからの直接酢酸エチル合成, *第12回触媒科学研究発表会,* 2021年6月.
1593. **野利本 剛, 空磨奈 伽, 平田 章, 堀 弘幸 :** 超好熱性アーキア Thermococcus kodakarensis 由来 tRNA メチル化酵素 Trm14 の機能解明 について, *第33回日本Archaea研究会,* 2021年7月.
1594. **横川 隆志, 能村 友一朗, 安田 旭宏, 尾木野 弘実, 日浦 恵太, 仲田 沙織, 岡 夏央, 安藤 香織, 河村 卓哉, 平田 章, 堀 弘幸, 大野 敏 :** アーキア特異的修飾ヌクレオシド， アーケオシンの生合成, *第33回日本Archaea研究会講演会,* 2021年7月.
1595. **中谷 是崇, 西山 賢一, 中尾 賢一, 佐藤 善輝, 羽田 裕貴, 鈴木 克明, 水野 清秀, 中島 礼 :** 徳島市中徳島町で掘削された第四系ボーリングの検討(速報), *日本第四紀学会講演要旨集,* 2021年8月.
1596. **西山 賢一, 後 誠介 :** 1946年昭和南海地震で発生した和歌山県田辺市本宮町の岩盤崩壊, *日本地質学会第128年度学術大会講演要旨,* 2021年9月.
1597. **秋山 大介, 川崎 祐, 岸本 豊, 真岸 孝一, 中村 浩一, 西野 克志, 何 長振, 伊藤 満 :** スピンギャップを持つ擬一次元系交代鎖BaCu2V2O8のNMR, *日本物理学会秋季大会,* 2021年9月.
1598. **佐藤 亜樹, 折橋 裕二, 中井 俊一, 新正 裕尚, A. Jose Naranjo, 高久 雄一, 淺原 良浩, 安間 了 :** アンデス南部火山地帯, 第四紀島弧火山岩類のマグマ成因:ホウ素を含む全岩化学組成の特徴からの制約, *日本地球化学会第68年会,* 2021年9月.
1599. **ウォリス サイモン, 青矢 睦月 :** 沈み込み境界プロセスを議論するためにどの蛇紋岩体が適切なのか?: 西南日本，三波川帯の例, *日本地質学会第128年学術大会予稿集,* T3-O-13, 2021年9月.
1600. **胡 淏, 一柳 浩輝, 中島 清彦, 小笠原 正道 :** 有機ホスフィン類におけるフェロセニル基の立体的/電子的影響の定量化, *第67回有機金属化学討論会,* 2021年9月.
1601. **榎本 裕貴, 一柳 浩輝, 胡 淏, 浦 康之, 小笠原 正道 :** 単一のジフェロセニルホスフィノ基を有するC1対称Binap誘導体の合成と応用, *第67回有機金属化学討論会,* 2021年9月.
1602. **田上 遥, 西本 昂司, 大路 健仁, 大木 靖弘, 小笠原 正道 :** 面不斉ビニルフェロセン類の立体選択的メタセシス不斉二量化反応, *第67回有機金属化学討論会,* 2021年9月.
1603. **峰 広嵩, 片田 将生, 山本 孝 :** 担持銅触媒によるエタノールからの直接酢酸エチル合成における担体ジルコニア結晶相の影響, *第128回触媒討論会,* 2021年9月.
1604. **中村 浩一, 富本 健介, 犬飼 宗弘, 森賀 俊広 :** チタン酸ナトリウムの局所構造とイオン運動におけるミリング効果, *日本物理学会2021年秋季大会 講演概要集,* 2021年9月.
1605. **萩原 亮, 戸波 勇人, 藤井 勇輔, 出口 博之, 久田 旭彦, 小山 晋之, 谷口 晴香, 松川 倫明 :** citrate-pyrolysis前駆体法を用いた単一相 R247系(R:Y or Pr)の合成手法, *日本物理学会講演概要集,* **Vol.76,** *No.2,* 2873, 2021年9月.
1606. **山本 祐平, 村瀬 遼亮, 中田 亮一, 永石 一弥, 今井 昭二 :** 遠隔山岳地域における湿性沈着中の鉛同位体比測定に基づく大気中鉛の発生源解析, *日本分析化学会第70年会,* 2021年9月.
1607. **西山 賢一, 後 誠介, 朽津 信明, 藤 隆宏, 白石 明香 :** 和歌山県新宮市九重地区で発生した土砂災害とその伝承, *日本応用地質学会令和3年度研究発表会講演論文集,* 2021年10月.
1608. **西山 賢一, 木村 一成, 加藤 弘徳 :** 中国四国地方における自然災害伝承費の分布と特徴, *日本応用地質学会中国四国支部,* 13-16, 2021年10月.
1609. **山崎 新太郎, 荒井 紀之, 西山 賢一, 矢野 真一郎, 丸谷 靖幸 :** 球磨地域の土砂災害2, *日本応用地質学会令和3年度研究発表会講演論文集,* 2021年10月.
1610. **森野 瑛介, 久田 旭彦, 髙木 拓海, 藤原 直樹 :** 組み込み式ガスケットを用いた対向アンビル型高圧装置の開発, *第59回高圧討論会講演要旨集,* **Vol.31,** 134, 2021年10月.
1611. **西山 賢一 :** 和歌山県新宮市に伝わる江戸時代の土砂災害慰霊碑の発見と活用, *防災学術連携体・特別シンポジウム 「防災教育と災害伝承への多様な視点 -東日本大震災から10年を経て-」,* 2021年11月.
1612. **山本 祐平 :** 山岳地域における湿性沈着中の鉛同位体比測定に基づく大気中鉛の発生源解析, *2021年日本化学会中国四国支部大会,* 2021年11月.
1613. **大路 健仁, 岡崎 願之晋, 劉 強, 安江 里紗, 吉田 和弘, 小笠原 正道 :** 面不斉フェロセン縮環ホスホン酸のエナンチオ選択的合成法の開発と応用, *2021年日本化学会中国四国支部大会,* 2021年11月.
1614. **小笠原 正道 :** キラルな遷移金属錯体の触媒的不斉合成, *東京都立大学大学院理学研究科 特別講演会(化学コロキウム),* 2021年11月.
1615. **野利本 剛, 空 磨奈伽, 平田 章, 堀 弘幸 :** 超好熱性アーキアThermococcus kodakarensis由来tRNA m2G6 メチル化酵素Trm14の 幅広い基質特異性, *第44回日本分子生物学会年会,* 2021年12月.
1616. **安間 了 :** 屋久島の成立とフィリッピン海プレートの移動, *屋久島学ソサエティー,* 2021年12月.
1617. **田中 康照, 犬飼 宗弘, 森賀 俊広, 河村 純一, 中村 浩一 :** チタン酸リチウムのミリングにともなう電気伝導挙動の変化, *第47回固体イオニクス討論会講演要旨集,* 156-157, 2021年12月.
1618. **朝倉 一恵, 横尾 頼子, 阪本 千尋, 北村 篤志, 金澤 康平, 安間 了, Shiva Mehrabani, 申 キチョル :** イランの降水に含まれる不溶性物質のPb同位体比にみられる地域的特徴, *第11回同位体環境学シンポジウム,* 2021年12月.
1619. **安間 了 :** 屋久島の中身はこうなっていた!花崗岩と堆積岩の地史, *屋久島大学,* 2021年12月.
1620. **山本 孝 :** ラボXAFS装置による固体触媒・環境試料の評価, *X線分析研究懇談会第276回例会,* 2022年1月.
1621. **平田 章 :** アーキアRNAスプライシング酵素の構造と機能に関する研究~タンパク質とRNAの共進化の仕組み~, *第145回徳島生物学会,* 2022年1月.
1622. **川崎 祐, 澤元 浩, 岸本 豊, 真岸 孝一, 松本 信洋, 永田 正一 :** スピネル型硫化物Cu(Ir1-xRhx)2S4の63Cu-NMR, *日本物理学会年次大会,* 2022年3月.
1623. **青矢 睦月, 小泉 比架留, 市川 裕一朗 :** 補正EDSデータによる眉山青色片岩の角閃石組成累帯構造:沈み込みP-T pathの屈曲, *変成岩などシンポジウム2022(オンライン),* 2022年3月.
1624. **森野 瑛介, 久田 旭彦, 髙木 拓海, 藤原 直樹 :** 物性測定用組み込みガスケット式対向アンビル高圧装置の開発, *日本物理学会講演概要集,* **Vol.77,** *No.1,* 2559, 2022年3月.
1625. **杉浦 輝, 真岸 孝一, 久田 旭彦, 川崎 祐, 雷 前坤, 松本 悠輝, 並木 孝洋, 西村 克彦 :** TmTr2Al20 (Tr = V, Ti) の NMR, *日本物理学会講演概要集,* **Vol.77,** *No.1,* 16pPSH96, 2022年3月.
1626. **小畠 美穂, 三好 德和, 上野 雅晴 :** Riccardin Cの全合成をモデルとした効率的なワンポットタンデムカップリング反応の研究-ユニット合成の簡便さを考慮した適切な炭素-炭素切断位置及び組み合わせる官能基の設計-, *日本化学会第102春季年会,オンライン開催,公演番号 K5-2am-04,* 2022年3月.
1627. **三好 德和 :** 高大連携事業を通じた理系人材育成, *日本化学会春季年会予稿集,* **Vol.2022,** B101-3, 2022年3月.
1628. **佐藤 晴紀, 宮西 孝一郎, 根来 誠, 香川 晃徳, 中村 浩一, 犬飼 宗弘 :** 光励起三重項状態の電子スピンを用いた混晶の室温核スピン超偏極, *日本化学会第102回春季年会,* 2022年3月.
1629. **安部 聖竜, 佐藤 晴紀, 宮西 孝一郎, 根来 誠, 香川 晃徳, 中村 浩一, 犬飼 宗弘 :** 室温核スピン超偏極を可能とする共晶の構造解析, *日本化学会第102回春季年会,* 2022年3月.
1630. **三好 亜季, 宮崎 泰彰, 久保 誠輝, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いる新しい形のシリルエーテル化反応の開発, *日本化学会第102春季年会,オンライン開催,公演番号 D202-4pm-07,* 2022年3月.
1631. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2021年10月.
1632. **西山 賢一, 山崎 新太郎 :** 洪水・斜面災害が発生した熊本県球磨・芦北・人吉地域の地形・地質学的背景, *令和2年7月九州豪雨災害調査団報告書,* 23-32, 2021年6月.
1633. **山崎 新太郎, 荒井 紀之, 西山 賢一, 矢野 真一郎, 丸谷 靖幸 :** 2020年熊本県南部豪雨災害による岩盤崩壊の特徴について, *令和2年7月九州豪雨災害調査団報告書,* 33-40, 2021年6月.
1634. **安間 了, Jaafar Jotheri :** 堆積物柱状試料と地形からみるメソポタミア下流域の堆積環境の変遷と塩害, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2021年度,* 154-171, 2022年3月.
1635. **下岡 順直, 安間 了, Jaafar Jotheri, 中川 清隆, 長島 秀樹, 平田 英隆 :** Tell Umm al-Aqaribの三日月型砂丘のルミネッセンス年代測定と風向風速解析, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2021年度,* 173-185, 2022年3月.
1636. **黒澤 正紀, 池端 慶, 安間 了, 西山 伸一 :** ヤシン・テペ遺跡出土の腐食青銅片の鉱物・化学的特徴, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2021年度,* 187-195, 2022年3月.
1637. **西山 賢一, 日本応用地質学会災害地質研究部会編 :** 土砂災害の疑問55, 成山堂書店, 東京, 2022年6月.
1638. **齋藤 有 :** ハイパーピクナイト, 朝倉書店, 2022年7月.
1639. **山本 孝 :** X線吸収端近傍構造(XANES)「触媒総合事典 第4章16節」, 朝倉書店, 東京, 2023年1月.
1640. **髙田 知紀, 西山 賢一, 薮内 佳順, 宇野 宏司, 瀬谷 今日子, 砂川 佳子 :** 地域防災と時間性, ユニオンプレス, 2023年2月.
1641. **Yu Nishida, Shiho Ohmori, Risa Kakizono, Kunpei Kawai, Miyu Namba, Kazuki Okada, Ryota Yamagami, Akira Hirata *and* Hiroyuki Hori :** Required Elements in tRNA for Methylation by the Eukaryotic tRNA (Guanine- N2-) Methyltransferase (Trm11-Trm112 Complex), *International Journal of Molecular Sciences,* **Vol.23,** *No.7,* 2022.
1642. **Takashi Yamamoto, Ayumi Yamashita, Nozomi Yokoyama, Akira Yukumoto *and* Shoji Imai :** Analysis of Leachate Copper from Sediment at a Small Abandoned Mine Site by Handy-type Liquid Electrode Plasma Atomic Emission Spectrometer, *ISIJ International,* **Vol.62,** *No.5,* 854-859, 2022.
1643. **Norikazu Miyoshi, Aki Miyoshi, Yasuaki Miyazaki, Shigeki Kubo *and* Masaharu Ueno :** Practical Method for Hydroxyl-Group Protection Using Strontium Metal and Readily Available Silyl Chlorides, *Chemical Communications,* **Vol.58,** *No.43,* 6312-6315, 2022.
1644. **Etsuo Uchida, Shinya Nagano, Sota Niki, Kou Yonezu, Yu Saitoh *and* Ki-Cheol Shin :** Geochemical and radiogenic isotopic signatures of granitic rocks in Chanthaburi and Chachoengsao provinces, southeastern Thailand: Implications for origin and evolution, *Journal of Asian Earth Sciences: X,* **Vol.8,** *No.100111,* 2022.
1645. **Takuya Kurihara, Munehiro Inukai *and* Motohiro Mizuno :** Slow CO2 Diffusion Governed by Steric Hindrance of Rotatory Ligands in Small Pores of a Metal-Organic Framework., *The Journal of Physical Chemistry Letters,* **Vol.13,** *No.30,* 7023-7028, 2022.
1646. **伊藤 博信, 長谷川 修一, 山中 稔, 西山 賢一, 石田 啓祐 :** 四国の高速道路切土法面における酸性土の分布と成因, *応用地質,* **Vol.63,** *No.3,* 96-111, 2022年.
1647. **Mutsuki Aoya :** Overprint of secondary Du folding in the Sambagawa metamorphic belt, SW Japan: implications for strain ellipsoids and Paleogene tectonics of the east-Eurasian margin, *The Island Arc,* **Vol.31,** DOI:10.1111/-iar.12463, 2022.
1648. **Koji Nishimoto, Haruka Taue, Takehito Ohji, Sayaka Funakoshi, Yasuhiro Ohki *and* Masamichi Ogasawara :** Diastereo- and Enantioselective Metathesis Dimerization/Kinetic Resolution of Racemic Planar-Chiral Vinylferrocenes, *Organic Letters,* **Vol.24,** 7355-7360, 2022.
1649. **Keigo Kumagai, Shota Sakai, Masaharu Ueno, Michiyo Kataoka, Shu Kobayashi *and* Kentaro Hanada :** Chlamydial Infection-Dependent Synthesis of Sphingomyelin as a Novel Anti-Chlamydial Target of Ceramide Mimetic Compounds, *International Journal of Molecular Sciences,* **Vol.23,** 14697, 2022.
1650. **Nishimura Madoka, Nakaya Yukari, Kashimoto Mao, Shoji Imai, Hideji TANAKA *and* Masaki Takeuchi :** Determination of trace perchlorate in river water by ion chromatography with online matrix removal and sample concentration, *Analytical Sciences,* 2023.
1651. **Yuhei Yamamoto, Kentaroh Oka, Tokoro Shunichi, Naomichi Nishii, Kikuchi Yoichi, Jun Nishimoto *and* Shoji Imai :** Investigation of the concentration ratios of anthropogenic metal elements in fresh snow at mountain area as a tracer for the discrimination between short and longrange transport contributions, *Analytical Sciences,* **Vol.39,** *No.1,* 679-687, 2023.
1652. **Miho Kobatake, Norikazu Miyoshi *and* Masaharu Ueno :** One-pot Tandem Coupling Method for the Short Step Formal Synthesis of Riccardin C, *Chemistry - A European Journal,* **Vol.29,** e202203805, 2023.
1653. **西山 賢一, 中尾 賢一, 辻野 泰之, 元山 茂樹, 石田 啓祐 :** 小松島市の地質と地形, *阿波学会紀要, No.64,* 1-10, 2023年.
1654. **A. Tsuneki, N. Watanabe, Ryo Anma, S. Jammo, Yu Saitoh *and* S. Ahmad :** Preliminary report of the Charmo (Jarmo) Prehistoric investigations, *Al -Rafidan,* **Vol.44,** 1-34, 2023.
1655. **常木 晃, 渡部 展也, 安間 了, サーリ ジャンモ, サーベル アハマド サーベル :** 肥沃な三日月地帯東部の新石器化-イラク·クルディスタン，スレマニ地域チャルモ遺跡の調査 (2022年), *第30回西アジア発掘調査報告会報告集,* 30-35, 2023年.
1656. **S. Abe, S Asami, M Eizuka, S Futagi, A Gando, Y Gando, T Gima, A Goto, T Hachiya, K. Hata, S Hayashida, K Hosokawa, K Ichimura, S Ieki, H Ikeda, K Inoue, K Ishidoshiro, Y. Kamei, N Kawada, Y Kishimoto, M Koga, M Kurasawa, N Maemura, T Mitsui, H Miyake, T Nakahata, K Nakamura, K Nakamura, R Nakamura, H Ozaki, T Sakai, H Sambonsugi, I Shimizu, J Shirai, K Shiraishi, A Suzuki, Y Suzuki, A Takeuchi, K Tamae, K Ueshima, H Watanabe, Y Yoshida, S Obara, AK Ichikawa, D Chernyak, A Kozlov, KZ Nakamura, S Yoshida, Y Takemoto, S Umehara, Ken-Ichi Fushimi, Kenta Kotera, Y Urano, BE Berger, BK Fujikawa, JG Learned, J Maricic, SN Axani, J Smolsky, Z Fu, LA Winslow, Y Efremenko, HJ Karwowski, DM Markoff, W Tornow, S DellOro, T ODonnell, JA Detwiler, S Enomoto, MP Decowski, C Grant, A Li *and* H Song :** Search for the Majorana nature of neutrinos in the inverted mass ordering region with KamLAND-Zen, *Physical Review Letters,* **Vol.130,** *No.5,* 2023.
1657. **伏見 賢一, Dmitry Chernyak, 江尻 宏泰, 畑 和実, 硲 隆太, T. Iida, H. Ikeda, K. Imagawa, K. Inoue, H. Ito, T. Kishimoto, M. Koga, 小寺 健太, A. Kozlov, K. Nakamura, 折戸 玲子, T. Shima, Y. Takemoto, S. Umehara, Y. Urano, 山本 祐平, K. Yasuda, S. Yoshida :** 高純度ヨウ化ナトリウムの結晶育成, *日本結晶成長学会誌,* **Vol.49,** *No.4,* 2023年1月.
1658. **西山 賢一 :** 徳島市の地形・地質, *地図中心, No.605,* 6-8, 2023年2月.
1659. **Yuji Orihashi, Aki Sato, Shunichi Nakai, Hironao Shinjoe, Jose Naranjo, Ryo Anma *and* Group Project CFRISTMASSY :** Magmagenesis of the Quaternary arc volcanic rocks, the Southern Volcanic Zone of the Andes: Constraints from characteristics of the bulk chemistry including boron contents, *XII South American Symposium on Isotope Geology,* Santiago, Jun. 2022.
1660. **Masamichi Ogasawara :** Diastereo- and Enantioselective Metathesis Dimerization/Kinetic Resolution of Racemic Planar-Chiral Vinylferrocenes, *29th International Conference on Organometallic Chemistry (ICOMC-29),* Praha, Jul. 2022.
1661. **KAKERU Masaoka, Haruka Taue *and* Masamichi Ogasawara :** Enantioselective Synthesis of Planar-Chiral Ferrocenes by Molybdenum-Catalyzed Metathesis Dimerization/Desymmetrization of Divinylferrocenes, *29th International Conference on Organometallic Chemistry (ICOMC-29),* Praha, Jul. 2022.
1662. **Takashi Yamamoto, Hirotaka Mine *and* Shoki Katata :** Effects of crystalline ZrO2 phase on direct ethyl acetate synthesis from ethanol over supported copper catalysts, *12th International Conference on Environmental Catalysis,* Suita, Aug. 2022.
1663. **Akihiko Hisada, Ko-ichi Magishi, Fujiwara Naoki, Uwatoko Yoshiya, Deng Guochu, Pomjakushina Ekaterina, Conder Kazimierz, Radheep Mohan Dinadhayalane, Thiyagarajan Raman, Esakkimuthu Sankaran *and* Arumugam Sonachalam :** NMR study under pressure on highly Ca-doped spin-ladder compound Sr1.4Ca12.6Cu24O41, *JPS Conf. Proc.,* **Vol.38,** 011048-1-011048-5, Sapporo, Aug. 2022.
1664. **Rina Fukuchi, Yoshiro Nishio, Rie Hori, Ryo Anma, Yuhji Yamamoto, Arito Sakaguchi, Lena Maeda, Saneatsu Saito *and* Kae Takahashi :** Potential of online webcasts of large Scientific Ocean Drilling Programs for University Education: a case study IODP Expedition 386 the R/V Kaimei, *IX GeoSciEd,* Shimane, Aug. 2022.
1665. **Masamichi Ogasawara :** Palladium-Catalyzed Asymmetric Synthesis of Axially Chiral Allenes, *Burapha University Special Seminar,* Chonburi, Feb. 2023.
1666. **Masamichi Ogasawara :** Palladium-Catalyzed Asymmetric Synthesis of Axially Chiral Allenes, *Chulalongkorn University Special Seminar,* Bangkok, Feb. 2023.
1667. **Yu Kawasaki, Hiro Mori, Yutaka Kishimoto, Ko-ichi Magishi, Koichi Nakamura, Zhangzhen He *and* Mitsuru Itoh :** NMR Study of Magnetic Structure and Spin Dynamics in α-CoV2O6, *JPS Conf.Proc.,* **Vol.38,** 011118-1-011118-6, Mar. 2023.
1668. **Ko-ichi Magishi, Mori Hiro, Sugiura Hikaru, Akihiko Hisada, Yu Kawasaki, Lei Qiankun, Matsumoto Yuki, Namiki Takahiro *and* Nishimura Katsuhiko :** NMR Study of Caged Compounds TmTr2Al20 (Tr = Ti, V), *JPS Conf.Proc.,* **Vol.38,** 011107-1-011107-6, Mar. 2023.
1669. **Masamichi Ogasawara, Haruka Taue, Masaoka Kakeru, Wakioka Masayuki *and* Ohki Yasuhiro :** Enantio- and Diastereoselective Metathesis Dimerization/Kinetic Resolution of Racemic Planar-Chiral Vinylphosphaferrocenes, *The 9th International Forum on Advanced Technologies (IFAT 2023),* Taipei, Mar. 2023.
1670. **Masaoka Kakeru, Taue Haruka, Higaki Tatsuya, Ohki Yasuhiro *and* Masamichi Ogasawara :** Enantioselective Metathesis Dimerization/Desymmetrization of Cs-Symmetric Divinylferrocenes, *The 9th International Forum on Advanced Technologies (IFAT 2023),* Taipei, Mar. 2023.
1671. **Mori Yuuki, Hitomi Tsukasa *and* Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Alkoxyallenes and Application in Addition Reactions, *The 9th International Forum on Advanced Technologies (IFAT 2023),* Taipei, Mar. 2023.
1672. **Masaharu Ueno :** Methods of Incorporating Catalysts into Microchannels Based on Structure and Function and Their Application to Organic Synthesis, *2023 International Workshop on Microfluidic Chemical Plant International University-Industry Collaboration for Innovation, March 13-14 2023, Hsinchu, Taiwan, Keynote Lecture,* Mar. 2023.
1673. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes, *National Taiwan Normal University Special Seminar,* Taipei, Mar. 2023.
1674. **西井 直道, 山本 祐平, 豊澤 大夢, 岡 健太郎, 所 竣一, 西本 潤, 菊地 洋一, 今井 昭二 :** 2020年度冬期に四国山岳, 瀬戸内沿岸, 広島県北部, 岩手県山岳で 採取した降雪, 降雨及び樹氷中の非水溶性物質の酸分解―ICP-MS 及びGFAAS法による無機微量元素の研究, *第82回分析化学討論会,* 2022年5月.
1675. **所 俊一, 山本 祐平, 中田 亮一, 村瀬 遼亮, 田中 稜真, 今井 昭二 :** ノビアスキレートPA-1固相抽出-二重収束型多重検知ICP-MS法 を用いた四国・中国地方において同時採取した夏季降雨中 の鉛同位体比による発生源の識別, *第82回分析化学討論会,* 2022年5月.
1676. **安間 了 :** 西アジア出土石材に関する分析とその手法, *日本文化の地質学的特質 研究発表会,* 2022年6月.
1677. **犬飼 宗弘 :** 共晶・混晶のトリプレットDNP, *DNP研究会,* 2022年7月.
1678. **Kawai Kumpei, Nishida Yu, Ohomori Shiho, Kakizono Risa, Namba Miyu, Okada Kazuki, Yamagami Ryota, 平田 章, Hori Hiroyuki :** Required Elements in tRNA for Methylation by the Eukaryotic tRNA (Guanine-N2-) Methyltransferase (Trm11-Trm112 Complex), *第23回日本RNA学会年会,* 2022年7月.
1679. **Liu Yancheng, Takagi Yuko, Sugijianto Milyadi, Nguyen Doung My Ken, 平田 章, Hori Hiroyuki, Ho Kiong :** Archaeal ATP-Dependent RNA Ligase Plays a Role in C/D Box sRNA Circularization and Ribosomal RNA Processing, *第23回日本RNA学会年会,* 2022年7月.
1680. **西山 賢一 :** 地盤情報データベースを用いた徳島県鳴門市周辺の地下地質, *日本地質学会第129年学術大会講演要旨,* 2022年9月.
1681. **政岡 翔, 田上 遥, 小笠原 正道 :** Cs-対称ジビニルフェロセン類の不斉メタセシス不斉二量化/非対称化反応, *第68回有機金属化学討論会,* 2022年9月.
1682. **浦野 雄介, 畑 和美, 折戸 玲子, 伏見 賢一, 他 :** PICOLON宇宙暗黒物質探索:単色中性子を用いたNaI(Tl)の消光因子測定2, *日本物理学会年次大会,* 2022年9月.
1683. **小笠原 正道, 田上 遥, 政岡 翔, 脇岡 正幸, 大木 靖弘 :** 面不斉ビニルホスファフェロセン類の立体選択的メタセシス不斉二量化反応, *第68回有機金属化学討論会,* 2022年9月.
1684. **西井 直道, 山本 祐平, 菊地 洋一, 今井 昭二 :** 降雪及び樹氷の融解水中の溶存態と懸濁態の微量元素, *日本分析化学年会第71回,* 2022年9月.
1685. **伏見 賢一, 小寺 健太, 折戸 玲子, 浦野 雄介, Chrnyak Dmitry, 江尻 宏泰, 岸本 忠史, 嶋 達志, 梅原 さおり, 吉田 斉, 畑 和実, 池田 晴雄, 井上 邦雄, 古賀 真之, 黒澤 俊介, 硲 隆太, 飯田 崇史, 今川 恭四郎, 保田 賢輔, 伊藤 博士, Alexandre Kozlov, 中村 健悟, 竹本 康浩 :** 超高純度無機シンチレーターによる宇宙・素粒子稀現象の研究, *公益社団法人 日本セラミックス協会 第35回秋季シンポジウム(徳島市),* **Vol.1T08,** 2022年9月.
1686. **山本 孝, 栗本 彰人 :** ガリウムイオン添加酸化ジルコニウム固溶体触媒のXRD/XAFS分析, *日本分析化学会第71年会,* 2022年9月.
1687. **中村 浩一, 田中 康照, 富本 健介, 犬飼 宗弘, 森賀 俊広 :** M-Ti 酸化物(M=Na, Li)の局所構造変化と電気伝導挙動, *日本物理学会2022年秋季大会 講演概要集,* 2022年9月.
1688. **杉浦 輝, 真岸 孝一, 久田 旭彦, 川崎 祐, 雷 前坤, 松本 悠輝, 並木 孝洋, 西村 克彦 :** TmTr2Al20 (Tr = V, Ti) の NMR Ⅱ, *日本物理学会講演概要集,* **Vol.77,** *No.2,* 2296, 2022年9月.
1689. **藤井 勇輔, 飯田 幹也, 萩原 亮, 出口 博之, 久田 旭彦, 松川 倫明 :** R2Ba4Cu7O15(R: Pr or Y)を合成するためのcitrate pyrolysis前駆体の生成手法, *日本物理学会講演概要集,* **Vol.77,** *No.2,* 2236, 2022年9月.
1690. **川口 皓大, 三明 瞭太, 藤井 孝徳, 出口 博之, 美藤 正樹, 萩原 亮, 久田 旭彦 :** YBCO(Y247 系)超伝導セラミクスにおけるグレイン間相転移, *日本物理学会講演概要集,* **Vol.77,** *No.2,* 2235, 2022年9月.
1691. **今井 昭二, 繁田 大地, 長崎 昌太, 上井 優佑, 奥田 桂子, 中尾 佑子, 山本 祐平 :** 四国の渓流水における非海塩性硫酸イオンの分布, *日本分析化学年会第71回,* 2022年9月.
1692. **山本 孝, 髻谷 圭, 片岡 卓哉, 塀内 聖奈 :** 低結晶性ZrW2O8の構造およびアルキル化活性, *第130回触媒討論会,* 2022年9月.
1693. **吉田 愛菜, 中村 光裕, 田井 章博 :** 高感度かつハイスループットなアスコルビン酸定量法, *日本農芸化学会2022年度中四国支部大会(第63回講演会),* 2022年9月.
1694. **西山 賢一, 木村 一成, 加藤 弘徳, 野々村 敦子 :** 中国四国地方に分布する自然災害伝承碑とその特徴, *日本応用地質学会令和4年度研究発表会講演論文集,* 139-140, 2022年10月.
1695. **小笠原 正道 :** 不斉炭素を持たないキラル化合物の立体制御 ~軸不斉アレンと面不斉メタロセンの触媒的不斉合成~, *熊本大学大学院 先端科学研究部 基礎科学部門 特別講演会,* 2022年10月.
1696. **西山 賢一, 露口 耕治, 大矢 基弘, 佐竹 一希, 川本 真由美 :** 愛媛県宇和島市に分布する崩壊堆積物に含まれる古土壌の14C年代, *日本応用地質学会中国四国支部令和4年度研究発表会発表論文集,* 35-38, 2022年11月.
1697. **佐藤 晴紀, 宮西 孝一郎, 根来 誠, 香川 晃徳, 西山 祐介, 中村 浩一, 犬飼 宗弘 :** 共晶法による生体分子のTriplet DNP, *第61回NMR討論会,* 2022年11月.
1698. **金子 宗平, 栗原 拓也, 水野 元博, 中村 浩一, 犬飼 宗弘 :** 多孔性配位高分子における二酸化炭素と水の 動的挙動と競争吸着, *第61回NMR討論会,* 2022年11月.
1699. **山本 孝, 峰 広嵩 :** ジルコニア担持銅触媒のXRD/XAFS分析およびエタノール転換反応活性, *第58回X線分析討論会,* 2022年11月.
1700. **三好 亜季, 谷 和紀, 三好 德和, 上野 雅晴 :** ワンポットタンデムカップリング反応によるポリアリール化合物の環境調和型効率的合成, *2022年日本化学会中国四国支部大会,広島大学(広島),講演番号2K-10,* 2022年11月.
1701. **川上 夕, 三好 德和, 上野 雅晴 :** セラミド輸送阻害剤HPA-12類縁体の効率的合成, *2022年日本化学会中国四国支部大会,広島大学(広島),講演番号2PB-10,* 2022年11月.
1702. **木村 恒輝, 三好 德和, 上野 雅晴 :** ビスマス塩を触媒とするアルデヒドの環境調和型酸化反応の開発, *2022年日本化学会中国四国支部大会,広島大学(広島),講演番号2PB-11,* 2022年11月.
1703. **河合 薫平, 野利本 剛, 空 磨奈伽, 山上 龍太, 平田 章, 堀 弘幸 :** 超好熱性アーキアThermococcus kodakarensisにおけるtRNAメチル化酵素Trm14の性質の解明, *第45回日本分子生物学会年会,* 2022年12月.
1704. **小笠原 正道 :** パラジウム触媒による軸不斉アレンの触媒的不斉合成 ~天然物アレンの不斉全合成への応用と不斉重合による光学活性アレンポリマー合成~, *第48回高分子講座,* 2022年12月.
1705. **田中 康照, 中村 浩一, 犬飼 宗弘, 森賀 俊広 :** チタン酸リチウムの電気伝導度における酸素欠損およびミリング効果, *第48回固体イオニクス討論会講演要旨集,* 196-197, 2022年12月.
1706. **髙木 拓海, 廣田 翔也, 久田 旭彦, 真岸 孝一, 藤原 直樹 :** 組み込み式ガスケットを用いた対向アンビル型高圧装置の開発Ⅱ, *第62回高圧討論会講演要旨集,* **Vol.32,** 69, 2022年12月.
1707. **小笠原 正道 :** キラルな遷移金属錯体の触媒的不斉合成, *京都大学化学研究所 附属元素科学国際研究センター,* 2023年1月.
1708. **澤元 浩, 川崎 祐, 岸本 豊, 真岸 孝一, 松本 信洋, 永田 正一 :** スピネル型硫化物Cu(Ir1-xRhx)2S4の63Cu-NMR II, *日本物理学会秋季大会,* 2023年3月.
1709. **青矢 睦月 :** 三波川帯における二次的なDu褶曲のオーバープリント: 歪み解析の解釈, *変成岩などシンポジウム(つくば),* 2023年3月.
1710. **金木 俊也, 纐纈 佑衣, 青矢 睦月, 中村 佳博, ウォリス サイモン :** Kouketsu et al. (2014) と Aoya et al. (2010) の炭質物ラマン温度計の自動化, *変成岩などシンポジウム(つくば),* 2023年3月.
1711. **齋藤 有 :** 泥の起源と行方:黒潮は砕屑粒子を遠洋域まで運べるのか?, *日本堆積学会20周年記念リレーセミナー,* **Vol.0,** *No.0,* 0, 2023年3月.
1712. **堀江 將仁, 澤野 恵理, 谷 友香里, 檜垣 達也, 大木 靖弘, 小笠原 正道 :** 新規不斉配位子の開発を基軸とするパラジウム触媒反応による軸不斉天然物アレンの不斉全合成, *日本化学会第103春季年会,* 2023年3月.
1713. **三好 亜季, 三好 德和, 上野 雅晴 :** 温度制御によるタンデムカップリング反応を用いたワンポット・ポリアリール化合物の環境調和型効率的合成, *日本化学会第103春季年会,東京理科大学(千葉),公演番号 P4-2am-33, 2023年3月.,* 2023年3月.
1714. **政岡 翔, 田上 遥, 檜垣 達也, 大木 靖弘, 小笠原 正道 :** Cs対称ジビニルフェロセン類のモリブデン触媒メタセシス二量化/不斉非対称化反応, *日本化学会第103春季年会,* 2023年3月.
1715. **渡邊 達哉, 大村 D. 聡, 上野 雅晴, 三好 德和 :** ストロンチウムを用いるアミド化合物の還元反応, *日本化学会第103春季年会,東京理科大学(千葉),公演番号 K404-4pm-06, 2023年3月.,* 2023年3月.
1716. **中村 浩一, 田中 康照, 犬飼 宗弘, 森賀 俊広 :** チタン酸リチウムの局所構造とリチウムイオン運動における酸素欠損およびミリングの効果, *2023年春季大会プログラム 講演概要集,* 2023年3月.
1717. **山本 孝 :** 固体表面の酸・塩基点とその触媒機能, *近畿化学協会触媒・表面部会 入門触媒科学セミナー,* 2022年10月.
1718. **西山 賢一, 鳥井 真之 :** 2016年熊本地震で地すべりが発生した阿蘇カルデラ・高野尾羽根火山北斜面における変形構造, *京都大学防災研究所「著しい変形を受けた地質地域の斜面災害評価に関する研究集会」,* 2022年11月.
1719. **西山 賢一 :** 1976年台風17号豪雨による土砂災害と雨量との関係, *第2回高知県・徳島県地すべり対策研究会,* 2023年1月.
1720. **犬飼 宗弘 :** 共晶を利用した生体分子の室温DNP, *金沢固体NMRセミナー2022,* 2023年1月.
1721. **西山 賢一 :** 石屋洞穴周辺の地形・地質的特徴, *石屋洞穴,長崎県埋蔵文化財センター調査報告書,* 59-63, 2023年3月.
1722. **安間 了, 佐野 貴司, 申 基澈, 昆 慶明, 松井 恵子 :** 携帯式蛍光X線分析用の標準岩石試料の整備, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2022年度,* 187-197, 2023年3月.
1723. **安間 了, 申 基澈, 渡辺 千香子, 辻 彰洋, 佐野 貴司, 齋藤 有, 中野 孝教, 横尾 頼子, 小泉 龍人, Mark Altaweel, Anke Marsh, Hashim Hama, Kamal Rasheed, Jaafar Jotheri :** スレマニ博物館所蔵の楔形文書粘土板胎土の化学組成, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2022年度,* 147-152, 2023年3月.
1724. **横尾 頼子, 堀井 彩衣, 濱口 弘平, 前田 哲弥, メラバニ シバ, 安間 了 :** イラン7都市の月別降水の主要イオン組成, *都市文明の本質:古代西アジアにおける都市の発生と変容の学際研究 研究成果報告2022年度,* 163-168, 2023年3月.
1725. **今井 昭二 :** 原子吸光分析法, 丸善出版, 2024年1月.
1726. **西山 賢一, 地学団体研究会編 :** 最新 地学事典, 平凡社, 2024年3月.
1727. **S Abe, A Aguasca-Cabot, Reiko Orito *and* 267 coauthors :** Multiwavelength study of the galactic PeVatron candidate LHAASO J2108+5157, *Astronomy & Astrophysics,* **Vol.673,** id.A75, 2023.
1728. **Satoshi Inouye, Kazuo Matsuda *and* Mitsuhiro Nakamura :** Enzymatic sulfation of coelenterazine by human cytosolic aryl sulfotransferase SULT1A1: identification of coelenterazine C2-benzyl monosulfate by LC/ESI-TOF-MS, *Biochemical and Biophysical Research Communications,* **Vol.665,** 133-140, 2023.
1729. **Haruki Sato, Koichiro Miyanishi, Makoto Negoro, Akinori Kagawa, Yusuke Nishiyama, Satoshi Horike, Koichi Nakamura *and* Munehiro Inukai :** Hyperpolarization of Biomolecules in Eutectic Crystals at Room Temperature Using Photoexcited Electrons, *The Journal of Physical Chemistry Letters,* **Vol.14,** *No.19,* 4560-4564, 2023.
1730. **Ko-ichi Magishi, Hikaru Sugiura, Akihiko Hisada, Yu Kawasaki, Qiankun Lei, Yuki Matsumoto, Takahiro Namiki *and* Katsuhiko Nishimura :** NMR Study of Caged Compounds TmTr2Al20 (Tr = Ti, V), *JPS Conference Proceedings,* **Vol.38,** 011107-1-011107-6, 2023.
1731. **Yu Kawasaki, Hiro Mori, Yutaka Kishimoto, Ko-ichi Magishi, Koichi Nakamura, Zhangzhen He *and* Mitsuru Itoh :** NMR Study of Magnetic Structure and Spin Dynamids in a-CoV2O6, *JPS Conference Proceedings,* **Vol.38,** 011118-1-011118-6, 2023.
1732. **Akihiko Hisada, Ko-ichi Magishi, Naoki Fujiwara, Yoshiya Uwatoko, Guochu Deng, Ekaterina Pomjakushina, Kazimierz Conder, Radheep Mohan Dinadhayalane, Raman Thiyagarajan, Sankaran Esakkimuthu *and* Sonachalam Arumugam :** NMR Study under pressure on highly Ca-doped spin-ladder compound Sr1.4Ca12.6Cu24O41, *JPS Conference Proceedings,* **Vol.38,** 011048-1-011048-6, 2023.
1733. **KAKERU Masaoka, Haruka Taue, Masayuki Wakioka, Yasuhiro Ohki *and* Masamichi Ogasawara :** Asymmetric Metathesis Dimerization/Kinetic Resolution of Racemic Planar-Chiral Vinylphosphaferrocenes, *Organometallics,* **Vol.42,** 1629-1638, 2023.
1734. **H. Azizi, N. Daneshvar, Y. Asahara, M. Minami *and* Ryo Anma :** The generation of Eocene mafic dike swarms during the exhumation of a core complex, Biarjmand area, NE Iran, *Journal of Petrology,* **Vol.64,** 1-18, 2023.
1735. **KAKERU Masaoka, Haruka Taue, Tatsuya Higaki, Yasuhiro Ohki *and* Masamichi Ogasawara :** Competitive Double Friedel-Crafts 2,5- and 1',2-Diacylation of Monophosphaferrocenes, *Organometallics,* **Vol.42,** 1667-1673, 2023.
1736. **Takashi Yamamoto, Hirotaka Mine, Shoki Katata *and* Taketo Tone :** Direct ethyl acetate synthesis from ethanol over amorphous-, monoclinic-, tetragonal ZrO2 supported copper catalysts prepared from the same zirconium source, *Applied Catalysis B: Environmental,* **Vol.327,** *No.15,* 122433, 2023.
1737. **F Acero, A Acharyya, Reiko Orito *and* 356 coauthors :** Sensitivity of the Cherenkov Telescope Array to spectral signatures of hadronic PeVatrons with application to Galactic Supernova Remnants, *Astroparticle Physics, No.150,* id.102850, 2023.
1738. **A Acharyya, R Adam, Reiko Orito *and* 333 coauthors :** Sensitivity of the Cherenkov Telescope Array to TeV photon emission from the Large Magellanic Cloud, *Monthly Notices of the Royal Astronomical Society,* **Vol.523,** *No.4,* 5353-5387, 2023.
1739. **Kazuhiro W. Makabe *and* Swalla W Makabe Hannah I Jensen Alexander C A Fodor Jennifer Hsu William R Jeffery Noriyuki Satoh Billie J Kazuhiro :** Cymric, a Maternal and Zygotic HTK-16-Like SHARK Family Tyrosine Kinase Gene, Is Disrupted in Molgula occulta, a Tailless Ascidian, *The Biological Bulletin,* **Vol.245,** *No.1,* 1-8, 2023.
1740. **Razieh Chaharlang, Jalil Ghalamghash, Yu Saitoh, Mihai N. Ducea *and* Axel K. Schmitt :** Sr Nd isotopes of Sabalan Volcano, NW Iran: insights into the origin of collisional adakites and geodynamic implications, *International Journal of Earth Sciences,* **Vol.112,** 2065-2080, 2023.
1741. **Ken-Ichi Fushimi, K. Imagawa, Y. Kishida, S. Kurosawa, Reiko Orito, A. Sakaue, Y. Takihira, S. Umehara, A. Yamaji, Yuhei Yamamoto, K. Yasuda *and* T. Yoshida :** Purification of CaF2 crystal for double beta decay experiments, *AIP Conference Proceedings,* **Vol.2908,** *No.1,* 2023.
1742. **J. C. Lissenberg, J. D. Condon, J. A. Smye *and* Ryo Anma :** Crystallization of superfast-spreading oceanic crust (ODP 1256D, Pacific Ocean): Constraints from zircon geochronology, *Geochemistry, Geophysics, Geosystems,* **Vol.24,** *No.10,* e2023GC010964, 2023.
1743. **H. Abe, K. Abe, Reiko Orito *and* 283 coauthors :** Observations of the Crab Nebula and Pulsar with the Large-sized Telescope Prototype of the Cherenkov Telescope Array, *The Astrophysical Journal,* **Vol.956,** *No.2 80,* 25, 2023.
1744. **K Abe, K Hiraide, N Kato, S Moriyama, M Nakahata, K Sato, H Sekiya, T Suzuki, Y Suzuki, A Takeda, BS Yang, Nam Young Kim, YD Kim, Yong Hamb Kim, Y Itow, K Martens, A Mason, M Yamashita, K Miuchi, Y Takeuchi, KB Lee, MK Lee, Y Fukuda, H Ogawa, K Ichimura, Y Kishimoto, K Nishijima, Ken-Ichi Fushimi, BD Xu, K Kobayashi *and* S Nakamura :** Direct dark matter searches with the full data set of XMASS-I, *Physical Review D,* **Vol.108,** *No.8,* 2023.
1745. **K. Abe, S. Abe, Reiko Orito *and* 273 coauthors :** Star tracking for pointing determination of Imaging Atmospheric Cherenkov Telescopes. Application to the Large-Sized Telescope of the Cherenkov Telescope Array, *Astronomy & Astrophysics,* **Vol.679,** 90, 2023.
1746. **H. Azizi, N. Daneshvar, G. Rafat, Y. Asahara, K. Horie, M. Takehara, Y. Kon, M. Minami *and* Ryo Anma :** O-Hf isotope ratios of Alvand S-type granite, western Iran, reveal crustal melting in an extensional regime, *Lithos,* **Vol.464-465,** *No.107437,* 2023.
1747. **H. Abe, K. Abe, Reiko Orito *and* 366 coauthors :** Performance of the joint LST-1 and MAGIC observations evaluated with Crab Nebula data, *Astronomy & Astrophysics,* **Vol.680,** 66, 2023.
1748. **K. Sakai, K. Fuke, K. Yoshimura, M. Sasaki, K. Abe, Reiko Orito *and* 13 coauthors :** Search for Antideuterons of Cosmic Origin Using the BESS-Polar II Magnetic-Rigidity Spectrometer, *Physical Review Letters,* **Vol.132,** *No.13,* 131001, 2023.
1749. **Yoh Kohno, Asako Ito, Aya Okamoto, Ryota Yamagami, Akira Hirata *and* Hiroyuki Hori :** Escherichia coli tRNA (Gm18) methyltransferase (TrmH) requires the correct localization of its methylation site (G18) in the D-loop for efficient methylation., *The Journal of Biochemistry,* **Vol.175,** *No.1,* 43-56, 2024.
1750. **Ko-ichi Magishi, Hikaru Sugiura, Akihiko Hisada, Yu Kawasaki *and* Kuwai Tomohiko :** NMR study of Si-substitution effect for Al in PrTi2Al20, *Interactions,* **Vol.245,** 11-1-11-8, 2024.
1751. **Akinobu NAKADA, Takayuki YOKOTA, Morihiko OGURA, Ninghui CHANG, Osamu TOMITA, Akinori SAEKI, Masamichi Ogasawara *and* Ryu ABE :** Metal-Free Carbazole-Thiophene Photosensitizers Designed for a Dye-Sensitized H2-Evolving Photocatalyst in Z-Scheme Water Splitting, *The Journal of Chemical Physics,* **Vol.160,** 044710, 2024.
1752. **Haruna Imazu, Kakeru Masaoka, Saki Uike *and* Masamichi Ogasawara :** Molybdenum-Catalyzed Enantioselective Ring-Closing Metathesis/Kinetic Resolution of Racemic Planar-Chiral 1,1'-Diallylferrocenes, *Catalysts,* **Vol.14,** 123, 2024.
1753. **Urano Yusuke, Ken-Ichi Fushimi, Hata Kazumi, Y. Kawai, S. Konishi, S. Kurosawa, K. Mukai, Y. Ogino, Reiko Orito *and* T. Sakabe :** Quenching factor measurement of low-energy Na recoils in ultra-pure NaI(Tl) crystal, *Proceedings of Science,* **Vol.441,** 087, 2024.
1754. **A. Tsuneki, A. S. Saber, N. Watanabe, Ryo Anma, S. Jammo, M. Makino, Y. Miyauchi, O. K. Lorentz, Y. Itahashi, M. Yoneda, M. Kurosawa *and* K. Ikehata :** Preliminary report of the Charmo (Jarmo) Prehistoric investigations, 2023, *Al -Rafidan,* **Vol.45,** 1-47, 2024.
1755. **羽田 裕貴, 中谷 是崇, 小松原 琢, 阿部 朋弥, 水野 清秀, 中島 礼, 西山 賢一, 中尾 賢一 :** 徳島県鳴門市里浦町の海岸に分布する大阪層群相当層の堆積年代, *地質調査総合センター速報,* **Vol.84,** 9-20, 2023年.
1756. **常木 晃, 渡部 展也, 安間 了, サーリ ジャンモ, 牧野 真理子, 宮内 優子, サーベル アハマド サーベル :** 肥沃な三日月地帯東部の新石器化-イラク·クルディスタン，スレマニ地域チャルモ遺跡の調査 (2023年), *第31回西アジア発掘調査報告会報告集,* 34-39, 2024年.
1757. **S. Abe, S Asami, M Eizuka, S Futagi, A Gando, Y Gando, T Gima, A Goto, T Hachiya, K. Hata, S Hayashida, K Hosokawa, K Ichimura, S Ieki, H Ikeda, K Inoue, K Ishidoshiro, Y. Kamei, N Kawada, Y Kishimoto, M Koga, M Kurasawa, N Maemura, T Mitsui, H Miyake, T Nakahata, K Nakamura, K Nakamura, R Nakamura, H Ozaki, T Sakai, H Sambonsugi, I Shimizu, J Shirai, K Shiraishi, A Suzuki, Y Suzuki, A Takeuchi, K Tamae, K Ueshima, H Watanabe, Y Yoshida, S Obara, AK Ichikawa, D Chernyak, A Kozlov, KZ Nakamura, S Yoshida, Y Takemoto, S Umehara, Ken-Ichi Fushimi, Kenta Kotera, Y Urano, BE Berger, BK Fujikawa, JG Learned, J Maricic, SN Axani, J Smolsky, Z Fu, LA Winslow, Y Efremenko, HJ Karwowski, DM Markoff, W Tornow, S DellOro, T ODonnell, JA Detwiler, S Enomoto, MP Decowski, C Grant, A Li *and* H Song :** First measurement of the strange axial coupling constant using neutral-current quasielastic interactions of atmospheric neutrinos at KamLAND, *Physical Review D,* **Vol.107,** *No.7,* 072006, 2023.
1758. **S. Abe, S Asami, M Eizuka, S Futagi, A Gando, Y Gando, T Gima, A Goto, T Hachiya, K. Hata, S Hayashida, K Hosokawa, K Ichimura, S Ieki, H Ikeda, K Inoue, K Ishidoshiro, Y. Kamei, N Kawada, Y Kishimoto, M Koga, M Kurasawa, N Maemura, T Mitsui, H Miyake, T Nakahata, K Nakamura, K Nakamura, R Nakamura, H Ozaki, T Sakai, H Sambonsugi, I Shimizu, J Shirai, K Shiraishi, A Suzuki, Y Suzuki, A Takeuchi, K Tamae, K Ueshima, H Watanabe, Y Yoshida, S Obara, AK Ichikawa, D Chernyak, A Kozlov, KZ Nakamura, S Yoshida, Y Takemoto, S Umehara, Ken-Ichi Fushimi, Kenta Kotera, Y Urano, BE Berger, BK Fujikawa, JG Learned, J Maricic, SN Axani, J Smolsky, Z Fu, LA Winslow, Y Efremenko, HJ Karwowski, DM Markoff, W Tornow, S DellOro, T ODonnell, JA Detwiler, S Enomoto, MP Decowski, C Grant, A Li *and* H Song :** Measurement of cosmic-ray muon spallation products in a xenon-loaded liquid scintillator with KamLAND, *Physical Review C,* **Vol.107,** *No.5,* 054612, 2023.
1759. **青矢 睦月 :** 瀬場谷岩体, *最新 地学事典 地学団体研究会編,* 2024年3月.
1760. **Takashi Yamamoto, Kei Tabusadani, Emina Kawaguchi *and* Takuya Kataoka :** Calcination temperature dependency of poorly crystallized ZrW2O8 on acidic property, *The 19th Korea-Japan Symposium on Catalysis,* Seoul, May 2023.
1761. **Ryo Anma, Jaafar Jotheri, Sachi Wakasa, Yorinao Shitaoka, Kicheol Shin, Takashi Sano *and* Yoriko Yokoo :** Late Quaternary climate changes and sedimentation in the Tigris-Euphrates watersheds: implication in the changing watercourses and early cities, *Symposium: Cities and Urbanization in West Asai and Egupt Shapes, Functions, and Ideology-,* Tsukuba, Jun. 2023.
1762. **Yuhei Yamamoto, Tokoro Shun-ichi, Murase Ryosuke, Nakada Ryoichi, Nagaishi Kazuya *and* Shoji Imai :** Application of MC-ICP-MS with solid extraction method using chelate column for trace lead isotopes in wet depositions, *Colloquium Spectroscopicum Internationale XLIII,* Jun. 2023.
1763. **Takashi Yamamoto *and* Kazunori Miyamoto :** XAFS study of thermal decomposition behaviour of platinum salt on catalyst support by laboratory-type spectrometer, *Colloquium Spectroscopicum Internationale XLIII & The 5th Asian Symposium on Laser Induced Breakdown Spectroscopy (CSI XLIII / ASLIBS2023),* Tokushima, Jun. 2023.
1764. **Masamichi Ogasawara :** Enantioselective Preparation of Planar-Chiral Ferrocenes: Asymmetric Metathesis Dimerization of Vinylferrocenes, *24th International Symposium on Olefin Metathesis and Related Chemistry (ISOM-24),* Bergen, Jul. 2023.
1765. **Munehiro Inukai, Sato Haruki, Miyanishi Koichiro, Negoro Makoto, Kagawa Akinori, Nishiyama Yusuke *and* Koichi Nakamura :** Hyperpolarization of biomolecules in eutectic crystals at room temperature using photo-excited electron, *EUROISMAR 2023,* Jul. 2023.
1766. **Munehiro Inukai, Sato Haruki, Miyanishi Koichiro, Negoro Makoto, Kagawa Akinori, Nishiyama Yusuke *and* Koichi Nakamura :** Hyperpolarization of biomolecules in eutectic crystals at room temperature using photo-excited electro, *IUCr 2023,* Aug. 2023.
1767. **Urano Yusuke, Hata Kazumi, Reiko Orito, S. Kurosawa, K. Fushimi, S. Konishi, K. Mukai, Y. Ogino, T. Sakabe *and* Y. Kawai :** Quenching factor measurement of low-energy Na recoils in ultra-pure NaI(Tl) crystal, *18th International Conference on Topics in Astroparticle and Underground Physics,* Aug. 2023.
1768. **Masamichi Ogasawara :** Palladium-Catalyzed Asymmetric Synthesis of Axially Chiral Allenes and Applications in Total Synthesis of Allenic Natural Products, *University of Ghana Special Seminar,* Accra, Oct. 2023.
1769. **Masamichi Ogasawara :** Palladium-Catalyzed Asymmetric Synthesis of Axially Chiral Allenes and Applications in Total Synthesis of Allenic Natural Products, *University for Development Studies Special Seminar,* Tamale, Oct. 2023.
1770. **Masamichi Ogasawara *and* Panida Surawatanawong :** Palladium-Catalyzed Synthesis of Allenes: Experimental Observations and Theoretical Rationale, *The Fifth Asian Workshop of Experiment and Theory in Quantum Beam Molecular Sciences (SSWIU2023),* Mito, Oct. 2023.
1771. **Hori Hiroyuki, Yamagami Ryota, Ishida Kazuo, Takuma Hiroyuki, Kusuba Hiroaki, Akira Hirata, Ochi Anna, Iwashita Chikako *and* Tomikawa Chie :** Regulatory Factors for tRNA modifications in Thermus thermophilus, *International Workshop on Neotechnologies for ThermusQ initiative,* Oct. 2023.
1772. **Ko-ichi Magishi, Sugiura Hikaru, Akihiko Hisada, Yu Kawasaki *and* Kuwai Tomohiko :** NMR study of Si-substitution effect for Al inPrTi2Al20, International Conference on Hyperfine Interactions and their Applications (HYPERFINE2023), *HYPERFINE2023,* Nara, Nov. 2023.
1773. **Masaoka Kakeru, Taue Haruka, Higaki Tatsuya, Ohki Yasuhiro *and* Masamichi Ogasawara :** Competitive Double Friedel-Crafts 2,5- and 1',2-Diacylation of Monophosphaferrocenes, *The 15th International Kyoto Conference on New Aspects of Organic Chemistry (IKCOC-15),* Kyoto, Nov. 2023.
1774. **MORI Yuki, Hitomi Tsukasa *and* Masamichi Ogasawara :** Palladium-Catalyzed Synthesis of Alkoxyallenes and Their Application in Regioselective Hydroelementation Reactions, *The 15th International Kyoto Conference on New Aspects of Organic Chemistry (IKCOC-15),* Kyoto, Nov. 2023.
1775. **HIBIKI Okabe *and* Masamichi Ogasawara :** Design, Synthesis, and Application of Planar-Chiral Skewed DPPF Derivatives, *The 15th International Kyoto Conference on New Aspects of Organic Chemistry (IKCOC-15),* Kyoto, Nov. 2023.
1776. **Masamichi Ogasawara :** Enantioselective Synthesis of Planar-Chiral Ferrocene Derivatives by Asymmetric Metathesis Dimerization of Vinylferrocenes, *The 15th International Kyoto Conference on New Aspects of Organic Chemistry (IKCOC-15),* Kyoto, Nov. 2023.
1777. **Masamichi Ogasawara :** Asymmetric Synthesis of Planar-Chiral Ferrocenes by Enantioselective Metathesis Dimerization of Vinylferrocenes, *The 16th International Conference on Cutting-Edge Organic Chemistry in Asia (ICCEOCA-16),* Singapore, Dec. 2023.
1778. **Masamichi Ogasawara :** Asymmetric Metathesis Dimerization of Vinylferrocenes: New Strategy for Preparing Enantiomerically Enriched Planar-Chiral Ferrocenes, *International Symposium on Catalysis and Fine Chemicals 2023 (C&FC2023),* Tokyo, Dec. 2023.
1779. **Masamichi Ogasawara :** Catalytic Enantioselective Synthesis of Planar-Chiral Transition-Metal Complexes, *Mahidol University Graduate Seminar in Chemistry,* Bangkok, Feb. 2024.
1780. **Yukinojo Koyama, Simon Wallis, Takayoshi Nagaya *and* Mutsuki Aoya :** Stress distribution in a subduction channel associated with rapid subduction: evidence from quartz piezometry applied to the Sanbagawa subduction-type metamorphic belt, *Abstract for Japan Geoscience Union (JpGU) meeting 2023,* **Vol.SCG45,** 31, May 2023.
1781. **松田 哲平, 山上 龍太, 鈴木 建夫, 平田 章, 堀 弘幸 :** tRNAの6位を2'-O-メチル化する新規 tRNAメチル化酵素の発見, *第64回日本生化学会・中国四国支部例会,* 2023年5月.
1782. **藤田 柊, 杉尾 譲, 河村 卓哉, 山上 龍太, 岡 夏央, 平田 章, 横川 隆志, 堀 弘幸 :** ArcS-RaSEA 複合体によるリジン転移反応の解析, *第64回日本生化学会・中国四国支部例会,* 2023年5月.
1783. **杉尾 譲, 山崎 颯太, 上田 隼也, 磯貝 亮, 松本 奈津美, 林 実, 山上 龍太, 平田 章, 冨川 千恵, 河村 卓哉, 堀 弘幸 :** 第3のtRNA 4-チオウリジン合成経路をThermoplasma acidophilumは持つ, *第35回日本Archaea研究会,* 2023年7月.
1784. **Fujita Shu, Sugio Yuzuru, Kawamura Takuya, Yamagami Ryota, Oka Natsuhisa, 平田 章, Yokogawa Takashi, Hori Hiroyuki :** Lysine-transfer reaction by the complex of ArcS and RaSEA for archaeosine biosynthesis in tRNA, *第24回日本RNA学会年会,* 2023年7月.
1785. **廣田 翔也, 髙木 拓海, 久田 旭彦, 真岸 孝一, 藤原 直樹 :** 組み込み式ガスケットを用いた物性測定用高圧装置の開発, *2023年度 応用物理・物理系学会 中国四国支部 合同学術講演会 講演予稿集,* 92, 2023年7月.
1786. **髙木 拓海, 久田 旭彦, 真岸 孝一, 齊藤 隆仁, 上床 美也 :** 鉄系梯子型化合物の合成および高圧下電気抵抗測定, *2023年度 応用物理・物理系学会 中国四国支部 合同学術講演会 講演予稿集,* 69, 2023年7月.
1787. **元田 秀樹, 政岡 翔, 小笠原 正道, 大西 敦 :** 耐溶剤型多糖系キラルカラムを用いた有効キラル配位子の迅速スクリーニング手法の開発, *日本プロセス化学会2023サマーシンポジウム,* 2023年8月.
1788. **羽田 裕貴, 中谷 是崇, 水野 清秀, 納谷 友規, 中島 礼, 西山 賢一 :** 徳島平野沿岸部における地下第四系層序と堆積環境, *日本地質学会第130 年学術大会講演集,* 2023年9月.
1789. **西山 賢一, 露口 耕治, 大矢 基弘, 佐竹 一希, 川本 真由美 :** 愛媛県宇和島市に分布する崩壊堆積物に含まれる古土壌の14C年代, *日本地質学会第130 年学術大会講演集,* 2023年9月.
1790. **山本 孝, 刀根 武大, 峰 広嵩 :** 銅ジルコニア触媒による気相酢酸エチル直接合成反応の担体焼成温度依存性, *第132回触媒討論会,* 2023年9月.
1791. **竹内 雅人, 三國 諒宏, 河口 笑奈, 山本 孝, 松岡 雅也 :** 近赤外分光法による酸化タングステン担持ジルコニア触媒の表面酸性質評価, *第132回触媒討論会,* 2023年9月.
1792. **政岡 翔, 田上 遥, 檜垣 達也, 大木 靖弘, 小笠原 正道 :** Competitive Double Friedel-Crafts 2,5- and 1',2-Diacylation of Monophosphaferrocenes, *第69回有機金属化学討論会,* 2023年9月.
1793. **所 竣一, 山本 祐平, 中田 亮一, 村瀬 遼亮, 今井 昭二 :** ノビアスキレートPA-1固相抽出法を用いた降雨及び降雪中鉛同位体分析の前処理方法の検討及びMC-ICP-MS法を用いた夏季降雨中鉛の発生源の識別, *日本分析化学会第72年会,* 2023年9月.
1794. **山本 祐平, 岡 健太郎, 今井 昭二 :** 真空凍結乾燥および分画抽出法を用いた凍結湿性沈着試料中の微量金属元素の溶解性の解析, *日本分析化学会第72年会,* 2023年9月.
1795. **森 勇綺, 人見 牧, 小笠原 正道 :** Palladium-Catalyzed Synthesis of Alkoxyallenes and Their Application in Regioselective Hydroelementation Reactions, *第69回有機金属化学討論会,* 2023年9月.
1796. **金木 俊也, 纐纈 佑衣, 青矢 睦月, 中村 佳博, ウォリス サイモン :** 炭質物ラマンスペクトルから経験温度を推定する際のピークフィッティングの自動化:Part 2, *日本地質学会第130年学術大会講演要旨,* T2-P1, 2023年9月.
1797. **小山 雪乃丞, ウォリス サイモン, 永冶 方敬, 青矢 睦月 :** 沈み込み境界の深部スロー地震発生領域における最大尊断応力分布とその時間変化:西南日本三波川沈み込み帯の例, *日本地質学会第130年学術大会講演要旨,* T2-O13, 2023年9月.
1798. **岡村 知樹, 中村 大輔, 青矢 睦月 :** 東五良津岩体のザクロ石グラニュライトの変成温度圧力条件, *日本地質学会第130年学術大会講演要旨,* T2-O8, 2023年9月.
1799. **澤元 浩, 川崎 祐, 岸本 豊, 真岸 孝一, 松本 信洋, 永田 正一 :** スピネル型硫化物 Cu(Ir1-xRhx)2S4 における金属絶縁体転移の NMR による研究, *日本物理学会年次大会,* 2023年9月.
1800. **真岸 孝一, 杉浦 輝, 久田 旭彦, 川崎 祐, 桑井 智彦 :** PrTi2(Al1-xSix)20 の NMR, *日本物理学会年次大会,* 2023年9月.
1801. **山本 翔太, 犬飼 宗弘, 中村 浩一 :** LiMn2O4のJahn-Tellerひずみにおけるミリング効果, *日本物理学会第78回年次大会講演要旨集,* 2023年9月.
1802. **中村 浩一, 山本 翔太, 田中 康照, 犬飼 宗弘, 村井 啓一郎, 森賀 俊広 :** 酸化物における格子ひずみとイオン運動, *日本物理学会第78回年次大会講演概要集,* 2023年9月.
1803. **齋藤 有 :** 半遠洋性泥質堆積物の供給源が示唆する黒潮の砕屑物長距離輸送, *日本地質学会第 130 年学術大会,* 2023年9月.
1804. **林 順司, 川上 竜巳, 平田 章, 金丸 芳, 大島 敏久, 櫻庭 春彦 :** 超好熱アーキア由来FAD依存性D-乳酸脱水素酵素のX線結晶構造解析, *日本農芸化学会中四国支部会2023年度中四国・西日本支部合同大会,* 2023年9月.
1805. **政岡 翔, 田上 遥, 檜垣 達也, 大木 靖弘, 小笠原 正道 :** モノホスファフェロセンの2,5-位及び1',2-位への競争的二重フリーデル・クラフツアシル化反応, *社会産業理工学研究交流会2023,* 2023年9月.
1806. **西山 賢一 :** 南海トラフ巨大地震に伴う津波の自然災害伝承碑とその活用, *日本応用地質学会令和5年度研究発表会講演論文集,* 111-112, 2023年10月.
1807. **西山 賢一 :** 徳島県三好市における土砂災害の歴史, *令和5年度全国治山林道協会長会議講演,* 2023年10月.
1808. **西山 賢一 :** 中国四国地方における自然災害伝承碑の分布とその活用, *日本応用地質学会中国四国支部設立30周年記念講演,* 2023年10月.
1809. **山本 孝, 刀根 武大, 峰 広嵩 :** ジルコニア担持銅触媒のXRD/XAFS分析 およびエタノール転換反応活性(2) ー担体焼成温度の影響ー, *第59回X線分析討論会,* 2023年10月.
1810. **山本 祐平 :** 遠隔地における大気エアロゾル中の微量元素分析, *プラズマ分光分析研究会第120回講演会,* 2023年10月.
1811. **廣田 翔也, 松下 かのん, 髙木 拓海, 久田 旭彦, 真岸 孝一, 藤原 直樹, 上床 美也 :** 組み込み式ガスケットを用いた対向アンビル型高圧装置の開発Ⅲ, *第64回高圧討論会講演要旨集 高圧力の科学と技術第33巻(2023年)特別号,* **Vol.33,** 120, 2023年11月.
1812. **犬飼 宗弘, 佐藤 晴紀, 宮西 孝一郎, 根来 誠, 香川 晃徳, 中村 浩一 :** 共結晶の超分子シントンを活用した薬剤のトリプレットDNP, *第62回NMR討論会,* 2023年11月.
1813. **佐藤 晴紀, 宮西 孝一郎, 根来 誠, 香川 晃徳, 中村 浩一, 犬飼 宗弘 :** 共結晶の構造チューニングによる脂肪族化合物のトリプレット DNP, *第62回NMR討論会,* 2023年11月.
1814. **竹内 雅人, 三國 諒宏, 河口 笑奈, 山本 孝, 松岡 雅也 :** 酸化タングステン担持ジルコニア触媒の表面酸性サイトに吸着したNH3, NH4+の分析, *第39回近赤外フォーラム,* 2023年11月.
1815. **Fujita Shu, Sugio Yuzuru, Kawamura Takuya, Yamagami Ryota, Oka Natsuhisa, 平田 章, Yokogawa Takashi, Hori Hiroyuki :** ArcS-RaSEA複合体によるLys転移反応の生化学解析, *第46回日本分子生物学会年会,* 2023年12月.
1816. **Kawai Kumpei, Norioto Go, Matsuda Teppei, Manaka Sora, Yamagami Ryota, 平田 章, Hori Hiroyuki :** Characterization of tRNA methyltransferase Trm14 from a hyper-thermophilic archaeon, Thermococcus kodakarensis, *第46回日本分子生物学会年会,* 2023年12月.
1817. **安間 了, 申 キチョル, 齋藤 有, 横尾 頼子, 渡辺 千香子, 辻 彰洋 :** メソポタミア氾濫原堆積物と粘土板胎土の元素・同位体比分析, *第2回人・モノ・自然シンポジウム,* 2023年12月.
1818. **和田 悠兵, 吉田 知洋, 横尾 頼子, 安間 了 :** 四国吉野川水系河川水のイオン組成・硫黄同位体比からみる硫黄の起源の推定と流域環境の関係, *第13回同位体環境学シンポジウム,* 2023年12月.
1819. **服部 喜嗣, 横尾 頼子, 濱口 弘平, 前田 哲弥, 安間 了, Shiva Mehrabani, 申 基澈 :** イラン・イラクにおける降水のイオン組成の地域別特徴, *第13回同位体環境学シンポジウム,* 2023年12月.
1820. **田島 美来, 上野 雅晴 :** カラム精製を用いない新規セラミド輸送タンパク質阻害剤HPCB-5のグラムスケール合成研究, *2023年度第2回 日本プロセス化学会東四国地区フォーラムセミナー, P-11,* 2024年1月.
1821. **三好 亜季, 大村 D. 聡, 上野 雅晴 :** 水中ワンポットタンデムカップリング反応を駆使したソルバトクロミズムを発現する機能性化合物の網羅的合成, *2023年度第2回 日本プロセス化学会東四国地区フォーラムセミナー, P-12,* 2024年1月.
1822. **小笠原 正道 :** 面不斉フェロセン類の不斉合成法の開発と応用, *東京都立大学大学院理学研究科 特別講演会,* 2024年1月.
1823. **西山 賢一 :** 佐那河内村の地形・地質に基づく土砂災害危険個所の分布, 2024年1月.
1824. **小笠原 正道 :** 有機合成手法による金属錯体の精密合成:面不斉遷移金属錯体の不斉合成と応用, *有機合成ミニシンポジウム2024,* 2024年1月.
1825. **政岡 翔, 脇岡 正幸, 大木 靖弘, 小笠原 正道 :** ホスフィン配位子の修飾に利用されている嵩高い置換アリール基の立体的/電子的パラメータの定量化, *日本化学会第104春季年会,* 2024年3月.
1826. **三好 亜季, 小畠 美穂, 三好 德和, 上野 雅晴 :** ワンポット・タンデムカップリング反応を駆使した天然物Riccardin類の効率的かつ網羅的合成法の開発, *日本化学会第104春季年会,日本大学理工学部(千葉),公演番号 E1143-2am-02,* 2024年3月.
1827. **三好 亜季, 大村 D. 聡, 三好 德和, 上野 雅晴 :** 環境調和型ワンポット・タンデムカップリング反応を用いた光機能性化合物の効率的合成及びその評価, *日本化学会第104春季年会,日本大学理工学部(千葉),公演番号 E1143-2pm-02,* 2024年3月.
1828. **上野 雅晴, 田島 美来, 三好 德和 :** 大量供給を志向した新規セラミド輸送タンパク質阻害剤HPCB-5の合成研究, *日本化学会第104春季年会,日本大学理工学部(千葉),公演番号 E1143-2pm-08,* 2024年3月.
1829. **今津 陽菜, 政岡 翔, 鵜池 紗希, 小笠原 正道 :** ラセミ体面不斉1,1'-ジアリルフェロセン類の不斉閉環メタセシス/速度論分割における不斉モリブデン触媒の影響, *日本化学会第104春季年会,* 2024年3月.
1830. **中村 浩一, 北島 葉月, 井藤 弘章, 犬飼 宗弘, 村井 啓一郎, 森賀 俊広 :** LiMPO4 (M=Fe, Mn)における格子ひずみとイオン拡散挙動, *2024年春季大会プログラム 講演概要集,* 2024年3月.
1831. **山本 翔太, 犬飼 宗弘, 中村 浩一, 山本 孝 :** リチウムマンガン酸化物における局所構造と電気伝導挙動, *2024年日本物理学会春季大会講演要旨集,* 2024年3月.
1832. **真岸 孝一, 久田 旭彦, 川崎 祐, 桑井 智彦 :** NdNb2Al20 の NMR, *日本物理学会春季大会,* 2207, 2024年3月.
1833. **渡邊 達哉, 上野 雅晴, 三好 德和 :** 金属ストロンチウムを用いるアミド化合物のアルデヒドへの還元反応, *日本化学会第104春季年会,日本大学理工学部(千葉),公演番号 E1111-4pm-03,* 2024年3月.
1834. **鎌田 知里, 下田 亮介, 上野 雅晴, 三好 德和 :** ストロンチウムアルコキシド触媒を用いる簡便かつ有用なエステル交換反応の開発, *日本化学会第104春季年会,日本大学理工学部(千葉),公演番号 E1111-4pm-04,* 2024年3月.
1835. **森 勇綺, 人見 牧, 小笠原 正道 :** パラジウム触媒を用いたアルコキシアレンの合成と立体選択的分子変換への応用, *日本化学会第104春季年会,* 2024年3月.
1836. **田井 章博, 吉田 愛菜, 中村 光裕, 古賀 武尊 :** アスコルビン酸の高感度ハイスループット定量法, *第169回ビタミンC研究委員会,* 2023年6月.
1837. **犬飼 宗弘 :** 配位高分子の固体NMR, *JEOL 分析機器 NMRユーザーズミーティング,* 2023年10月.
1838. **西山 賢一 :** 徳島県における 自然災害伝承碑の分布とその活用, *第3回高知県・徳島県地すべり対策研究会,* 2024年2月.
1839. **犬飼 宗弘 :** 共結晶を活用したトリプレットDNPマトリクスの開発, *第2回DNP研究会,* 2024年3月.
1840. **西山 賢一 :** 福井洞窟の地質調査について, *史跡福井洞窟保存活用計画,* 105-106, 2024年3月.
1841. **安間 了 :** 石材考古学における可搬型蛍光X線分析装置の利用と問題点, *国際日本文化研究センター共同研究報告書, No.200,* 49-54, 2024年3月.
1842. **Takashi Yamamoto :** 2.9 Pre-edge structure, selection rules and quadrupole contributions, "International Tables for Crystallography Volume I: X-ray Absorption Spectroscopy and Related Techniques", John Wiley & Sons, Inc., Oct. 2024.
1843. **西山 賢一, 福井洞窟ミュージアム, 岡山理科大学 :** 空から見た洞窟遺跡, 雄山閣, 東京, 2024年10月.
1844. **Parisa Talebi, Rossella Greco, Takashi Yamamoto, Mahdiyeh Zeynali, Saeid Asgharizadeh *and* Wei Cao :** Hierarchical Nickel Carbonate Hydroxide Nanostructures for Photocatalytic Hydrogen Evolution from Water Splitting, *Materials Advances,* **Vol.5,** *No.7,* 2968-2973, 2024.
1845. **Shunsuke Endo, Yui Kouketsu *and* Mutsuki Aoya :** Sanbagawa subduction: what went in, how deep, and how hot did it get?, *Elements,* **Vol.20,** *No.2,* 77-82, 2024.
1846. **Takuya Kurihara, Yue Souri, Munehiro Inukai *and* Motohiro Mizuno :** CO2-induced gate-opening structural transition process of a porous coordination polymer revealed by solid-state 13C NMR, *Chemical Communications,* **Vol.60,** *No.38,* 5074-5077, 2024.
1847. **Wang Qingyu, Son Ka, Pietropaolo Adriana, Fortino Mariagrazia, Masamichi Ogasawara, Ohji Takehito, Shimoda Shuhei, Bando Masayoshi *and* Nakano Tamaki :** Distinctive Chiral Conformations Induced to Poly(naphthalene-1,4-diyl) by Helix-sense-selective Polymerization and Circularly Polarized Light Irradiation, *Chemistry - A European Journal,* **Vol.30,** e202304275-(1-6), 2024.
1848. **今井 昭二, 山本 祐平, 長崎 昌太, 黒谷 功, 奥田 桂子, 中尾 佑子 :** 四国山岳地域における渓流水中硫酸イオンの長距離輸送および国内起源に基づく地質環境画分と大気降下物画分の推計法, *分析化学,* **Vol.73,** *No.4, 5,* 131-140, 2024年.
1849. **Munehiro Inukai, Haruki Sato, Koichiro Miyanishi, Makoto Negoro, Akinori Kagawa, Yuta Hori, Yasuteru Shigeta, Takuya Kurihara *and* Koichi Nakamura :** Cocrystalline Matrices for Hyperpolarization at Room Temperature Using Photoexcited Electrons, *Journal of the American Chemical Society,* **Vol.146,** *No.21,* 14539-14545, 2024.
1850. **Filipp Temerov, Rossella Greco, Joran Celis, Salvador Eslava, Weimin Wang, Takashi Yamamoto *and* Wei Cao :** Activating 2D MoS2 by loading 2D Cu-S nanoplatelets for improved visible light photocatalytic hydrogen evolution, drug degradation, and CO2 reduction, *Results in Materials,* **Vol.22,** 100569, 2024.
1851. **Takashi Yamamoto *and* Kazunori Miyamoto :** Decomposition and migration behavior of Pt salt on Al2O3 and SiO2 upon calcination studied by XAFS and UV-Visible spectroscopy using a laboratory-type spectrometer, *Spectrochimica Acta. Part B: Atomic Spectroscopy,* **Vol.217,** 106974, 2024.
1852. **Shunya Kaneki, Yui Kouketsu, Mutsuki Aoya, Yoshihiro Nakamura, Simon Wallis, Yusuke Shimura *and* Ken Yamaoka :** An automatic peak deconvolution code for Raman spectra of carbonaceous material and a revised geothermometer for intermediate- to moderately high-grade metamorphism, *Progress in Earth and Planetary Science,* **Vol.11,** 35, 2024.
1853. **Nomura Kotohiro, Mancuso Raffaella, Cai Zhengguo, Dagorne Samuel, Eisen S. Moris, Gonsalvi Luca, Kotora Martin, Lee Yeoul Bun, Liu Shaofeng, Martins Margarida Luísa, Ohkuma Takeshi, Pombeiro Armando, Ragaini Fabio, Redshaw Carl, Visseaux Marc, Wu Zongquan, Yoshida Hiroto *and* Masamichi Ogasawara :** Exclusive Papers of the Editorial Board Members and Topical Advisory Panel Members of Catalysts in Section Catalysis in Organic and Polymer Chemistry, *Catalysts,* **Vol.14,** 407, 2024.
1854. **S. Abe, J. Abhir, Reiko Orito *and* 563 coauthors :** Dark matter line searches with the Cherenkov Telescope Array, *Journal of Cosmology and Astroparticle Physics,* **Vol.7,** 047, 2024.
1855. **Shu Fujita, Yuzuru Sugio, Takuya Kawamura, Ryota Yagamai, Natsuhisa Oka, Akira Hirata, Takashi Yokogawa *and* Hiroyuki Hori :** ArcS from Thermococcus kodakarensis transfers L-lysine to preQ0 nucleoside derivatives as minimum substrate RNAs., *The Journal of Biological Chemistry,* **Vol.300,** *No.8,* 107505, 2024.
1856. **Akihiro Takamatsu, Masatsugu Oishi, Shimpei Gohda, Hiroki Takemura, Konosuke Mitsushio, Satoshi Sugano, Takashi Yamamoto, Toshiaki Ina, Haruo Kishimoto *and* Takaaki Sakai :** Characteristics of Ag-doped LaMnO3 perovskite oxide and its application as a solid oxide fuel cell cathode, *Materials Advances,* 2024.
1857. **山本 孝, 宮本 一範 :** 白金化合物L3吸収端XANESホワイトライン強度評価法と酸化数―実験室型装置による観察―, *鉄と鋼,* **Vol.110,** *No.12,* 973-980, 2024年.
1858. **Akira Hirata *and* Bunzo Mikami :** Structural insight into sugar-binding modes of microbial ß-amylase, *Biochemical and Biophysical Research Communications,* **Vol.733,** *No.12,* 150695, 2024.
1859. **Schilling E. Manuel, Plissart Gaëlle, Akizawa Norikatsu, Lin Kuan-Yu, Corgne Alexandre, Alvear Maite, González Emilio, Marín Catalina, Walker J. Richard, Ryo Anma, Warren Jessica, Ishikawa Akira, Prigent Cecile, González Victoria, Rivera Ana, Martínez Felipe *and* González-Jiménez María José :** Magmatic and tectonic processes recorded by mantle rocks of the Taitao ophiolite (6 Ma), southern Chile, *Goldschmidt meeting,* 2024.
1860. **山本 孝 :** 白金化合物および担持金属塩熱分解時のL3吸収端XANESホワイトライン強度, *第84回分析化学討論会,* 2024年5月.
1861. **Fujita Shu, Sugio Yuzuru, Kawamura Takuya, Yamgami Ryota, Oka Natsuhida, 平田 章, Yokogawa Takashi, Hori Hiroyuki :** Lysine-transfer reaction by the complex of ArcS and RaSEA for archaeosine biosynthesis in tRNA, *第25回日本RNA学会年会,* 2024年6月.
1862. **Sugio Yuzuru, Yamasaki Sota, Ueda Junya, Isogai Ryo, Matsumoto Natsumi, Hayashi Minoru, Yamagami Ryota, 平田 章, Tomikawa Chie, Yokogawa Takashi, Hori Hiroyuki :** The third biosynthesis pathway of 4-thiouridine in tRNA, *第25回日本RNA学会年会,* 2024年6月.
1863. **松田 哲平, 山上 龍太, 鈴木 健夫, 平田 章, 堀 弘幸 :** Thermococcus kodakarensis のtRNAの6位を2'-O-メチル化する新規tRNAメチル化酵素の発見, *第36回日本Archaea研究会,* 2024年7月.
1864. **藤田 柊, 杉尾 譲, 河村 卓哉, 山上 龍太, 岡 夏央, 平田 章, 横川 隆志, 堀 弘幸 :** アーケオシン合成系第二段階目酵素・ArcSの基質RNA特異性, *第36回日本Archaea研究会,* 2024年7月.
1865. **宮田 侑奈, 河村 卓哉, 山上 龍太, 堀 弘幸, 平田 章 :** 極小アーキアARMAN-2 tRNAイントロン切断酵素ε₂ EndAの触媒反応機構, *第36回日本Archaea研究会,* 2024年7月.
1866. **倪 俊豪, 金井 保, 福岡 香里, 杉野 洋誉, 堀 弘幸, 平田 章 :** アーキア転写制御因子Tgrのマルトトリオース認識機構, *第36回日本Archaea研究会,* 2024年7月.
1867. **生島 くるみ, 櫻井 樹, 藤田 柊, 杉尾 譲, 河村 卓哉, 山上 龍太, 増田 太郎, 岡 夏央, 横川 隆志, 堀 弘幸, 平田 章 :** アーケオシン合成におけるLys転移酵素ArcSのX線結晶構造, *第36回日本Archaea研究会,* 2024年7月.
1868. **松下 かのん, 髙木 拓海, 久田 旭彦, 真岸 孝一, 上床 美也 :** Ba1-xKxFe2Se3 の合成および高圧下電気抵抗測定, *2024年度 応用物理・物理系学会 中国四国支部 合同学術講演会 講演予稿集,* 118, 2024年7月.
1869. **佐々木 亮太, 真岸 孝一, 久田 旭彦, 川崎 祐, 桑井 智彦 :** 磁場に鈍感な相転移を示す SmNb2Al20 の NMR による研究, *2024年度 応用物理・物理系学会 中国四国支部 合同学術講演会 講演予稿集,* 119, 2024年7月.
1870. **山本 孝 :** 白金化合物および担持金属塩熱分解時のL3吸収端XANESホワイトライン強度, *DV-Xα研究協会 DV-Xα講演会,* 2024年9月.
1871. **小山 雪乃丞, ウォリス サイモン, 永冶 方敬, 青矢 睦月 :** 沈み込み帯プレート境界にかかる最大剪断応力の推定と沈み込み帯熱モデルの再検討:後期白亜紀, 三波川沈み込み帯の例, *日本地質学会第131年学術大会講演要旨,* T1-O11, 2024年9月.
1872. **西山 賢一, 鳥井 真之 :** 熊本県阿蘇火山山麓に分布するアースフロー堆積物の14C年代, *日本地質学会学術大会講演要旨,* 2024年9月.
1873. **金木 俊也, 纐纈 佑衣, 青矢 睦月, 中村 佳博, ウォリス サイモン, 志村 侑亮, 山岡 健 :** 炭質物ラマンスペクトルから岩石の最高被熱温度を推定するための自動ピーク分離コードの紹介と使用者への実践的な助言, *日本地質学会第131年学術大会講演要旨,* T1-O12, 2024年9月.
1874. **井上 雄介, 安藤 徹, 柴田 伊廣, 竹本 帝人, 道家 涼介, 殿谷 梓, 中尾 賢一, 西山 賢一, 馬場 俊孝, 日色 知也, 山﨑 新太郎, 山田 芳恵, 横山 光, 長谷川 修一 :** 第23回地震火山地質こどもサマースクール in 吉野川「妖怪と探る吉野川 のヒミツ」でこどもたちが発見したこと, *日本地震学会予稿集,* 2024年10月.
1875. **野々村 敦子, 多田 智貴, 柳内 守, 殿谷 梓, 古谷 勇人, 西山 賢一, 長谷川 修一 :** 中山間地域における地形解析による豪雨時危険個所の可視化, *日本応用地質学会令和6年度研究発表会講演論文集,* 77-78, 2024年10月.
1876. **山本 孝 :** 異なるプロファイル関数により評価した白金化合物 L3 吸収端 XANES のホワイトライン強度, *第60回X線分析討論会,* 2024年10月.
1877. **山本 孝 :** Chemical Bonds / おもしろワクワク化学の世界 '23 徳島化学展, *化学と教育,* **Vol.72,** *No.4,* 173, 2024年.