

微生物関係論文

2022年以降公開分

No	部署分類	大分類	小分類	公開年	論文区分	論文タイトル	著者	雑誌・本	キーワード	論文フリーアクセス	URL
1	感染症部	細菌	カルバペネム耐性菌	2022	英文査読論文	Characterization of a Novel Plasmid in <i>Serratia marcescens</i> Harboring <i>bla</i> _{GES-5} Isolated from a Nosocomial Outbreak in Japan	Noriko Nakanishi, Shoko Komatsu, Tomotada Iwamoto, Ryohei Nomoto	J. Hosp. Infect. 121: 128-131, 2022.	Serratia marcescens Carbapenemase blaGES-5	はい	https://pubmed.ncbi.nlm.nih.gov/34906601/
2	感染症部	細菌	S. parvus	2022	英文査読論文	Complete Genome Sequences of Four <i>Streptococcus parvus</i> Strains Obtained from Saliva of Domestic Pigs in Japan	Ryohei Nomoto, Kasumi Ishida-Kuroki, Mari Tohya, Ichiro Nakagawa, Tsutomu Sekizaki	Microbiol Resour Announc. 2022.	Streptococcus parvus, Genome	はい	https://journals.asm.org/doi/10.1128/mra.01245-21
3	感染症部	細菌	S. ruminantium	2022	英文査読論文	Complete Genome Sequences of Three <i>Streptococcus ruminantium</i> Strains Obtained from Endocarditis Lesions of Cattle in Japan.	Ryohei Nomoto, Kasumi Ishida-Kuroki, Mari Tohya, Ichiro Nakagawa, Tsutomu Sekizaki	Microbiol Resour Announc. 2022.	Streptococcus ruminantium, Genome	はい	https://journals.asm.org/doi/10.1128/mra.01248-21
4	感染症部	細菌	薬剤耐性菌	2022	英文査読論文	Antibiotic Resistance in Non-Typhoidal <i>Salmonella enterica</i> Strains Isolated from Chicken Meat in Indonesia	Minori Takaichi, Kayo Osawa, Ryohei Nomoto, Norike Nakanishi, Masanori Kameoka, Makiko Miura, Katsumi Shigemura, Shohiro Kinoshita, Koichi Kitagawa, Atsushi Uda, Takayuki Miyara, Ni Made Mertaniasih, Usman Hadi, Dadik	Pathogens 2022, 11, 543.	non-typhoidal Salmonella enterica (NTS); Indonesia;	はい	https://www.mdpi.com/2076-0817/11/5/543
5	感染症部	ウイルス	SARS-CoV-2	2022	英文査読論文	Comparison of RT-PCR, RT-LAMP, and antigen quantification assays for the detection of SARS-CoV-2	Takayuki Wada, Eriko Kage-Nakadai and Yoshikazu Nishikawa	Jpan. J. Infect. Dis. 2022, 75, 249-253	antigen; pig; seroepidemiology; zoonosis	はい	https://www.jstage.jst.go.jp/article/yoken/advpub/0/advpub_JJID.2021.476/_article-char/ja/
6	感染症部	細菌	非結核性抗酸菌	2022	英文査読論文	Genomic features of <i>Mycobacterium avium</i> subsp. <i>hominissuis</i> isolated from pigns Japan	Tetsuya Komatsu, Kenji Ohya, Atsushi Ota, Yukiko Nishiuchi, Hirokazu Yano, Kayoko Matsuo, Justice Opare Odoi, Shota Suganuma, Kotaro Sawai, Akemi Hasebe, Tetsuo Asai, Tokuma Yanai, Hideto Fukushima, Takayuki Wada, Shiomi	Gigabyte, 2021, 1-12	Mycobacterium avium, Genome	はい	https://gigabytejournal.com/articles/33
7	感染症部	感染症情報	新型コロナ	2022	その他	2度のパンデミックを経験して	岩本朋忠	公衆衛生情報 2022.6, 20-21	パンデミック	いいえ	
8	感染症部	ウイルス	SARS-CoV-2	2022	英文査読論文	SARS-CoV-2 RNA in Wastewater Was Highly Correlated With the Number of COVID-19 Cases During the Fourth and Fifth Pandemic Wave in Kobe City, Japan	Yoshihiko Tanimoto, Erika Ito, Sonoko Miyamoto, Ai Mori, Ryohei Nomoto, Noriko Nakanishi, Naohiro Oka, Takao Morimoto, Tomotada Iwamoto	Frontiers in Microbiology 2022, 13, 892447	COVID-19; SARS-CoV-2, sewage, wastewater	はい	https://pubmed.ncbi.nlm.nih.gov/35756040/
9	感染症部	ウイルス	E型肝炎ウイルス	2022	英文査読論文	Isolation and genome sequencing of hepatitis 1 E virus genotype 1 imported from India to Japan	Sakura Kobayashi, Ai Mori, Ryuichi Sugiyama, Tian-Cheng Li, Yoshiki Fujii, Keigo Yato, Mami Matsuda, Tomoyuki Shiota, Masaya Katsunata, Tomotada Iwamoto, Masamichi Muramatsu, Ryosuke Suzuki	Jpan. J. Infect. Dis. 2022, online	Hepatitis E Virus Genotype 1	はい	https://www.jstage.jst.go.jp/article/yoken/75/6/75_JJID.2022.127/_pdf/-char/en
10	感染症部	細菌	薬剤耐性菌	2022	英文査読論文	Genetic characteristics of azithromycin-resistant <i>Neisseria gonorrhoeae</i> collected in Hyogo, Japan during 2015-2019	Makiko Miura, Katsumi Shigemura, Kayo Osawa, Noriko Nakanishi, Ryohei Nomoto, Reo Onishi, Hiroyuki Yoshida, Tom Sawamura, Shuih-Bin Fang, Yi-Te Chiang, Shian-ying Sung, Kuan-cho Chen, Takayuki Miyara and Masato Fuisawa	Journal of Medical Microbiology 2022;71:001533	23S rRNA, azithromycin resistance, mtrN <i>Neisseria</i>	はい	https://www.microbiologyresearch.org/content/journal/jmm/10.1099/jmm.0.001533
11	感染症部	細菌	レジオネラ	2023	英文査読論文	Evaluation of <i>Legionella pneumophila</i> SGUT serotypes isolated from bath water using multiplex-PCR serotyping assay	Shoko Komatsu, Shinobu Tanaka, Noriko Nakanishi	Jpan. J. Infect. Des. 2022, online	Legionella pneum	はい	https://www.jstage.jst.go.jp/article/yoken/76/1/76_JJID.2022.397/_article
12	感染症部	細菌	カンピロバクター	2022	英文査読論文	Quantitative detection and genetic characterization of thermotolerant <i>Campylobacter</i> spp. in fresh chicken meats at retail in Japan	Hiroshi Asakura, Shiori Yamamoto, Kazuhiro Yamada, Jun Kawase, Hiromi Nakamura, Kou-ichiro Abe, Yoshimasa Sasaki, Tetsuya Ikeda, Ryohei Nomoto	Front. Microbiol. 13:1014212.	retailed poultry meat, contamination level Japan;	はい	https://www.frontiersin.org/journals/microbiology/articles/10.3389/fmicb.2022.1014212/full
13	感染症部	細菌	レジオネラ	2022	英文査読論文	Investigation of a <i>Legionella pneumophila</i> Outbreak at a Bath Facility in Japan Using Whole-Genome Sequencing of Isolates from Clinical and Environmental Samples.	Noriko Nakanishi *, Shoko Komatsu, Shinobu Tanaka, Kengo Mukai and Ryohei Nomoto	Microorganisms 2023, 11, 28.	Legionella pneumophila; Legionnaires'	はい	https://pubmed.ncbi.nlm.nih.gov/36677320/
14	感染症部	細菌	結核	2022	その他	当院における外国生まれ結核患者の検討	滝口純司、李正道、岩林正明、横田真、橋本梨花、網本久敬、金子正博、藤井宏、富岡洋海、藤山理世、楠信也、有川健太郎、岩本朋忠	Kobe City Hosp Bull 61:13-17, 2022	結核	いいえ	
15	感染症部	細菌	大腸菌	2022	英文査読論文	Seroepidemiological survey on pigs and cattle for novel K88 (F4)-like colonisation factor detected in human enterotoxigenic <i>Escherichia coli</i>	Yoshihiko Tanimoto, Miyoko Inoue, Kana Komatsu, Atsuyuki Odani,	Epidemiology and Infection 150 2021-12-13	Cattle; enterotoxigenic Escherichia coli; K88	はい	https://www.cambridge.org/core/services/aop-cambridge-core/content/view/CC3E58314AA3F941CC6552C66549C909/S0950268821002697a.pdf/seroepidemiological-survey-on-pigs-and-cattle-for-novel-k88-f4-like-colonisation-factor-detected-in-human-enterotoxigenic-escherichia-coli.pdf
16	感染症部	ウイルス	サボウイルス	2023	英文査読論文	Prevalence of sapovirus and astrovirus in pediatric infectious gastroenteritis surveillance in Kobe City, Japan, during 2016-2019	Takeshi Hanafusa, Kentaro Arikawa, Yoshihiko Tanimoto *	Jpan. J. Infect. Dis., 76, 246-250, 2023	sapovirus, astrov	はい	https://www.jstage.jst.go.jp/article/yoken/76/4/76_JJID.2023.037/_article/-char/en
17	感染症部	感染症情報	進行性多巣性白質脳症	2023	英文査読論文	Nationwide Laboratory Surveillance of Progressive Multifocal Leukoencephalopathy in Japan: Fiscal Years 2011-2020	Kazuo Nakamichi, (略: 16名), Souichi Nukuzuma, (略: 7名) and Masayuki Saijo	Viruses 15, 968, 2023	JC virus; cerebrospinal fluid; laboratory surveillance;	はい	https://pubmed.ncbi.nlm.nih.gov/37112948/
18	感染症部	細菌	非結核性抗酸菌	2023	英文査読論文	Unique genomic sequences in a novel <i>Mycobacterium avium</i> subsp. <i>hominissuis</i> lineage enable fine scale transmission route tracing during pig movement	Kayoko Matsuo, Justice Opare Odoi, Shota Suganuma, Kotaro Sawai, Akemi Hasebe, Tetsuo Asai, Tokuma Yanai, Hideto Fukushima, Takayuki Wada, Shiomi Yoshida, Toshihiro Ito, Kentaro Arikawa, Mikihiro Kawai, Manabu Ato, Anthony D. Baughn, Tomotada Iwamoto, Fumito Maruyama	One Health 16 (2023) 100559	MAH VNTR Draft genome sequences Pig	はい	https://www.sciencedirect.com/science/article/pii/S2352771423000794
19	感染症部	ウイルス	新型コロナ	2023	その他	神戸市における新型コロナウイルス感染症のゲノムサーベイランス<第4波で経験>	岩本朋忠、中西典子、有川健太郎、谷本佳彦、野本竜平	日本公衆衛生協会 新型コロナウイルス感染症対応記録 p276-279:2023年3月	新型コロナ	いいえ	

20	感染症部	細菌	ヒアルロン酸分解菌	2023	英文査読論文	Isolation and identification of hyaluronandegrading bacteria from Japanese fecal microbiota	Hazuki Akazawa, Itsuko Fukuda, Haruna Kaneda, Shoichi Yoda, Mamoru Kinura, Ryohei Nomoto, Shuji Ueda, Yasuhito Shirai, Ro Osawa	PLoS ONE 18(5):e0284517, 2023	Hyaluronan, human gut microbiota	はい	https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0284517
21	感染症部	細菌	結核	2023	その他	全ゲノム解析時代を迎えた結核分子疫学の潮流	岩本朋忠	公衆衛生 vol.87 434-440	結核、ゲノム	いいえ	
22	感染症部	細菌	侵襲性肺炎球菌	2023	IASR	神戸市における侵襲性肺炎球菌感染症由来12F型の分子疫学解析 (IASR vol.44 8-9: 2023)	中西典子 米澤武志 田中 忍 瀨 夏樹 岩本朋忠 野本竜平	IASR vol.44 8-9: 2023	侵襲性肺炎球菌, 分子疫学	はい	https://www.niid.go.jp/niid/ja/typhi-m/iasr-reference/2606-related-articles/related-articles-515/11770-515r05.html
23	感染症部	細菌	結核	2023	英文査読論文	Transmission dynamics variability of lineage 2 Mycobacterium tuberculosis strains in Kobe, Japan, determined using population-based whole-genome sequencing analysis	Tomotada Iwamoto, Kentaro Arikawa, Yoshiro Murase, Tsuyoshi Sekizuka, Makoto Kuroda, Yukiko Nishiuchi, Nobuya Kusunoki, Riyo Fujiyama, Satoshi Mitarai	Infection, Genetics and Evolution 114 (2023) 105495	Tuberculosis Heterogeneity in transmission dynamics	はい	https://doi.org/10.1016/j.meegid.2023.105495
24	感染症部	細菌	レジオネラ	2023	英文査読論文	Prevalence and genetic distribution of Legionella spp. in public bath facilities in Kobe City, Japan	Shoko Komatsu, Shinobu Tanaka and Noriko Nakanishi	Journal of Water and Health Vol 21 No 11, 1727	bath water, Legionella, Legionella antimicrobial	はい	https://iwaponline.com/jwh/article/21/11/1727/98275/Prevalence-and-genetic-distribution-of-Legionella
25	感染症部	細菌	薬剤耐性菌	2024	英文査読論文	Metagenome and Resistome Analysis of Beta-Lactam-Resistant Bacteria Isolated from River Waters in Surabaya, Indonesia	Ryoneer Nomoto, Kayo Osawa, Shohiro Kinoshita, Koichi Kitagawa, Noriko Nakanishi, Rosantia Sarassari, Dadik Raharjo, Masato Fujisawa, Kuntaman Kuntaman, Toshiro Shirakawa	Microorganisms 2024, 12, 199.	resistance genes (ARGs); antimicrobial	はい	https://doi.org/10.3390/microorganisms12010199