

微生物関係論文

2020年以降公開分

| No | 部署分類 | 公開年 | 論文区分 | 論文タイトル | 著者 | 雑誌・本 |
|----|------|------|--------|---|---|--|
| 1 | 感染症部 | 2020 | 英文査読論文 | Bacterial population kinetics in heteroresistant <i>Mycobacterium tuberculosis</i> harbouring rare resistance-conferring mutations in <i>gyrA</i> and <i>rpoB</i> imply an epistatic interaction of mutations in a pre-XDR-TB patient | Shiomi Yoshida, Tomotada Iwamoto, Kentaro Arikawa, Tsuyoshi Sekizuka, Makoto Kuroda, Yoshikazu Inoue, Satoshi Mitarai, Taisuke Tsuji, Kazunari Tsuyuguchi and Katsuhiro Suzuki | J. Antimicrob. Chemother. 75:1722–1725, 2020 |
| 2 | 感染症部 | 2020 | 英文査読論文 | Two new cases of pulmonary infection by <i>Mycobacterium shigaense</i> , Japan | Shiomi Yoshida, Tomotada Iwamoto, Takehiko Kobayashi, Ryohei Nomoto, Yoshikazu Inoue, Kazunari Tsuyuguchi, and Katsuhiro Suzuki | Emerg. Infect. Dis. 26:2728–2732, 2020. |
| 3 | 感染症部 | 2020 | 英文査読論文 | Higher genome mutation rates of Beijing lineage of <i>Mycobacterium tuberculosis</i> during human infection | Mariko Hakamata, Hayato Takihara, Tomotada Iwamoto, Aki Tamari, Atsushi Hashimoto, Takahiro Tanaka, Shaban A. Kaboso, Gebremichal Gebretsadik, Aleksandr Iljinov, Akira Yokoyama, Yuriko Ozeki, Akihito Nishiyama, Yoshitaka Tateishi, Hiroshi Moro, Toshiaki Kikuchi, Shujiro Okuda and Sohkichi Matsumoto | Sci. Reports. 10:17997, 2020. |
| 4 | 感染症部 | 2020 | 英文査読論文 | A genome epidemiological study of SARS-CoV-2 introduction into Japan | Tsuyoshi Sekizuka, Kentaro Itokawa, Masanori Hashino, Tetsuro Kawano-Sugaya, Rina Tanaka, Koji Yatsu, Asami Ohnishi, Keiko Goto, Hiroyuki Tsukagoshi, Hayato Ehara, Kenji Sadamasu, Masakatsu Taira, Shinichiro Shibata, Ryohei Nomoto, Satoshi Hiroi, Miho Toho, Tomoe Shimada, Tamano Matsui, Tomimasa Sunagawa, Hajime Kamiya, Yuichiro Yahata, Takuya Yamagishi, Motoi Suzuki, Takaji Wakita, and Makoto Kuroda | mSphere 5:e00786–20, 2020. |
| 5 | 感染症部 | 2020 | 英文査読論文 | Agglutination of human polyomaviruses by using a tetravalent glycocluster as a cross-linker | Makoto Ogata, Takashi Onoda, Ami Koizumi, Yuhei Tokunaga, Isao Ohta, Souichi Nukuzuma, Enoch Y. Park, Taichi Usui, and Tetsuro Suzuki | ACS Omega 5:21940–21947, 2020 |
| 6 | 感染症部 | 2021 | 英文査読論文 | Characterization of JC Polyomavirus Derived from COS-IMRb Cells | Souichi Nukuzuma, Chiyoko Nukuzuma, Masanori Kameoka, Shigeki Sugiura, Kazuo Nakamichi, Takafumi Tasaki, Koushi Hidaka, and Tsutomu Takegami | Jpn. J. Infect. Dis. 74:48–53, 2021. |
| 7 | 感染症部 | 2021 | 英文査読論文 | The novel coronavirus pandemic and the state of the epidemic in Kobe, Japan | Noriko Nakanishi and Yoshio Iijima | J. Disaster Res. 16:84–87, 2021. |
| 8 | 感染症部 | 2021 | 英文査読論文 | False-positive results in SARS-CoV-2 antigen test with rhinovirus-A infection. | Shogo Otake, Sonoko Miyamoto, Ai Mori, Tomotada Iwamoto, Masashi Kasai | Pediatric. Int. 63:1135–1137, 2021. |
| 9 | 感染症部 | 2021 | 英文査読論文 | Whole-Genome Sequencing of Shiga toxin-producing <i>Escherichia coli</i> OX18 from a fatal hemolytic uremic syndrome case. | Kenichi Lee, Atsushi Iguchi, Kazuhiro Uda, Sohshi Matsumura, Isao Miyairi, Kenji Ishikura, Makoto Ohnishi, Junji Seto, Kanako Ishikawa, Noriko Konishi, Hiromi Obata, Ichiro Furukawa, Hiromi Nagaoka, Hirotaka Morinushi, Natsuki Hama, Ryohei Nomoto, Hiroshi Nakajima, Hideaki Kariya, Mitsuhiro Hamasaki, Sunao Iyoda | Emerg. Infect. Dis. 27:1509–1512, 2021. |

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| 10 | 感染症部 | 2021 | 英文査読論文 | A discernable increase in the severe acute respiratory syndrome coronavirus 2 R.1 lineage carrying an E484K spike protein mutation in Japan | Tsuyoshi Sekizuka, Kentaro Itokawa, Masanori Hashino, Kazuhiro Okubo, Asami Ohnishi, Keiko Goto, Hiroyuki Tsukagoshi, Hayato Ehara, Ryohei Nomoto, Makoto Ohnishi, Makoto Kuroda, Virus Diagnosis Group (NIID Toyama), COVID-19 Genomic Surveillance Network in Japan (COG-JP) | Infect. Genet. Evol. 94:105013, 2021. |
| 11 | 感染症部 | 2021 | 英文査読論文 | Occurrence of carriage of multidrug resistant Enterobacteriaceae among pregnant women in the primary health center and hospital Setting in Surabaya, Indonesia | Siti R. O. Sulikah, Miratul Hasanah, Wahyu Setyarini, Hari Parathon, Koichi Kitagawa, Noriko Nakanishi, Ryohei Nomoto, Kayo Osawa, Shohiro Kinoshita, Itaru Hirai, Toshiro Shirakawa, and Kuntaman Kuntaman Siti R. O. Sulikah, Miratul Hasanah, Wahyu Setyarini, Hari Parathon, Koichi Kitagawa, Noriko Nakanishi, Ryohei Nomoto, Kayo Osawa, Shohiro Kinoshita, Itaru Hirai, Toshiro Shirakawa, and Kuntaman Kuntaman | Microbial Drug Resistance ●: ●-●, 2021. |
| 12 | 感染症部 | 2022 | 英文査読論文 | Characterization of a Novel Plasmid in <i>Serratia marcescens</i> Harbouiring <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. |
| 13 | 感染症部 | 2022 | 英文査読論文 | Complete Genome Sequences of Four <i>Streptococcus parasuis</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. |
| 14 | 感染症部 | 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. |
| 15 | 感染症部 | 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, Noriko Nakanishi, Masanori Kameoka, Makiko Miura, Katsumi Shigemura, Shohiro Kinoshita, Koichi Kitagawa, Atsushi Uda, Takayuki Miyara, Ni Made Mertaniasih, Usman Hadi, Dadik Raharjo, Ratna Yulistiani, Masato Fujisawa, Kuntaman Kuntaman, Toshiro Shirakawa | Pathogens 2022, 11, 543. |
| 16 | 感染症部 | 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 | Jpn. J. Infect. Dis. 2022, 75, 249-253 |
| 17 | 感染症部 | 2022 | 英文査読論文 | Genomic features of <i>Mycobacterium avium</i> subsp. <i>hominissuis</i> isolated from pigs in Japan | Tetsuya Komatsu, Kenji Ohya, Atsushi Ota, Yukiko Ishiuchi, Hirokazu Yano, Kayoko Matsuo, Justice Opere Odoi, Shota Saganuma, Kotaro Sawai, Akemi Hasebe, Tetsuo Asai, Tokuma Yanai, Hideto Fukushi, Takayuki Wada, Shiomi Yoshida, Toshihiro Ito, Kentaro Arikawa, Mikihiko Kawai, Manabu Ato, Anthony D Baughn, Tomotada Iwamoto and Fumito Maruyama | Gigabyte, 2021, 1-12 |
| 18 | 感染症部 | 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 |
| 19 | 感染症部 | 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 |
| 20 | 感染症部 | 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 Katsumata, Tomotada Iwamoto, Masamichi Muramatsu, Ryosuke Suzuki | Jpn. J. Infect. Dis. 2022, on-line |

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| 21 | 感染症部 | 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, Toru Sawamura, Shiu-Bin Fang, Yi-Te Chiang, Shian-ying Sung, Kuan-cho Chen, Takayuki Miyara and Masato Fujisawa | Journal of Medical Microbiology 2022;71:001533 |
| 22 | 感染症部 | 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 | Jpn. J. Infect. Dis. 2022, online |
| 23 | 感染症部 | 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. |
| 24 | 感染症部 | 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. |
| 25 | 感染症部 | 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 |
| 26 | 感染症部 | 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 |
| 27 | 感染症部 | 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 *. | Jpn. J. Infect. Dis., 76, 246-250, 2023 |
| 28 | 感染症部 | 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 Sajio | Viruses 15, 968, 2023 |
| 29 | 感染症部 | 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 Fukushi, Takayuki Wada, Shiomji Yoshida, Toshihiro Ito, Kentaro Arikawa, Mikihiko Kawai, Manabu Ato, Anthony D. Baughn, Tomotada Iwamoto, Fumito Maruyama | One Health 16 (2023) 100559 |
| 30 | 感染症部 | 2023 | その他 | 神戸市における新型コロナ感染症のゲノムサーベイランス<第4波で経験した> | 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 |
| 31 | 感染症部 | 2023 | 英文査読論文 | Isolation and identification of hyaluronan-degrading bacteria from Japanese fecal microbiota | Hazuki Akazawa, Itsuko Fukuda, Haruna Kaneda, Shoichi Yoda, Mamoru Kimura,Ryohei Nomoto, Shuji Ueda, Yasuhito Shirai, Ro Osawa | PLoS ONE 18(5):e0284517, 2023 |
| 32 | 感染症部 | 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 |
| 33 | 感染症部 | 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 濱 夏樹 岩本朋忠 野本竜平 | IASR vol.44 8-9: 2023 |

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| 36 | 感染症部 | 2024 | 英文査読論文 | Metagenome and Resistome Analysis of Beta-Lactam-Resistant Bacteria Isolated from River Waters in Surabaya, Indonesia | Ryohei Nomoto, Kayo Osawa , Shohiro Kinoshita, Koichi Kitagawa, Noriko Nakanishi, Rosantia Sarassari, Dadik Raharjo, Masato Fujisawa, Kuntaman Kuntaman, Toshiro Shirakawa | Microorganisms 2024, 12, 199. |
| 37 | 感染症部 | 2024 | 日本語査読論文 | 尿中抗原検査により診断し得た <i>Legionella longbeachae</i> によるレジオネラ肺炎の1例 | 池成拓哉、池町真実、中西典子、小松頌子、田中忍、松岡佑、徳重康介、多田公英、二村絢子、大戸美穂、中野茉生、小池千裕、竹川啓史 | 日本臨床微生物学会雑誌 34 (3): 214-221, 2024 |
| 38 | 感染症部 | 2024 | IASR | 循環式気泡浴槽を感染源としたレジオネラ症発生事例における菌株解析—神戸市 | 小松頌子、田中忍、野本竜平、向井健悟、中西典子 | IASR Vol. 45 p111-112, 2024 |
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| 42 | 感染症部 | 2024 | 日本語査読論文 | HIV スクリーニング検査偽陽性およびHIV-1 陽性検体を用いた新規HIV-1/2 抗体確認検査法の有用性の検討 | 河上麻美代、北村有里恵、伊藤 仁、黒木倫士郎、青木 均、中澤 桢哉、柴田伸一郎、野本 竜平、仁平 稔、柿田 徹也、藤原 卓士、三宅 啓文、長島 真美、貞升 健志、吉村 和久 | 日本エイズ学会誌 26 : 158–162 |
| 43 | 感染症部 | 2024 | 英文査読論文 | <i>Mycobacterium tuberculosis</i> is less likely to acquire pathogenic mutations during latent infection than during active disease | Asami Osugi, Aki Tamaru, Takashi Yoshiyama, <u>Tomotada Iwamoto</u> , Satoshi Mitarai, Yoshiro Murase | Microbiol Spectr. 2024 Jul 2;12(7):e0428923. |