

研究成果の刊行に関する一覧表

雑誌

- 1) Kakita T, Iyoda S, Taira H, Kudeken T, Akeda Y, Ohnishi M, Nakamura K. Closed genome sequences of 39 Shiga toxin-producing *Escherichia coli* strains isolated from healthy cattle in Okinawa, Japan. *Microbiol Resour Announc*. 2025 13;14(11):e0081625.
- 2) Shimabukuro S, Kobayashi T, Yokoyama E, Ando N, Kikuchi T, Momose F, Kageyama T, Matsuoka H, Ishioka T, Hirai S. Establishment of autoclave conditions for disaccharide-containing liquid media using spectrophotometry and high-performance liquid chromatography. *Journal of Microbiological Methods*. 2026; 245: 107487.
- 3) 泉谷秀昌、李謙一、伊豫田淳、明田幸宏：2024 年に分離された腸管出血性大腸菌の MLVA 法による解析。IASR、第 46 巻、103-104、2025 年 5 月
- 4) Kubomura A, Lee K, Iyoda S, Akeda Y. 2025. Complete genome sequence of three atypical diarrheagenic *Escherichia coli* O166:H15 strains from foodborne outbreaks. *Microbiol Resour Announc* 14:e0074625.
- 5) Ooka T, Arai S, Lee K, Gotoh Y, Kubomura A, Imuta N, Hara-Kudo Y, Iyoda S, Hayashi T, Nishi J. 2025. Prevalence, sequence diversity, and amplification of an IS-associated enterotoxin gene, *astA*, in *Escherichia coli*. *Front Microbiol* 16:1635769.
- 6) Imai Y, Okuno M, Hoshiko Y, Kaneko H, Sato T, Noguchi A, Lee K, Yamamoto T, Nakayama-Imahiji H, Kuwahara T, Iyoda S, Ogura Y. 2025. Genomics of Shiga toxin-producing *Escherichia coli* from asymptomatic carriers in Japan highlights risk-adaptive control. *Commun Biol* 9.
- 7) Morita M, Lee K, Sugiyama A, Kojima N, Kawai Y, Izumiya H, Akeda Y. 2026. A case of laboratory-acquired *Salmonella* Typhi infection due to phage typing in Japan: whole-genome sequencing confirms the source of infection. *Access Microbiol* 8.