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Metagenome-based diversity analyses suggest a strong locality signal for bacterial communities associated with oyster aquaculture farms in Ofunato Bay

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Atsushi Kobiyama^a, Kazuho Ikeo^a, Md. Shaheed Reza^{a,1}, Jonaira Rashid ^a, Yuichiro Yamada^a, Yuri Ikeda^a, Daisuke Ikeda^a, Nanami Mizusawa^a, Shigeru Sato^a, Takehiko Ogata^a, Mitsuru Jimbo^a, Toshiaki Kudo^a, Shinnosuke Kaga^{b,2}, Shiho Watanabe^b, Kimiaki Naiki^b, Yoshimasa Kaga^b, Katsuhiko Mineta^c, Vladimir Bajic^c, Takashi Gojobori^{c*} and Shugo Watabe^{a,*}

^aKitasato University School of Marine Biosciences, Minami-ku, Sagamihara, Kanagawa 252-0373, Japan;

^bIwate Fisheries Technology Center, Kamaishi, Iwate 026-0001, Japan;

^c King Abdullah University of Science and Technology (KAUST), Computational Bioscience Research Center (CBRC), Thuwal 23955-6900, Kingdom of Saudi Arabia

*Corresponding authors:

Professor Shugo Watane, Department of Marine Biochemistry, Kitasato University School of Marine Biosciences, Minami-ku, Sagamihara, Kanagawa 252-0373, Japan. E-mail: swatabe@kitasato-u.ac.jp

Professor Takashi Gojobori, Computational Bioscience Research Center and Biological and Environmental Science and Engineering, King Abdullah University of Science and Technology, Thuwal 23955-6900, Kingdom of Saudi Arabia Email: takashi.gojobori@kaust.edu.sa

Present addresses: ¹Department of Fisheries Technology, Bangladesh Agricultural University, Mymensingh-2202, Bangladesh; ²Ofunato Fisheries Promotion Center, Iwate Prefectural Government, Ofunato, Iwate 022-8502, Japan.

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