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Addendum: Comparative Genomic Analysis of the Class *Epsilonproteobacteria* and Proposed Reclassification to Epsilonbacteraeota (phyl. nov.)

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Waite DW, Vanwonterghem I, Rinke C, Parks DH, Zhang Y, Takai K, Sievert SM, Simon J, Campbell BJ, Hanson TE, Woyke T, Klotz MG and Hugenholtz P (2018) Addendum: Comparative Genomic Analysis of the Class Epsilonproteobacteria and Proposed Reclassification to Epsilonbacteraeota (phyl. nov.). Front. Microbiol. 9:772. doi: 10.3389/fmicb.2018.00772 ¹ Australian Centre for Ecogenomics, School of Chemistry and Molecular Biosciences, The University of Queensland, St Lucia, QLD, Australia, ² Department of Cell and Molecular Biology, College of the Environment and Life Sciences, University of Rhode Island, Kingston, RI, United States, ³ Department of Subsurface Geobiological Analysis and Research (D-SUGAR), Japan Agency for Marine-Earth Science and Technology, Yokosuka, Japan, ⁴ Department of Biology, Woods Hole Oceanographic Institution, Woods Hole, MA, United States, ⁵ Microbial Energy Conversion and Biotechnology, Department of Biology, Technische Universität Darmstadt, Darmstadt, Germany, ⁶ Department of Biological Sciences, Life Science Facility, Clemson University, Clemson, SC, United States, ⁷ School of Marine Science and Policy, College of Earth, Ocean, and Environment, University of Delaware, Newark, DE, United States, ⁸ Delaware Biotechnology Institute, University of Delaware, Newark, DE, United States, ⁹ Department of Energy, Joint Genome Institute, Walnut Creek, CA, United States, ¹⁰ School of Molecular Biosciences, College of Veterinary Medicine, Washington State University, Pullman, WA, United States, ¹¹ State Key Laboratory of Marine Environmental Science, Institute of Marine Microbes and Ecospheres, College of Ocean and Earth Sciences, Xiamen University, Xiamen, China

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An addendum on

Comparative Genomic Analysis of the Class *Epsilonproteobacteria* and Proposed Reclassification to Epsilonbacteraeota (phyl. nov.)

by Waite, D. W., Vanwonterghem, I., Rinke, C., Parks, D. H., Zhang, Y., Takai, K., et al. (2017). Front. Microbiol. 8:682. doi: 10.3389/fmicb.2017.00682

In our original publication, we proposed the phylum name Epsilonbacteraeota according to a proposal to modify Rule 8 of the International Code of Nomenclature of Prokaryotes, under which the suffix *–aeota* would be used to denote prokaryotic phyla (Oren et al., 2015). An addendum to this proposal was recently made whereby the shorter suffix *–ota*, instead of *–aeota*, be added to the stem of the name of one of the contained classes (Whitman et al., 2018). In accordance with this amendment, and the requirement that a class name is derived from a type genus (Oren et al., 2015), we propose to replace the name Epsilonbacteraeota with Campylobacterota. This change does not affect subordinate ranks of the phylum.

Campylobacterota (Cam.py.lo.bac.ter.o'ta. N.L. neut. n. *Campylobacter* type genus of the type order of the type class of the phylum; suff. *-ota*, proposed ending to denote a phylum; N.L. neut. pl. n. Campylobacterota the phylum of the class *Campylobacteria*).

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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