



## OPEN ACCESS

EDITED AND REVIEWED BY  
Eric Altermann,  
Massey University, New Zealand

\*CORRESPONDENCE  
Shanquan Wang  
✉ wangshanquan@mail.sysu.edu.cn

SPECIALTY SECTION  
This article was submitted to  
Microbiotechnology,  
a section of the journal  
Frontiers in Microbiology

RECEIVED 30 November 2022  
ACCEPTED 05 December 2022  
PUBLISHED 15 December 2022

CITATION  
Wang S, He J, Shen C and  
Manefield MJ (2022) Editorial:  
Organohalide respiration: New  
findings in metabolic mechanisms and  
bioremediation applications, Volume II.  
*Front. Microbiol.* 13:1112309.  
doi: 10.3389/fmicb.2022.1112309

COPYRIGHT  
© 2022 Wang, He, Shen and  
Manefield. This is an open-access  
article distributed under the terms of  
the [Creative Commons Attribution  
License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use, distribution  
or reproduction in other forums is  
permitted, provided the original  
author(s) and the copyright owner(s)  
are credited and that the original  
publication in this journal is cited, in  
accordance with accepted academic  
practice. No use, distribution or  
reproduction is permitted which does  
not comply with these terms.

# Editorial: Organohalide respiration: New findings in metabolic mechanisms and bioremediation applications, Volume II

Shanquan Wang<sup>1\*</sup>, Jianzhong He<sup>2</sup>, Chaofeng Shen<sup>3</sup> and  
Michael J. Manefield<sup>4</sup>

<sup>1</sup>Environmental Microbiome Research Center, School of Environmental Science and Engineering, Guangdong Provincial Key Laboratory of Environmental Pollution Control and Remediation Technology, Southern Marine Science and Engineering Guangdong Laboratory (Zhuhai), Sun Yat-sen University, Guangzhou, China, <sup>2</sup>Department of Civil and Environmental Engineering, National University of Singapore, Singapore, Singapore, <sup>3</sup>Department of Environmental Engineering, College of Environmental and Resource Sciences, Zhejiang University, Hangzhou, China, <sup>4</sup>School of Civil and Environmental Engineering, University of New South Wales, Sydney, NSW, Australia

## KEYWORDS

organohalide respiration, *Dehalococcoides*, electron transport chain, reductive dehalogenase, microbiome, bioremediation

## Editorial on the Research Topic

[Organohalide respiration: New findings in metabolic mechanisms and bioremediation applications, Volume II](#)

The massive production and use of organohalides resulted in their worldwide contamination in soil, sediment and other environmental matrices (He et al., 2021). Organohalide-respiring bacteria (OHRB)-mediated reductive dehalogenation not only represents a promising solution for remediation of sites contaminated by organohalides (Jugder et al., 2016; Atashgahi et al., 2018), but involves element cycling in both terrestrial and marine environments (Horna-Gray et al., 2022; Xu et al., 2022). Recent research progress in characterizing major OHRB and crystal structures of key functional enzymes provided critical insights into organohalide respiration of OHRB (Bommer et al., 2014; Payne et al., 2015; Kublik et al., 2016; Wang et al., 2018; Picott et al., 2022). Nonetheless, there are still many puzzles to be resolved for better mechanistic understanding and bioremediation applications. Therefore, this Research Topic was formulated in two volumes to solicit manuscripts related to organohalide-respiring bacteria, reductive dehalogenase (RDase) and associated electron transport chain, dehalogenating microbiome, and organohalide bioremediation. Given the success of Volume I of this Research Topic and the rapidly evolving subject area, Volume II was launched for the publication of new research findings and updated information. We selected four manuscripts for publication after a rigorous peer review process.

## Organohalide-respiring bacteria

In a *Dehalococcoides*-containing enrichment culture, Zhao et al. reported extensive and even complete debromination of two commonly used polybrominated diphenyl ethers (PBDEs, i.e., BDE47 and BDE183). In addition, the debromination extent and rate of BDE183 could be enhanced by amendment of the BDE47. This study provides knowledge on new capabilities of *Dehalococcoides* and its potential in bioremediation of sites contaminated by both DBE47 and BDE183.

## RDases and associated electron transport chains

Reductive dehalogenase is the key enzyme to catalyze halogen removal from organohalides. Based on both transcription and translation analyses, Cimmino et al. deciphered the stoichiometry of *pceABCT* individual gene products in OHRB of Firmicutes. Notably, in contrast to a previously proposed model, results showed the formation of a membrane-bound PceA<sub>2</sub>B that could be devoid of PceC. These results provide unprecedented insight into the electron-accepting complex in PCE-dechlorinating OHRB of the phylum of Firmicutes.

## Organohalide bioremediation

Bioelectrochemical systems (BES) hold great potential for bioremediation of sites co-contaminated by organohalides and heavy metals. Matturro et al. employed both 16S rRNA gene amplicon sequencing and metagenomic analyses to elucidate the microbial interactions among *Dehalococcoides*, *Methanobrevibacter* and *Methanobacterium* for the efficient dechlorination of trichloroethene (TCE) and reduction of Cr(VI) in a BES. In addition, at sites contaminated with chlorinated ethenes, abiotic factors (e.g., iron sulfide minerals) could determine the fate of chloroethenes by affecting organohalide respiration of OHRB. Li et al. reported that FeS enhanced *Dehalococcoides*-mediated reductive dechlorination of TCE by formation of FeS nanoparticles and up-regulation of *tceA* transcription. These results could guide efficient bioremediation of sites contaminated by chlorinated ethenes and other contaminants.

With the success of the two volumes of this Research Topic, we would like to thank all the authors and reviewers for their valuable contributions. These papers significantly improve

our understanding in organohalide-respiring bacteria and their electron transport chains, as well as in dehalogenating microbiome and bioremediation implications. Notably, several research gaps were also highlighted in this Research Topic, and awaited future studies: (1) contribution of microbial reductive dehalogenation to attenuation of organohalides in natural environments; (2) cycling of organohalides in varied environmental matrices and associated functional microorganisms and enzymes; (3) reciprocal interactions of the commonly co-existing abiotic processes with the OHRB-mediated reductive dehalogenation process. We hope that this collection of reviews and original research articles will be helpful for researchers and engineers seeking information on organohalide respiration and bioremediation applications.

## Author contributions

All authors listed have made a substantial, direct, and intellectual contribution to the work and approved it for publication.

## Funding

Topic editors acknowledge support of NSFC Grants 41922049 and 42161160306 to SW, the Ministry of Education in Singapore under Academic Research Fund Tier 2 Project MOE-00003301 to JH, NSFC Grant 21876149 to CS, and Australian Research Council Discovery Project DP190103640 to MM.

## Conflict of interest

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.

## Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

## References

- Atashgahi, S., Häggblom, M. M., and Smidt, H. (2018). Organohalide respiration in pristine environments: implications for the natural halogen cycle. *Environ. Microbiol.* 20, 934–948. doi: 10.1111/1462-2920.14016
- Bommer, M., Kunze, C., Fessler, J., Schubert, T., Diekert, G., Dobbek, H., et al. (2014). Structural basis for organohalide respiration. *Science*. 346, 455–458. doi: 10.1126/science.1258118
- He, H., Li, Y., Shen, R., Shim, H., Zeng, Y., Zhao, S., et al. (2021). Environmental occurrence and remediation of emerging organohalides: a review. *Environ. Pollut.* 290, 118060. doi: 10.1016/j.envpol.2021.118060
- Horna-Gray, I., Lopez, N. A., Ahn, Y., Saks, B., Girer, N., Hentschel, U., et al. (2022). *Desulfoluna* spp. form a cosmopolitan group of anaerobic dehalogenating bacteria widely distributed in marine sponges. *FEMS Microbiol. Ecol.* 98, fiac063. doi: 10.1093/femsec/fiac063
- Jugder, B. E., Ertan, H., Bohl, S., Lee, M., Marquis, C. P., Manefield, M., et al. (2016). Organohalide respiring bacteria and reductive dehalogenases: key tools in organohalide bioremediation. *Front. Microbiol.* 7, 249. doi: 10.3389/fmicb.2016.00249
- Kublik, A., Deobald, D., Hartwig, S., Schiffmann, C. L., Andrades, A., von Bergen, M., et al. (2016). Identification of a multi-protein reductive dehalogenase complex in *Dehalococcoides mccartyi* strain CBDB1 suggests a protein-dependent respiratory electron transport chain obviating quinone involvement. *Environ. Microbiol.* 18, 3044–3056. doi: 10.1111/1462-2920.13200
- Payne, K. A., Quezada, C. P., Fisher, K., Dunstan, M. S., Collins, F. A., Sjuts, H., et al. (2015). Reductive dehalogenase structure suggests a mechanism for B12-dependent dehalogenation. *Nature*. 517, 513–516. doi: 10.1038/nature13901
- Picott, K. J., Flick, R., and Edwards, E. A. (2022). Heterologous expression of active dehalobacter respiratory reductive dehalogenases in *Escherichia coli*. *Appl. Environ. Microbiol.* 88, e0199321. doi: 10.1128/aem.01993-21
- Wang, S., Qiu, L., Liu, X., Xu, G., Siegert, M., Lu, Q., et al. (2018). Electron transport chains in organohalide-respiring bacteria and bioremediation implications. *Biotechnol. Adv.* 36, 1194–1206. doi: 10.1016/j.biotechadv.2018.03.018
- Xu, G., Zhang, N., Zhao, X., Chen, C., Zhang, C., He, J., et al. (2022). Offshore marine sediment microbiota respire structurally distinct organohalide pollutants. *Environ. Sci. Technol.* 56, 3065–3075. doi: 10.1021/acs.est.1c06680