Organohalide-respiring bacteria in polluted urban rivers employ novel bifunctional reductive dehalogenases to dechlorinate polychlorinated biphenyls and tetrachloroethene

Lan Qiu,[†] Wenwen Fang,[†] Haozheng He,[†] Zhiwei Liang,[†] Yangyue Zhan,[†] Qihong Lu,[†] Dawei Liang,[‡] Zhili He,[†] Bixian Mai,[§] Shanquan Wang^{*,†}

⁺ Environmental Microbiomics 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 510275

^{*} Beijing Key Laboratory of Bio-inspired Energy Materials and Devices, School of Space & Environment, Beihang University, Beijing, China 100191

[§] State Key Laboratory of Organic Geochemistry and Guangdong Key Laboratory of Environmental Protection and Resources Utilization, Guangzhou Institute of Geochemistry, Chinese Academy of Sciences, Guangzhou, China 510640

The supplementary information include 4 supplementary figures and 5 supplementary tables.



Figure S1. PCE dechlorination in the 174 microcosms. Blank boxes represent no detection of PCE dechlorination activity after three months of incubation. Samples in red font represent no PCB180 dechlorination activity.



Figure S2. Preference of *meta*- and *para*-chlorines in reductive dechlorination of PCB180 in the 135 PCB/PCE-dechlorinating sediment-free cultures (A) and characterized *Dehalococcoides mccartyi* strains (B). * A mixed culture containing *D. mccartyi* 195.







- Bacteria, Chloroflexi, Dehalococcoidia, Dehalococcoidales, Dehalococcoidaceae, Dehalococcoides
- Bacteria, Proteobacteria, Deltaproteobacteria, Desulfuromonadales, Geobacteraceae, Geobacter

Figure S3. Microbial community composition of (A) 135 PCB/PCE-dechlorinating microcosms and (B)29 PCE-dechlorinating microcosms without PCB dechlorination activity. Top 20 most abundant genus and known OHRBs (average relative abundance > 0.01%) were included in the figure.



Figure S4. Predicted microbial community function of the 135 PCB/PCE-dechlorinating microcosms. S6