

CURRICULUM VITAE

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Publications

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2. Li Wang, Xiang Xiong, Xuesong Luo, Wenli Chen, Shilin Wen, Boren Wang, Chengrong Chen, Qiaoyun Huang* 2020 Aggregational differentiation of ureolytic microbes in an Ultisol under long-term organic and chemical fertilizations. *Science of The Total Environment*, 716: 137103.
3. Xuesong Luo, Hang Qian, Li Wang, Shun Han, Shilin Wen, Boren Wang, Qiaoyun Huang*, Wenli Chen* 2019 Fertilizer types shaped the microbial guilds driving the dissimilatory nitrate reduction to ammonia process in a Ferralic Cambisol. *Soil Biology & Biochemistry*, 141: 107677
4. Yujie Xiao, Huizhong Liu, Meina He, Liang Nie, Hailing Nie, Wenli Chen* and Qiaoyun Huang* 2019 A crosstalk between c-di-GMP and cAMP in regulating transcription of GesA, a diguanylate cyclase involved in swimming motility in *Pseudomonas putida*. *Environmental Microbiology*, 22(1):142-157.
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7. Shaozu Xu, Yonghui Xing, Song Liu, Qiaoyun Huang* & Wenli Chen* 2019 Role of novel bacterial *Raoultella* sp. strain X13 in plant growth promotion and cadmium bioremediation in soil. *Appl Microbiol Biotechnol*. DOI 10.1007/s00253-019-09700-7
8. Shaozu Xu, Xuesong Luo, Yonghui Xing, Song Liu, Qiaoyun Huang, Wenli Chen* 2019 Complete genome sequence of *Raoultella* sp. strain X13, a promising cell factory for the

synthesis of CdS quantum dots. 3 Biotech. 9:120
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9. Shun Han, Xiang Xiong, Xuesong Luo, Luyang Zeng, Dan Wei, Wenli Chen*, Qiaoyun Huang* 2018 Fertilization rather than aggregate size fractions shape the nitrite-oxidizing microbial community in a Mollisol. *Soil Biology & Biochemistry*, 124:179-183. (DOI:10.1016/j.soilbio.2018.06.015).
10. Shun Han, Luyang Zeng, Xuesong Luo, Xiang Xiong, Shilin Wen, Boren Wang, Wenli Chen*, Qiaoyun Huang* 2018 Shifts in Nitrobacter- and Nitrospira-like nitrite-oxidizing bacterial communities under long-term fertilization practices. *Soil Biology & Biochemistry*, 124:118-125
11. Li Wang, Xuesong Luo, Hao Liao, Wenli Chen*, Dan Wei, Peng Cai, Qiaoyun Huang* 2018 Ureolytic microbial community is modulated by fertilization regimes and particle-size fractions in a Black soil of Northeastern China. *Soil Biology and Biochemistry* 116:171-178
12. Yujie Xiao, Wenjing Zhu, Huizhong Liu, Hailing Nie, Wenli Chen*, Qiaoyun Huang* 2018 FinR Regulates Expression of nicC and nicX Operons, Involved in Nicotinic Acid Degradation in *Pseudomonas putida* KT2440. *Appl Environ Microbiol.* 84(20):e01210-18
13. Shun Han, Xiang Li, Xuesong Luo, Shilin Wen, Wenli Chen* and Qiaoyun Huang* 2018 Nitrite-Oxidizing Bacteria Community Composition and Diversity Are Influenced by Fertilizer Regimes, but Are Independent of the Soil Aggregate in Acidic Subtropical Red Soil. *Front. Microbiol.*,<https://doi.org/10.3389/fmicb.2018.00885>
14. Hao Liao, Yuchen Zhang, Qinyan Zuo, Binbin Du, Wenli Chen*, Dan Wei, Qiaoyun Huang*. 2018 Contrasting responses of bacterial and fungal communities to aggregate-size fractions and long-term fertilizations in soils of northeastern China. *Sci Total Environ* 635:784-792.
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25. Ming Li, Xue Tian, Rong-Zi Liu, Wen-Li Chen, Peng Cai, Xing-Min Rong, Ke Dai, and Qiao-Yun Huang. 2014 Combined Application of Rice Straw and Fungus *Penicillium Chrysogenum* to Remediate Heavy-Metal-Contaminated Soil. *Soil and Sediment Contamination: An International Journal*. 23(3): 328-338
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