Yi Xiao

School of Life Sciences and Biotechnology, Shanghai Jiao Tong University Xitong Building (Room C404), 800 Dongchuan Road, Shanghai 200240, P.R. China Phone: +86 02134208601 Email: yi_xiao@sjtu.edu.cn

Research Summary

Yi Xiao's research interests focus on developing biotechnology tools for basic research and applied research. We design and construct genetically-encoded biosensors and CRISPR-associated tools. Using these tools, we modify cell genomes and reprogram cell behaviors to explore metabolism and enhance biosynthesis.

2019- A member of the editorial board of Applied and Environmental Microbiology

Education

Ph.D. (Microbiology) Wuhan Institute of Virology, Chinese Academy of Sciences, China, 2008

B.S. (Biotechnology) Huazhong University of Science and Technology, China, 2003

Employment

Jan. 2017-Current, Associate Professor Shanghai Jiao Tong University

Aug. 2010-Nov. 2016, Research Scientist and Postdoc. Washington University in St. Louis, USA

Sep. 2008-Jul. 2010, Postdoc. South China Sea Institute of Oceanology, Chinese Academy of Sciences, China

Publications

25, Chen-Guang Liu, **Yi Xiao**, Xiao-Xia Xia, Xin-Qing Zhao, Liangcai Peng, Penjit Srinophakun, Feng-Wu Bai. (2019) Cellulosic ethanol production: Progress, challenges and strategies for solutions. **Biotechnology Advances** 37(3):491-504.

24, Ruilian Yao[#], Di Liu[#], Xiao Jia, Yuan Zheng, Wei Liu, **Yi Xiao**^{*}. (2018) CRISPR-Cas9/Cas12a biotechnology and application in bacteria. Synthetic and Systems Biotechnology. 3(3):135-149

23, Juan Xia, ChenGuang Liu, XinQing Zhao, **Yi Xiao**, XiaoXia Xia, FengWu Bai. Contribution of cellulose synthesis, formation of fibrils and their entanglement to the self-flocculation of Zymomonas mobilis. *Biotechnology and Bioengineering* 115(11):2714-2725

22, **Yi Xiao**, Shaojie Wang, Sarah Rommelfanger, Andrea Balassy, Carlos Barba-Ostria, Pengfei Gu, Jonathan M. Galazka, Fuzhong Zhang^{*}. (2018) Developing a Cas9-based tool to engineer native plasmids in Synechocystis sp. PCC 6803. *Biotechnology and Bioengineering* 115(9):2305-2314.

21, **Yi Xiao**, Wen Jiang, Fuzhong Zhang. (2017) Developing a genetically encoded, cross-species biosensor for detecting ammonium and regulating biosynthesis of cyanophycin. *ACS Synthetic Biology*. 6 (10), pp 1807–1815

20, Gayle J. Bentley, Wen Jiang, Linda P. Guaman, **Yi Xiao**, Fuzhong Zhang. (2016) Engineering Escherichia coli to produce branched-chain fatty acids in high percentages. *Metabolic Engineering* 38:148–158

19, **Yi Xiao**, Christopher H. Bowen, Di Liu, Fuzhong Zhang. (2016) Exploiting nongenetic cellto-cell variation for enhanced biosynthesis. *Nature Chemical Biology* 12, 339–344

18, Di Liu, **Yi Xiao**, Bradley S. Evans, and Fuzhong Zhang. (2015) Negative Feedback Regulation of Fatty Acid Production Based on a Malonyl-CoA Sensor–Actuator. *ACS Synthetic Biology* 4(2):132-140.

17, Wen Jiang, Yanfang Jiang, Gayle J. Bentley, Di Liu, **Yi Xiao**, Fuzhong Zhang. (2015) Enhanced production of branched-chain fatty acids by replacing β-ketoacyl-(acyl-carrier-protein) synthase III (FabH). *Biotechnology and Bioengineering*. 112, 1613-1622

16, Lian He[#], **Yi Xiao**[#], Nikodimos Gebreselassie[#], Fuzhong Zhang, Maciek R. Antoniewicz, Yinjie J. Tang and Lifeng Peng. (2014) Central metabolic responses to the overproduction of fatty acids in Escherichia coli based on 13C-metabolic flux analysis. *Biotechnology and Bioengineering*. 111(3):575-85.

15, **Yi Xiao**, Zhenhua Ruan, Zhiguo Liu, Stephen G. Wu, Arul M. Varman, Yan Liu, Yinjie J. Tang. (2013) Engineering *Escherichia coli* to convert acetic acid to free fatty acids. *Biochemical Engineering Journal* 76:60-69)

14, Zhiguo Liu, Zhenhua Ruan, **Yi Xiao**, Yi Yu, Yinjie J. Tang, Wei Liao, Yan Liu. (2013) Integration of sewage sludge digestion with advanced biofuel synthesis. *Bioresource Technology* 132:166-170.

13, Arul Varman, **Yi Xiao**, Himadri Pakrasi, Yinjie J. Tang. (2013) Metabolic engineering of *Synechocystis* 6803 for isobutanol production. *Applied and Environmental Microbiology* 79:908-914.

12, **Yi Xiao**[#], Xueyang Feng[#], Arul M. Varman[#], Lian He, Huifeng Yu, Yinjie J. Tang. (2012) Kinetic modeling and isotopic investigation of isobutanol fermentation by two engineered *Escherichia coli* strains. *Industrial & Engineering Chemistry Research (ACS Publications)* 51:15855–15863.

11, **Yi Xiao**, Ting-Ting Liu, Hui Dai, Jun-Jie Zhang, Hong Liu, Huiru Tang, David J. Leak, and Ning-Yi Zhou. (2012) OnpA, an unusual flavin-dependent monooxygenase containing a cytochrome b5 Domain. *Journal of Bacteriology*. 194: 1342–1349

10, **Yi Xiao**[#], Sumei Li[#], Siwen Niu, Liang Ma, Guangtao Zhang, Haibo Zhang, Gaiyun Zhang, Jianhua Ju, and Changsheng Zhang. (2011) Characterization of tiacumicin B biosynthetic gene cluster affording diversified tiacumicin analogues and revealing a tailoring dihalogenase. *Journal of the American Chemical Society*. 133:1092-1105.

9, Arul Varman, **Yi Xiao**, Effendi Leonard. Yinjie J. Tang. (2011) Statistics-based model for prediction of chemical biosynthesis yield from *Saccharomyces cerevisiae*. *Microbial Cell Factories*. 10:45

8, Siwen Niu, Tao Hu, Sumei Li, **Yi Xiao**, Liang Ma, Guangtao Zhang, Haibo Zhang, Xiaohong Yang, Jianhua Ju, Changsheng Zhang. (2011) Characterization of a sugar-O-methyltransferase TiaS5 affords new Tiacumicin analogues with improved antibacterial properties and reveals substrate promiscuity. *Chembiochem*. 12:1740-1748.

7, Liping Gong, **Yi Xiao**, Qiang Liu, Sumei Li, Changsheng Zhang, Jinsong Liu. (2011). Preliminary X-ray crystallographic analysis of the glycosyltransferase from a marine *Streptomyces* species. *Acta Crystallographica Section F*. 67:136-139

6, **Yi Xiao**, Sumei Li, Liang Ma, Jianhua Ju, and Changsheng Zhang. (2010) Genetic manipulation system for tiacumicin producer *Dactylosporangium aurantiacum* NRRL 18085. *Wei Sheng Wu Xue Bao*. 50:1014-1022 (In Chinese)

5, Ying Yin, **Yi Xiao**, Hai-Zhou Liu, Fuhua Hao, Simon Rayner, Huiru Tang and Ning-Yi Zhou. (2010) Characterization of catabolic meta-nitrophenol nitroreductase from *Cupriavidus necator* JMP134. *Applied Microbiology and Biotechnology*. 87: 2077-2085

4, Qing Wei, Hong Liu, Jun-Jie Zhang, Song-He Wang, **Yi Xiao**, Ning-Yi Zhou. (2010) Characterization of a *para*-nitrophenol catabolic cluster in *Pseudomonas* sp. strain NyZ402 and construction of an engineered strain capable of simultaneously mineralizing both *para*- and *ortho*-nitrophenols. *Biodegradation*. 21: 575-584

3, Jun-Jie Zhang, Hong Liu, **Yi Xiao**, Xian-En Zhang, and Ning-Yi Zhou. (2009) Identification and characterization of catabolic *para*-nitrophenol 4-monooxygenase and *para*-benzoquinone reductase from *Pseudomonas* sp. strain WBC-3. *Journal of Bacteriology*. 191: 2703-2710

2, **Yi Xiao**, Jun-Jie Zhang, Hong Liu, Ning-Yi Zhou. (2007) Molecular characterization of a novel *ortho*-nitrophenol catabolic gene cluster in *Alcaligenes* sp. strain NyZ215. *Journal of Bacteriology*. 189:6587-6593

1, **Yi Xiao**[#], Jian-Feng Wu[#], Hong Liu, Shu-Jun Wang, Shuang-Jiang Liu, Ning-Yi Zhou. (2006) Characterization of genes involved in the initial reactions of 4-chloronitrobenzene degradation in *Pseudomonas putida* ZWL73. *Applied Microbiology and Biotechnology*. 73:166-171.