

张驰

手机：13606704530 出生年月：1989.09 民族：汉

邮箱：sidelinejudge@hotmail.com 性别：男

职称：助理研究员 政治面貌：中共党员



教育背景

200809-201207	中山大学	生命科学学院生态学系	理学学士
201209-201507	中国科学院大学	沈阳应用生态研究所	理学硕士
	论文《大豆 <i>GmIDA</i> 基因功能研究与转录组数据分析》		
	合作导师：崔敏龙研究员		
201509-201907	美国田纳西大学	赫伯特农学院植物科学系	哲学博士
	论文《Functional Study of Plant SABATH Methyltransferases in the Biosynthesis of Methyl Cinnamate, Juvenile Hormone III and Methyl Gibberellins》		
	合作导师：Feng Chen 教授		

工作经历

201309-201507	沈阳应用生态研究所	研究助理
201509-201907	美国田纳西大学	赫伯特农学院植物科学系 研究助理
201910-202203	西湖大学	理学院化学生物学实验室 博士后/助理研究员
202204-202501	哥本哈根大学	植物和环境系 博士后

论文发表

Wang, W., Zhang, C., Guo, H., & Chen, F. (2025). A novel auxin methyltransferase of the SABATH family for phenylacetic acid methylation is conserved in potato and tomato. *Plant physiology and biochemistry : PPB*, 224, 109972. Advance online publication.
<https://doi.org/10.1016/j.plaphy.2025.109972>

Liang, F., Xie, Y., Zhang, C., Zhao, Y., Motawia, M. S., Kampranis, S. C. (2025). Elucidation of the final steps in Taxol biosynthesis and its biotechnological production. *Nature Synthesis*, <https://doi.org/10.1038/s44160-025-00800-z>

Chen, L., Gao, R., Wei, G., Luo, S., Köllner, T. G., Xu, H., Jiang, Y., Zhang, C., Chen, X., Dickschat, J. S., Gershenzon, J., Li, S., Chang, Y., & Chen, F. (2025). Microbial-type terpene synthases significantly contribute to the terpene profile of glandular trichomes of the fern *Dryopteris fragrans* (L.). *The Plant journal : for cell and molecular biology*, 121(6), e70079. <https://doi.org/10.1111/tpj.70079>

Magnus, N., von Reuss, S. H., Braack, F., Zhang, C., Baer, K., Koch, A., Hampe, P. L., Sutour, S., Chen, F., & Piechulla, B. (2023). Non-canonical Biosynthesis of the Brexane-Type Bishomosesquiterpene Chlororaphen through Two Consecutive Methylation Steps in *Pseudomonas chlororaphis* O6 and *Variovorax boronicumulans* PHE5-4. *Angewandte Chemie (International ed. in English)*, 62(29), e202303692.

Chen, X., Nowicki, M., Wadl, P. A., Zhang, C., Köllner, T. G., Payá-Milans, M., Huff, M. L., Staton, M. E., Chen, F., & Trigiano, R. N. (2023). Chemical profile and analysis of biosynthetic pathways and genes of volatile terpenes in *Pityopsis ruthii*, a rare and endangered flowering plant. *PloS one*, 18(6), e0287524.

Zhang, W., Jiang, Y., Chen, F., Guan, Z., Wei, G., Chen, X., Zhang, C., Köllner, T. G., Chen, S., Chen, F., & Chen, F. (2022). Dynamic regulation of volatile terpenoid production and emission from *Chrysanthemum morifolium* capitula. *Plant physiology and biochemistry : PPB*, 182, 11 – 21.

Chen, S., *Zhang, C., & Zhang, L. (2022). Investigation of the Molecular Landscape of Bacterial Aromatic Polyketides by Global Analysis of Type II Polyketide Synthases. *Angewandte Chemie (International ed. in English)*, 61(24), e202202286. (*Co-first authors)

Jiang, Y., Liu, G., Zhang, W., Zhang, C., Chen, X., Chen, Y., Yu, C., Yu, D., Fu J., Chen, F., 2021. Biosynthesis and emission of methyl hexanoate, the major constituent of floral scent of a night-blooming water lily *Vicoria cruziana*. *Phytochemistry*. 191:112899

Piechulla, B., Zhang, C., Eisenschmidt-Bönn, D., Chen, F., Magnus, N., 2021. Non-canonical substrates for terpene synthases in bacteria are synthesized by a new family of methyltransferases. FEMS Microbiol Rev. fuab024.

Zhang, C., Chaiprasongsuk, M., Chanderbali, A.S., Chen, X., Fu, J., Soltis, D.E., Chen, F., 2020. Origin and evolution of a gibberellin-deactivating enzyme GAMT. Plant Direct. 4(12):e00287.

Zhang, L., Chen, F., Zhang, X., Li, Z., Zhao, Y., Lohaus, R., Chang, X., Dong, W., Ho, S.Y.W., Liu, X., Song, A., Chen, J., Guo, W., Wang, Z., Zhuang, Y., Wang, H., Chen, X., Hu, J., Liu, Y., Qin, Y., Wang, K., Dong, S., Liu, Y., Zhang, S., Yu, X., Wu, Q., Wang, L., Yan, X., Jiao, Y., Kong, H., Zhou, X., Yu, C., Chen, Y., Li, F., Wang, J., Chen, W., Chen, X., Jia, Q., Zhang, C., Jiang, Y., Zhang, W., Liu, G., Fu, J., Chen, F., Ma, H., Van de Peer, Y., Tang, H., 2020. The water lily genome and the early evolution of flowering plants. Nature. 577(7788):79-84.

Wei, G., Eberl, F., Chen, X., Zhang, C., Unsicker, S.B., Köllner, T.G., Gershenzon, J., Chen, F., 2020. Evolution of isoprenyl diphosphate synthase-like terpene synthases in fungi. Sci Rep. 10(1):14944

Zhang, C., Chen, X., Crandall-Stotler, B., Guo, H., Qian, P., Köllner, T. and Chen, F., 2019. Biosynthesis of methyl (*E*)-cinnamate in the liverwort *Conocephalum salebrosum* and evolution of cinnamic acid methyltransferase. Phytochemistry. 164, 50-59.

*Chaiprasongsuk, M., *Zhang, C., Qian, P., Chen, X., Li, G., Trigiano, R. N., Guo, H. and Chen, F., 2018. Biochemical characterization in Norway spruce (*Picea abies*) of SABATH methyltransferases that methylate phytohormones. Phytochemistry. 149, 146-154. (*Co-first authors)

Zhang, C., Gao, Z., Dong, Y. and Cui, M., 2015. Transcriptome analysis of inflorescences from four soybean cultivars. Chinese Journal of Ecology. 34, 3391-3396. (*In Chinese*)

Li, Y., Gao, Z., Zhang, C., Li, N. and Liu, C., 2015. Research progress on the molecular regulation mechanism of anthocyanin biosynthesis pathway. Chinese Journal of Ecology. 34, 2937-2942. (*In Chinese*)

Song, Q., Liu, C., Gao, Z., Li, Y., Zhang, C. and Cui, M., 2014. Optimal temperature for hsp18.2 promoter in gene expression of anthocyanin biosynthesis of *Antirrhinum majus*. Chinese Journal of Ecology. 33, 2436-2441. (*In Chinese*)

参会报告

“Biochemical Characterization of SABATH Methyltransferases in Liverworts” iMOSS 2018, St. Petersburg, Florida, June 2018.

参研项目

时间	项目名称	资助
2024-2026	真菌中非典型萜类合成	诺和诺德基金
2023-2025	紫杉醇的生物合成	丹麦国家研究基金
2019-2022	《聚酮合酶组装线改造》	西湖大学
2018-2019	《phytosensor》	DARPA
2015-2019	《Functional and Evolutionary Study of SABATH methyltransferase》	UTAgResearch
2015-2017	《大豆中编码 PIP 结构域的 GmIDL 基因在侧根发育中的功能研究》	国家自然科学基金
2015-2017	《GmPHRs 基因在大豆根应答低磷胁迫过程中的功能分析》	国家自然科学基金
2014-2017	《大豆根系中应答重金属 Cd 胁迫的基因表达模式的研究》	国家自然科学基金

专业服务

《Horticulture Research》, 《Plant Direct》 审稿人

获奖荣誉

AgResearch Travel Award, UT Institute of Agriculture, 2018

院级优秀团支书, 2008, 2010

研究兴趣

天然产物通路发掘

微生物与植物互作

大数据与人工智能

酶的进化与功能改造

个人爱好

跑山、足球、水上运动、西方文学