TAO YAN(严涛)

Lecturer at College of Agronomy, Hunan Agricultural University (HUAU), Changsha, China. My Ph.D research work covers a range of issues: Population Genetics Evolution and Ecotype Divergence Analysis of *Brassica napus*, Genome-wide Association Study (GWAS) of Agronomic Traits. Currently, I am interested in Transposable Elements Insertion Polymorphisms (TIPs) in Crop Population and genetic basis such as SV, CNV and TIPs etc. Now my research focus on Crop Stress and Improvement.

I am broadly interested in bioinformatics, data integration and visualization.





▲ Download a PDF of this CV

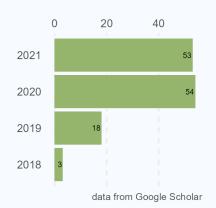
CONTACT

- **▼** tyan@zju.edu.cn
- TaoYan
- github.com/YTLogos
- Ø taoyan.netlify.app
- **%** yt056410
- **(86)** 13372566428

Citation = 128

H-index = 4

110-index = 3



Last updated on 2021-08-19.

2015 2014 2014 2013		Second-class Scholarship and Merit Student Zhejiang University Third-class Scholarship and Merit Student Zhejiang University → Hangzhou, CN PLIDI ICATIONS
2021	•	PUBLICATIONS BnaGVD: A genomic variation database of rapeseed (Brassica napus) Plant and Cell Physiology. 2021, 62(2):378–383. • Yan, T., Yao, Y., Wu, D., Jiang, L.* • First author • Impact Factor = 5.516
	•	Genome-wide association study reveals a patatin-like lipase relating to the reduction of seed oil content in Brassica napus BMC Plant Biology. 2021, 21(6). · Haoyi Wang, Qian Wang, Haksong Pak, Tao Yan, Mingxun Chen, Xiaoyang Chen, Dezhi Wu and Lixi Jiang* · Impact Factor = 4.960
2020	•	Genome-wide association study reveals new genes involved in leaf trichome formation in polyploid oilseed rape (Brassica napus L.) Plant, Cell & Environment. 2020, 43(3):675-691. • Xuan, L.*, Yan, T.*, Lu, L., Zhao, X., Wu, D., Hua, S., Jiang, L.* • Co-first author • Impact Factor = 7.791
		BnaSNPDB: An interactive web portal for the efficient retrieval and analysis of SNPs among 1,007 rapeseed accessions Computational and Structural Biotechnology Journal. 2020, 18:2766-2773. • Yan, T., Wang, Q., Maodzeka, A., Wu, D., Jiang, L.* • First author • Impact Factor = 7.409
2019		Whole-genome resequencing of a world-wide collection of rapeseed accessions reveals genetic basis of their ecotype divergence <i>Molecular Plant</i> . 2019, 12(1):30-43. • Wu, D., Liang, Z., <i>Yan, T.</i> , Xu, Y., Xuan, L., Tang, J., Zhou, G., Lohwasser, U., Hua, S., Wang, H., Chen, X., Wang, Q., Zhu, L., Maodzeka, A., Hussain, N., Li, Z., Li, X., Shamsi, I.H., Jilani, G., Wu, L., Zheng, H., Zhang, G., Chalhoub, B., Shen, L., Yu, H., Jiang, L.* • Impact Factor = 16.357

2018

Effect of high night temperature on storage lipids and transcriptome changes in developing seeds of oilseed rape

Journal of Experimental Botany. 2018, 69(7):1721-1733.

- · Zhou, L., Yan, T., Chen, X., Li, Z., Wu, D., Hua, S., Jiang, L.*
- Impact Factor = 7.860
- TRANSPARENT TESTA 4-mediated flavonoids negatively affect embryonic fatty acid biosynthesis in Arabidopsis

Plant, Cell & Environment. 2018, 41(12):2773-2790.

- · Xuan, L., Zhang, C., *Yan, T.*, Wu, D., Hussain, N., Li, Z., Chen, M., Pan, J., Jiang, L.*
- · Impact Factor = 7.791

CONFERENCE PROCEEDINGS

2020

Construction and utilization of a core germplasm of Brassica napus

第一届全国作物学科博士生论坛, Oct 2020

♀ Taiyuan, CN

·获取优秀奖

2019

GWAS reveals new genes involved in leaf trichome formation in polyploid oilseed rape (Brassica napus L.)

第十二届长三角作物学博士生论坛, Oct 2019

Yangzhou, CN

·获取优秀奖