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学习和工作经历	<p>2007–2011 本科，农学院，南京农业大学（金善宝实验班--植物生产）</p> <p>2011–2015 硕博连读，园艺学院，南京农业大学</p> <p>2013–2014 联合培养博士，自然与科技学院，丹麦奥胡斯大学</p> <p>2014–2015 联合培养博士，自然与科技学院，丹麦奥胡斯大学</p> <p>2016–2018 助理研究员，蔬菜研究所，江苏省农业科学院</p> <p>2018–2021 助理教授，自然与科技学院，丹麦奥胡斯大学</p> <p>2021– 副教授，园艺学院，南京农业大学</p> <p>主要研究方向为非生物胁迫下植物的生理生化及分子调控机制。通过耐逆野生番茄材料的筛选，我们从多角度联合分析阐明了植物耐逆（尤其是复合胁迫下）机制。同时，结合胁迫记忆、外源激素调控、CO₂加富等手段，提出了有效增强植物应对逆境的新见解。</p>			
科研项目	<p>[1] The effect of elevated CO₂ concentration and exogenous melatonin on tomatoes at combined heat and drought (复合胁迫下 CO₂ 上调和外源褪黑素对番茄生长发育的影响), Aarhus University Research Foundation, 在研, 主持。</p> <p>[2] Biosubstrate, Bio based growth media for plant production (基于生物的植物生长基质), GUDP (丹麦绿色发展示范项目), 在研, 主持。</p> <p>[3] 国家自然科学基金青年基金项目, 31601745, 番茄响应高温干旱胁迫的生理生化和环状 RNAs 调控机制解析, 结题, 主持。</p> <p>[4] 江苏省自然科学基金青年基金项目, BK20160579, 高温干旱胁迫下番茄生理生化和环状 RNAs 调控机制解析, 结题, 主持。</p> <p>[5] 江苏省农业科学院院基金, 基于混池分组分析法的番茄花青素遗传特点分析及分子标记开发, 结题, 主持。</p> <p>[6] 欧盟 Interreg 项目, Salfar-耐盐作物关键生长调控机理研究, 在研, 子课题主持。</p> <p>[7] DFF-Forskningsprojekt1 (tematisk forskning)/DFF-Research Project1 (Thematic Research), Impact of plant-based diet on the</p>			

	<p>consumption of health promoting microRNA' s, 在研, 参加。</p> <p>[8] MUDP, ' Waste to Value – Biofertilizer' med Baccess A/S,在研, 参加。</p> <p>[9] 省农业重大品种创制, 优质多抗高产设施番茄重大新品种创制,在研, 参加。</p> <p>[10] 国家重点研发计划, 茄科蔬菜优质多抗设施新品种培育,结题, 参加。</p> <p>[11] 江苏省农业科学院院基金, 番茄杂种优势群划分及性状相关性。结题, 参加。</p>
<p>发表论文</p>	<p>[1] <u>Rong Zhou</u>[*], et al. (2020). The alleviation of photosynthetic damage in tomato under drought and cold stress by high CO₂ and melatonin. International Journal of Molecular Sciences, 21, 5587.</p> <p>[2] <u>Rong Zhou</u>[*], et al. (2020). Genotype-dependent responses of chickpea to high temperature and moderately increased light. Plant Physiology and Biochemistry, 154: 353-359.</p> <p>[3] <u>Rong Zhou</u>[*], et al. (2020). Interactive effects of elevated CO₂ concentration and combined heat and drought stress on tomato photosynthesis. BMC Plant Biology, 20: 1-12.</p> <p>[4] <u>Rong Zhou</u>[*], et al. (2020). Combined high light and heat stress induced complex response in tomato with better leaf cooling after heat priming. Plant Physiology and Biochemistry, 151: 1-9.</p> <p>[5] <u>Rong Zhou</u>[*], et al. (2020). Unique miRNAs and their targets in tomato leaf responding to combined drought and heat stress. BMC Plant Biology, 20: 107.</p> <p>[6] <u>Rong Zhou</u>[*], et al. (2020). High throughput sequencing of circRNAs in tomato leaf responding to multiple stress of drought and heat. Horticultural Plant Journal, 6: 34-38.</p> <p>[7] <u>Rong Zhou</u>[*], et al. (2019). Physiological analysis and transcriptome sequencing reveal the effects of combined cold and drought on tomato leaf. BMC Plant Biology, 19: 377.</p> <p>[8] <u>Rong Zhou</u>, et al. (2019). Oxidative damage and antioxidant mechanism in tomatoes responding to drought and heat stress. Acta Physiol Plant, 41: 20-30.</p> <p>[9] <u>Rong Zhou</u>, et al. (2019). Physiological response of tomatoes at drought, heat and their combination followed by recovery, Physiol Plant, 165: 144-154. Highly cited paper</p> <p>[10] <u>Rong Zhou</u>, et al. (2019). Genome-wide identification of circRNAs in tomato seeds in response to high temperature. Biol Plantarum, 63: 97-103.</p>

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- [14] **Rong Zhou**, et al. (2017). Drought stress had a predominant effect over heat stress on three tomato cultivars subjected to combined stress. *BMC Plant Biol*, 17: 24-36. **Highly cited paper**
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- [16] **Rong Zhou**, et al. (2016). Identification of miRNAs and their targets in tomato at moderately and acutely elevated temperatures by high-throughput sequencing and degradome analysis. *Sci Rep*, 6: 33777-33789.
- [17] **Rong Zhou**, et al. (2015). Screening and validation of tomato genotypes under heat stress using F_v/F_m to reveal the physiological mechanism of heat tolerance. *Environ Exp Bot*, 118: 1-11.
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