

1. Huang et al. (2014) Adaptive divergence with gene flow in incipient speciation of *Miscanthus floridulus/sinensis* complex (Poaceae). *Plant J*, 80, 834-847.
2. Huang et al. (2014) Deciphering mycorrhizal fungi in cultivated *Phalaenopsis* microbiome with next-generation sequencing of multiple barcodes. *Fungal Diversity*, 66, 77-88.
3. Tran et al. (2019) Revegetation on abandoned salt ponds relieves the seasonal fluctuation of soil microbiomes. *BMC Genomics*, 20, 478.