

				
	<p>1. Ding Lian[#], Zhao Kunkun[#], Zhang Xue[#], Song Aiping, Su Jiangshuo, Hu Yueheng, Zhao Wenqian, Jiang Jiafu, Chen Fadi*. (2019) Comprehensive characterization of a floral mutant reveals the mechanism of hooked petal morphogenesis in <i>Arabidopsis thaliana</i>. <i>Plant Cell</i> 31(12): 2325-2340. IF_{5years} = 7.658</p> <p>2. Ding Lian[#], Song Aiping[#], Zhang Xue, Li Song, Su Jiangshuo, Xia Weikang, Zhao Kunkun, Zhao Wenqian, Guan Yunxiao, Fang Weimin, Chen Sumei, Jiang Jiafu, Chen Fadi* (2020) The core regulatory networks and hub genes in <i>Arabidopsis thaliana</i>. <i>Plant Cell</i> 32(12): 3315-3328. IF_{5years} = 7.658</p>			

	<p>regulating flower development in <i>Chrysanthemum morifolium</i>. . 103 (6):669-688. IF_{5years} = 4.065</p> <p>3. Zhao Kunkun[#], Ding Lian[#], Xia Weikang, Zhao Wenqian, Zhang Xue, Jiang Jiafu, Chen Sumei, Chen Fadi* (2020) Characterization of an APETALA1 and a FRUITFUL-like homolog in chrysanthemum. 272:109518. IF_{5years} = 2.844</p> <p>4. Wang Jingjing[#], Guan Yunxiao[#], Ding Lian, Li Peiling, Zhao Wenqian, Jiang Jiafu, Chen Sumei, Chen Fadi* (2019) The CmTCP20 gene regulates petal elongation growth in <i>Chrysanthemum morifolium</i>. 280:248-257. IF_{5years} = 4.253</p> <p>5. Wang J, Wang H, Ding L, Song A, Shen F, Jiang J, Chen S, Chen F* (2017) Transcriptomic and hormone analyses reveal mechanisms underlying petal elongation in <i>Chrysanthemum morifolium</i> ‘Jinba’. 93 (6):593-606. IF_{5years} = 4.065</p> <p>6. Ding Lian; Yan Shuangshuang; Jiang Li; Liu Meiling; Zhang Juan; Zhao Jianyu; Zhao Wensheng; Han Yingyan; Wang Qian.; Zhang Xiaolan*; HANABA TARANU regulates the shoot apical meristem and leaf development in cucumber (), , 2015, 66: 7075-7087. IF_{5years} = 7.011</p> <p>7. Ding Lian[#]; Yan Shuangshuang[#]; Jiang Li[#]; Zhao Wensheng; Ning Kang; Zhao Jianyu; Liu Xiaofeng; Zhang Juan; Wang Qian; Zhang Xiaolan*; HANABA TARANU (HAN) bridges meristem and organ primordia boundaries through PINHEAD, JAGGED, BLADE-ON-PETIOLE2 and CYTOKININ OXIDASE 3 during flower development in , , 2015, 11(9): e1005479. IF_{5years} = 5.857</p> <p>8. Zhao Jianyu[#]; Li Yanqiang[#]; Ding Lian[#]; Yan Shuangshuang; Liu Meiling; Jiang Li; Zhao Wensheng; Wang Qian; Yan Liqiang; Liu Renyi; Zhang Xiaolan*; Phloem transcriptome signatures underpin the physiological differentiation of the pedicel, stalk and fruit of cucumber (), , 2015, 57: 19-34. IF_{5years} = 4.799</p>