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科研项目				
发表论文	1. Wang YH, Jiang B, Dymerski R, et al (2021) Quantitative trait loci for horticulturally important traits defining the Sikkim cucumber, <i>Cucumis sativus</i> var. <i>sikkimensis</i> . <i>Theor Appl Genet</i> 134:229–247. 2. Wang YH, Bo K, Gu X, Pan P, Li Y, Chen J, Wen C, Ren ZH, Ren HZ, Chen X, Grumet R, Weng Y(2020). Molecularly Tagged Genes and Quantitative Trait Loci in Cucumber - And recommendation of controlled vocabulary for QTL mapping. <i>Horticulture Research</i> 7 (3) 3. Wang YH, Tan J, Wu Z, Wen C, VandenLangenberg K, Wehner TC, Thornton A, Zheng X, Owens K, Hoeft E, Kraan PAG, Suelmann J, Pan J, Weng Y (2019) STAYGREEN, STAY HEALTHY: a loss-of-susceptibility mutation in the <i>STAYGREEN</i> gene provides durable, broad-spectrum disease resistances for over 50 years of US cucumber production. <i>New Phytologist</i> 221:415–430. 4. Wang YH, VandenLangenberg K, Wen CL, Wehner TC, Weng Y (2018) QTL mapping of downy and powdery mildew resistances in PI 197088 cucumber with genotyping-by- sequencing in RIL population. <i>Theoretical and Applied Genetics</i> (131): 597. 5. Wang YH, VandenLangenberg K, Wehner TC, Kraan PAG, Suelmann J, Zheng XY, Owens K, Weng Y (2016) QTL mapping for downy mildew resistance in cucumber inbred line WI7120 (PI 330628). <i>Theoretical and Applied Genetics</i> (129): 1493. 6. Wang YH, VandenLangenberg K, Wehner TC, Weng Y. 2014. QTLs for Downy mildew resistance and their association with LRR-RLK resistance gene analogs in cucumber. <i>Proc Cucurbitaceae</i> 2014, pp 17-20.			

7. Pan, Y., **Wang, Y.**, McGregor, C., Liu, S., Luan, F., Gao, M. and Weng, Y. (2020) Genetic architecture of fruit size and shape variation in cucurbits: a comparative perspective. *Theor. Appl. Genet.*, 133, 1–21. Available at: <https://doi.org/10.1007/s00122-019-03481-3>.
8. Li, Z., Han, Y., Niu, H., **Wang, Y.**, Jiang, B. and Weng, Y. (2020) Gynoecy instability in cucumber (*Cucumis sativus* L.) is due to unequal crossover at the copy number variation-dependent *Femaleness* (*F*) locus. *Hortic. Res.*, 7. Available at: <http://dx.doi.org/10.1038/s41438-020-0251-2>.
9. Pan, Y., Wen, C., Han, Y., **Wang, Y.**, Li, Y., Li, S., Cheng, X. and Weng, Y. (2020) QTL for horticulturally important traits associated with pleiotropic andromonoecy and carpel number loci, and a paracentric inversion in cucumber. *Theor. Appl. Genet.*, 133, 2271–2290. Available at: <https://doi.org/10.1007/s00122-020-03596-y>.
10. Wen, C., Zhao, W., Liu, W., Yang, L., **Wang, Y.**, et al. (2019) CsTFL1 inhibits determinate growth and terminal flower formation through interaction with CsNOT2a in cucumber. *Development*, 146.
11. Rett-Cadman, S., Colle, M., Mansfeld, B., Barry, C.S., **Wang, Y.**, Weng, Y., Gao, L., Fei, Z. and Grumet, R. (2019) QTL and Transcriptomic Analyses Implicate Cuticle Transcription Factor <i>SHINE</i> as a Source of Natural Variation for Epidermal Traits in Cucumber Fruit. *Front. Plant Sci.*, 10, 1–16.
12. Yagcioglu, M., Jiang, B., Wang, P., **Wang, Y.**, Ellialtioglu, S.S. and Weng, Y. (2019) QTL mapping of low temperature germination ability in cucumber. *Euphytica*, 215, 84. Available at: <http://link.springer.com/10.1007/s10681-019-2408-3>.
13. Pan, J., Tan, J., **Wang, Y.**, Zheng, X., Owens, K., Li, D., Li, Y. and Weng, Y. (2018) STAYGREEN (CsSGR) is a candidate for the anthracnose (*Colletotrichum orbiculare*) resistance locus cla in Gy14 Cucumber. *Theor. Appl. Genet.*
14. Bo, K., Wang, H., Pan, Y., Behera, T., Pandey, S., Wen, C., **Wang, Y.**, et al. (2016) *SHORT HYPOCOTYL1* encodes a SMARCA3-Like chromatin remodeling factor regulating elongation. *Plant Physiol.*, 172, 1273–1292. Available at: <http://www.plantphysiol.org/lookup/doi/10.1104/pp.16.00501>.
15. Weng, Y., Colle, M., **Wang, Y.**, Yang, L., Rubinstein, M., Sherman, A., Ophir, R. and Grumet, R. (2015) QTL mapping in multiple populations and development stages reveals dynamic quantitative trait loci for fruit size in cucumbers of different market classes. *Theor. Appl. Genet.*, 128, 1747–1763.