

## 胡大刚

发布时间: 2017-03-22 浏览次数: 3781



胡大刚, 1986年出生, 博士, 园艺科学与工程学院副教授, 教育部“长江学者和创新团队发展计划”创新团队成员, 山东农业大学“杰出青年”基金获得者。自博士毕业以来, 一直在山东农业大学园艺科学与工程学院从事教学和科研工作; 在此期间, 曾赴各农业名校参观学习, 并到各地(包括河南郑州和西南大学柑桔研究所等)作技术指导, 现被聘为“泰安市科普志愿者专家服务团成员”; “中国细胞生物学学会专业分会成员”; 在国际及国内重要学术会议上作学术报告10余次; 讲授《园艺最新研究进展》和《园艺植物分子生物学实验》等课程; 主持国家重点研发子课题1项; 国家和山东省自然基金各1项; 山东农业大学杰出青年基金和科技创新基金各1项; 作物遗传与种质创新国家重点实验室开放基金课题1项; 近5年发表学术论著30余篇, 发表期刊包括SCI和国内核心期刊, 累积影响因子70左右。其中第一作者论文8篇(JCR一区4篇, 二区2篇, 三区2篇), 主要发表在PLOS Genetics、New Phytologist、Plant Physiology和Plant Journal等国外著名植物学期刊上; 通讯作者论文18篇(JCR一区1篇, 二区4篇, 三区2篇), 主要发表在Horticulture Research、Plant Science、BMC Plant Biology和Frontiers in Plant Science等国外著名植物学期刊上。所发表第一和通讯作者论文自2014年以后被SCI论文他引共计238次, 参编教材4部, 授权发明专利1项; 担任Journal of Agricultural and Food Chemistry (JCR一区), planta (JCR二区) 及DNA and Cell Biology (JCR三区) 等期刊的审稿人。

### 研究方向简介:

1、苹果果实品质的形成与调控。从苹果果实的糖酸代谢和运输入手, 研究了它们的合成、运输和液泡贮存; 然后, 从糖酸积累相关基因的环境和内部信号响应切入, 揭示了适度胁迫等促进糖酸积累、改善果实品质的机理。

2、苹果果实品质保持的机理与调控。研究采后苹果果实果肉软化和风味保持的调控机理, 并研发配套的苹果果实保鲜新技术; 同时揭示采后苹果果实病害发生的分子机理, 寻求降低采后果实病害的调控措施, 最终减少采后苹果果实的腐烂率。

### 近3年发表论文:

1. **Hu DG<sup>1</sup>**, Yu JQ, Han PL, Xie XB, Sun CH, Zhang QY, Wang JH, Hao YJ\*. The regulatory module MdPUB29-MdbHLH3 connects ethylene biosynthesis with fruit quality in apple. *New Phytologist*, DOI.org/10.1111/nph.15511.
2. Sun MH<sup>1</sup>, Ma QJ<sup>1</sup>, **Hu DG<sup>1</sup>**, Zhu XP, You CX, Shu HR, Hao YJ\* (2018) The glucose sensor MdHXK1 phosphorylates a tonoplast Na<sup>+</sup>/H<sup>+</sup> exchanger to improve salt tolerance. *Plant Physiology*, 176(4): 2977-2990. (Cite: 2)
3. Sun CH, Yu JQ, Duan Xi, Wang JH, Zhang QY, Gu KD, **Hu DG\***, Zheng CS\*. (2018) The MADS transcription factor CmANR1 positively modulates root system development by directly regulating *CmPIN2* in chrysanthemum. *Horticulture Research*, 5, 52. (Cite: 0)

4. Yu JQ, Wang JH, Sun CH, Zhang QY, **Hu DG\***, Hao YJ\*. (2018) Ectopic expression of the apple nucleus-encoded thylakoid protein MdY3IP1 triggers early-flowering and enhanced salt-tolerance in *Arabidopsis thaliana*. BMC Plant Biology, 18, 18. (Cite: 1)
5. Sun CH, Yu JQ, Wen LZ, Guo YH, Sun X, Hao YJ, **Hu DG\***, Zheng CS\*. (2018) Chrysanthemum MADS-box transcription factor *CmANR1* modulates lateral root development via homo-/heterodimerization to influence auxin accumulation in *Arabidopsis*. Plant Science, 266, 27-36. (Cite: 2)
6. Qi CH, Zhao XY, Jiang H, Liu HT, Wang YX, **Hu DG\***, Hao YJ\*. (2018) Molecular cloning and functional identification of an apple flagellin receptor *MdFLS2* gene. Journal of Integrative Agriculture, 17(0): 60345-7. (Cite: 0)
7. **Hu DG**, Li YY, Zhang QY, Li M, Sun CH, Yu JQ, Hao YJ\*. (2017) R2R3-MYB transcription factor MdMYB73 is involved in malate accumulation and vacuolar acidification in apple. The Plant Journal, 91, 443-454. (Cite: 6)
8. Sun CH, Yu JQ, **Hu DG\***. (2017) Nitrate: A Crucial Signal during Lateral Roots Development. Frontiers in Plant Science, 8, 485. (Cite: 15)
9. Sun MH, Ma QJ, Liu X, Zhu XP, **Hu DG\***, Hao YJ\*. (2017) Molecular cloning and functional characterization of *MdNHX1* reveals its involvement in salt tolerance in apple calli and *Arabidopsis*. Scientia Horticulturae, 215(2017), 126-133. (Cite: 2)
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11. Zhang QY, Liu XJ, Liu X, Wang JH, Yu JQ, **Hu DG\***, Hao YJ\*. (2017) Genome-Wide Identification, Characterization, and Expression Analysis of Calmodulin-Like Proteins (CMLs) in Apple. Horticultural Plant Journal, DOI:<https://doi.org/10.1016/j.hpj.2017.12.002>. (Cite: 0)
12. Sun CH, Zhang QY, Ming Li, Yu JQ, **Hu DG\***. (2017) A Universal Method of Protoplast Isolation and Transient Expression in Plants. Asian Journal of Life Sciences, 2017, AJLS-105.
13. Liu XJ, An XH, Liu X, **Hu DG**, Wang XF, You CX, Hao YJ\*. (2017) MdSnRK1.1 interacts with MdJAZ18 to regulate sucrose-induced anthocyanin and proanthocyanidin accumulation in apple. Journal of Experimental Botany, 68(11), 2977-2990.
14. Ma QJ, Sun MH, Lu J, Liu YJ, **Hu DG**, Hao YJ\*. (2017) Transcription factor AREB2 is involved in soluble sugar accumulation by activating sugar transporter and amylase genes. Plant Physiology, 174(4), 2348-2362.
15. **Hu DG**, Sun CH, Ma QJ, You CX, Cheng L, Hao YJ\*. (2016) MdMYB1 regulates anthocyanin and malate accumulation by directly facilitating their transport into vacuoles in apples. Plant Physiology, 170(3), 1315-1330. (Cite: 52)
16. **Hu DG**, Sun CH, Zhang QY, An JP, You CX, Hao YJ\*. (2016) Glucose sensor MdHXK1 phosphorylates and stabilizes MdbHLH3 to promote anthocyanin biosynthesis in apple. PLOS Genetics, 12(8), e1006273. (Cite: 24)
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18. **Hu DG**, Sun CH, Sun MH, Hao YJ\*. (2016) MdSOS2L1 phosphorylates MdVHA-B1 to modulate malate accumulation in response to salinity in apple. Plant cell Reports, 35(3), 705-718. (Cite: 3)
19. Ma QJ, Sun MH, Liu YJ, Lu J, **Hu DG\***, Hao YJ\*. (2016) Molecular cloning and functional characterization of the apple sucrose transporter gene *MdSUT2*. Plant Physiology and Biochemistry, 109, 442-451. (Cite: 5)
20. Sun CH, Zhang QY, Sun MH, **Hu DG\***. (2016) MdSOS2L1 forms a complex with MdMYB1 to control vacuolar pH by transcriptionally regulating MdVHA-B1 in apples. Plant Signaling & Behavior, 11(3), e1146846.
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