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2018年联合工程实验室发表文章汇总

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2018-08-22 发布于 发表文章 栏目中 浏览次数 79

[1] Y. Peng, L. Chen, S. Li, Y. Zhang, R. Xu, Z. Liu, W. Liu, J. Kong, X. Huang, Y. Wang, B. Cheng, L. Zheng BAK1 interact with G proteins and regulate sugar-responsive growth and development in *Arabidopsis*, 9 (2018). doi:10.1038/s41467-018-03884-8

[2] Y. Jiang, Q. Zheng, L. Chen, Y. Liang, J. Wu, Ectopic overexpression of maize heat shock transcript ZmHsf04 confers increased thermo and salt-stress tolerance in transgenic *Arabidopsis*, **Acta Physiol. P** doi:10.1007/s11738-017-2587-2.

[3] W. Chai, X. Peng, B. Liu, J. Wang, Z. Zhu, Y. Liu, K. Zhao, B. Cheng, W. Si, H. Jiang, Comparative Genome Re-sequencing and Expression Profile Analysis of Nucleobase Cation Symporter 2 (NCS2) C **Front. Plant Sci.** 9 (2018). doi:10.3389/fpls.2018.00856.

[4] N. Song, Z. Xu, J. Wang, Q. Qin, H. Jiang, W. Si, X. Li, Genome-wide analysis of maize CONSTANS-L and expression profiling under light/dark and abscisic acid treatment, **Gene.** 673 (201 10.1016/j.gene.2018.06.032.

[5] W. Chai, W. Si, W. Ji, Q. Qin, M. Zhao, H. Jiang, Genome-Wide Investigation and Expression Prof Transcription Factors in Foxtail Millet *Setaria italica*, **BioMed Res. Int.** 2018 (2018) 1–18. doi:10.1155/201

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