



七、发表的研究论文

序号	通讯作者	论文题目	刊物名称/发表日期	影响因子
1	蒋才富	Environmentally responsive genome-wide accumulation of de novo Arabidopsis thaliana mutations and epimutations	<i>Genome Research</i> , 2014, 24:1821-1829	14.927
2	张舒群	Mitogen-activated protein kinase cascades in signaling plant growth and development.	<i>Trends in Plant Science</i> , 2014, doi: 10.1016/j.tplants.2014.10.001	14.22
3	毛传藻	Rice SPX1 and SPX2 inhibit phosphate starvation responses through interacting with PHR2 in a phosphate-dependent manner	<i>Proc Natl Acad Sci USA.</i> , 2014, 111: 14953-14958	10.727
4	巩志忠	REPRESSOR of SILENCING5 encodes a member of the small heat shock protein family and is required for DNA demethylation in Arabidopsis	<i>The Plant Cell</i> , 2014, 26:2660-2675	10.656
5	郭岩	Inhibition of the <i>Arabidopsis</i> salt Overly Sensitive pathway by 14-3-3 proteins	<i>The Plant Cell</i> , 2014, 26:1166-1182	10.656
6	毛同林	Arabidopsis microtubule-destabilizing protein 25 functions in pollen tube growth by severing actin filaments	<i>The Plant Cell</i> , 2014, 26: 325-339	10.656



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7	王毅	The Os-AKT1 channel is critical for K ⁺ uptake in rice roots and is modulated by the rice CBL1-CIPK23 complex	<i>The Plant Cell</i> , 2014, 26: 3387-3402	10.656
8	吴平	SPX4 negatively regulates phosphate signaling and homeostasis through its interaction with PHR2 in rice	<i>The Plant Cell</i> , 2014, 26: 1586-1597	10.656
9	巩志忠	ABA-mediated ROS in mitochondria regulate root meristem activity by controlling PLETHORA expression in Arabidopsis	<i>PLoS Genetics</i> , 2014, 10: e1004791.	8.901
10	苏震	PNRD: A plant non-coding RNA database	<i>Nucleic Acids Research</i> , 2014, doi:10.1093/nar/gku1162	8.378
11	陈益芳	Arabidopsis WRKY45 transcription factor activates <i>PHOSPHATE TRANSPORTER1;1</i> expression in response to phosphate starvation	<i>Plant Physiology</i> , 2014, 164: 2020-2029	7.908
12	郭岩	A calcium-independent activation of the Arabidopsis SOS2-like protein kinase24 by its interacting SOS3-like calcium binding protein1	<i>Plant Physiology</i> , 2014, 164:2197-2206	7.908
13	李颖章	Histone H2B monoubiquitination is involved in regulating the dynamics of microtubules during the defense response to <i>Verticillium dahliae</i> toxins in Arabidopsis	<i>Plant Physiology</i> , 2014, 164:1857-1865	7.908



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14	郑绍建	Xyloglucan endotransglucosylase-hydrolase17 Interacts with xyloglucan endotransglucosylase-hydrolase31 to confer xyloglucan endotransglucosylase action and affect Aluminum sensitivity in Arabidopsis	<i>Plant Physiology</i> , 2014, 165:1566-1574	7.908
15	郑绍建	<i>TRICHOME BIREFRINGENCE-LIKE27</i> affects aluminum sensitivity by modulating the O-acetylation of xyloglucan and aluminum-binding capacity in Arabidopsis	<i>Plant Physiology</i> , 2014, 166:181-189	7.908
16	陈益芳	Arabidopsis RAV1 transcription factor, phosphorylated by SnRK2 kinases, regulates the expressions of <i>ABI3</i> , <i>ABI4</i> , and <i>ABI5</i> during seed germination and early seedling development	<i>The Plant Journal</i> , 2014, 80: 654-668	7.535
17	巩志忠	Abscisic acid inhibits root growth in Arabidopsis through ethylene biosynthesis	<i>The Plant Journal</i> , 2014 79:44-55	7.535
18	孙传清	<i>TOND1</i> confers tolerance to nitrogen deficiencies in rice	<i>The Plant Journal</i> , 2014 Dec.2 published online	7.535
19	叶 德	The ARID-HMG DNA-binding protein AtHMGB15 is required for pollen tube growth in Arabidopsis thaliana	<i>The Plant Journal</i> , 2014, 79, 741-756	7.535



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20	蒋德安	OsMOGS is required for N-glycan formation and auxin-mediated root development in rice (<i>Oryza sativa</i> L.)	<i>The Plant Journal</i> , 2014, 78: 632–645	7.535
21	蒋德安	OsABCB14 functions in auxin transport and iron homeostasis in rice (<i>Oryza sativa</i> .L).	<i>The Plant Journal</i> , 2014, 79:106-117.	7.535
22	郑绍建	Transcription factor WRKY46 regulates osmotic stress responses and stomatal movement independently in <i>Arabidopsis</i> .	<i>The Plant Journal</i> , 2014, 79:13-27	7.535
23	郑绍建	WRKY41 controls <i>Arabidopsis</i> seed dormancy via direct regulation of ABI3 transcript levels not downstream of ABA.	<i>The Plant Journal</i> , 2014, 79:810-823	7.535
24	任东涛	Activation of MKK9-MPK3/MPK6 enhances phosphate acquisition in <i>Arabidopsis thaliana</i>	<i>New Phytologist</i> , 2014, 203,1146-1160	7.289
25	杨淑华	<i>Arabidopsis</i> HSP90 protein modulates RPP4-mediated temperature-dependent cell death and defense responses	<i>New Phytologist</i> , 2014, 202: 1320-1334.	7.289
26	蒋德安	Auxin response factor (OsARF12), a novel regulator for phosphate homeostasis in rice (<i>Oryza sativa</i>)	<i>New Phytologist</i> , 2014, 201: 91–103	7.289
27	李召虎	Histone Lysine methyltransferase SDG8 is involved in brassinosteroid regulated gene expression in <i>Arabidopsis thaliana</i>	<i>Molecular Plant</i> , 2014, 7:1303-1315	6.348



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28	张舒群	Regulation of ethylene biosynthesis and signaling by protein kinases and phosphatases	<i>Molecular Plant</i> , 2014, 7,6,939-942	6.348
29	徐娟	Reactive oxygen species in signaling the transcriptional activation of WIPK expression in tobacco	<i>Plant Cell & Environment</i> , 2014, 37,7,1614-1625	6.242
30	杨建立	Identification of early Al-responsive genes in rice bean (<i>Vigna umbellata</i>) roots provides new clues to molecular mechanisms of Al toxicity and tolerance	<i>Plant Cell & Environment</i> , 2014, 37:1586-1597	6.242
31	韩玉珍	Tudor-SN, a component of stress granules, regulates growth under salt stress by modulating <i>GA20ox3</i> mRNA levels in <i>Arabidopsis</i>	<i>Journal of Experimental Botany</i> , 2014, 65, 20, 5933–5944	6.019
32	寿惠霞	Mutation in xyloglucan 6-xylosyltransferase results in abnormal root hair development in <i>Oryza sativa</i>	<i>Journal of Experimental Botany</i> , 2014, doi:10.1093/jxb/eru189	6.019
33	吴平	The paralogous SPX3 and SPX5 genes redundantly modulate Pi homeostasis in rice	<i>Journal of Experimental Botany</i> , 2014, 65: 859-870	6.019
34	杨淑华	Cold signal transduction and its interplay with phytohormones during cold acclimation	<i>Plant & Cell Physiology</i> , 2014, doi:10.1093/pcp/pcu115	4.972



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35	陈其军	A CRISPR/Cas9 toolkit for multiplex genome editing in plants	<i>BMC Plant Biology</i> , 2014, 14:327	4.758
36	苏震	SFGD: a comprehensive platform for mining functional information from soybean transcriptome data and its use in identifying acyl-lipid metabolism pathways	<i>BMC Genomics</i> , 2014, 15:271	4.5
37	李岩	The circular F-Actin bundles provide a track for turnaround and bidirectional movement of mitochondria in Arabidopsis root hair	<i>PLoS ONE</i> , 2014, 9(3): e91501	4.015
38	李岩	The C-terminus of AtGRIP is crucial for its self-association and for targeting to Golgi stacks in Arabidopsis	<i>PLoS ONE</i> , 2014, 9(6): e98963	4.015
39	李召虎	The phytotoxin coronatine induces abscission-related gene expression and boll ripening during defoliation of cotton	<i>PLoS ONE</i> , 2014, 9(5): e97652	4.015
40	刘国琴	Arabidopsis voltage-dependent anion channel 1 (AtVDAC1) is required for female development and maintenance of mitochondrial functions related to energy-transaction	<i>PLoS ONE</i> , 2014, 9(9):e106941	4.015
41	张明才	The effect of mepiquat chloride on elongation of cotton (<i>Gossypium hirsutum</i> L.) internode is associated with low concentration of gibberellic acid	<i>Plant Science</i> , 2014, 225:15-23	3.785



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42	肖兴国	Genome-wide analysis of <i>AP2/ERF</i> family genes from <i>Lotus corniculatus</i> shows <i>LcERF054</i> enhances salt tolerance	Functional & integrative Genomics , 2014, 14:453-466	3.217
43	田晓莉	Plant growth regulation enhanced potassium uptake and use efficiency in cotton	Field Crops Research , 2014, 163,109-118	2.957
44	张明才	Tillage practices affect biomass and grain yield through regulating root growth, root-bleeding sap and nutrients uptake in summer maize	Field Crops Research , 2014, 157:89-97	2.957
45	田晓莉	Functional characterization of <i>GhAKT1</i> , a novel Shaker-like K^+ channel gene involved in K^+ uptake from cotton (<i>Gossypium hirsutum</i>)	Gene , 2014, 545: 61-71	2.246