

Han Hu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2068257/publications.pdf>

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9
papers

2,647
citations

1307366

7
h-index

1474057

9
g-index

9
all docs

9
docs citations

9
times ranked

7271
citing authors

#	ARTICLE	IF	CITATIONS
1	Functional redundancy of <i>OsPIN1</i> paralogous genes in regulating plant growth and development in rice. <i>Plant Signaling and Behavior</i> , 2022, 17, 2065432.	1.2	3
2	<i>OsCLT1</i> , a CRT-like transporter 1, is required for glutathione homeostasis and arsenic tolerance in rice. <i>New Phytologist</i> , 2016, 211, 658-670.	3.5	75
3	Integration of summary data from GWAS and eQTL studies predicts complex trait gene targets. <i>Nature Genetics</i> , 2016, 48, 481-487.	9.4	1,757
4	GIPS: A Software Guide to Sequencing-Based Direct Gene Cloning in Forward Genetics Studies. <i>Plant Physiology</i> , 2016, 170, 1929-1934.	2.3	18
5	The paralogous <i>SPX3</i> and <i>SPX5</i> genes redundantly modulate Pi homeostasis in rice. <i>Journal of Experimental Botany</i> , 2014, 65, 859-870.	2.4	88
6	Expression of PIN Genes in Rice (<i>Oryza sativa</i> L.): Tissue Specificity and Regulation by Hormones. <i>Molecular Plant</i> , 2009, 2, 823-831.	3.9	185
7	Regulation of <i>OsSPX1</i> and <i>OsSPX3</i> on Expression of <i>OsSPX</i> domain Genes and Pi Starvation Signaling in Rice. <i>Journal of Integrative Plant Biology</i> , 2009, 51, 663-674.	4.1	119
8	QTLNetwork: mapping and visualizing genetic architecture of complex traits in experimental populations. <i>Bioinformatics</i> , 2008, 24, 721-723.	1.8	396
9	Genetic analysis for brix weight per stool and its component traits in sugarcane (<i>Saccharum</i>) Tj ETQq1 1 0.784314 $\frac{rgBT}{Overlock}$ 10		