## Peng Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The genome of the pear ( <i>Pyrus bretschneideri</i> Rehd.). Genome Research, 2013, 23, 396-408.	2.4	832
2	Spermidine oxidase-derived H2O2 regulates pollen plasma membrane hyperpolarization-activated Ca2+-permeable channels and pollen tube growth. Plant Journal, 2010, 63, 1042-1053.	2.8	182
3	LNK1 and LNK2 Are Transcriptional Coactivators in the <i>Arabidopsis</i> Circadian Oscillator. Plant Cell, 2014, 26, 2843-2857.	3.1	148
4	Genome-wide identification and comparative analysis of the heat shock transcription factor family in Chinese white pear (Pyrus bretschneideri) and five other Rosaceae species. BMC Plant Biology, 2015, 15, 12.	1.6	138
5	Phosphatidic Acid Counteracts S-RNase Signaling in Pollen by Stabilizing the Actin Cytoskeleton. Plant Cell, 2018, 30, 1023-1039.	3.1	101
6	A Na+/Ca2+ Exchanger-like Protein (AtNCL) Involved in Salt Stress in Arabidopsis. Journal of Biological Chemistry, 2012, 287, 44062-44070.	1.6	81
7	Genome-wide characterization, evolution, and expression analysis of the leucine-rich repeat receptor-like protein kinase (LRR-RLK) gene family in Rosaceae genomes. BMC Genomics, 2017, 18, 763.	1.2	62
8	Gene-expression profile of developing pollen tube of Pyrus bretschneideri. Gene Expression Patterns, 2016, 20, 11-21.	0.3	40
9	<i>COR27</i> and <i>COR28</i> encode nighttime repressors integrating <i>Arabidopsis</i> circadian clock and cold response. Journal of Integrative Plant Biology, 2017, 59, 78-85.	4.1	39
10	Identification and testing of reference genes for gene expression analysis in pollen of Pyrus bretschneideri. Scientia Horticulturae, 2015, 190, 43-56.	1.7	34
11	Physiological and Nutritional Responses of Pear Seedlings to Nitrate Concentrations. Frontiers in Plant Science, 2018, 9, 1679.	1.7	33
12	Genome-wide identification and comparative analysis of the cation proton antiporters family in pear and four other Rosaceae species. Molecular Genetics and Genomics, 2016, 291, 1727-1742.	1.0	32
13	Characterization of the pectin methyl-esterase gene family and its function in controlling pollen tube growth in pear (Pyrus bretschneideri). Genomics, 2020, 112, 2467-2477.	1.3	27
14	Mitochondrial dysfunction mediated by cytoplasmic acidification results in pollen tube growth cessation in <i>Pyrus pyrifolia</i> . Physiologia Plantarum, 2015, 153, 603-615.	2.6	18
15	Characterization of Dof family in Pyrus bretschneideri and role of PbDof9.2 in flowering time regulation. Genomics, 2020, 112, 712-720.	1.3	18
16	LNK1 and LNK2 recruitment to the evening element require morning expressed circadian related MYB-like transcription factors. Plant Signaling and Behavior, 2015, 10, e1010888.	1.2	17
17	Evolutionary and Expression Analysis Provides Evidence for the Plant Glutamate-like Receptors Family is Involved in Woody Growth-related Function. Scientific Reports, 2016, 6, 32013.	1.6	16
18	Phylogenetic and expression analysis of the magnesium transporter family in pear, and functional verification of <i>PbrMGT7</i> in pear pollen. Journal of Horticultural Science and Biotechnology, 2018, 93, 51-63.	0.9	14

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19	Identification and functional characterization of SOC1-like genes in Pyrus bretschneideri. Genomics, 2020, 112, 1622-1632.	1.3	13
20	Characterization of the pectin methylesterase inhibitor gene family in Rosaceae and role of PbrPMEI23/39/41 in methylesterified pectin distribution in pear pollen tube. Planta, 2021, 253, 118.	1.6	13
21	Genome-wide identification and expression analysis of the <i>OSCA</i> gene family in <i>Pyrus bretschneideri</i> . Canadian Journal of Plant Science, 2018, 98, 918-929.	0.3	12
22	PbrROP1/2-elicited imbalance of cellulose deposition is mediated by a CrRLK1L-ROPGEF module in the pollen tube of <i>Pyrus</i> . Horticulture Research, 2022, 9, .	2.9	8
23	PbCOL8 is a clock-regulated flowering time repressor in pear. Tree Genetics and Genomes, 2017, 13, 1.	0.6	7
24	PbGLR3.3 Regulates Pollen Tube Growth in the Mediation of Ca2+ Influx in Pyrus bretschneideri. Journal of Plant Biology, 2018, 61, 217-226.	0.9	7
25	Comprehensive genomic analysis of the RNase T2 gene family in Rosaceae and expression analysis in Pyrus bretschneideri. Plant Systematics and Evolution, 2020, 306, 1.	0.3	7
26	The unique evolutionary pattern of the Hydroxyproline-rich glycoproteins superfamily in Chinese white pear (Pyrus bretschneideri). BMC Plant Biology, 2018, 18, 36.	1.6	6
27	PbrSLAH3 is a nitrate-selective anion channel which is modulated by calcium-dependent protein kinase 32 in pear. BMC Plant Biology, 2019, 19, 190.	1.6	6
28	Phylogenetic and Expression Analysis of Pear Yellow Stripe-Like Transporters and Functional Verification of PbrYSL4 in Pear Pollen. Plant Molecular Biology Reporter, 2016, 34, 737-747.	1.0	3
29	PbrPOE21 inhibits pear pollen tube growth in vitro by altering apical reactive oxygen species content. Planta, 2020, 252, 43.	1.6	3
30	Identification and function analysis of fasciclin-like arabinogalactan protein family genes in pear (Pyrus bretschneideri). Plant Systematics and Evolution, 2021, 307, 1.	0.3	3
31	Identification and expression analysis of the PbrMLO gene family in pear, and functional verification of PbrMLO23. Journal of Integrative Agriculture, 2021, 20, 2410-2423.	1.7	2
32	Cellulose accumulation mediated by <scp>PbrCSLD5</scp> , a cellulose synthaseâ€like protein, results in cessation of pollen tube growth in <i>Pyrus bretschneideri</i> . Physiologia Plantarum, 2022, 174, e13700.	2.6	2
33	Characterization and Functional Explorations of O-glycosylation Enzymes SECRET AGENT and SPINDLY in Pyrus bretschneideri. Journal of Plant Biology, 0, , 1.	0.9	0