## **Zhouxin Shen**

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5978373/publications.pdf

Version: 2024-02-01

ors

#	Article	IF	CITATIONS
1	Nitrate triggered phosphoproteome changes and a PIN2 phosphosite modulating root system architecture. EMBO Reports, 2021, 22, e51813.	4.5	22
2	Wnt5a Induces ROR1 to Interact Grb2 to Enhance Ras Activation in Chronic Lymphocytic Leukemia. Blood, 2021, 138, 247-247.	1.4	1
3	Dynamic regulation of Pep-induced immunity through post-translational control of defence transcript splicing. Nature Plants, 2020, 6, 1008-1019.	9.3	40
4	Genetic elucidation of interconnected antibiotic pathways mediating maize innate immunity. Nature Plants, 2020, 6, 1375-1388.	9.3	52
5	Auxin Induces Widespread Proteome Remodeling in Arabidopsis Seedlings. Proteomics, 2019, 19, 1900199.	2.2	10
6	LIKE SEX4 1 Acts as a $\hat{i}^2$ -Amylase-Binding Scaffold on Starch Granules during Starch Degradation. Plant Cell, 2019, 31, 2169-2186.	6.6	26
7	Wnt5a induces ROR1 to recruit cortactin to promote breast-cancer migration and metastasis. Npj Breast Cancer, 2019, 5, 35.	5.2	18
8	The Second Site Modifier, Sympathy for the ligule, Encodes a Homolog of Arabidopsis ENHANCED DISEASE RESISTANCE4 and Rescues the Liguleless narrow Maize Mutant. Plant Cell, 2019, 31, 1829-1844.	6.6	17
9	Magnesium Flux Modulates Ribosomes to Increase Bacterial Survival. Cell, 2019, 177, 352-360.e13.	28.9	77
10	Quantitative Early Auxin Root Proteomics Identifies GAUT10, a Galacturonosyltransferase, as a Novel Regulator of Root Meristem Maintenance. Molecular and Cellular Proteomics, 2019, 18, 1157-1170.	3.8	29
11	Multiple genes recruited from hormone pathways partition maize diterpenoid defences. Nature Plants, 2019, 5, 1043-1056.	9.3	60
12	Discovery, Biosynthesis and Stress-Related Accumulation of Dolabradiene-Derived Defenses in Maize. Plant Physiology, 2018, 176, 2677-2690.	4.8	94
13	Wnt5a induces ROR1 to recruit DOCK2 to activate Rac1/2 in chronic lymphocytic leukemia. Blood, 2018, 132, 170-178.	1.4	36
14	The Dictyostelium GSK3 kinase GlkA coordinates signal relay and chemotaxis in response to growth conditions. Developmental Biology, 2018, 435, 56-72.	2.0	6
15	Integration of omic networks in a developmental atlas of maize. Science, 2016, 353, 814-818.	12.6	411
16	The small GTPases Ras and Rap1 bind to and control TORC2 activity. Scientific Reports, 2016, 6, 25823.	3.3	47
17	Identification of Evening Complex Associated Proteins in Arabidopsis by Affinity Purification and Mass Spectrometry. Molecular and Cellular Proteomics, 2016, 15, 201-217.	3.8	170
18	Wnt5a Induces ROR1 to Complex with HS1, Which Undergoes Tyrosine Phosphorylation and Contributes to Planar-Cell-Polarity Migration in Chronic Lymphocytic Leukemia. Blood, 2016, 128, 301-301.	1.4	2

#	Article	IF	CITATIONS
19	Wnt5a Induces Association of ROR1 with 14-3-3î¶ to Enhance Chemotaxis and Proliferation in Chronic Lymphocytic Leukemia. Blood, 2016, 128, 349-349.	1.4	1
20	Stemness factor Sall4 is required for DNA damage response in embryonic stem cells. Journal of Cell Biology, 2015, 208, 513-520.	5.2	50
21	A High-Resolution Tissue-Specific Proteome and Phosphoproteome Atlas of Maize Primary Roots Reveals Functional Gradients along the Root Axes. Plant Physiology, 2015, 168, 233-246.	4.8	64
22	Plastid-produced interorgannellar stress signal MEcPP potentiates induction of the unfolded protein response in endoplasmic reticulum. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 6212-6217.	7.1	82
23	The matrix protein Fibulin-5 is at the interface of tissue stiffness and inflammation in fibrosis. Nature Communications, 2015, 6, 8574.	12.8	64
24	Parallel Proteomic and Phosphoproteomic Analyses of Successive Stages of Maize Leaf Development. Plant Cell, 2013, 25, 2798-2812.	6.6	94
25	Reconstruction of protein networks from an atlas of maize seed proteotypes. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, E4808-17.	7.1	174
26	Plant elicitor peptides are conserved signals regulating direct and indirect antiherbivore defense. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 5707-5712.	7.1	179
27	ROR1 Can Interact With TCL1 and Enhance Leukemogenesis In EÂμ-TCL1 Transgenic Mice. Blood, 2013, 122, 868-868.	1.4	0
28	Use of high-throughput LC–MS/MS proteomics technologies in drug discovery. Drug Discovery Today: Technologies, 2006, 3, 301-306.	4.0	6
29	Differential Expression Profile of the Proteome and Transcriptome in Aggressive and Indolent Chronic Lymphocytic Leukemia Blood, 2005, 106, 2101-2101.	1.4	0