## Xiuren Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1907987/publications.pdf

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41 4,846 23 39 g-index

43 43 43 43 5994

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	scInTime: A Computational Method Leveraging Single-Cell Trajectory and Gene Regulatory Networks to Identify Master Regulators of Cellular Differentiation. Genes, 2022, 13, 371.	2.4	4
2	PRP4KA phosphorylates SERRATE for degradation via 20 <i>S</i> proteasome to fine-tune miRNA production in <i>Arabidopsis</i> Science Advances, 2022, 8, eabm8435.	10.3	16
3	The R-loop influences miRNA birth place. Nature Plants, 2022, 8, 320-321.	9.3	4
4	RNA architecture influences plant biology. Journal of Experimental Botany, 2021, 72, 4144-4160.	4.8	12
5	HASTY moves to chromatin for miRNA production. Molecular Plant, 2021, 14, 364-365.	8.3	2
6	The epigenetic factor FVE orchestrates cytoplasmic SGS3-DRB4-DCL4 activities to promote transgene silencing in <i>Arabidopsis</i> . Science Advances, 2021, 7, .	10.3	11
7	In vitro Reconstitution Assays of Arabidopsis 20S Proteasome. Bio-protocol, 2021, 11, e3967.	0.4	O
8	Identification and Quantification of Small RNAs. Methods in Molecular Biology, 2021, 2200, 225-254.	0.9	2
9	Multiple Quality Control Mechanisms in the ER and TGN Determine Subcellular Dynamics and Salt-Stress Tolerance Function of KORRIGAN1. Plant Cell, 2020, 32, 470-485.	6.6	21
10	Degradation of SERRATE via ubiquitin-independent 20S proteasome to survey RNA metabolism. Nature Plants, 2020, 6, 970-982.	9.3	32
11	Site-specific and substrate-specific control of accurate mRNA editing by a helicase complex in trypanosomes. Rna, 2020, 26, 1862-1881.	3 <b>.</b> 5	9
12	Lack of endoplasmic reticulum quality control (ERQC) promotes tonoplast (TP) targeting of KORRIGAN 1 (KOR1). Plant Signaling and Behavior, 2020, 15, 1744348.	2.4	0
13	$\hat{l}^2C1$ protein encoded in geminivirus satellite concertedly targets MKK2 and MPK4 to counter host defense. PLoS Pathogens, 2019, 15, e1007728.	4.7	49
14	Genome-wide probing RNA structure with the modified DMS-MaPseq in Arabidopsis. Methods, 2019, 155, 30-40.	3.8	17
15	Transactivator: A New Face of Arabidopsis AGO1. Developmental Cell, 2018, 44, 277-279.	<b>7.</b> 0	2
16	Genome-Wide Investigation of the Role of MicroRNAs in Desiccation Tolerance in the Resurrection Grass Tripogon Ioliiformis. Plants, 2018, 7, 68.	3.5	8
17	SWI2/SNF2 ATPase CHR2 remodels pri-miRNAs via Serrate to impede miRNA production. Nature, 2018, 557, 516-521.	27.8	106
18	Actions of plant Argonautes: predictable or unpredictable?. Current Opinion in Plant Biology, 2018, 45, 59-67.	7.1	46

#	Article	lF	Citations
19	The Trojan Horse of the Plant Kingdom. Cell Host and Microbe, 2018, 24, 1-3.	11.0	24
20	Tomato leaf curl Yunnan virus-encoded C4 induces cell division through enhancing stability of Cyclin D 1.1 via impairing NbSKl·-mediated phosphorylation in Nicotiana benthamiana. PLoS Pathogens, 2018, 14, e1006789.	4.7	93
21	Arabidopsis Serrate Coordinates Histone Methyltransferases ATXR5/6 and RNA Processing Factor RDR6 to Regulate Transposon Expression. Developmental Cell, 2018, 45, 769-784.e6.	7.0	50
22	KETCH1 imports HYL1 to nucleus for miRNA biogenesis in <i>Arabidopsis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 4011-4016.	7.1	70
23	The functions of plant small <scp>RNA</scp> s in development and in stress responses. Plant Journal, 2017, 90, 654-670.	5.7	198
24	Tough GC beats transgene silencing. Nature Plants, 2017, 3, 850-851.	9.3	3
25	Salt Stress and CTD PHOSPHATASE-LIKE4 Mediate the Switch between Production of Small Nuclear RNAs and mRNAs. Plant Cell, 2017, 29, 3214-3233.	6.6	13
26	Small RNA-Sequencing Links Physiological Changes and RdDM Process to Vegetative-to-Floral Transition in Apple. Frontiers in Plant Science, 2017, 8, 873.	3 <b>.</b> 6	27
27	RISC-interacting clearing 3'- 5' exoribonucleases (RICEs) degrade uridylated cleavage fragments to maintain functional RISC in Arabidopsis thaliana. ELife, 2017, 6, .	6.0	48
28	Arabidopsis AGO3 predominantly recruits 24-nt small RNAs to regulate epigenetic silencing. Nature Plants, 2016, 2, 16049.	9.3	64
29	Trehalose Accumulation Triggers Autophagy during Plant Desiccation. PLoS Genetics, 2015, 11, e1005705.	3.5	94
30	Spatiotemporal Sequestration of miR165/166 by Arabidopsis Argonaute10 Promotes Shoot Apical Meristem Maintenance. Cell Reports, 2015, 10, 1819-1827.	6.4	106
31	In vitro Reconstitution Assay of miRNA Biogenesis by Arabidopsis DCL1. Bio-protocol, 2015, 5, .	0.4	2
32	Geminivirus-encoded TrAP suppressor inhibits the histone methyltransferase SUVH4/KYP to counter host defense. ELife, 2015, 4, e06671.	6.0	92
33	Bidirectional processing of pri-miRNAs with branched terminal loops byÂArabidopsisÂDicer-like1. Nature Structural and Molecular Biology, 2013, 20, 1106-1115.	8.2	133
34	Argonautes compete for miR165/166 to regulate shoot apical meristem development. Current Opinion in Plant Biology, $2012, 15, 652-658$ .	7.1	59
35	Arabidopsis Argonaute 10 Specifically Sequesters miR166/165 to Regulate Shoot Apical Meristem Development. Cell, 2011, 145, 242-256.	28.9	420
36	Deep sequencing of small RNAs specifically associated with Arabidopsis AGO1 and AGO4 uncovers new AGO functions. Plant Journal, 2011, 67, 292-304.	5.7	114

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37	Cucumber mosaic virus-encoded 2b suppressor inhibits Arabidopsis Argonaute1 cleavage activity to counter plant defense. Genes and Development, 2006, 20, 3255-3268.	5.9	589
38	Tomato is a highly effective vehicle for expression and oral immunization with Norwalk virus capsid protein. Plant Biotechnology Journal, 2006, 4, 419-432.	8.3	113
39	Agrobacterium-mediated transformation of Arabidopsis thaliana using the floral dip method. Nature Protocols, 2006, 1, 641-646.	12.0	1,758
40	Bean Yellow Dwarf Virus replicons for high-level transgene expression in transgenic plants and cell cultures. Biotechnology and Bioengineering, 2006, 93, 271-279.	3.3	66
41	The AIP2 E3 ligase acts as a novel negative regulator of ABA signaling by promoting ABI3 degradation. Genes and Development, 2005, 19, 1532-1543.	5.9	369