

Chao Di

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12194391/publications.pdf>

Version: 2024-02-01

16
papers

14,465
citations

706676

14
h-index

1051228

16
g-index

17
all docs

17
docs citations

17
times ranked

37575
citing authors

#	ARTICLE	IF	CITATIONS
1	U1 snRNP regulates cancer cell migration and invasion in vitro. <i>Nature Communications</i> , 2020, 11, 1.	5.8	12,921
2	A Complex of U1 snRNP with Cleavage and Polyadenylation Factors Controls Telescripting, Regulating mRNA Transcription in Human Cells. <i>Molecular Cell</i> , 2019, 76, 590-599.e4.	4.5	72
3	U1 snRNP Telescripting: Suppression of Premature Transcription Termination in Introns as a New Layer of Gene Regulation. <i>Cold Spring Harbor Perspectives in Biology</i> , 2019, 11, a032235.	2.3	53
4	U1 snRNP Telescripting Roles in Transcription and Its Mechanism. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2019, 84, 115-122.	2.0	17
5	U1 snRNP telescripting regulates a size- and function-stratified human genome. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 993-999.	3.6	93
6	Critical roles of long noncoding RNAs in <i>Drosophila</i> spermatogenesis. <i>Genome Research</i> , 2016, 26, 1233-1244.	2.4	164
7	JAZ7 negatively regulates dark-induced leaf senescence in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2016, 67, 751-762.	2.4	113
8	Systematic study of novel lncRNAs in different gastrointestinal cancer cells. <i>Discovery Medicine</i> , 2016, 21, 159-71.	0.5	9
9	Rice transcriptome analysis to identify possible herbicide quinclorac detoxification genes. <i>Frontiers in Genetics</i> , 2015, 6, 306.	1.1	28
10	CLIPdb: a CLIP-seq database for protein-RNA interactions. <i>BMC Genomics</i> , 2015, 16, 51.	1.2	210
11	A common set of distinct features that characterize noncoding RNAs across multiple species. <i>Nucleic Acids Research</i> , 2015, 43, 104-114.	6.5	63
12	Comparative analysis of the transcriptome across distant species. <i>Nature</i> , 2014, 512, 445-448.	13.7	289
13	Characterization of stress-responsive lncRNAs in <i>Arabidopsis thaliana</i> by integrating expression, epigenetic and structural features. <i>Plant Journal</i> , 2014, 80, 848-861.	2.8	264
14	Down-Regulation of OsSPX1 Causes High Sensitivity to Cold and Oxidative Stresses in Rice Seedlings. <i>PLoS ONE</i> , 2013, 8, e81849.	1.1	28
15	Gene Expression Profiles Deciphering Rice Phenotypic Variation between Nipponbare (Japonica) and 93-11 (Indica) during Oxidative Stress. <i>PLoS ONE</i> , 2010, 5, e8632.	1.1	52
16	Increased expression of <i>OsSPX1</i> enhances cold/subfreezing tolerance in tobacco and <i>Arabidopsis thaliana</i> . <i>Plant Biotechnology Journal</i> , 2009, 7, 550-561.	4.1	89