

Aiwei Wu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9705196/publications.pdf>

Version: 2024-02-01

11
papers

386
citations

1307594

7
h-index

1372567

10
g-index

11
all docs

11
docs citations

11
times ranked

837
citing authors

#	ARTICLE	IF	CITATIONS
1	Co-LncRNA: investigating the lncRNA combinatorial effects in GO annotations and KEGG pathways based on human RNA-Seq data. Database: the Journal of Biological Databases and Curation, 2015, 2015, .	3.0	107
2	The correlations of tumor mutational burden among single-region tissue, multi-region tissues and blood in non-small cell lung cancer. , 2019, 7, 98.		53
3	Identification of module biomarkers from the dysregulated ceRNAâ€“ceRNA interaction network in lung adenocarcinoma. Molecular BioSystems, 2015, 11, 3048-3058.	2.9	50
4	Intratumor heterogeneity comparison among different subtypes of non-small-cell lung cancer through multi-region tissue and matched ctDNA sequencing. Molecular Cancer, 2019, 18, 7.	19.2	48
5	Extensive ceRNAâ€“ceRNA interaction networks mediated by miRNAs regulate development in multiple rhesus tissues. Nucleic Acids Research, 2016, 44, gkw587.	14.5	46
6	Integrating analysis reveals microRNA-mediated pathway crosstalk among Crohn's disease, ulcerative colitis and colorectal cancer. Molecular BioSystems, 2014, 10, 2317.	2.9	27
7	DOT1L complex regulates transcriptional initiation in human erythroleukemic cells. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	26
8	Functional dissection of virusâ€“human crosstalk mediated by miRNAs based on the VmiReg database. Molecular BioSystems, 2015, 11, 1319-1328.	2.9	12
9	Characterizing Genes with Distinct Methylation Patterns in the Context of Protein-Protein Interaction Network: Application to Human Brain Tissues. PLoS ONE, 2013, 8, e65871.	2.5	8
10	TOX4 facilitates promoter-proximal pausing and C-terminal domain dephosphorylation of RNA polymerase II in human cells. Communications Biology, 2022, 5, 300.	4.4	6
11	Integrative analysis reveals clinically relevant molecular fingerprints in pancreatic cancer. Molecular Therapy - Nucleic Acids, 2021, 26, 11-21.	5.1	3