Likun Wang

List of Publications by Year
in descending order

13 papers	4,111 citations	933447 10 h-index	1125743 13 g-index
13 all docs	13 docs citations	13 times ranked	6805 citing authors

#	Article	IF	Citations
1	DEGseq: an R package for identifying differentially expressed genes from RNA-seq data. Bioinformatics, 2010, 26, 136-138.	4.1	3,728
2	Systematic analysis of gene expression patterns associated with postmortem interval in human tissues. Scientific Reports, 2017, 7, 5435.	3.3	80
3	ASEB: a web server for KAT-specific acetylation site prediction. Nucleic Acids Research, 2012, 40, W376-W379.	14.5	72
4	Characterization and Prediction of Lysine (K)-Acetyl-Transferase Specific Acetylation Sites. Molecular and Cellular Proteomics, 2012, 11, M111.011080.	3.8	49
5	Application of Whole Exome and Targeted Panel Sequencing in the Clinical Molecular Diagnosis of 319 Chinese Families with Inherited Retinal Dystrophy and Comparison Study. Genes, 2018, 9, 360.	2.4	47
6	A mutation in ADIPOR1 causes nonsyndromic autosomal dominant retinitis pigmentosa. Human Genetics, 2016, 135, 1375-1387.	3.8	44
7	A Brief Review on Software Tools in Generating Chou's Pseudo-factor Representations for All Types of Biological Sequences. Protein and Peptide Letters, 2018, 25, 822-829.	0.9	33
8	Observations on novel splice junctions from RNA sequencing data. Biochemical and Biophysical Research Communications, 2011, 409, 299-303.	2.1	16
9	UltraPse: A Universal and Extensible Software Platform for Representing Biological Sequences. International Journal of Molecular Sciences, 2017, 18, 2400.	4.1	16
10	cisPath: an R/Bioconductor package for cloud users for visualization and management of functional protein interaction networks. BMC Systems Biology, 2015, 9, S1.	3.0	12
11	XGEM: Predicting Essential miRNAs by the Ensembles of Various Sequence-Based Classifiers With XGBoost Algorithm. Frontiers in Genetics, 2022, 13, 877409.	2.3	6
12	SoftPanel: a website for grouping diseases and related disorders for generation of customized panels. BMC Bioinformatics, 2016, 17, 153.	2.6	4
13	SGII: Systematic Identification of Essential lncRNAs in Mouse and Human Genome With lncRNA-Protein-Protein Heterogeneous Interaction Network. Frontiers in Genetics, 2022, 13, 864564.	2.3	4