

Wen-Yun Yang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12085474/publications.pdf>

Version: 2024-02-01

12
papers

816
citations

1478505

6
h-index

1588992

8
g-index

12
all docs

12
docs citations

12
times ranked

2878
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrating Functional Data to Prioritize Causal Variants in Statistical Fine-Mapping Studies. PLoS Genetics, 2014, 10, e1004722.	3.5	475
2	A model-based approach for analysis of spatial structure in genetic data. Nature Genetics, 2012, 44, 725-731.	21.4	147
3	Identification of causal genes for complex traits. Bioinformatics, 2015, 31, i206-i213.	4.1	72
4	Accounting for Population Structure in Gene-by-Environment Interactions in Genome-Wide Association Studies Using Mixed Models. PLoS Genetics, 2016, 12, e1005849.	3.5	61
5	Leveraging reads that span multiple single nucleotide polymorphisms for haplotype inference from sequencing data. Bioinformatics, 2013, 29, 2245-2252.	4.1	23
6	Spatial Localization of Recent Ancestors for Admixed Individuals. G3: Genes, Genomes, Genetics, 2014, 4, 2505-2518.	1.8	19
7	A Comparative Study on Feature Extraction from Protein Sequences for Subcellular Localization Prediction. , 2006, , .		8
8	CLASSIFICATION OF PROTEIN SEQUENCES BASED ON WORD SEGMENTATION METHODS. , 2007, , .		7
9	Incorporating cellular sorting structure for better prediction of protein subcellular locations. Journal of Experimental and Theoretical Artificial Intelligence, 2011, 23, 79-95.	2.8	2
10	A Spatial-Aware Haplotype Copying Model with Applications to Genotype Imputation. Lecture Notes in Computer Science, 2014, , 371-384.	1.3	1
11	STRING KERNELS WITH FEATURE SELECTION FOR SVM PROTEIN CLASSIFICATION. , 2007, , .		1
12	A Spatial Haplotype Copying Model with Applications to Genotype Imputation. Journal of Computational Biology, 2015, 22, 451-462.	1.6	0