

Sharlee Climer

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8488957/publications.pdf>

Version: 2024-02-01

22
papers

361
citations

840776

11
h-index

888059

17
g-index

26
all docs

26
docs citations

26
times ranked

372
citing authors

#	ARTICLE	IF	CITATIONS
1	COVID-19 and the differential dilemma. <i>Patterns</i> , 2021, 2, 100260.	5.9	2
2	Connecting the dots: The boons and banes of network modeling. <i>Patterns</i> , 2021, 2, 100374.	5.9	3
3	Linking crop traits to transcriptome differences in a progeny population of tetraploid potato. <i>BMC Plant Biology</i> , 2020, 20, 120.	3.6	18
4	Network Modeling of Complex Data Sets. <i>Methods in Molecular Biology</i> , 2020, 2096, 197-215.	0.9	5
5	Eyeing the patterns: Data visualization using doubly-seriated color heatmaps. <i>Advances in Computers</i> , 2020, 119, 121-156.	1.6	1
6	Parallel accelerated Custom Correlation Coefficient calculations for genomics applications. <i>Parallel Computing</i> , 2019, 84, 15-23.	2.1	12
7	Attacking the Opioid Epidemic: Determining the Epistatic and Pleiotropic Genetic Architectures for Chronic Pain and Opioid Addiction. , 2018, , .		29
8	Phytobiome and Transcriptional Adaptation of <i>Populus deltoides</i> to Acute Progressive Drought and Cyclic Drought. <i>Phytobiomes Journal</i> , 2018, 2, 249-260.	2.7	23
9	Latitudinal Clines of the Human Vitamin D Receptor and Skin Color Genes. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 1251-1266.	1.8	23
10	The complete parsimony haplotype inference problem and algorithms based on integer programming, branch-and-bound and Boolean satisfiability. <i>Journal of Discrete Algorithms</i> , 2016, 37, 68-83.	0.7	1
11	Moving from capstones toward cornerstones: successes and challenges in applying systems biology to identify mechanisms of autism spectrum disorders. <i>Frontiers in Genetics</i> , 2015, 6, 301.	2.3	14
12	Human gephyrin is encompassed within giant functional noncoding yin-yang sequences. <i>Nature Communications</i> , 2015, 6, 6534.	12.8	15
13	Allele-Specific Network Reveals Combinatorial Interaction That Transcends Small Effects in Psoriasis GWAS. <i>PLoS Computational Biology</i> , 2014, 10, e1003766.	3.2	25
14	A Custom Correlation Coefficient (CCC) Approach for Fast Identification of Multi-SNP Association Patterns in Genome-Wide SNPs Data. <i>Genetic Epidemiology</i> , 2014, 38, 610-621.	1.3	38
15	SplittingHeirs. , 2010, , .		2
16	NETWORK ANALYSIS OF EPISTATIC INTERACTIONS AMONG FOUR MYOCARDIAL FATTY ACID METABOLISM CANDIDATE GENES MODULATING HYPERTENSIVE HEART DISEASE. <i>Journal of the American College of Cardiology</i> , 2010, 55, A131.E1226.	2.8	0
17	How frugal is mother nature with haplotypes?. <i>Bioinformatics</i> , 2009, 25, 68-74.	4.1	10
18	Complete Parsimony Haplotype Inference Problem and Algorithms. <i>Lecture Notes in Computer Science</i> , 2009, , 337-348.	1.3	3

#	ARTICLE	IF	CITATIONS
19	Cut-and-solve: An iterative search strategy for combinatorial optimization problems. Artificial Intelligence, 2006, 170, 714-738.	5.8	60
20	Take a walk and cluster genes. , 2004, , .		12
21	Local Lines: A linear time line detector. Pattern Recognition Letters, 2003, 24, 2291-2300.	4.2	18
22	Image database indexing using JPEG coefficients. Pattern Recognition, 2002, 35, 2479-2488.	8.1	38