

Anna Ferrer-Admetlla

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11301922/publications.pdf>

Version: 2024-02-01

10
papers

1,467
citations

1040056

9
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

3151
citing authors

#	ARTICLE	IF	CITATIONS
1	An Approximate Markov Model for the Wright-Fisher Diffusion and Its Application to Time Series Data. <i>Genetics</i> , 2016, 203, 831-846.	2.9	70
2	Influenza Virus Drug Resistance: A Time-Sampled Population Genetics Perspective. <i>PLoS Genetics</i> , 2014, 10, e1004185.	3.5	126
3	On Detecting Incomplete Soft or Hard Selective Sweeps Using Haplotype Structure. <i>Molecular Biology and Evolution</i> , 2014, 31, 1275-1291.	8.9	335
4	Signatures of Environmental Genetic Adaptation Pinpoint Pathogens as the Main Selective Pressure through Human Evolution. <i>PLoS Genetics</i> , 2011, 7, e1002355.	3.5	464
5	Human pseudogenes of the ABO family show a complex evolutionary dynamics and loss of function. <i>Glycobiology</i> , 2009, 19, 583-591.	2.5	12
6	A variant in the gene FUT9 is associated with susceptibility to placental malaria infection. <i>Human Molecular Genetics</i> , 2009, 18, 3136-3144.	2.9	11
7	A Natural History of FUT2 Polymorphism in Humans. <i>Molecular Biology and Evolution</i> , 2009, 26, 1993-2003.	8.9	209
8	Evolutionary analysis of genes of two pathways involved in placental malaria infection. <i>Human Genetics</i> , 2008, 123, 343-357.	3.8	6
9	Balancing Selection Is the Main Force Shaping the Evolution of Innate Immunity Genes. <i>Journal of Immunology</i> , 2008, 181, 1315-1322.	0.8	173
10	SNP analysis to results (SNPator): a web-based environment oriented to statistical genomics analyses upon SNP data. <i>Bioinformatics</i> , 2008, 24, 1643-1644.	4.1	61