## Amin Ardeshirdavani

List of Publications by Year in descending order

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1040056 1199594 12 799 9 12 citations g-index h-index papers 17 17 17 1804 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Effective diagnosis of genetic disease by computational phenotype analysis of the disease-associated genome. Science Translational Medicine, 2014, 6, 252ra123.	12.4	223
2	Associations of Disease-Modifying Therapies With COVID-19 Severity in Multiple Sclerosis. Neurology, 2021, 97, e1870-e1885.	1.1	168
3	eXtasy: variant prioritization by genomic data fusion. Nature Methods, 2013, 10, 1083-1084.	19.0	153
4	Candidate gene prioritization with Endeavour. Nucleic Acids Research, 2016, 44, W117-W121.	14.5	111
5	Towards practical privacy-preserving genome-wide association study. BMC Bioinformatics, 2018, 19, 537.	2.6	32
6	<i>Beegle:</i> from literature mining to disease-gene discovery. Nucleic Acids Research, 2016, 44, e18-e18.	14.5	30
7	pBRIT: gene prioritization by correlating functional and phenotypic annotations through integrative data fusion. Bioinformatics, 2018, 34, 2254-2262.	4.1	24
8	NGS-Logistics: federated analysis of NGS sequence variants across multiple locations. Genome Medicine, 2014, 6, 71.	8.2	16
9	NGS-Logistics : Federated analysis of NGS sequence variants across multiple locations. Genome Medicine, 2014, 6, 71.	8.2	13
10	Multiple Sclerosis Data Alliance $\hat{a} \in A$ global multi-stakeholder collaboration to scale-up real world data research. Multiple Sclerosis and Related Disorders, 2021, 47, 102634.	2.0	11
11	Galahad: a web server for drug effect analysis from gene expression. Nucleic Acids Research, 2015, 43, W208-W212.	14.5	8
12	Systematic evaluation of NIPT aneuploidy detection software tools with clinically validated NIPT samples. PLoS Computational Biology, 2021, 17, e1009684.	3.2	6