

Brad Gulko

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

9
papers

912
citations

7
h-index

10
g-index

10
ext. papers

1,202
ext. citations

31.8
avg, IF

4.45
L-index

#	Paper	IF	Citations
9	Bayesian inference of ancient human demography from individual genome sequences. <i>Nature Genetics</i> , 2011 , 43, 1031-4	36.3	387
8	Fast, scalable prediction of deleterious noncoding variants from functional and population genomic data. <i>Nature Genetics</i> , 2017 , 49, 618-624	36.3	178
7	A method for calculating probabilities of fitness consequences for point mutations across the human genome. <i>Nature Genetics</i> , 2015 , 47, 276-83	36.3	175
6	Genome-wide inference of natural selection on human transcription factor binding sites. <i>Nature Genetics</i> , 2013 , 45, 723-9	36.3	95
5	Is a super-enhancer greater than the sum of its parts?. <i>Nature Genetics</i> , 2016 , 49, 2-3	36.3	44
4	An evolutionary framework for measuring epigenomic information and estimating cell-type-specific fitness consequences. <i>Nature Genetics</i> , 2019 , 51, 335-342	36.3	16
3	An inferred fitness consequence map of the rice genome. <i>Nature Plants</i> , 2020 , 6, 119-130	11.5	14
2	Fast, scalable prediction of deleterious noncoding variants from functional and population genomic data		2
1	How Much Information is Provided by Human Epigenomic Data? An Evolutionary View		1