

Li Song

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10789484/publications.pdf>

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10
papers

2,464
citations

933447

10
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

4585
citing authors

#	ARTICLE	IF	CITATIONS
1	Centrifuge: rapid and sensitive classification of metagenomic sequences. <i>Genome Research</i> , 2016, 26, 1721-1729.	5.5	1,025
2	Rcorrector: efficient and accurate error correction for Illumina RNA-seq reads. <i>GigaScience</i> , 2015, 4, 48.	6.4	381
3	Thousands of exon skipping events differentiate among splicing patterns in sixteen human tissues. <i>F1000Research</i> , 2013, 2, 188.	1.6	372
4	Thousands of exon skipping events differentiate among splicing patterns in sixteen human tissues. <i>F1000Research</i> , 2013, 2, 188.	1.6	293
5	Lighter: fast and memory-efficient sequencing error correction without counting. <i>Genome Biology</i> , 2014, 15, 509.	8.8	201
6	CLASS2: accurate and efficient splice variant annotation from RNA-seq reads. <i>Nucleic Acids Research</i> , 2016, 44, e98-e98.	14.5	70
7	Hybrid assembly with long and short reads improves discovery of gene family expansions. <i>BMC Genomics</i> , 2017, 18, 541.	2.8	51
8	CLASS: constrained transcript assembly of RNA-seq reads. <i>BMC Bioinformatics</i> , 2013, 14, S14.	2.6	30
9	A multi-sample approach increases the accuracy of transcript assembly. <i>Nature Communications</i> , 2019, 10, 5000.	12.8	30
10	Genomic differences between nasal <i>Staphylococcus aureus</i> from hog slaughterhouse workers and their communities. <i>PLoS ONE</i> , 2018, 13, e0193820.	2.5	11