

# Peter Edge

## 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.

6

papers

510

citations

4

h-index

6

g-index

6

ext. papers

763

ext. citations

13.2

avg, IF

4.27

L-index

#	Paper	IF	Citations
6	Widespread contribution of transposable elements to the innovation of gene regulatory networks. <i>Genome Research</i> , <b>2014</b> , 24, 1963-76	9.7	251
5	HapCUT2: robust and accurate haplotype assembly for diverse sequencing technologies. <i>Genome Research</i> , <b>2017</b> , 27, 801-812	9.7	161
4	Longshot enables accurate variant calling in diploid genomes from single-molecule long read sequencing. <i>Nature Communications</i> , <b>2019</b> , 10, 4660	17.4	69
3	Ultraaccurate genome sequencing and haplotyping of single human cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2017</b> , 114, 12512-12517	11.5	27
2	Longshot: accurate variant calling in diploid genomes using single-molecule long read sequencing		1
1	InPhaDel: integrative shotgun and proximity-ligation sequencing to phase deletions with single nucleotide polymorphisms. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, e111	20.1	1