

# Han Fang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4315792/publications.pdf>

Version: 2024-02-01

10  
papers

2,974  
citations

1039406

9  
h-index

1372195

10  
g-index

12  
all docs

12  
docs citations

12  
times ranked

7849  
citing authors

#	ARTICLE	IF	CITATIONS
1	Scikit-ribo Enables Accurate Estimation and Robust Modeling of Translation Dynamics at Codon Resolution. <i>Cell Systems</i> , 2018, 6, 180-191.e4.	2.9	41
2	Accurate detection of complex structural variations using single-molecule sequencing. <i>Nature Methods</i> , 2018, 15, 461-468.	9.0	1,175
3	Whole genome sequencing of one complex pedigree illustrates challenges with genomic medicine. <i>BMC Medical Genomics</i> , 2017, 10, 10.	0.7	15
4	GenomeScope: fast reference-free genome profiling from short reads. <i>Bioinformatics</i> , 2017, 33, 2202-2204.	1.8	1,183
5	Indel variant analysis of short-read sequencing data with Scalpel. <i>Nature Protocols</i> , 2016, 11, 2529-2548.	5.5	99
6	Genome-wide variant analysis of simplex autism families with an integrative clinical-bioinformatics pipeline. <i>Journal of Physical Education and Sports Management</i> , 2015, 1, a000422.	0.5	6
7	TAF1 Variants Are Associated with Dysmorphic Features, Intellectual Disability, and Neurological Manifestations. <i>American Journal of Human Genetics</i> , 2015, 97, 922-932.	2.6	101
8	Reducing INDEL calling errors in whole genome and exome sequencing data. <i>Genome Medicine</i> , 2014, 6, 89.	3.6	144
9	Accurate de novo and transmitted indel detection in exome-capture data using microassembly. <i>Nature Methods</i> , 2014, 11, 1033-1036.	9.0	194
10	Integrating precision medicine in the study and clinical treatment of a severely mentally ill person. <i>PeerJ</i> , 2013, 1, e177.	0.9	12