## Han Fang

List of Publications by Year in descending order

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HAN FANC

#	Article	IF	CITATIONS
1	Scikit-ribo Enables Accurate Estimation and Robust Modeling of Translation Dynamics at Codon Resolution. Cell Systems, 2018, 6, 180-191.e4.	6.2	41
2	Accurate detection of complex structural variations using single-molecule sequencing. Nature Methods, 2018, 15, 461-468.	19.0	1,175
3	Whole genome sequencing of one complex pedigree illustrates challenges with genomic medicine. BMC Medical Genomics, 2017, 10, 10.	1.5	15
4	GenomeScope: fast reference-free genome profiling from short reads. Bioinformatics, 2017, 33, 2202-2204.	4.1	1,183
5	Indel variant analysis of short-read sequencing data with Scalpel. Nature Protocols, 2016, 11, 2529-2548.	12.0	99
6	Genome-wide variant analysis of simplex autism families with an integrative clinical-bioinformatics pipeline. Journal of Physical Education and Sports Management, 2015, 1, a000422.	1.2	6
7	TAF1 Variants Are Associated with Dysmorphic Features, Intellectual Disability, and Neurological Manifestations. American Journal of Human Genetics, 2015, 97, 922-932.	6.2	101
8	Reducing INDEL calling errors in whole genome and exome sequencing data. Genome Medicine, 2014, 6, 89.	8.2	144
9	Accurate de novo and transmitted indel detection in exome-capture data using microassembly. Nature Methods, 2014, 11, 1033-1036.	19.0	194
10	Integrating precision medicine in the study and clinical treatment of a severely mentally ill person. PeerJ, 2013, 1, e177.	2.0	12