

# Yang Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/221988/publications.pdf>

Version: 2024-02-01

10  
papers

819  
citations

1307594

7  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

3878  
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparative Analysis of Brain and Fat Body Gene Splicing Patterns in the Honey Bee, <i>Apis mellifera</i> . <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1055-1063.	1.8	5
2	Allele-Specific Quantification of Structural Variations in Cancer Genomes. <i>Cell Systems</i> , 2016, 3, 21-34.	6.2	41
3	Single-molecule analysis reveals widespread structural variation in multiple myeloma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 7689-7694.	7.1	43
4	UnSplicer: mapping spliced RNA-seq reads in compact genomes and filtering noisy splicing. <i>Nucleic Acids Research</i> , 2014, 42, e25-e25.	14.5	3
5	Transcriptional regulation of autophagy by an FXR-CREB axis. <i>Nature</i> , 2014, 516, 108-111.	27.8	342
6	Search for chromosome rearrangements: New approaches toward discovery of novel translocations in head and neck squamous cell carcinoma. <i>Head and Neck</i> , 2013, 35, 831-835.	2.0	2
7	TrueSight: a new algorithm for splice junction detection using RNA-seq. <i>Nucleic Acids Research</i> , 2013, 41, e51-e51.	14.5	31
8	RNA interference knockdown of <i>DNA methyl-transferase 3</i> affects gene alternative splicing in the honey bee. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 12750-12755.	7.1	237
9	A <i>de novo</i> metagenomic assembly program for shotgun DNA reads. <i>Bioinformatics</i> , 2012, 28, 1455-1462.	4.1	42
10	FusionHunter: identifying fusion transcripts in cancer using paired-end RNA-seq. <i>Bioinformatics</i> , 2011, 27, 1708-1710.	4.1	73