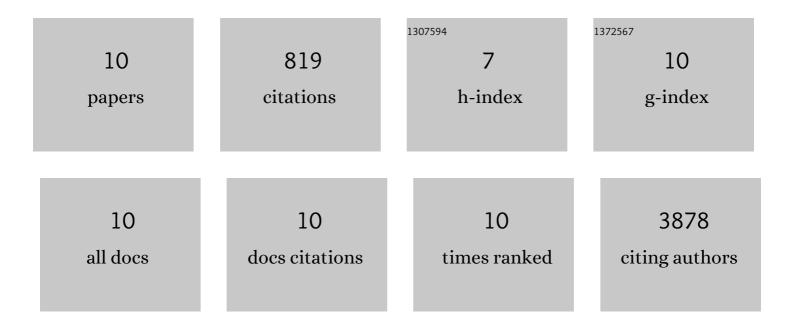


List of Publications by Year in descending order

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