

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Integrative analysis of 111 reference human epigenomes. Nature, 2015, 518, 317-330.	27.8	5,653
2	SRSF1-Regulated Alternative Splicing in Breast Cancer. Molecular Cell, 2015, 60, 105-117.	9.7	290
3	Systematic discovery of regulated and conserved alternative exons in the mammalian brain reveals NMD modulating chromatin regulators. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 3445-3450.	7.1	131
4	OLego: fast and sensitive mapping of spliced mRNA-Seq reads using small seeds. Nucleic Acids Research, 2013, 41, 5149-5163.	14.5	116
5	SpliceTrap: a method to quantify alternative splicing under single cellular conditions. Bioinformatics, 2011, 27, 3010-3016.	4.1	86
6	Chd5 orchestrates chromatin remodelling during sperm development. Nature Communications, 2014, 5, 3812.	12.8	82
7	Distinct and Predictive Histone Lysine Acetylation Patterns at Promoters, Enhancers, and Gene Bodies. G3: Genes, Genomes, Genetics, 2014, 4, 2051-2063.	1.8	39
8	Direct cloning of double-stranded RNAs from RNase protection analysis reveals processing patterns of C/D box snoRNAs and provides evidence for widespread antisense transcript expression. Nucleic Acids Research, 2011, 39, 9720-9730.	14.5	35
9	BSSF: a fingerprint based ultrafast binding site similarity search and function analysis server. BMC Bioinformatics, 2010, 11, 47.	2.6	27
10	DrugViz: a Cytoscape plugin for visualizing and analyzing small molecule drugs in biological networks. Bioinformatics, 2008, 24, 2117-2118.	4.1	14
11	SHEETSPAIR: A Database of Amino Acid Pairs in Protein Sheet Structures. Data Science Journal, 2007, 6, S589-S595.	1.3	7
12	ERPDB: An Integrated Database of ERP Data for Neuroinformatics Research. Data Science Journal, 2007, 6, S743-S748.	1.3	1
13	Wavelet Analysis of ERP Recordings for Dual Tasks in Man. , 2008, , 631-635.		0