

Periannan Senapathy

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10859287/publications.pdf>

Version: 2024-02-01

10
papers

3,138
citations

1039406

9
h-index

1372195

10
g-index

10
all docs

10
docs citations

10
times ranked

3146
citing authors

#	ARTICLE	IF	CITATIONS
1	RNA splice junctions of different classes of eukaryotes: sequence statistics and functional implications in gene expression. <i>Nucleic Acids Research</i> , 1987, 15, 7155-7174.	6.5	2,322
2	[16] Splice junctions, branch point sites, and exons: Sequence statistics, identification, and applications to genome project. <i>Methods in Enzymology</i> , 1990, 183, 252-278.	0.4	637
3	Distribution and consensus of branch point signals in eukaryotic genes: a computerized statistical analysis. <i>Nucleic Acids Research</i> , 1990, 18, 3015-3015.	6.5	81
4	EuSplice: a unified resource for the analysis of splice signals and alternative splicing in eukaryotic genes. <i>Bioinformatics</i> , 2007, 23, 1815-1823.	1.8	25
5	Mechanism of coupling periodate-oxidized nucleosides to proteins. <i>FEBS Letters</i> , 1985, 190, 337-341.	1.3	14
6	Introns and the origin of protein-coding genes. <i>Science</i> , 1995, 268, 1366-1367.	6.0	14
7	RoBuST: an integrated genomics resource for the root and bulb crop families Apiaceae and Alliaceae. <i>BMC Plant Biology</i> , 2010, 10, 161.	1.6	14
8	AspAlt: A tool for inter-database, inter-genomic and user-specific comparative analysis of alternative transcription and alternative splicing in 46 eukaryotes. <i>Genomics</i> , 2009, 94, 48-54.	1.3	13
9	Targeted Genome-Wide Enrichment of Functional Regions. <i>PLoS ONE</i> , 2010, 5, e11138.	1.1	11
10	Origination of the Split Structure of Spliceosomal Genes from Random Genetic Sequences. <i>PLoS ONE</i> , 2008, 3, e3456.	1.1	7