

MiklÅ³s CsÅ±rÅ¶s

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10101357/publications.pdf>

Version: 2024-02-01

12
papers

872
citations

1040056

9
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

1744
citing authors

#	ARTICLE	IF	CITATIONS
1	Count: evolutionary analysis of phylogenetic profiles with parsimony and likelihood. <i>Bioinformatics</i> , 2010, 26, 1910-1912.	4.1	393
2	A Detailed History of Intron-rich Eukaryotic Ancestors Inferred from a Global Survey of 100 Complete Genomes. <i>PLoS Computational Biology</i> , 2011, 7, e1002150.	3.2	160
3	Streamlining and Large Ancestral Genomes in Archaea Inferred with a Phylogenetic Birth-and-Death Model. <i>Molecular Biology and Evolution</i> , 2009, 26, 2087-2095.	8.9	114
4	Extremely Intron-Rich Genes in the Alveolate Ancestors Inferred with a Flexible Maximum-Likelihood Approach. <i>Molecular Biology and Evolution</i> , 2008, 25, 903-911.	8.9	55
5	Malin: maximum likelihood analysis of intron evolution in eukaryotes. <i>Bioinformatics</i> , 2008, 24, 1538-1539.	4.1	47
6	Reconsidering the significance of genomic word frequencies. <i>Trends in Genetics</i> , 2007, 23, 543-546.	6.7	31
7	In search of lost introns. <i>Bioinformatics</i> , 2007, 23, i87-i96.	4.1	27
8	Rapid Homology Search with Neighbor Seeds. <i>Algorithmica</i> , 2007, 48, 187-202.	1.3	14
9	Splice Sites Seldom Slide: Intron Evolution in Oomycetes. <i>Genome Biology and Evolution</i> , 2016, 8, 2340-2350.	2.5	14
10	Non-identifiability of identity coefficients at biallelic loci. <i>Theoretical Population Biology</i> , 2014, 92, 22-29.	1.1	9
11	Statistical Alignment of Retropseudogenes and Their Functional Paralogs. <i>Molecular Biology and Evolution</i> , 2005, 22, 2457-2471.	8.9	8
12	Gain-loss-duplication models for copy number evolution on a phylogeny: Exact algorithms for computing the likelihood and its gradient. <i>Theoretical Population Biology</i> , 2022, 145, 80-94.	1.1	0