

Kingdom-Wide Analysis of Fungal Protein-Coding and Patterns of Adaptive Evolution

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Tempo of Degeneration Across Independently Evolved Nonrecombining Regions. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	9
4	Evolution and codon usage bias of mitochondrial and nuclear genomes in <i>Aspergillus</i> section <i>Flavi</i> . <i>G3: Genes, Genomes, Genetics</i> , 2023, 13, .	1.8	5
5	Composition and Codon Usage Pattern Results in Divergence of the Zinc Binuclear Cluster (Zn(II) ₂ Cys ₆) Sequences among Ascomycetes Plant Pathogenic Fungi. <i>Journal of Fungi (Basel)</i> , 2023, 9, 1076.	1.0	1
6	The Evolution of tRNA Copy Number and Repertoire in Cellular Life. <i>Genes</i> , 2023, 14, 27.	2.4	7
7	De Novo Assembly and Characterization of the Transcriptome of an Omnivorous Camel Cricket (<i>Tachycines meditationis</i>). <i>International Journal of Molecular Sciences</i> , 2023, 24, 4005.	4.1	0
8	Analysis of synonymous codon usage patterns in mitochondrial genomes of nine <i>Amanita</i> species. <i>Frontiers in Microbiology</i> , 2023, 14, .	3.5	17
9	Codon Optimization Improves the Prediction of Xylose Metabolism from Gene Content in Budding Yeasts. <i>Molecular Biology and Evolution</i> , 2023, 40, .	8.9	6
12	Till death do us pair: Co-evolution of plant–necrotroph interactions. <i>Current Opinion in Plant Biology</i> , 2023, 76, 102457.	7.1	2