

TigrScan and GlimmerHMM: two open source ab initio

Bioinformatics

20, 2878-2879

DOI: [10.1093/bioinformatics/bth315](https://doi.org/10.1093/bioinformatics/bth315)

Citation Report

#	ARTICLE	IF	CITATIONS
1	An empirical analysis of training protocols for probabilistic gene finders. BMC Bioinformatics, 2004, 5, 206.	1.2	23
2	Effective statistical features for coding and non-coding DNA sequence classification for yeast, C. elegans and human. International Journal of Bioinformatics Research and Applications, 2005, 1, 181.	0.1	8
3	Excitatory cortical neurons form fine-scale functional networks. Nature, 2005, 433, 868-873.	13.7	553
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5	Genomic and Proteomic Evidence for a Second Family of Dense Core Granule Cargo Proteins in Tetrahymena thermophila. Journal of Eukaryotic Microbiology, 2005, 52, 291-297.	0.8	29
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23	Genome Sequence of Babesia bovis and Comparative Analysis of Apicomplexan Hemoprotozoa. PLoS Pathogens, 2007, 3, e148.	2.1	335
24	Following the Viterbi Path to Deduce Flagellar Actin-Interacting Proteins of Leishmania spp.: Report on Cofilins and Twinfilins. AIP Conference Proceedings, 2007, , .	0.3	3
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463	A chromosome-scale reference genome for <i>Giardia intestinalis</i> WB. <i>Scientific Data</i> , 2020, 7, 38.	2.4	69
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467	The sequence and de novo assembly of the wild yak genome. <i>Scientific Data</i> , 2020, 7, 66.	2.4	16
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471	Chromosome-level assemblies of multiple <i>Arabidopsis</i> genomes reveal hotspots of rearrangements with altered evolutionary dynamics. <i>Nature Communications</i> , 2020, 11, 989.	5.8	162
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525	The genome of <i>Tripterygium wilfordii</i> and characterization of the celastrol biosynthesis pathway. <i>GigaByte</i> , 0, 2021, 1-32.	0.0	3
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528	Genome annotation of disease-causing microorganisms. <i>Briefings in Bioinformatics</i> , 2021, 22, 845-854.	3.2	13
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549	Novel PSSM-Based Approaches for Gene Identification Using Support Vector Machine. <i>Journal of Information Technology Research</i> , 2021, 14, 152-173.	0.3	3

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560	An Atlas of Genomic Resources for Studying Rosaceae Fruits and Ornamentals. <i>Frontiers in Plant Science</i> , 2021, 12, 644881.	1.7	5
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584	The chromosome-level <i>Stevia</i> genome provides insights into steviol glycoside biosynthesis. <i>Horticulture Research</i> , 2021, 8, 129.	2.9	35
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594	Genomic Analysis of <i>Sarcomyxa edulis</i> Reveals the Basis of Its Medicinal Properties and Evolutionary Relationships. <i>Frontiers in Microbiology</i> , 2021, 12, 652324.	1.5	6
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608	Improved draft reference genome for the Glassy-winged Sharpshooter (<i>Homalodisca</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 102 Td (v	0.8	9
610	Genome assembly of <i>Scorias spongiosa</i> and comparative genomics provide insights into ecological adaptation of honeydew-dependent sooty mould fungi. <i>Genomics</i> , 2021, 113, 2189-2198.	1.3	2

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1021	Genome Resource of <i>Rhizoctonia solani</i> anastomosis group 4 strain AG4-JY, a pathomycete of sheath blight of foxtail millet. <i>Plant Disease</i> , 0, .	0.7	0
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1036	Identification of sex-linked marker and candidate sex determination gene in ornamental fish, African scat (<i>Scatophagus tetracanthus</i>). <i>Aquaculture</i> , 2023, 563, 739023.	1.7	1

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