

Resource Summary Report

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ExaBayes

RRID:SCR_024019

Type: Tool

Proper Citation

ExaBayes (RRID:SCR_024019)

Resource Information

URL: <https://cme.h-its.org/exelixis/web/software/exabayes/>

Proper Citation: ExaBayes (RRID:SCR_024019)

Description: Software package for Bayesian tree inference. Used for large-scale analyses on computer clusters.

Synonyms: exabayes

Resource Type: software resource, software toolkit

Defining Citation: [PMID:25135941](#)

Keywords: Bayesian tree inference, large-scale analyses, computer clusters,

Availability: Free, Available for download, Freely available

Resource Name: ExaBayes

Resource ID: SCR_024019

Alternate IDs: OMICS_10303

Old URLs: <https://sources.debian.org/src/exabayes/>

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260821T194247+0000

Ratings and Alerts

No rating or validation information has been found for ExaBayes.

No alerts have been found for ExaBayes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Kovacs TGL, et al. (2024) Dating in the Dark: Elevated Substitution Rates in Cave Cockroaches (Blattodea: Nocticolidae) Have Negative Impacts on Molecular Date Estimates. *Systematic biology*, 73(3), 532.

Rossitto De Marchi B, et al. (2020) Genetic diversity and SNP's from the chloroplast coding regions of virus-infected cassava. *PeerJ*, 8, e8632.

Gan HM, et al. (2019) Two reads to rule them all: Nanopore long read-guided assembly of the iconic Christmas Island red crab, *Gecarcoidea natalis* (Pocock, 1888), mitochondrial genome and the challenges of AT-rich mitogenomes. *Marine genomics*, 45, 64.

Twort VG, et al. (2019) New Zealand Tree and Giant W?t? (Orthoptera) Transcriptomics Reveal Divergent Selection Patterns in Metabolic Loci. *Genome biology and evolution*, 11(4), 1293.

Platt RN, et al. (2018) Conflicting Evolutionary Histories of the Mitochondrial and Nuclear Genomes in New World Myotis Bats. *Systematic biology*, 67(2), 236.

Branstetter MG, et al. (2017) Phylogenomic Insights into the Evolution of Stinging Wasps and the Origins of Ants and Bees. *Current biology : CB*, 27(7), 1019.

Yerramsetty P, et al. (2016) Evolution of RLSB, a nuclear-encoded S1 domain RNA binding protein associated with post-transcriptional regulation of plastid-encoded rbcL mRNA in vascular plants. *BMC evolutionary biology*, 16(1), 141.