Resource Summary Report

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Chromaseq

RRID:SCR_005587
Type: Tool

Proper Citation

Chromaseq (RRID:SCR_005587)

Resource Information

URL: http://mesquiteproject.org/packages/chromaseq/

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Description: A software package in Mesquite that processes chromatograms, makes contigs, base calls, etc., using in part the programs Phred and Phrap.

Abbreviations: Chromaseq

Synonyms: Chromaseq: a package for processing chromatograms and sequence data in Mesquite

Resource Type: software resource

Keywords: chromatogram, sequence, mesquite

Funding: NSF EF-0531754

Availability: Acknowledgement required

Resource Name: Chromaseq

Resource ID: SCR_005587

Alternate IDs: OMICS_01017

Record Creation Time: 20220129T080231+0000

Record Last Update: 20260725T190609+0000

Ratings and Alerts

No rating or validation information has been found for Chromaseq.

No alerts have been found for Chromaseq.

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 nidm-terms .

Martinez-Hernandez F, et al. (2023) Genetic Variability of the Internal Transcribed Spacer and Pyruvate:Ferredoxin Oxidoreductase Partial Gene of Trichomonas vaginalis from Female Patients. Microorganisms, 11(9).

Kavanaugh DH, et al. (2021) Phylogeny of the supertribe Nebriitae (Coleoptera, Carabidae) based on analyses of DNA sequence data. ZooKeys, 1044, 41.

Huang YL, et al. (2020) Effect of Host, Environment and Fungal Growth on Fungal Leaf Endophyte Communities in Taiwan. Journal of fungi (Basel, Switzerland), 6(4).

Villanueva-Garcia C, et al. (2017) Clarifying the Cryptic Host Specificity of Blastocystis spp. Isolates from Alouatta palliata and A. pigra Howler Monkeys. PloS one, 12(1), e0169637.

Brunet T, et al. (2016) The evolutionary origin of bilaterian smooth and striated myocytes. eLife, 5.

Maddison WP, et al. (2016) Phylogenetic placement of the unusual jumping spider Depreissia Lessert, and a new synapomorphy uniting Hisponinae and Salticinae (Araneae, Salticidae). ZooKeys(549), 1.

Kanda K, et al. (2015) Successful Recovery of Nuclear Protein-Coding Genes from Small Insects in Museums Using Illumina Sequencing. PloS one, 10(12), e0143929.