Resource Summary Report

Generated by NIF on Apr 27, 2025

Chromas

RRID:SCR_000598

Type: Tool

Proper Citation

Chromas (RRID:SCR_000598)

Resource Information

URL: http://technelysium.com.au/?page_id=13

Proper Citation: Chromas (RRID:SCR_000598)

Description: Software ideal for the most basic of sequencing projects, where assembly of

multiple sequences is not required.

Abbreviations: Chromas

Resource Type: commercial organization, software resource

Keywords: sequencing, bio.tools

Funding:

Availability: Commercial license, Free trial

Resource Name: Chromas

Resource ID: SCR_000598

Alternate IDs: OMICS_01016, biotools:chromas

Alternate URLs: https://bio.tools/chromas

Record Creation Time: 20220129T080202+0000

Record Last Update: 20250426T055421+0000

Ratings and Alerts

No rating or validation information has been found for Chromas.

No alerts have been found for Chromas.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 14 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Sonnemann HM, et al. (2024) Placental co-transcriptional activator Vestigial-like 1 (VGLL1) drives tumorigenesis via increasing transcription of proliferation and invasion genes. Frontiers in oncology, 14, 1403052.

Loginova OA, et al. (2023) First report of Orthostrongylus sp. (Nematoda: Protostrongylidae) in wild reindeer (Rangifer tarandus) from the Taimyr, Russia: Nearctic parasites in a Palearctic host. Parasitology research, 122(2), 685.

Trivieri N, et al. (2022) Growth factor independence underpins a paroxysmal, aggressive Wnt5aHigh/EphA2Low phenotype in glioblastoma stem cells, conducive to experimental combinatorial therapy. Journal of experimental & clinical cancer research: CR, 41(1), 139.

Mazurek S, et al. (2021) Disruption of RING and PHD Domains of TRIM28 Evokes Differentiation in Human iPSCs. Cells, 10(8).

Carraturo F, et al. (2018) Comparative assessment of the quality of commercial black and green tea using microbiology analyses. BMC microbiology, 18(1), 4.

da Silva EC, et al. (2017) Cryptococcus gattii molecular type VGII infection associated with lung disease in a goat. BMC veterinary research, 13(1), 41.

Shum P, et al. (2017) Harnessing mtDNA variation to resolve ambiguity in 'Redfish' sold in Europe. PeerJ, 5, e3746.

Illera JC, et al. (2017) Factors governing the prevalence and richness of avian haemosporidian communities within and between temperate mountains. PloS one, 12(9), e0184587.

De Castro O, et al. (2017) What is in your cup of tea? DNA Verity Test to characterize black and green commercial teas. PloS one, 12(5), e0178262.

Grunin M, et al. (2016) Transcriptome Analysis on Monocytes from Patients with Neovascular Age-Related Macular Degeneration. Scientific reports, 6, 29046.

De Castro O, et al. (2016) Disentangling Phylogenetic Relationships in a Hotspot of Diversity: The Butterworts (Pinguicula L., Lentibulariaceae) Endemic to Italy. PloS one, 11(12), e0167610.

Wang S, et al. (2015) A Single-Nucleotide Polymorphism of TaGS5 Gene Revealed its Association with Kernel Weight in Chinese Bread Wheat. Frontiers in plant science, 6, 1166.

Shum P, et al. (2015) Three-dimensional post-glacial expansion and diversification of an exploited oceanic fish. Molecular ecology, 24(14), 3652.

Avin FA, et al. (2014) Molecular divergence and species delimitation of the cultivated oyster mushrooms: integration of IGS1 and ITS. TheScientificWorldJournal, 2014, 793414.