

Resource Summary Report

Generated by [NIF](#) on Aug 28, 2026

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., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: Chromas

Resource Type: commercial organization, software resource

Keywords: sequencing, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Chromas

Resource ID: SCR_000598

Alternate IDs: OMICS_01016, biotools:chromas

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

License: Commercial license

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260820T054312+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 16 mentions in open access literature.

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

Wang M, et al. (2024) Genomic insights into Neolithic founding paternal lineages around the Qinghai-Xizang Plateau using integrated YanHuang resource. *iScience*, 27(12), 111456.

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 O, et al. (2024) *Fasciola hepatica*: Updates on egg morphology, host range, and distribution. *Food and waterborne parasitology*, 36, e00237.

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 Wnt5a^{High}/EphA2^{Low} 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.

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.

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.

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.