

Resource Summary Report

Generated by T1D on Sep 2, 2026

ChromasPro

RRID:SCR_000229

Type: Tool

Proper Citation

ChromasPro (RRID:SCR_000229)

Resource Information

URL: http://technelysium.com.au/?page_id=27

Proper Citation: ChromasPro (RRID:SCR_000229)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 31,2023. Software which is able to assemble data from 454 and Illumina next-generation sequencers, with up to 100,000 sequences if 2 Gb RAM is available.

Resource Type: data acquisition software, data processing software, software application, software resource

Keywords: genome, sequence, dna, assemble, data, illumina, next gen sequence, next generation

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ChromasPro

Resource ID: SCR_000229

Alternate IDs: OMICS_01808

License: Licenses for purchase. Options for packages of licenses or single-user installations. Acceptance of license terms allows for use of software for up to 30 days, after which payment must be made to the vendor.

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260829T182021+0000

Ratings and Alerts

No rating or validation information has been found for ChromasPro.

No alerts have been found for ChromasPro.

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 [T1D](#).

Salehipour P, et al. (2023) CRISPR-Based Fluorescent Reporter (CBFR) Assay for Sensitive, Specific, Inexpensive, and Visual Detection of a Specific EGFR Exon 19 Deletion in NSCLC. *Molecular biotechnology*, 65(5), 807.

Gattoni G, et al. (2022) Genomic study and lipidomic bioassay of *Leeuwenhoekiella parthenopeia*: A novel rare biosphere marine bacterium that inhibits tumor cell viability. *Frontiers in microbiology*, 13, 1090197.

Testini C, et al. (2019) Myc-dependent endothelial proliferation is controlled by phosphotyrosine 1212 in VEGF receptor-2. *EMBO reports*, 20(11), e47845.

Diawara I, et al. (2018) Analysis of amino acid motif of penicillin-binding proteins 1a, 2b, and 2x in invasive *Streptococcus pneumoniae* nonsusceptible to penicillin isolated from pediatric patients in Casablanca, Morocco. *BMC research notes*, 11(1), 632.

Alari V, et al. (2018) Generation of the Rubinstein-Taybi syndrome type 2 patient-derived induced pluripotent stem cell line (IAli001-A) carrying the EP300 exon 23 stop mutation c.3829A>?T, p.(Lys1277*). *Stem cell research*, 30, 175.

Pawlik A, et al. (2016) Purification and characterization of laccase from *Sinorhizobium meliloti* and analysis of the lacc gene. *International journal of biological macromolecules*, 92, 138.

Clemente A, et al. (2015) Increased STAT3 phosphorylation on CD27(+) B-cells from common variable immunodeficiency disease patients. *Clinical immunology (Orlando, Fla.)*, 161(2), 77.

Amer S, et al. (2015) Molecular characterization of *Echinococcus granulosus sensu lato* from farm animals in Egypt. *PloS one*, 10(3), e0118509.

Tongo M, et al. (2015) Near full-length HIV type 1M genomic sequences from Cameroon : Evidence of early diverging under-sampled lineages in the country. *Evolution, medicine, and public health*, 2015(1), 254.

Okeke MI, et al. (2014) Molecular characterization and phylogenetics of Fennoscandian cowpox virus isolates based on the p4c and atip genes. *Virology journal*, 11, 119.

Tajer Mohammad Ghazvini P, et al. (2014) Isolation and Characterization of a Novel Magnetotactic Bacterium From Iran: Iron Uptake and Producing Magnetic Nanoparticles in

Alphaproteobacterium MTB-KTN90. Jundishapur journal of microbiology, 7(9), e19343.

Cardoso L, et al. (2014) Molecular and histopathological detection of *Hepatozoon canis* in red foxes (*Vulpes vulpes*) from Portugal. *Parasites & vectors*, 7, 113.

Wang L, et al. (2013) Concurrent infections of *Giardia duodenalis*, *Enterocytozoon bieneusi*, and *Clostridium difficile* in children during a cryptosporidiosis outbreak in a pediatric hospital in China. *PLoS neglected tropical diseases*, 7(9), e2437.

Scott AP, et al. (2011) Recombination mapping of the susceptibility region for sporadic inclusion body myositis within the major histocompatibility complex. *Journal of neuroimmunology*, 235(1-2), 77.