Resource Summary Report

Generated by RRID on Sep 10, 2026

OpenBioinformatics.org

RRID:SCR_002229
Type: Tool

Proper Citation

OpenBioinformatics.org (RRID:SCR_002229)

Resource Information

URL: http://www.openbioinformatics.org/

Proper Citation: OpenBioinformatics.org (RRID:SCR_002229)

Description: An open bioinformatis software repository with no tie to any organization or institution. Contact them to host your software.

Abbreviations: OpenBioinformatics.org

Synonyms: OpenBioinformatics.org: An open bioinformatis software repository

Resource Type: software repository, software resource

Keywords: bioinformatics, software

Availability: Free, Available for download, Freely available

Resource Name: OpenBioinformatics.org

Resource ID: SCR_002229

Alternate IDs: nlx_155524

License: Open unspecified license

Record Creation Time: 20220129T080212+0000

Record Last Update: 20260905T133226+0000

Ratings and Alerts

No rating or validation information has been found for OpenBioinformatics.org.

No alerts have been found for OpenBioinformatics.org.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Moradi MH, et al. (2022) Genome-wide evaluation of copy gain and loss variations in three Afghan sheep breeds. Scientific reports, 12(1), 14286.

Yamasaki M, et al. (2020) Sensitivity to gene dosage and gene expression affects genes with copy number variants observed among neuropsychiatric diseases. BMC medical genomics, 13(1), 55.

Bushehri A, et al. (2019) Identification of PROS1 as a Novel Candidate Gene for Juvenile Retinitis Pigmentosa. International journal of molecular and cellular medicine, 8(3), 179.

Yao G, et al. (2019) Identification of a novel mutation of FGFR3 gene in a large Chinese pedigree with hypochondroplasia by next-generation sequencing: A case report and brief literature review. Medicine, 98(4), e14157.

Dorjbal B, et al. (2019) Hypomorphic caspase activation and recruitment domain 11 (CARD11) mutations associated with diverse immunologic phenotypes with or without atopic disease. The Journal of allergy and clinical immunology, 143(4), 1482.

Staninova-Stojovska M, et al. (2019) Molecular Basis of Inherited Colorectal Carcinomas in the Macedonian Population: An Update. Balkan journal of medical genetics : BJMG, 22(2), 5.

Chen Y, et al. (2018) A novel compound heterozygous variant of the SLC12A3 gene in Gitelman syndrome pedigree. BMC medical genetics, 19(1), 17.

Jeong JE, et al. (2017) The association between the nicotinic acetylcholine receptor ?4 subunit gene (CHRNA4) rs1044396 and Internet gaming disorder in Korean male adults. PloS one, 12(12), e0188358.

Todd JN, et al. (2015) Genetic Evidence for a Causal Role of Obesity in Diabetic Kidney Disease. Diabetes, 64(12), 4238.

Scheidecker S, et al. (2015) Mutations in TUBGCP4 alter microtubule organization via the ?-tubulin ring complex in autosomal-recessive microcephaly with chorioretinopathy. American journal of human genetics, 96(4), 666.

Liu Z, et al. (2010) MixHMM: inferring copy number variation and allelic imbalance using SNP arrays and tumor samples mixed with stromal cells. PloS one, 5(6), e10909.