

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

## OpenBioinformatics.org

RRID:SCR\_002229

Type: Tool

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### Proper Citation

OpenBioinformatics.org (RRID:SCR\_002229)

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### Resource Information

**URL:** <http://www.openbioinformatics.org/>

**Proper Citation:** OpenBioinformatics.org (RRID:SCR\_002229)

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

**Abbreviations:** OpenBioinformatics.org

**Synonyms:** OpenBioinformatics.org: An open bioinformatics 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

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### 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 [PRECISE-TBI](#) .

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  $\gamma 4$  subunit gene (CHRNA4) rs1044396 and Internet gaming disorder in Korean male adults. PloS one, 12(12), e0188358.

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

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

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.