

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

ChroMoS

RRID:SCR_008320

Type: Tool

Proper Citation

ChroMoS (RRID:SCR_008320)

Resource Information

URL: <http://epicenter.immunbio.mpg.de/services/chromos/>

Proper Citation: ChroMoS (RRID:SCR_008320)

Description: Combines genetic and epigenetic data to facilitate SNP classification, prioritization and prediction of their functional effect.

Abbreviations: ChroMoS

Synonyms: Chromatin Modified SNPs

Resource Type: software resource

Defining Citation: [PMID:23782616](#)

Keywords: Chromatin, modified SNPs, SNPs, genetic and epigenetic data combination

Availability: Free, Freely available

Resource Name: ChroMoS

Resource ID: SCR_008320

Alternate IDs: OMICS_00145

Record Creation Time: 20220129T080246+0000

Record Last Update: 20260725T190651+0000

Ratings and Alerts

No rating or validation information has been found for ChroMoS.

No alerts have been found for ChroMoS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Cheng Z, et al. (2020) From Folk Taxonomy to Species Confirmation of Acorus (Acoraceae): Evidences Based on Phylogenetic and Metabolomic Analyses. Frontiers in plant science, 11, 965.

Xiao X, et al. (2017) Molecular mechanisms underlying noncoding risk variations in psychiatric genetic studies. Molecular psychiatry, 22(4), 497.

Skibola CF, et al. (2014) Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. American journal of human genetics, 95(4), 462.

Edwards SL, et al. (2013) Beyond GWASs: illuminating the dark road from association to function. American journal of human genetics, 93(5), 779.

Barizzone N, et al. (2013) Rare variants in the TREX1 gene and susceptibility to autoimmune diseases. BioMed research international, 2013, 471703.