Resource Summary Report

Generated by RRID on Sep 10, 2026

CRLMM

RRID:SCR_012580
Type: Tool

Proper Citation

CRLMM (RRID:SCR_012580)

Resource Information

URL: http://www.bioconductor.org/packages/2.12/bioc/html/crlmm.html

Proper Citation: CRLMM (RRID:SCR_012580)

Description: Genotype Calling and Copy Number Analysis tool for Affymetrix SNP 5.0 and 6.0 and Illumina arrays.

Abbreviations: CRLMM

Resource Type: software resource

Resource Name: CRLMM

Resource ID: SCR_012580

Alternate IDs: OMICS_00717

Record Creation Time: 20220129T080311+0000

Record Last Update: 20260905T132725+0000

Ratings and Alerts

No rating or validation information has been found for CRLMM.

No alerts have been found for CRLMM.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Zorn S, et al. (2025) Obesity due to MC4R deficiency is associated with reduced cholesterol, triglycerides and cardiovascular disease risk. Nature medicine, 31(12), 4180.

Hamdi Y, et al. (2018) A genome wide SNP genotyping study in the Tunisian population: specific reporting on a subset of common breast cancer risk loci. BMC cancer, 18(1), 1295.

Jmel H, et al. (2018) Pharmacogenetic landscape of Metabolic Syndrome components drug response in Tunisia and comparison with worldwide populations. PloS one, 13(4), e0194842.

Matovinovic E, et al. (2017) Genome-wide linkage and association analysis of primary open-angle glaucoma endophenotypes in the Norfolk Island isolate. Molecular vision, 23, 660.

Bros??us J, et al. (2016) Relapsed diffuse large B-cell lymphoma present different genomic profiles between early and late relapses. Oncotarget, 7(51), 83987.

Mather KA, et al. (2016) Genome-wide significant results identified for plasma apolipoprotein H levels in middle-aged and older adults. Scientific reports, 6, 23675.

Benton MC, et al. (2015) Serum bilirubin concentration is modified by UGT1A1 haplotypes and influences risk of type-2 diabetes in the Norfolk Island genetic isolate. BMC genetics, 16, 136.

Benton MC, et al. (2015) A Phenomic Scan of the Norfolk Island Genetic Isolate Identifies a Major Pleiotropic Effect Locus Associated with Metabolic and Renal Disorder Markers. PLoS genetics, 11(10), e1005593.

Mather KA, et al. (2015) Investigating the genetics of hippocampal volume in older adults without dementia. PloS one, 10(1), e0116920.

Kanthaswamy S, et al. (2013) Identifying human-rhesus macaque gene orthologs using heterospecific SNP probes. Genomics, 101(1), 30.