

Resource Summary Report

Generated by [Kravitz](#) on Sep 18, 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: 20260912T195750+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 [Kravitz](#) .

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.

Broschus 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.

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

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.

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