

# Resource Summary Report

Generated by [dkNET](#) on Aug 24, 2026

## CRLMM

RRID:SCR\_012580

Type: Tool

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### Proper Citation

CRLMM (RRID:SCR\_012580)

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### 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:** 20260815T182431+0000

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### Ratings and Alerts

No rating or validation information has been found for CRLMM.

No alerts have been found for CRLMM.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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### Usage and Citation Metrics

We found 10 mentions in open access literature.

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

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.