# **Resource Summary Report**

Generated by <u>NIF</u> on Apr 28, 2025

# **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

Funding:

Resource Name: CRLMM

Resource ID: SCR\_012580

Alternate IDs: OMICS\_00717

Record Creation Time: 20220129T080311+0000

Record Last Update: 20250420T014615+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 9 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

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 openangle glaucoma endophenotypes in the Norfolk Island isolate. Molecular vision, 23, 660.

Broséus 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.