

Resource Summary Report

Generated by [nidm-terms](#) on Aug 9, 2026

ILLUMINUS

RRID:SCR_000388

Type: Tool

Proper Citation

ILLUMINUS (RRID:SCR_000388)

Resource Information

URL: <https://github.com/wtsi-npg/Illuminus>

Proper Citation: ILLUMINUS (RRID:SCR_000388)

Description: A fast and accurate algorithm for assigning single nucleotide polymorphism (SNP) genotypes to microarray data from the Illumina BeadArray technology.

Abbreviations: Illuminus

Synonyms: Illuminus: the genotype calling algorithm

Resource Type: software resource, software application

Defining Citation: [PMID:17846035](#)

Keywords: gene, genetic, genomic, c++, single nucleotide polymorphism, genotype, microarray, illumina beadarray, illumina

Availability: Free, Available for download, Freely available

Resource Name: ILLUMINUS

Resource ID: SCR_000388

Alternate IDs: OMICS_00726, nlx_154408

Alternate URLs: <http://www.sanger.ac.uk/resources/software/illuminus/>

Old URLs: <http://www.sanger.ac.uk/science/tools/illuminus>

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260808T190446+0000

Ratings and Alerts

No rating or validation information has been found for ILLUMINUS.

No alerts have been found for ILLUMINUS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.