

# Resource Summary Report

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## 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 application, software resource

**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:** 20260905T133221+0000

### Ratings and Alerts

No rating or validation information has been found for ILLUMINUS.

No alerts have been found for ILLUMINUS.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.