

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

Bycom

RRID:SCR_000659

Type: Tool

Proper Citation

Bycom (RRID:SCR_000659)

Resource Information

URL: <http://sourceforge.net/projects/bycom/>

Proper Citation: Bycom (RRID:SCR_000659)

Description: A software which can perform methylcytosine calling from BS-seq (WGBS and RRBS), and permits either unmapped reads (FASTQ) or mapped reads (SAM/BAM) to be used as the input data. Certain SNPs (C>A/G) can also be selected in the output.

Resource Type: software resource

Defining Citation: [PMID:25255082](#)

Keywords: methylcytosine, bs-seq, fastq, sam, bam, snps, snp, wgbs, rrbs, sorftw

Availability: Free, Available for download, Freely available

Resource Name: Bycom

Resource ID: SCR_000659

Alternate IDs: OMICS_00594

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260725T191024+0000

Ratings and Alerts

No rating or validation information has been found for Bycom.

No alerts have been found for Bycom.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.