

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

BlindCall

RRID:SCR_001280

Type: Tool

Proper Citation

BlindCall (RRID:SCR_001280)

Resource Information

URL: <http://cbcb.umd.edu/~hcorrada/secgen/>

Proper Citation: BlindCall (RRID:SCR_001280)

Description: Software for ultra-fast base-calling of second-generation sequencing data by blind deconvolution.

Abbreviations: BlindCall

Resource Type: software resource

Defining Citation: [PMID:24413520](#)

Keywords: base-calling, second-generation sequencing, blind deconvolution

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: BlindCall

Resource ID: SCR_001280

Alternate IDs: OMICS_02216

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260808T185727+0000

Ratings and Alerts

No rating or validation information has been found for BlindCall.

No alerts have been found for BlindCall.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Wang B, et al. (2017) An adaptive decorrelation method removes Illumina DNA base-calling errors caused by crosstalk between adjacent clusters. Scientific reports, 7, 41348.