

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

JointSLM

RRID:SCR_010823

Type: Tool

Proper Citation

JointSLM (RRID:SCR_010823)

Resource Information

URL: <http://seqanswers.com/wiki/JointSLM>

Proper Citation: JointSLM (RRID:SCR_010823)

Description: Copy number estimation from read depth information.

Abbreviations: JointSLM

Resource Type: software resource

Resource Name: JointSLM

Resource ID: SCR_010823

Alternate IDs: OMICS_00346

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260725T190703+0000

Ratings and Alerts

No rating or validation information has been found for JointSLM.

No alerts have been found for JointSLM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Luo F, et al. (2019) A systematic evaluation of copy number alterations detection methods on real SNP array and deep sequencing data. BMC bioinformatics, 20(Suppl 25), 692.

Dharanipragada P, et al. (2018) iCopyDAV: Integrated platform for copy number variations-Detection, annotation and visualization. PloS one, 13(4), e0195334.