

Resource Summary Report

Generated by [NIF](#) on Aug 11, 2026

quantsmooth

RRID:SCR_001271

Type: Tool

Proper Citation

quantsmooth (RRID:SCR_001271)

Resource Information

URL: <http://bioconductor.org/packages/release/bioc/html/quantsmooth.html>

Proper Citation: quantsmooth (RRID:SCR_001271)

Description: Software package for quantile smoothing and genomic visualization of array data.

Abbreviations: quantsmooth

Resource Type: software resource

Defining Citation: [PMID:15572474](#)

Keywords: copy number variation, visualization, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: quantsmooth

Resource ID: SCR_001271

Alternate IDs: OMICS_02067, BioTools:quantsmooth, biotools:quantsmooth

Alternate URLs: <https://bio.tools/quantsmooth>, <https://bio.tools/quantsmooth>, <https://bio.tools/quantsmooth>

License: GNU General Public License, v3

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260808T185725+0000

Ratings and Alerts

No rating or validation information has been found for quantsmooth.

No alerts have been found for quantsmooth.

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 [NIF](#) .

Law R, et al. (2014) Biallelic truncating mutations in FMN2, encoding the actin-regulatory protein Formin 2, cause nonsyndromic autosomal-recessive intellectual disability. American journal of human genetics, 95(6), 721.