

# Resource Summary Report

Generated by [dkNET](#) on Apr 26, 2025

## [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

**Funding:**

**Availability:** GNU General Public License, v2

**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>

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20250420T014024+0000

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## Ratings and Alerts

No rating or validation information has been found for quantsmooth.

No alerts have been found for quantsmooth.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.