

Resource Summary Report

Generated by RRID on Aug 2, 2026

CiLiQuant

RRID:SCR_019319

Type: Tool

Proper Citation

CiLiQuant (RRID:SCR_019319)

Resource Information

URL: <https://github.com/almorlio/CiLiQuant>

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Description: Software tool to separate junction reads based on their linear or circular origin. Only non ambiguous junction reads are used to compare relative linear and circular transcript abundance.

Resource Type: software resource, software application, data processing software

Keywords: RNA, splicing, circular origin, separate junction reads, circular transcript abundance, linear transcript abundance, compare, bio.tools

Funding: FWO ;
Special Research Fund UGent ;
Stichting Tegen Kanker ;
Kom Op Tegen Kanker (Stand Up To Cancer) ;
European Union's Horizon 2020

Availability: Free, Available for download, Freely available

Resource Name: CiLiQuant

Resource ID: SCR_019319

Alternate IDs: biotools:ciliquant

Alternate URLs: <https://bio.tools/ciliquant>

License: BSD 3-Clause License

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260801T190846+0000

Ratings and Alerts

No rating or validation information has been found for CiLiQuant.

No alerts have been found for CiLiQuant.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.