Resource Summary Report

Generated by dkNET on May 18, 2025

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: 20250513T062048+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.