# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 21, 2025

# CQN

RRID:SCR 001786

Type: Tool

## **Proper Citation**

CQN (RRID:SCR\_001786)

### **Resource Information**

URL: http://www.bioconductor.org/packages/2.13/bioc/html/cqn.html

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**Description:** A normalization tool for RNA-Seq data, implementing the conditional quantile

normalization method.

**Abbreviations:** CQN

**Synonyms:** Conditional Quantile Normalization

**Resource Type:** software resource

**Defining Citation: PMID:22285995** 

**Keywords:** rna-seq, differential expression, preprocessing, bio.tools

**Funding:** 

Availability: Artistic License, v3

Resource Name: CQN

Resource ID: SCR\_001786

Alternate IDs: OMICS 01949, biotools:cqn

Alternate URLs: https://bio.tools/cqn

**Record Creation Time:** 20220129T080209+0000

Record Last Update: 20250420T014039+0000

## **Ratings and Alerts**

No rating or validation information has been found for CQN.

No alerts have been found for CQN.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.

Hosseinzadeh L, et al. (2024) The androgen receptor interacts with GATA3 to transcriptionally regulate a luminal epithelial cell phenotype in breast cancer. Genome biology, 25(1), 44.

Park MH, et al. (2019) CCN1 interlinks integrin and hippo pathway to autoregulate tip cell activity. eLife, 8.

de la Torre-Ubieta L, et al. (2018) The Dynamic Landscape of Open Chromatin during Human Cortical Neurogenesis. Cell, 172(1-2), 289.

Grand Moursel L, et al. (2018) Brain Transcriptomic Analysis of Hereditary Cerebral Hemorrhage With Amyloidosis-Dutch Type. Frontiers in aging neuroscience, 10, 102.

Hsu SC, et al. (2017) The BET Protein BRD2 Cooperates with CTCF to Enforce Transcriptional and Architectural Boundaries. Molecular cell, 66(1), 102.

Amabile A, et al. (2016) Inheritable Silencing of Endogenous Genes by Hit-and-Run Targeted Epigenetic Editing. Cell, 167(1), 219.