### **Resource Summary Report**

Generated by dkNET on Apr 24, 2025

# **TurboNorm**

RRID:SCR\_012963 Type: Tool

#### **Proper Citation**

TurboNorm (RRID:SCR\_012963)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/2.12/bioc/html/TurboNorm.html

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**Description:** Software providing a fast scatterplot smoother suitable for microarray normalization based on B-splines with second-order difference penalty. Functions for microarray normalization of single-colour data i.e. Affymetrix/Illumina and two-colour data supplied as marray MarrayRaw-objects or limma RGList-objects are available.

Abbreviations: TurboNorm

Resource Type: software resource

Funding:

Resource Name: TurboNorm

Resource ID: SCR\_012963

Alternate IDs: OMICS\_00856

Record Creation Time: 20220129T080313+0000

Record Last Update: 20250420T014626+0000

#### **Ratings and Alerts**

No rating or validation information has been found for TurboNorm.

No alerts have been found for TurboNorm.

### Data and Source Information

Source: SciCrunch Registry

## Usage and Citation Metrics

We have not found any literature mentions for this resource.