

Resource Summary Report

Generated by [NIF](#) on Aug 30, 2026

FDM

RRID:SCR_006733

Type: Tool

Proper Citation

FDM (RRID:SCR_006733)

Resource Information

URL: <http://csbio-linux001.cs.unc.edu/nextgen/software/FDM/>

Proper Citation: FDM (RRID:SCR_006733)

Description: A graph-based statistical method to detect differential transcription using RNA-seq data.

Abbreviations: FDM

Resource Type: software resource

Resource Name: FDM

Resource ID: SCR_006733

Alternate IDs: OMICS_01332

Record Creation Time: 20220129T080237+0000

Record Last Update: 20260829T182241+0000

Ratings and Alerts

No rating or validation information has been found for FDM.

No alerts have been found for FDM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Singh D, et al. (2011) FDM: a graph-based statistical method to detect differential transcription using RNA-seq data. Bioinformatics (Oxford, England), 27(19), 2633.