

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[FX](#)

RRID:SCR_011902

Type: Tool

Proper Citation

FX (RRID:SCR_011902)

Resource Information

URL: <http://fx.gmi.ac.kr/>

Proper Citation: FX (RRID:SCR_011902)

Description: A user-Frendly RNA-Seq gene eXpression analysis tool, empowered by the concept of cloud-computing.

Abbreviations: FX

Resource Type: software resource

Defining Citation: [PMID:22257667](#)

Keywords: mapreduce/hadoop

Resource Name: FX

Resource ID: SCR_011902

Alternate IDs: OMICS_01403

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260801T190423+0000

Ratings and Alerts

No rating or validation information has been found for FX.

No alerts have been found for FX.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Yu P, et al. (2016) Single-cell Transcriptome Study as Big Data. Genomics, proteomics & bioinformatics, 14(1), 21.

Chen J, et al. (2013) Translational biomedical informatics in the cloud: present and future. BioMed research international, 2013, 658925.

Lin YC, et al. (2013) Enabling large-scale biomedical analysis in the cloud. BioMed research international, 2013, 185679.

Dai L, et al. (2012) Bioinformatics clouds for big data manipulation. Biology direct, 7, 43.