

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

D-EE

RRID:SCR_019058

Type: Tool

Proper Citation

D-EE (RRID:SCR_019058)

Resource Information

URL: <https://github.com/ShaoKunAn/D-EE>

Proper Citation: D-EE (RRID:SCR_019058)

Description: Software tool for distributed dimensionality reduction and visualization. Distributed software for visualizing intrinsic structure of large scale single cell data written in C language. Its distributed storage and distributed computation technique allows efficiently analyze large scale single cell data at cost of constant time speedup.

Resource Type: data visualization software, data analysis software, software application, software resource, data processing software

Keywords: Distributed dimensionality reduction, dimensionality reduction, distributed storage, distributed computation, large scale data, single cell data, data, , bio.tools

Availability: Free, Available for download, Freely available

Resource Name: D-EE

Resource ID: SCR_019058

Alternate IDs: biotools:d-ee

Alternate URLs: <https://bio.tools/d-ee>

License: GNU GPLv3

Record Creation Time: 20220129T080343+0000

Record Last Update: 20260801T042825+0000

Ratings and Alerts

No rating or validation information has been found for D-EE.

No alerts have been found for D-EE.

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 [PRECISE-TBI](#) .

An S, et al. (2020) D-EE: Distributed software for visualizing intrinsic structure of large-scale single-cell data. GigaScience, 9(11).