

# Resource Summary Report

Generated by [NIF](#) on Aug 21, 2026

## D-EE

RRID:SCR\_019058

Type: Tool

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### Proper Citation

D-EE (RRID:SCR\_019058)

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### 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 analysis software, data processing software, data visualization software, software application, software resource

**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:** 20260821T194154+0000

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### Ratings and Alerts

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

No alerts have been found for D-EE.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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