

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

2s-LCA

RRID:SCR_025613

Type: Tool

Proper Citation

2s-LCA (RRID:SCR_025613)

Resource Information

URL: <https://academic.oup.com/biostatistics/article/23/4/1200/6561796>

Proper Citation: 2s-LCA (RRID:SCR_025613)

Description: Software for multi-omic data integration. Used to jointly decompose multiple biologically related experimental data sets with biological and technological relationships that can be structured into the decomposition.

Synonyms: two-stage Linked Component Analysis

Resource Type: software application, software resource

Defining Citation: [PMID:35358296](#)

Keywords: multi-omic data integration, two-stage linked component analysis, jointly decompose biologically related experimental data sets, decomposition,

Resource Name: 2s-LCA

Resource ID: SCR_025613

Alternate URLs: <https://github.com/CHuanSite/SJD>

Record Creation Time: 20240815T053252+0000

Record Last Update: 20260810T040848+0000

Ratings and Alerts

No rating or validation information has been found for 2s-LCA.

No alerts have been found for 2s-LCA.

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 [RRID](#) .

Sonthalia S, et al. (2026) NeMO Analytics: a compendium of transcriptomic data for the exploration of neocortical development. Nature neuroscience, 29(4), 978.