

Resource Summary Report

Generated by [dkNET](#) on Aug 17, 2026

BootstrappingWithoutReplacement

RRID:SCR_017673

Type: Tool

Proper Citation

BootstrappingWithoutReplacement (RRID:SCR_017673)

Resource Information

URL: <https://github.com/shanglicheng/BootstrappingWithoutReplacement>

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Description: Software tool to dig out more robust and reliable differentially expressed genes between two groups. Samples from different groups will be re-sampled randomly associated with total number of samples.

Synonyms: Bootstrapping coupled with DESeq2

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

Keywords: Differentially, expressed, gene, two, group, randomly, associated, sample

Availability: Free, Freely available

Resource Name: BootstrappingWithoutReplacement

Resource ID: SCR_017673

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260817T040609+0000

Ratings and Alerts

No rating or validation information has been found for BootstrappingWithoutReplacement.

No alerts have been found for BootstrappingWithoutReplacement.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Arango D, et al. (2022) Direct epitranscriptomic regulation of mammalian translation initiation through N4-acetylcytidine. *Molecular cell*, 82(15), 2797.

Aguila J, et al. (2021) Spatial RNA Sequencing Identifies Robust Markers of Vulnerable and Resistant Human Midbrain Dopamine Neurons and Their Expression in Parkinson's Disease. *Frontiers in molecular neuroscience*, 14, 699562.

Khan SA, et al. (2021) Probing the signaling requirements for naive human pluripotency by high-throughput chemical screening. *Cell reports*, 35(11), 109233.