

# Resource Summary Report

Generated by [NIF](#) on Aug 9, 2026

## BlockClust

RRID:SCR\_003347

Type: Tool

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### Proper Citation

BlockClust (RRID:SCR\_003347)

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### Resource Information

**URL:** <https://github.com/bgruening/galaxytools/tree/master/workflows/blockclust>

**Proper Citation:** BlockClust (RRID:SCR\_003347)

**Description:** Software for efficient clustering and classification of non-coding RNAs from short read RNA-seq profiles.

**Resource Type:** software resource

**Defining Citation:** [PMID:24931994](#)

**Availability:** Free, Available for download, Freely available

**Resource Name:** BlockClust

**Resource ID:** SCR\_003347

**Alternate IDs:** OMICS\_04641

**Record Creation Time:** 20220129T080218+0000

**Record Last Update:** 20260808T185756+0000

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### Ratings and Alerts

No rating or validation information has been found for BlockClust.

No alerts have been found for BlockClust.

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### 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 [NIF](#) .

Hamar R, et al. (2025) The zebrafish (Danio rerio) snoRNAome. NAR genomics and bioinformatics, 7(1), lqaf013.