

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

[rna-stability](#)

RRID:SCR_019259

Type: Tool

Proper Citation

rna-stability (RRID:SCR_019259)

Resource Information

URL: <https://github.com/nch-igm/rna-stability>

Proper Citation: rna-stability (RRID:SCR_019259)

Description: Software tool as parallel processing framework for large scale generation of secondary RNA structures and folding statistics for transcriptome of any species.

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

Keywords: Secondary RNA structures generation, large scale generation, RNA structures, transcriptome folding statistics, , bio.tools

Availability: Free, Freely available

Resource Name: rna-stability

Resource ID: SCR_019259

Alternate IDs: biotools:rna-stability

Alternate URLs: <https://bio.tools/rna-stability>

License: BSD 3-Clause

Record Creation Time: 20220129T080344+0000

Record Last Update: 20260912T200108+0000

Ratings and Alerts

No rating or validation information has been found for rna-stability.

No alerts have been found for rna-stability.

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 [SPARC SAWG](#) .

Gaither JBS, et al. (2021) Synonymous variants that disrupt messenger RNA structure are significantly constrained in the human population. GigaScience, 10(4).