

# Resource Summary Report

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## ProbRNA

RRID:SCR\_001288

Type: Tool

### Proper Citation

ProbRNA (RRID:SCR\_001288)

### Resource Information

**URL:** <http://yiplab.cse.cuhk.edu.hk/probrna/>

**Proper Citation:** ProbRNA (RRID:SCR\_001288)

**Description:** Software for computational identification of protein binding sites on RNAs using high-throughput RNA structure-probing data.

**Abbreviations:** ProbRNA

**Resource Type:** software resource

**Defining Citation:** [PMID:24376038](#)

**Keywords:** high-throughput sequencing, probe, rna structure, rna, bio.tools

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** ProbRNA

**Resource ID:** SCR\_001288

**Alternate IDs:** biotools:probrna, OMICS\_02195

**Alternate URLs:** <https://bio.tools/probrna>

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260919T194938+0000

### Ratings and Alerts

No rating or validation information has been found for ProbRNA.

No alerts have been found for ProbRNA.

## 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 [nidm-terms](#) .

Ge P, et al. (2015) Computational analysis of RNA structures with chemical probing data. Methods (San Diego, Calif.), 79-80, 60.