

Resource Summary Report

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

RNAcontext

RRID:SCR_000179

Type: Tool

Proper Citation

RNAcontext (RRID:SCR_000179)

Resource Information

URL: <http://www.cs.toronto.edu/~hilal/rnacontext/>

Proper Citation: RNAcontext (RRID:SCR_000179)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31, 2025. Motif finding software suited for using large-scale RNA-binding affinity datasets to determine the relative binding preferences of RNA-binding proteins (RBPs) for a wide range of RNA sequences and structures. The tool is also implemented in a website.

Abbreviations: RNAcontext

Resource Type: web service, production service resource, software resource, data access protocol, analysis service resource, service resource

Defining Citation: [PMID:20617199](#)

Keywords: rna-binding protein, motif, rna sequence, rna structure, rna, binding preference, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: RNAcontext

Resource ID: SCR_000179

Alternate IDs: OMICS_02253, biotools:rnacontext

Alternate URLs: <https://bio.tools/rnacontext>

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260808T185715+0000

Ratings and Alerts

No rating or validation information has been found for RNAcontext.

No alerts have been found for RNAcontext.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Pan X, et al. (2018) Prediction of RNA-protein sequence and structure binding preferences using deep convolutional and recurrent neural networks. BMC genomics, 19(1), 511.

Han Y, et al. (2016) Integrating Epigenomics into the Understanding of Biomedical Insight. Bioinformatics and biology insights, 10, 267.