# Resource Summary Report

Generated by dkNET on May 11, 2025

## University of Freiburg Bioinformatics Group: Servers and Web-Applications

RRID:SCR_008256
Type: Tool

### Proper Citation

University of Freiburg Bioinformatics Group: Servers and Web-Applications (RRID:SCR_008256)

### Resource Information

**URL:** http://www.bioinf.uni-freiburg.de/Software

**Proper Citation:** University of Freiburg Bioinformatics Group: Servers and Web-Applications (RRID:SCR_008256)

**Description:** This resource takes you to the Server and Web-Applications of the University of Freiburg Bioinformatics Group. Sponsors: This resource is supported by University of Freiburg. Keywords: Server, Web application, Bioinformatics, Software,

**Synonyms:** Freiburg Bioinformatics Group

**Resource Type:** data or information resource, portal, topical portal

**Funding:**

**Resource Name:** University of Freiburg Bioinformatics Group: Servers and Web-Applications

**Resource ID:** SCR_008256

**Alternate IDs:** nif-0000-30196

**Record Creation Time:** 20220129T080246+0000

**Record Last Update:** 20250508T065130+0000

### Ratings and Alerts

No rating or validation information has been found for University of Freiburg Bioinformatics Group: Servers and Web-Applications.

No alerts have been found for University of Freiburg Bioinformatics Group: Servers and Web-Applications.

## Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 8 mentions in open access literature.

**Listed below are recent publications.** The full list is available at dkNET.


Kleinkauf R, et al. (2015) antaRNA--Multi-objective inverse folding of pseudoknot RNA using ant-colony optimization. BMC bioinformatics, 16, 389.

Wright PR, et al. (2014) CopraRNA and IntaRNA: predicting small RNA targets, networks and interaction domains. Nucleic acids research, 42(Web Server issue), W119.

Mann M, et al. (2014) Memory-efficient RNA energy landscape exploration. Bioinformatics (Oxford, England), 30(18), 2584.

Mann M, et al. (2014) Atom mapping with constraint programming. Algorithms for molecular biology : AMB, 9(1), 23.

Silva IJ, et al. (2013) An RpoS-dependent sRNA regulates the expression of a chaperone involved in protein folding. RNA (New York, N.Y.), 19(9), 1253.

Marín RM, et al. (2012) Optimal use of conservation and accessibility filters in microRNA target prediction. PloS one, 7(2), e32208.

Li W, et al. (2012) Predicting sRNAs and their targets in bacteria. Genomics, proteomics & bioinformatics, 10(5), 276.

Richter AS, et al. (2012) Accessibility and conservation: general features of bacterial small RNA-mRNA interactions? RNA biology, 9(7), 954.