

Resource Summary Report

Generated by [NIF](#) on Aug 10, 2026

[wapRNA](#)

RRID:SCR_013292

Type: Tool

Proper Citation

wapRNA (RRID:SCR_013292)

Resource Information

URL: <http://waprna.big.ac.cn/>

Proper Citation: wapRNA (RRID:SCR_013292)

Description: A free web-based application for the processing of high-throughput RNA-Seq data from next generation sequencing (NGS) platforms, such as Genome Analyzer of Illumina Inc. (Solexa) and SOLiD of Applied Biosystems (SOLiD).

Abbreviations: wapRNA

Resource Type: software resource

Defining Citation: [PMID:21896507](#)

Resource Name: wapRNA

Resource ID: SCR_013292

Alternate IDs: OMICS_00370

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260808T190009+0000

Ratings and Alerts

No rating or validation information has been found for wapRNA.

No alerts have been found for wapRNA.

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](#) .

Jin J, et al. (2012) Mechanism analysis of acid tolerance response of bifidobacterium longum subsp. longum BBMN 68 by gene expression profile using RNA-sequencing. PloS one, 7(12), e50777.