

Resource Summary Report

Generated by [NIF](#) on Aug 19, 2026

Sno/scaRNAbase

RRID:SCR_007938

Type: Tool

Proper Citation

Sno/scaRNAbase (RRID:SCR_007938)

Resource Information

URL: <https://omictools.com/sno-scarabase-tool>

Proper Citation: Sno/scaRNAbase (RRID:SCR_007938)

Description: A curated database for small nucleolar RNAs and small cajal body-specific RNAs. It presents sno/scaRNA-associated genetic and functional data and provides access to several other database sources via web-accessible search interfaces. Consisting of 1979 sno/scaRNA records obtained from 85 organisms, sno/scaRNAbase is a combination of systematic literature curation and annotation effort. small nucleolar RNA, small cajal body-specific RNA

Synonyms: Sno/scaRNAbase

Resource Type: database, data or information resource

Keywords: scarna, small cajal body-specific rna, small nucleolar rna, snorna

Resource Name: Sno/scaRNAbase

Resource ID: SCR_007938

Alternate IDs: nif-0000-03475

Old URLs: <http://bioinfo.fudan.edu.cn/snoRNAbase.nsf>

Record Creation Time: 20220129T080244+0000

Record Last Update: 20260819T044219+0000

Ratings and Alerts

No rating or validation information has been found for Sno/scaRNAbase.

No alerts have been found for Sno/scaRNAbase.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Deschamps-Francoeur G, et al. (2014) Identification of discrete classes of small nucleolar RNA featuring different ends and RNA binding protein dependency. Nucleic acids research, 42(15), 10073.

Chorev M, et al. (2013) Computational identification of functional introns: high positional conservation of introns that harbor RNA genes. Nucleic acids research, 41(11), 5604.

Motsch N, et al. (2012) MicroRNA profiling of Epstein-Barr virus-associated NK/T-cell lymphomas by deep sequencing. PloS one, 7(8), e42193.

Scott MS, et al. (2012) Human box C/D snoRNA processing conservation across multiple cell types. Nucleic acids research, 40(8), 3676.