# **Resource Summary Report**

Generated by <u>dkNET</u> on May 18, 2025

# Sno/scaRNAbase

RRID:SCR\_007938 Type: Tool

**Proper Citation** 

Sno/scaRNAbase (RRID:SCR\_007938)

### **Resource Information**

URL: https://omictools.com/sno-scarnabase-tool

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**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

Funding:

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: 20250517T055843+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 <u>dkNET</u>.

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.

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

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