Resource Summary Report

Generated by ASWG on Aug 4, 2026

SSCprofiler

RRID:SCR_001282
Type: Tool

Proper Citation

SSCprofiler (RRID:SCR_001282)

Resource Information

URL: http://mirna.imbb.forth.gr/SSCprofiler.html

Proper Citation: SSCprofiler (RRID:SCR_001282)

Description: Tool which can be used to identify novel miRNA gene candidates in the human genome.

Abbreviations: SSCprofiler

Synonyms: Sequence Structure and Conservation profiler

Resource Type: data analysis service, service resource, analysis service resource, production service resource

Defining Citation: PMID:19324892

Keywords: microrna, gene, genome, sequence, structure, conservation

Availability: Free, Freely available

Resource Name: SSCprofiler

Resource ID: SCR_001282

Alternate IDs: OMICS_02055

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260804T043119+0000

Ratings and Alerts

No rating or validation information has been found for SSCprofiler.

No alerts have been found for SSCprofiler.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Lachinani L, et al. (2023) The oncogene Musashi1 encodes novel miRNAs in breast cancer. Scientific reports, 13(1), 13710.

Rezaei H, et al. (2021) MicroRNA and Hemophilia-A Disease: Bioinformatics Prediction and Experimental Analysis. Cell journal, 23(3), 341.

Akhtar MM, et al. (2016) Bioinformatic tools for microRNA dissection. Nucleic acids research, 44(1), 24.