

Resource Summary Report

Generated by RRID on Aug 2, 2026

DSAP

RRID:SCR_013352

Type: Tool

Proper Citation

DSAP (RRID:SCR_013352)

Resource Information

URL: <http://dsap.cgu.edu.tw/>

Proper Citation: DSAP (RRID:SCR_013352)

Description: A web server designed to provide a total solution to analyze small RNAs sequencing data generated by SOLEXA., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: DSAP

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

Defining Citation: [PMID:20478825](#)

Keywords: bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: DSAP

Resource ID: SCR_013352

Alternate IDs: biotools:dsap, OMICS_00357

Alternate URLs: <https://bio.tools/dsap>

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260801T191007+0000

Ratings and Alerts

No rating or validation information has been found for DSAP.

No alerts have been found for DSAP.

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 [RRID](#).

Chen Y, et al. (2024) DNA framework signal amplification platform-based high-throughput systemic immune monitoring. *Signal transduction and targeted therapy*, 9(1), 28.

Edwards CM, et al. (2024) An ascending vagal sensory-central noradrenergic pathway modulates retrieval of passive avoidance memory. *bioRxiv : the preprint server for biology*.

Tyagi A, et al. (2018) Genome-wide discovery of tissue-specific miRNAs in clusterbean (*Cyamopsis tetragonoloba*) indicates their association with galactomannan biosynthesis. *Plant biotechnology journal*, 16(6), 1241.

Huang JY, et al. (2017) Shrimp miR-10a Is Co-opted by White Spot Syndrome Virus to Increase Viral Gene Expression and Viral Replication. *Frontiers in immunology*, 8, 1084.

Wan Y, et al. (2016) Transcriptome analysis reveals a ribosome constituents disorder involved in the RPL5 downregulated zebrafish model of Diamond-Blackfan anemia. *BMC medical genomics*, 9, 13.

Bai J, et al. (2015) NGSmirPlant: comprehensive characterization of the small RNA transcriptomes of plants. *Protein & cell*, 6(6), 397.

Xiong Q, et al. (2014) Characterization of miRNomes in acute and chronic myeloid leukemia cell lines. *Genomics, proteomics & bioinformatics*, 12(2), 79.

Huang PJ, et al. (2012) Identification of putative miRNAs from the deep-branching unicellular flagellates. *Genomics*, 99(2), 101.