

Resource Summary Report

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miRTarVisPlus

RRID:SCR_017103

Type: Tool

Proper Citation

miRTarVisPlus (RRID:SCR_017103)

Resource Information

URL: <http://sehilyi.com/mirtarvisplus/>

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Description: Web based interactive visual analytics tool for miRNA target predictions and integrative analyses of multiple prediction results. Used for sequence based miRNA target prediction by exploiting miRNA-mRNA expression profile data.

Abbreviations: miRTarVis+

Synonyms: mirtarvisplus

Resource Type: service resource, data access protocol, data analysis service, software resource, web service, production service resource, analysis service resource

Defining Citation: [PMID:28600227](#)

Keywords: interactive, visual, analytical, miRNA, target, prediction, analysis, mRNA, expression, data

Funding: National Research Foundation of Korea ;
Hankuk University of Foreign Studies Research Fund

Availability: Free, Freely available

Resource Name: miRTarVisPlus

Resource ID: SCR_017103

Alternate URLs: <http://hcil.snu.ac.kr/research/mirtarvisplus>

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260731T043006+0000

Ratings and Alerts

No rating or validation information has been found for miRTarVisPlus.

No alerts have been found for miRTarVisPlus.

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 [kravitz2](#) .

Awoyemi T, et al. (2024) MicroRNA analysis of medium/large placenta extracellular vesicles in normal and preeclampsia pregnancies. *Frontiers in cardiovascular medicine*, 11, 1371168.

Bi S, et al. (2020) Bioinformatics analysis of key genes and miRNAs associated with Stanford type A aortic dissection. *Journal of thoracic disease*, 12(9), 4842.

Yang L, et al. (2019) Clinical Features and MicroRNA Expression Patterns Between AML Patients With DNMT3A R882 and Frameshift Mutations. *Frontiers in oncology*, 9, 1133.