

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

[stMMR](#)

RRID:SCR_025601

Type: Tool

Proper Citation

stMMR (RRID:SCR_025601)

Resource Information

URL: <https://github.com/nayu0419/stMMR>

Proper Citation: stMMR (RRID:SCR_025601)

Description: Software tool for spatial domain identification from spatially resolved transcriptomics with multi-modal feature representation.

Resource Type: software resource, source code, software application

Defining Citation: [DOI:10.1101/2024.02.22.581503](https://doi.org/10.1101/2024.02.22.581503)

Keywords: spatial domain identification, spatially resolved transcriptomics, multi-modal feature representation,

Funding: National Natural Science Foundation of China ;
Natural Science Foundation of Shandong Province

Availability: Free, Available for download, Freely available,

Resource Name: stMMR

Resource ID: SCR_025601

License: MIT license

Record Creation Time: 20240808T053249+0000

Record Last Update: 20260804T043910+0000

Ratings and Alerts

No rating or validation information has been found for stMMR.

No alerts have been found for stMMR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Zhang D, et al. (2024) stMMR: accurate and robust spatial domain identification from spatially resolved transcriptomics with multimodal feature representation. GigaScience, 13.