

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

[Reads2Map](#)

RRID:SCR_023593

Type: Tool

Proper Citation

Reads2Map (RRID:SCR_023593)

Resource Information

URL: <https://github.com/Cristianetaniguti/Reads2Map>

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Description: Software package as collection of WDL workflows to build linkage maps from sequencing reads. WDL workflows to build genetic maps from sequencing reads.

Resource Type: software toolkit, software resource

Defining Citation: [DOI:10.1101/2022.11.24.517847](https://doi.org/10.1101/2022.11.24.517847)

Keywords: Genetic Map, Linkage analysis, Variant Calling, Dosage Calling

Funding: National Council for Scientific and Technological Development 313269/2021-1; USDA National Institute of Food and Agriculture (NIFA); 2020-51181-32156 Bill and Melinda Gates Foundation; OPP1213329 European Union's Horizon 2020

Availability: Free, Available for download, Freely available

Resource Name: Reads2Map

Resource ID: SCR_023593

License: MIT license

Record Creation Time: 20230523T050222+0000

Record Last Update: 20260810T040842+0000

Ratings and Alerts

No rating or validation information has been found for Reads2Map.

No alerts have been found for Reads2Map.

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](#) .

Taniguti CH, et al. (2022) Developing best practices for genotyping-by-sequencing analysis in the construction of linkage maps. GigaScience, 12.