

Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

Marvel

RRID:SCR_017621

Type: Tool

Proper Citation

Marvel (RRID:SCR_017621)

Resource Information

URL: <https://github.com/schloi/MARVEL>

Proper Citation: Marvel (RRID:SCR_017621)

Description: Software set of tools that facilitate overlapping, patching, correction and assembly of noisy long reads.

Resource Type: alignment software, data processing software, image analysis software, software application, software resource, software toolkit

Keywords: Overlapping, patching, correction, assembly, noisy, long, read, sequence, align

Availability: Free, Available for download, Freely available

Resource Name: Marvel

Resource ID: SCR_017621

License: <https://github.com/schloi/MARVEL/blob/master/LICENSE>

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260829T182811+0000

Ratings and Alerts

No rating or validation information has been found for Marvel .

No alerts have been found for Marvel .

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Meyer A, et al. (2021) Giant lungfish genome elucidates the conquest of land by vertebrates. *Nature*, 590(7845), 284.

Wang W, et al. (2020) The draft nuclear genome assembly of *Eucalyptus pauciflora*: a pipeline for comparing de novo assemblies. *GigaScience*, 9(1).