

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

matemaker

RRID:SCR_017199

Type: Tool

Proper Citation

matemaker (RRID:SCR_017199)

Resource Information

URL: <https://github.com/josephryan/matemaker>

Proper Citation: matemaker (RRID:SCR_017199)

Description: Software tool to make artificial mate pairs from long sequences for scaffolding.

Synonyms: matemaker v1.0.0

Resource Type: data analysis software, data processing software, sequence analysis software, software application, software resource, standalone software

Keywords: artificial, mate, pair, long, sequence, scaffolding, genomics, genome, assembly

Availability: Free, Available for download, Freely available

Resource Name: matemaker

Resource ID: SCR_017199

License: GNU GPL v3

Record Creation Time: 20220129T080334+0000

Record Last Update: 20260912T195849+0000

Ratings and Alerts

No rating or validation information has been found for matemaker.

No alerts have been found for matemaker.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Dur??n-Fuentes JA, et al. (2025) Repeatome diversity in sea anemone genomics (Cnidaria: Actiniaria) based on the Actiniaria-REPlib library. BMC genomics, 26(1), 473.

Vargas AM, et al. (2024) Morphological and dietary changes encoded in the genome of Beroe ovata, a ctenophore-eating ctenophore. NAR genomics and bioinformatics, 6(2), lqae072.

Ketchum RN, et al. (2020) Population Genomic Analyses of the Sea Urchin Echinometra sp. EZ across an Extreme Environmental Gradient. Genome biology and evolution, 12(10), 1819.

Ohdera A, et al. (2019) Box, stalked, and upside-down? Draft genomes from diverse jellyfish (Cnidaria, Acraspeda) lineages: Alatina alata (Cubozoa), Calvadosia cruxmelitensis (Staurozoa), and Cassiopea xamachana (Scyphozoa). GigaScience, 8(7).