

Resource Summary Report

Generated by [T1D](#) on Aug 10, 2026

HaploMaker

RRID:SCR_021928

Type: Tool

Proper Citation

HaploMaker (RRID:SCR_021928)

Resource Information

URL: <https://github.com/mfuzan/HaploMaker>

Proper Citation: HaploMaker (RRID:SCR_021928)

Description: Software pipeline for haplotype assembly of diploid genomes using paired end or PacBio reads.

Resource Type: software resource

Keywords: haplotype assembly, diploid genomes, paired end reads, PacBio reads

Availability: Free, Available for download, Freely available

Resource Name: HaploMaker

Resource ID: SCR_021928

Record Creation Time: 20220421T050137+0000

Record Last Update: 20260808T190221+0000

Ratings and Alerts

No rating or validation information has been found for HaploMaker.

No alerts have been found for HaploMaker.

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 [T1D](#) .

Fruzangohar M, et al. (2022) HaploMaker: An improved algorithm for rapid haplotype assembly of genomic sequences. GigaScience, 11.