

Resource Summary Report

Generated by [ASWG](#) on Aug 24, 2026

[ABMapper](#)

RRID:SCR_010242

Type: Tool

Proper Citation

ABMapper (RRID:SCR_010242)

Resource Information

URL: <http://hkbic.cuhk.edu.hk/software/abmapper>

Proper Citation: ABMapper (RRID:SCR_010242)

Description: A portable, easy-to-use package for spliced alignment, junction site detection, and reads mapping. The core module was written in C++ and wrapped in PERL scripts.

Abbreviations: ABMapper

Synonyms: ABMapper: A suffix-array based spliced alignment tool

Resource Type: software resource

Resource Name: ABMapper

Resource ID: SCR_010242

Alternate IDs: OMICS_01238

Record Creation Time: 20220129T080257+0000

Record Last Update: 20260815T182351+0000

Ratings and Alerts

No rating or validation information has been found for ABMapper.

No alerts have been found for ABMapper.

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 [ASWG](#) .

Li MW, et al. (2013) Silicon era of carbon-based life: application of genomics and bioinformatics in crop stress research. International journal of molecular sciences, 14(6), 11444.