

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[PALMapper](#)

RRID:SCR_011466

Type: Tool

Proper Citation

PALMapper (RRID:SCR_011466)

Resource Information

URL: <http://www.raetschlab.org/suppl/palmapper>

Proper Citation: PALMapper (RRID:SCR_011466)

Description: Computes both spliced and unspliced alignments at high accuracy while taking advantage of base quality information and splice site predictions.

Abbreviations: PALMapper

Resource Type: software resource

Resource Name: PALMapper

Resource ID: SCR_011466

Alternate IDs: OMICS_01245

Record Creation Time: 20220129T080304+0000

Record Last Update: 20260801T190409+0000

Ratings and Alerts

No rating or validation information has been found for PALMapper.

No alerts have been found for PALMapper.

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 [RRID](#) .

Dimitrakopoulos C, et al. (2021) Multi-omics data integration reveals novel drug targets in hepatocellular carcinoma. BMC genomics, 22(1), 592.

Imprialou M, et al. (2017) Genomic Rearrangements in Arabidopsis Considered as Quantitative Traits. Genetics, 205(4), 1425.

Dubin MJ, et al. (2015) DNA methylation in Arabidopsis has a genetic basis and shows evidence of local adaptation. eLife, 4, e05255.

Wolfe AL, et al. (2014) RNA G-quadruplexes cause eIF4A-dependent oncogene translation in cancer. Nature, 513(7516), 65.