

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 10, 2026

QPALMA

RRID:SCR_008791

Type: Tool

Proper Citation

QPALMA (RRID:SCR_008791)

Resource Information

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

Proper Citation: QPALMA (RRID:SCR_008791)

Description: An alignment tool targeted to align spliced reads produced by Next Generation sequencing platforms such as Illumina Solexa or 454.

Abbreviations: QPALMA

Synonyms: QPALMA: Optimal Spliced Alignments of Short Sequence Reads

Resource Type: software resource

Resource Name: QPALMA

Resource ID: SCR_008791

Alternate IDs: OMICS_01248

Record Creation Time: 20220129T080249+0000

Record Last Update: 20260808T185920+0000

Ratings and Alerts

No rating or validation information has been found for QPALMA.

No alerts have been found for QPALMA.

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 [PRECISE-TBI](#) .

Alser M, et al. (2021) Technology dictates algorithms: recent developments in read alignment. Genome biology, 22(1), 249.