

Resource Summary Report

Generated by T1D on Sep 5, 2026

OptimiR

RRID:SCR_024149

Type: Tool

Proper Citation

OptimiR (RRID:SCR_024149)

Resource Information

URL: <https://github.com/FlorianThibord/OptimiR>

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Description: Software miRSeq data alignment workflow. Used to integrate genetic information to assess the impact of variants on miRNA expression. Used for integrating genome wide genotype data into miRNA sequence alignment analysis.

Synonyms: optimir

Resource Type: software library, software resource, software toolkit

Defining Citation: [PMID:30819774](https://pubmed.ncbi.nlm.nih.gov/30819774/)

Keywords: miRSeq data alignment workflow, integrate genetic information, assess impact of variants, miRNA expression, integrating genome wide genotype data, miRNA sequence alignment analysis,

Availability: Free, Available for download, Freely available,

Resource Name: OptimiR

Resource ID: SCR_024149

Alternate IDs: OMICS_33123

Alternate URLs: <https://sources.debian.org/src/optimir/>

License: GPL-3.0 license

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260904T000121+0000

Ratings and Alerts

No rating or validation information has been found for OptimIR.

No alerts have been found for OptimIR.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.