

Resource Summary Report

Generated by [nidm-terms](#) on Jul 30, 2026

InsertionMapper

RRID:SCR_004163

Type: Tool

Proper Citation

InsertionMapper (RRID:SCR_004163)

Resource Information

URL: <http://sourceforge.net/projects/insertionmapper/>

Proper Citation: InsertionMapper (RRID:SCR_004163)

Description: A pipeline tool for the identification of targeted sequences from multidimensional high throughput sequencing data. It consists of four independently working modules: Data Preprocessing, Database Modeling, Dimension Deconvolution and Element Mapping. This pipeline tool is applicable to scenarios requiring analysis of the tremendous output of short reads produced in NGS sequencing experiments of targeted genome sequences.

Abbreviations: InsertionMapper

Resource Type: software resource

Defining Citation: [PMID:24090499](#)

Keywords: high throughput sequencing, dna sequence, next generation sequencing, bio.tools

Availability: Acknowledgement requested, GNU General Public License

Resource Name: InsertionMapper

Resource ID: SCR_004163

Alternate IDs: OMICS_01547, biotools:insertionmapper

Alternate URLs: <https://bio.tools/insertionmapper>

Record Creation Time: 20220129T080223+0000

Record Last Update: 20260725T190547+0000

Ratings and Alerts

No rating or validation information has been found for InsertionMapper.

No alerts have been found for InsertionMapper.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.