

# Resource Summary Report

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## InsertionMapper

RRID:SCR\_004163

Type: Tool

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### Proper Citation

InsertionMapper (RRID:SCR\_004163)

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### 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

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### Ratings and Alerts

No rating or validation information has been found for InsertionMapper.

No alerts have been found for InsertionMapper.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.