

# Resource Summary Report

Generated by [NIF](#) on Apr 17, 2025

## Batman-Seq

RRID:SCR\_000048

Type: Tool

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### Proper Citation

Batman-Seq (RRID:SCR\_000048)

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### Resource Information

**URL:** <http://sourceforge.net/projects/batman-seq/>

**Proper Citation:** Batman-Seq (RRID:SCR\_000048)

**Description:** A fast BWT-based short reads mapping tools which uses additional statistical method to model error profile of the sequencing experiment.

**Abbreviations:** Batman-Seq

**Synonyms:** Basic Alignment Tool for MAny Nucleotides

**Resource Type:** software resource

**Keywords:** c++

**Funding:**

**Resource Name:** Batman-Seq

**Resource ID:** SCR\_000048

**Alternate IDs:** OMICS\_00651

**Record Creation Time:** 20220129T080159+0000

**Record Last Update:** 20250410T064504+0000

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### Ratings and Alerts

No rating or validation information has been found for Batman-Seq.

No alerts have been found for Batman-Seq.

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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.