

# Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 18, 2026

## Sequgio

RRID:SCR\_008867

Type: Tool

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### Proper Citation

Sequgio (RRID:SCR\_008867)

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### Resource Information

**URL:** <http://r-forge.r-project.org/projects/sequgio/>

**Proper Citation:** Sequgio (RRID:SCR\_008867)

**Description:** An algorithm to estimate isoforms expression from RNA-seq data based on a model that doesn't assume uniform distribution of count within transcripts.

**Abbreviations:** Sequgio

**Resource Type:** software resource

**Defining Citation:** [PMID:24307704](#)

**Resource Name:** Sequgio

**Resource ID:** SCR\_008867

**Alternate IDs:** OMICS\_01290

**Record Creation Time:** 20220129T080249+0000

**Record Last Update:** 20260912T195710+0000

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### Ratings and Alerts

No rating or validation information has been found for Sequgio.

No alerts have been found for Sequgio.

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### Data and Source Information

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

## Usage and Citation Metrics

We have not found any literature mentions for this resource.