## **Resource Summary Report**

Generated by dkNET on Apr 30, 2025

# **Sequgio**

RRID:SCR\_008867

Type: Tool

### **Proper Citation**

Sequgio (RRID:SCR\_008867)

#### **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

**Funding:** 

Resource Name: Sequgio

Resource ID: SCR\_008867

Alternate IDs: OMICS\_01290

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

**Record Last Update:** 20250420T014443+0000

### **Ratings and Alerts**

No rating or validation information has been found for Sequgio.

No alerts have been found for Sequgio.

## Data and Source Information

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.