

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

## SeqGene

RRID:SCR\_011861

Type: Tool

### Proper Citation

SeqGene (RRID:SCR\_011861)

### Resource Information

**URL:** <http://sourceforge.net/apps/mediawiki/seqgene/?title=SeqGene>

**Proper Citation:** SeqGene (RRID:SCR\_011861)

**Description:** An open-source software for mining next-gen sequencing datasets, focusing on post-alignment quality control, SNP and indel identification and annotation, RNA expression quantification, etc.

**Abbreviations:** SeqGene

**Resource Type:** software resource

**Availability:** Open unspecified license

**Resource Name:** SeqGene

**Resource ID:** SCR\_011861

**Alternate IDs:** OMICS\_01134

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260725T190726+0000

### Ratings and Alerts

No rating or validation information has been found for SeqGene.

No alerts have been found for SeqGene.

### Data and Source Information

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

## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [nidm-terms](#) .

Alloza I, et al. (2020) BIRC6 Is Associated with Vulnerability of Carotid Atherosclerotic Plaque. International journal of molecular sciences, 21(24).

Devaiah BN, et al. (2016) BRD4 is a histone acetyltransferase that evicts nucleosomes from chromatin. Nature structural & molecular biology, 23(6), 540.