

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 12, 2025

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

Funding:

Availability: Open unspecified license

Resource Name: SeqGene

Resource ID: SCR_011861

Alternate IDs: OMICS_01134

Record Creation Time: 20220129T080307+0000

Record Last Update: 20250410T070213+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 [FDI Lab - SciCrunch.org](#).

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.