## **Resource Summary Report**

Generated by <u>dkNET</u> on Apr 28, 2025

# **SeqEM**

RRID:SCR\_002021 Type: Tool

**Proper Citation** 

SeqEM (RRID:SCR\_002021)

#### **Resource Information**

URL: http://hihg.med.miami.edu/software-download/seqem-version-1.0

Proper Citation: SeqEM (RRID:SCR\_002021)

**Description:** Online tool for utilizing a genotype calling algorithm for next-generation sequence data.

**Resource Type:** data processing software, sequence analysis software, software resource, data analysis software, algorithm resource, web application, software application

Defining Citation: PMID:20861027

Keywords: genotype, algorithm, sequence, rna, dna, bio.tools

Funding:

Availability: Available for download, Public

Resource Name: SeqEM

Resource ID: SCR\_002021

Alternate IDs: OMICS\_00074, biotools:seqem

Alternate URLs: https://bio.tools/seqem

Record Creation Time: 20220129T080211+0000

Record Last Update: 20250428T052917+0000

### **Ratings and Alerts**

No rating or validation information has been found for SeqEM.

No alerts have been found for SeqEM.

#### Data and Source Information

Source: SciCrunch Registry

### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>dkNET</u>.

Goovaerts T, et al. (2018) A comprehensive overview of genomic imprinting in breast and its deregulation in cancer. Nature communications, 9(1), 4120.