# **Resource Summary Report**

Generated by dkNET on Apr 29, 2025

# **Genestack**

RRID:SCR\_011885

Type: Tool

### **Proper Citation**

Genestack (RRID:SCR\_011885)

#### Resource Information

URL: http://www.genestack.com

Proper Citation: Genestack (RRID:SCR\_011885)

**Description:** A universal collaborative platform for bioinformatics application development that allows users to store and share large data sets securely within and across organizations, with free access to public data from major databases. The platform includes open-source and proprietary genomics applications, working together independent of file formats. For developers an SDK, APIs and a marketplace are provided.

**Abbreviations:** Genestack

Resource Type: service resource

**Keywords:** data sharing, data management, sequencing, quality control, exome, genome, variant call, somatic mutation, rna-seq, chip-seq, data analysis, visualization, computing, big data, next-generation sequencing

**Funding:** 

Resource Name: Genestack

Resource ID: SCR\_011885

Alternate IDs: OMICS\_01218

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

Record Last Update: 20250420T014601+0000

### **Ratings and Alerts**

No rating or validation information has been found for Genestack.

No alerts have been found for Genestack.

#### **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 dkNET.

Rauluseviciute I, et al. (2019) DNA methylation data by sequencing: experimental approaches and recommendations for tools and pipelines for data analysis. Clinical epigenetics, 11(1), 193.

Overton NLD, et al. (2018) Mutations in EEA1 are associated with allergic bronchopulmonary aspergillosis and affect phagocytosis of Aspergillus fumigatus by human macrophages. PloS one, 13(3), e0185706.