HTSeq
RRID:SCR_005514
Type: Tool

Proper Citation

HTSeq (RRID:SCR_005514)

Resource Information

URL: http://htseq.readthedocs.io/en/release_0.9.1/

Description: A Python package that provides infrastructure to process data from high-throughput sequencing assays. While the main purpose of HTSeq is to allow you to write your own analysis scripts, customized to your needs, there are also a couple of stand-alone scripts for common tasks that can be used without any Python knowledge.

Resource Name: HTSeq

Proper Citation: HTSeq (RRID:SCR_005514)

Resource Type: Resource, software resource, software application, standalone software, data processing software, authoring tool

Keywords: python, high-throughput sequencing assay

Resource ID: SCR_005514

Parent Organization: European Molecular Biology Laboratory

Availability: GNU General Public License v3 or later

Website Status: Last checked up

Alternate IDs: OMICS_01053

Old URLs: http://www-huber.embl.de/users/anders/HTSeq/

Abbreviations: HTSeq
Ratings and Alerts

No rating or validation information has been found for HTSeq.
No alerts have been found for HTSeq.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 3149 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](https://www.ncbi.nlm.nih.gov/pmc/tools/embor/).


Cardoso Alves L, et al. (2020) Non-apoptotic TRAIL function modulates NK cell activity during viral infection. EMBO reports, 21(1), e48789.


