HTSeq
RRID:SCR_005514
Type: Tool

Proper Citation

HTSeq (RRID:SCR_005514)

Resource Information

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

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

Keywords: python, high-throughput sequencing assay, bio.tools

Parent Organization: European Molecular Biology Laboratory

Availability: GNU General Public License v3 or later

Website Status: Last checked up

Abbreviations: HTSeq

Resource Name: HTSeq

Resource ID: SCR_005514

Alternate IDs: OMICS_01053, biotools:htseq

Alternate URLs: https://bio.tools/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 4085 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


