

Resource Summary Report

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htseq-count

RRID:SCR_011867

Type: Tool

Proper Citation

htseq-count (RRID:SCR_011867)

Resource Information

URL: <http://www-huber.embl.de/users/anders/HTSeq/doc/count.html>

Proper Citation: htseq-count (RRID:SCR_011867)

Description: Script distributed with the HT-Seq Python framework for processing RNA-seq or DNA-seq data., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: htseq-count

Synonyms: Counting reads in features

Resource Type: software resource

Defining Citation: [PMID:23975260](#)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: htseq-count

Resource ID: SCR_011867

Alternate IDs: OMICS_01162

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260725T190726+0000

Ratings and Alerts

No rating or validation information has been found for htseq-count.

No alerts have been found for htseq-count.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2587 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Avanessian SC, et al. (2026) IL2/IL15 Signaling Induces NK Cell Production of FLT3LG, Augmenting Anti-PD-1 Immunotherapy. *Cancer immunology research*, 14(1), 122.

Jing J, et al. (2026) Recombinant tissue plasminogen activator protects neurons after intracerebral hemorrhage through activating the PI3K/AKT/mTOR pathway. *Neural regeneration research*, 21(4), 1574.

Canato S, et al. (2025) Survivin Promotes Stem Cell Competence for Skin Cancer Initiation. *Cancer discovery*, 15(2), 427.

Culbert BM, et al. (2025) Effects of environmental salinity on global and endocrine-specific transcriptomic profiles in the caudal neurosecretory system of salmonid fishes. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 39(6), e70477.

Foo CZ, et al. (2025) Differential Chromatin Accessibility, Gene Expression, and mRNA Splicing Between Developing Cochlear Inner and Outer Hair Cells. *Journal of the Association for Research in Otolaryngology : JARO*.

Mokhtari A, et al. (2025) Using multiomic integration to improve blood biomarkers of major depressive disorder: a case-control study. *EBioMedicine*, 113, 105569.

Bestehorn A, et al. (2025) Cytoplasmic mRNA decay controlling inflammatory gene expression is determined by pre-mRNA fate decision. *Molecular cell*, 85(4), 742.

Engel K, et al. (2025) Segment specific loss of NFAT5 function in the kidneys is sufficient to induce a global kidney injury like phenotype. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 39(2), e70352.

Fonseca-Pereira D, et al. (2025) The metabolic sensor LKB1 regulates ILC3 homeostasis and mitochondrial function. *Cell reports*, 44(4), 115456.

Zeiler C, et al. (2025) Protocol to distinguish pre-mRNA from mRNA in RNA-protein interaction studies. *STAR protocols*, 6(3), 103967.

Ayabe H, et al. (2025) Cellular crosstalk mediated by TGF-?? drives epithelial-mesenchymal transition in patient-derived multi-compartment biliary organoids. *Nature communications*, 16(1), 6575.

Laplane P, et al. (2025) Effect of MisMatch repair deficiency on metastasis occurrence in a syngeneic mouse model. *Neoplasia (New York, N.Y.)*, 62, 101145.

Li Y, et al. (2025) Integration of network pharmacology and transcriptomics to reveal the ROS/NLRP3/Caspase-1/GSDMD-mediated mechanism of baicalin in alleviating contrast-induced acute kidney injury. *Naunyn-Schmiedeberg's archives of pharmacology*.

Ren L, et al. (2025) Genomes reveal pervasive distant hybridization in nature among cyprinid fishes. *GigaScience*, 14.

Treaster MJ, et al. (2025) A Y-linked duplication of anti-Mullerian hormone is the sex determination gene in threespine stickleback. *bioRxiv : the preprint server for biology*.

Zhao Y, et al. (2025) An atlas of transcriptomic changes in human immune cells driven by 364 endogenous and gut-microbiota-derived metabolites. *Scientific reports*, 15(1), 13814.

Hsieh YH, et al. (2025) Telomeric repeat-containing RNA increases in aged human cells. *Nucleic acids research*, 53(13).

Mayorek N, et al. (2025) L-2-hydroxyglutarate regulates centromere and heterochromatin conformation in the male germline. *PLoS genetics*, 21(7), e1011785.

Singh I, et al. (2025) Epigenetic targeting of MECOM/KRAS axis by JIB-04 impairs tumorigenesis and cisplatin resistance in MECOM-amplified ovarian cancer. *Cell death discovery*, 11(1), 326.

Chen Y, et al. (2025) Non-disruptive 3D profiling of combinations of epigenetic marks in single cells. *bioRxiv : the preprint server for biology*.