Resource Summary Report

Generated by NIF on Aug 30, 2026

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: 20260829T182410+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 2601 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF .

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.

Zhao S, et al. (2026) RNF25 confers mRNA damage tolerance by curbing activation of the integrated stress response. Molecular cell, 86(7), 1275.

Kinget L, et al. (2026) Prognostic and Predictive Value of the Clearseq1-4 Tumor Microenvironment Classification in Localized and Metastatic Clear-Cell Renal Cell Carcinoma. Cancer research communications, 6(4), 884.

Harris Snell P, et al. (2026) Coordinated regulation of mRNA translation and stability by ZC3H7A and ZC3H7B RNA-binding proteins. Cell reports, 45(6), 117511.

Mukherjee S, et al. (2026) A single Citrobacter rodentium infection in Pink1 knockout and wild-type mice leads to regional blood-brain-barrier perturbation and limited microglial activation without dopamine neuron axon terminal loss. PLoS pathogens, 22(6), e1014315.

Choubey P, et al. (2026) Cohesin complex cooperates with PU.1 at super-enhancers to regulate the differentiation and identity of conventional dendritic cells. Cell reports, 45(5), 117331.

van de Grift YBC, et al. (2026) GRHL and PGR control WNT4 expression via 3D looping of conserved and species-specific enhancers. Cell reports, 45(7), 117575.

Kaur H, et al. (2026) Dynamicity of Topoisomerase 1 binding at super-enhancers regulates TLR-responsive gene expression program in macrophages. Cell reports, 45(7), 117604.

Schmitt AM, et al. (2026) Spatially-resolved subtype progression reveals metabolic vulnerabilities in pancreatic ductal adenocarcinoma. Molecular cancer, 25(1).

Ferreira S, et al. (2026) Inactivation of CDKN2AARF Promotes p53-Independent Remodeling of the PDAC Tumor Microenvironment. Cancer research, 86(13), 3109.

Kryeziu K, et al. (2026) Patient-derived organoids from metastatic colorectal cancer mirror tumor heterogeneity and predict patient survival and drug sensitivity. Cell reports. Medicine, 7(6), 102840.

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.

Gilad N, et al. (2026) Chronic activation of p38? in skeletal muscle causes necrotic changes, but also abolishes expression of MK2, MK3, and MKK6 and the muscle recovers. The Journal of biological chemistry, 302(4), 111338.

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.

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.

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

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

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