

Resource Summary Report

Generated by SPARC SAWG on Sep 15, 2026

cutadapt

RRID:SCR_011841

Type: Tool

Proper Citation

cutadapt (RRID:SCR_011841)

Resource Information

URL: <http://code.google.com/p/cutadapt/>

Proper Citation: cutadapt (RRID:SCR_011841)

Description: Software tool that removes adapter sequences from DNA sequencing reads., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: cutadapt

Resource Type: software resource

Defining Citation: [DOI:10.14806/ej.17.1.200](https://doi.org/10.14806/ej.17.1.200)

Keywords: bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: cutadapt

Resource ID: SCR_011841

Alternate IDs: OMICS_01086, biotools:cutadapt

Alternate URLs: <https://bio.tools/cutadapt>

Old URLs: <https://sources.debian.org/src/cutadapt/>

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260912T195740+0000

Ratings and Alerts

No rating or validation information has been found for cutadapt.

No alerts have been found for cutadapt.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 7088 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

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

Nitsch S, et al. (2026) H4K16 acylations destabilize chromatin architecture and facilitate transcriptional response during metabolic perturbations. *Molecular cell*, 86(1), 24.

Diaz-Jimenez D, et al. (2026) Timing of Glucocorticoid Treatment Dictates Glucocorticoid Receptor Actions Modulating the NLRP3-Inflammasome Activation in Macrophages. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 40(3), e71510.

Lehmann AE, et al. (2026) Aneuploidy promotes transient stress adaptation and metabolic flexibility in the human fungal pathogen *Aspergillus fumigatus*. *Current biology : CB*, 36(13), 3258.

Hasegawa K, et al. (2026) Age-related decline in nuclear envelope LINC complex drives neuronal aging via axon initial segment dysfunction. *EMBO reports*, 27(13), 3788.

Lu P, et al. (2026) Bromodomain-containing protein 4 knockdown promotes neuronal ferroptosis in a mouse model of subarachnoid hemorrhage. *Neural regeneration research*, 21(2), 715.

Liu YY, et al. (2026) Assessing whether rectal swabs reflect appendiceal microbiota profiles in acute appendicitis: a 16S rRNA-based comparative study. *Gut pathogens*, 18(1), 10.

Li Y, et al. (2026) BOLL-Containing Aggregates Mediate the Translational Regulation During Human Oogenesis. *Cell proliferation*, 59(4), e70181.

Tao H, et al. (2026) NSD1-Mediated PPAR γ Methylation Enhances PTEN Activity to Suppress Glycolysis and Tumor Progression in Endometrial Cancer. *Cancer research*, 86(10), 2464.

Oluwatola O, et al. (2026) Transdermal bicarbonate buffer therapy increases intratumoral pH and elicits antitumor responses in bladder cancer. *Frontiers in immunology*, 17, 1706250.

Hoang CV, et al. (2026) Genomic and biological characterization of lytic phages infecting *Pseudomonas syringae* associated with almond bacterial blast. *Scientific reports*, 16(1).

Friske JD, et al. (2026) A Conserved Enhancer Locus in Extrachromosomal DNA and Homogeneously Staining Regions Activates MYC Transcription in Group 3 Medulloblastoma. *Cancer research*, 86(13), 3160.

Xiangwen W, et al. (2026) A De Novo Splicing Mutation of SRP72 in Bone Marrow Failure Syndrome Type 1: Case Report and Review of the Literature. *Molecular genetics & genomic medicine*, 14(1), e70168.

Barajas JM, et al. (2026) Tandem Duplications in UBTF Create XPO1-Dependent Nuclear Export Signals that Reveal a Leukemic Therapeutic Dependency. *Blood cancer discovery*, 7(1), 51.

Williams EC, et al. (2026) Expression of endogenous retroviral elements is associated with extracellular matrix remodeling in prostate cancer. *Mobile DNA*, 17(1), 1.

Doré G, et al. (2026) First draft genome of the decaploid species, *Ludwigia grandiflora* subsp. *hexapetala*, validated through gene expression. *GigaByte (Hong Kong, China)*, 2026, gigabyte176.

Zhang W, et al. (2026) A novel duplication frameshift mutation in the BAG3 gene in a patient with dilated cardiomyopathy. *BMC cardiovascular disorders*, 26(1).

Palimirmo FS, et al. (2026) Genomic Insights and Matrilineal Evolution Reveal Potential Dispersal Patterns of Chitala Species (Osteoglossomorpha: Notopteridae) in the Sundaland Biodiversity Hotspot. *Ecology and evolution*, 16(3), e73257.

Isoyama S, et al. (2026) Cooperative Roles of Class IA PI3K Isoforms in Translocation-Related Sarcoma Cell Survival and Proliferation. *Cancer research communications*, 6(4), 976.

Lee M, et al. (2026) LINE-1 Locus Transcription Nucleates Oncogenic Chromatin Architecture. *Cancer discovery*, 16(4), 800.