Resource Summary Report

Generated by dkNET on Apr 25, 2025

Flexbar

RRID:SCR_013001
Type: Tool

Proper Citation

Flexbar (RRID:SCR_013001)

Resource Information

URL: http://sourceforge.net/projects/flexbar/

Proper Citation: Flexbar (RRID:SCR_013001)

Description: Flexible barcode and adapter removal for sequencing platforms.

Abbreviations: Flexbar

Resource Type: software resource

Defining Citation: DOI:10.3390/biology1030895

Keywords: bio.tools

Funding:

Resource Name: Flexbar

Resource ID: SCR_013001

Alternate IDs: biotools:flexbar

Alternate URLs: https://bio.tools/flexbar, https://sources.debian.org/src/flexbar/

Record Creation Time: 20220129T080313+0000

Record Last Update: 20250420T014628+0000

Ratings and Alerts

No rating or validation information has been found for Flexbar.

No alerts have been found for Flexbar.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 260 mentions in open access literature.

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

Domínguez M, et al. (2025) Genomics Reveal Population Structure and Intergeneric Hybridization in an Endangered South American Bird: Implications for Management and Conservation. Ecology and evolution, 15(1), e70820.

Oo JA, et al. (2025) Long non-coding RNAs direct the SWI/SNF complex to cell type-specific enhancers. Nature communications, 16(1), 131.

Xu M, et al. (2025) BASAL: a universal mapping algorithm for nucleotide base-conversion sequencing. Nucleic acids research, 53(2).

Brezski A, et al. (2024) A systematic analysis of circRNAs in subnuclear compartments. RNA biology, 21(1), 1.

Tian R, et al. (2024) Sirenian genomes illuminate the evolution of fully aquatic species within the mammalian superorder afrotheria. Nature communications, 15(1), 5568.

G MK, et al. (2024) Efficacy of Plyometric and TheraBand FlexBar Exercises in Tennis Elbow Patients: A Comparative Study. Cureus, 16(6), e61525.

Kim KQ, et al. (2024) eIF4F complex dynamics are important for the activation of the integrated stress response. Molecular cell, 84(11), 2135.

Reichel M, et al. (2024) ALBA proteins facilitate cytoplasmic YTHDF-mediated reading of m6A in Arabidopsis. The EMBO journal, 43(24), 6626.

La Ferlita A, et al. (2024) tRFUniverse: A comprehensive resource for the interactive analyses of tRNA-derived ncRNAs in human cancer. iScience, 27(2), 108810.

Gurgul A, et al. (2024) Cannabidiol (CBD) modulates the transcriptional profile of ethanol-exposed human dermal fibroblast cells. Journal of applied genetics, 65(4), 773.

Alvarez-Martinez M, et al. (2024) Blimp-1 and c-Maf regulate immune gene networks to protect against distinct pathways of pathobiont-induced colitis. Nature immunology, 25(5),

886.

Lin D, et al. (2024) Transcriptome and proteome profiling reveals TREM2-dependent and -independent glial response and metabolic perturbation in an Alzheimer's mouse model. The Journal of biological chemistry, 300(11), 107874.

Ghareeb AFA, et al. (2024) Host transcriptome response to heat stress and Eimeria maxima infection in meat-type chickens. PloS one, 19(2), e0296350.

Arifuzzaman M, et al. (2024) Dietary fiber is a critical determinant of pathologic ILC2 responses and intestinal inflammation. The Journal of experimental medicine, 221(5).

Kanwal N, et al. (2024) GPATCH4 regulates rRNA and snRNA 2'-O-methylation in both DHX15-dependent and DHX15-independent manners. Nucleic acids research, 52(4), 1953.

Kelbert M, et al. (2024) The zinc-finger transcription factor Sfp1 imprints specific classes of mRNAs and links their synthesis to cytoplasmic decay. eLife, 12.

Verheyden NA, et al. (2024) A high-resolution map of functional miR-181 response elements in the thymus reveals the role of coding sequence targeting and an alternative seed match. Nucleic acids research, 52(14), 8515.

Núñez-Lillo G, et al. (2024) A First Omics Data Integration Approach in Hass Avocados to Evaluate Rootstock-Scion Interactions: From Aerial and Root Plant Growth to Fruit Development. Plants (Basel, Switzerland), 13(5).

Li Y, et al. (2024) Landscape of RNA pseudouridylation in archaeon Sulfolobus islandicus. Nucleic acids research, 52(8), 4644.

Tzur Y, et al. (2024) Ribosomal protein L24 mediates mammalian microRNA processing in an evolutionarily conserved manner. Cellular and molecular life sciences : CMLS, 81(1), 55.