**Flexbar**

**RRID:** SCR_013001  
**Type:** Tool

**Proper Citation**

Flexbar (RRID:SCR_013001)

**Resource Information**

**URL:** [http://sourceforge.net/projects/flexbar/](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](https://doi.org/10.3390/biology1030895)

**Keywords:** bio.tools

**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:** 20240424T182929+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 236 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](http://FDI-Lab-SciCrunch.org).


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.

Arnold B, et al. (2024) hGRAD: A versatile "one-fits-all" system to acutely deplete RNA binding proteins from condensates. The Journal of cell biology, 223(2).


Rotinsulu DA, et al. (2023) Molecular Features and Antimicrobial Susceptibilities of Streptococcus equi ssp. equi Isolates from Strangles Cases in Indonesia. Veterinary sciences, 10(1).


Das A, et al. (2023) The RNA-binding protein RBP42 regulates cellular energy metabolism in mammalian-infective Trypanosoma brucei. mSphere, 8(5), e0027323.


Varma E, et al. (2023) Translational control of Ybx1 expression regulates cardiac function in response to pressure overload in vivo. Basic research in cardiology, 118(1), 25.


Mendes LW, et al. (2023) Impact of the fungal pathogen Fusarium oxysporum on the taxonomic and functional diversity of the common bean root microbiome. Environmental microbiome, 18(1), 68.

Jolly AJ, et al. (2023) Redistribution of the chromatin remodeler Brg1 directs smooth muscle-derived adventitial progenitor-to-myofibroblast differentiation and vascular fibrosis. JCI insight, 8(9).