

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 17, 2025

AS-Peak

RRID:SCR_000380

Type: Tool

Proper Citation

AS-Peak (RRID:SCR_000380)

Resource Information

URL: <http://sourceforge.net/projects/as-peak/>

Proper Citation: AS-Peak (RRID:SCR_000380)

Description: A software that utilizes a peak detection algorithm to identify RNA-protein binding sites.

Abbreviations: ASPeak

Resource Type: software resource

Defining Citation: [PMID:23929032](https://pubmed.ncbi.nlm.nih.gov/23929032/)

Keywords: rna, peak detection, protein, binding, site, algorithm, analysis

Funding:

Availability: Open Source, Freely Available

Resource Name: AS-Peak

Resource ID: SCR_000380

Alternate IDs: OMICS_00566

Alternate URLs: <https://omictools.com/aspeak-tool>

License: GNU General Public License version 3.0

Record Creation Time: 20220129T080201+0000

Record Last Update: 20250410T064554+0000

Ratings and Alerts

No rating or validation information has been found for AS-Peak.

No alerts have been found for AS-Peak.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Mikedis MM, et al. (2020) DAZL mediates a broad translational program regulating expansion and differentiation of spermatogonial progenitors. eLife, 9.

Kucukural A, et al. (2013) ASPeak: an abundance sensitive peak detection algorithm for RIP-Seq. Bioinformatics (Oxford, England), 29(19), 2485.