Atac
RRID:SCR_015980
Type: Tool

Proper Citation
Atac (RRID:SCR_015980)

Resource Information

URL: http://kmer.sourceforge.net
Proper Citation: Atac (RRID:SCR_015980)
Description: Alignment analysis software tool for comparative mapping between two genome assemblies or between two different genomes. It can cache intermediate results to speed a comparisons of multiple sequences.

Resource Type: Resource, image analysis software, data analysis software, data processing software, alignment software, software application, sequence analysis software, software resource

Keywords: software, tool, DNA, sequence, analysis, aligning, genome, compare, mapping, assembly, bio.tools

Availability: Free, Freely available, Available for download

Website Status: Last checked up

Resource Name: Atac
Resource ID: SCR_015980
Alternate IDs: biotools:atac
Alternate URLs: https://bio.tools/atac
No rating or validation information has been found for Atac.

No alerts have been found for Atac.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 259 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.


Li Y, et al. (2020) R-loops coordinate with SOX2 in regulating reprogramming to pluripotency. Science advances, 6(24), eaba0777.


Korchynska S, et al. (2020) Life-long impairment of glucose homeostasis upon prenatal exposure to psychostimulants. The EMBO journal, 39(1), e100882.


