Resource Summary Report

Generated by NIF on May 15, 2025

BS Seeker

RRID:SCR_005641
Type: Tool

Proper Citation

BS Seeker (RRID:SCR_005641)

Resource Information

URL: http://pellegrini.mcdb.ucla.edu/BS_Seeker/BS_Seeker.html

Proper Citation: BS Seeker (RRID:SCR_005641)

Description: Software which performs accurate and fast mapping of bisulfite-treated short reads. Supplementary information and examples are provided on the site.

Synonyms: Bisulfite Sequence Seeker

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

Defining Citation: PMID:20416082

Keywords: bisulfite sequencing, sequence analysis software, short read, sequence mapping, bio.tools

Funding:

Availability: Free, Available for download, Freely available

Resource Name: BS Seeker

Resource ID: SCR_005641

Alternate IDs: OMICS_00578, biotools:bs_seeker

Alternate URLs: https://bio.tools/bs_seeker

Record Creation Time: 20220129T080231+0000

Record Last Update: 20250514T061344+0000

Ratings and Alerts

No rating or validation information has been found for BS Seeker.

No alerts have been found for BS Seeker.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Lowe MG, et al. (2022) EED is required for mouse primordial germ cell differentiation in the embryonic gonad. Developmental cell, 57(12), 1482.

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.

Chen PY, et al. (2010) BS Seeker: precise mapping for bisulfite sequencing. BMC bioinformatics, 11, 203.