

Resource Summary Report

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

Defining Citation: [PMID:20416082](#)

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

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: 20260811T043236+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.