

Resource Summary Report

Generated by SPARC SAWG on Sep 12, 2026

BAIT

RRID:SCR_000511

Type: Tool

Proper Citation

BAIT (RRID:SCR_000511)

Resource Information

URL: <http://sourceforge.net/projects/bait/>

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Description: Software to create strand inheritance plots in data derived from the Strand-Seq sequencing protocol. The software is designed to be flexible with a range of species, and basic template folders can be called to read in species-specific data.

Abbreviations: BAIT

Synonyms: BAIT - Software to help analyse Strand-Seq data

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

Defining Citation: [PMID:24028793](https://pubmed.ncbi.nlm.nih.gov/24028793/)

Keywords: create strand inheritance plots, strand-seq, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: BAIT

Resource ID: SCR_000511

Alternate IDs: biotools:bait, OMICS_01531

Alternate URLs: <https://bio.tools/bait>

License: BSD License

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260912T195510+0000

Ratings and Alerts

No rating or validation information has been found for BAIT.

No alerts have been found for BAIT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [SPARC SAWG](#) .

Hills M, et al. (2013) BAIT: Organizing genomes and mapping rearrangements in single cells. Genome medicine, 5(9), 82.