

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

BAIT

RRID:SCR_000511

Type: Tool

Proper Citation

BAIT (RRID:SCR_000511)

Resource Information

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

Proper Citation: BAIT (RRID:SCR_000511)

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 processing software, data analysis software, software resource, software application, data visualization software

Defining Citation: [PMID: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: 20260805T044009+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 [RRID](#) .

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