

Resource Summary Report

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

CBaSE

RRID:SCR_027765

Type: Tool

Proper Citation

CBaSE (RRID:SCR_027765)

Resource Information

URL: <https://weghornlab.org/software.html>

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Description: Software tool which derives gene-specific probabilistic estimates of the strength of negative and positive selection in cancer.

Synonyms: Cancer Bayesian SElection estimation

Resource Type: software application, software resource, source code

Defining Citation: [PMID:29106416](#)

Keywords: Cancer Genes, SNV, indel, gene-specific probabilistic estimates, strength of negative and positive selection, cancer

Funding: NCI U54 CA143874;
NIMH R01 MH101244;
NIGMS R01 GM078598

Availability: Free, Available for download, Freely available

Resource Name: CBaSE

Resource ID: SCR_027765

Alternate URLs: <https://github.com/weghornlab/CBaSE>,
<http://genetics.bwh.harvard.edu/cbase>

Record Creation Time: 20251211T070811+0000

Record Last Update: 20260805T044618+0000

Ratings and Alerts

No rating or validation information has been found for CBaSE.

No alerts have been found for CBaSE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.