

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 18, 2026

TREESELECT

RRID:SCR_009422

Type: Tool

Proper Citation

TREESELECT (RRID:SCR_009422)

Resource Information

URL: <http://www.hsph.harvard.edu/faculty/alkes-price/software/>

Proper Citation: TREESELECT (RRID:SCR_009422)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 6th, 2023. Software package for inferring natural selection from unusual population differentiation between closely related populations. (entry from Genetic Analysis Software)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: TREESELECT

Resource ID: SCR_009422

Alternate IDs: nlx_154682

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260912T200246+0000

Ratings and Alerts

No rating or validation information has been found for TREESELECT.

No alerts have been found for TREESELECT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Rowan TN, et al. (2021) Powerful detection of polygenic selection and evidence of environmental adaptation in US beef cattle. PLoS genetics, 17(7), e1009652.

Blackburn HD, et al. (2017) A fine structure genetic analysis evaluating ecoregional adaptability of a Bos taurus breed (Hereford). PloS one, 12(5), e0176474.

Decker JE, et al. (2016) Origins of cattle on Chirikof Island, Alaska, elucidated from genome-wide SNP genotypes. Heredity, 116(6), 502.

Mersha TB, et al. (2015) Mapping asthma-associated variants in admixed populations. Frontiers in genetics, 6, 292.