

# Resource Summary Report

Generated by [kravitz2](#) on Sep 17, 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 [kravitz2](#) .

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.