

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

HTSNPER

RRID:SCR_009242

Type: Tool

Proper Citation

HTSNPER (RRID:SCR_009242)

Resource Information

URL: <http://www.chgb.org.cn/htSNPer/htSNPer.html>

Proper Citation: HTSNPER (RRID:SCR_009242)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 4, 2023. Java program for characterizing the haplotype block structure and selecting haplotype tagging SNP (entry from Genetic Analysis Software)

Abbreviations: HTSNPER

Synonyms: Haplotype Tagging SNPer

Resource Type: software application, software resource

Defining Citation: [PMID:15740612](#)

Keywords: gene, genetic, genomic, c++, jvm, ms-windows, linux

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: HTSNPER

Resource ID: SCR_009242

Alternate IDs: nlx_154403

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260727T040447+0000

Ratings and Alerts

No rating or validation information has been found for HTSNPER.

No alerts have been found for HTSNPER.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Cuervo-Alarcon L, et al. (2021) A candidate gene association analysis identifies SNPs potentially involved in drought tolerance in European beech (*Fagus sylvatica* L.). *Scientific reports*, 11(1), 2386.

Ding K, et al. (2005) htSNPer1.0: software for haplotype block partition and htSNPs selection. *BMC bioinformatics*, 6, 38.