

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

## HTSNPER

RRID:SCR\_009242

Type: Tool

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### Proper Citation

HTSNPER (RRID:SCR\_009242)

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### 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:** 20260905T133247+0000

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### Ratings and Alerts

No rating or validation information has been found for HTSNPER.

No alerts have been found for HTSNPER.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

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.