

Resource Summary Report

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

BPPH

RRID:SCR_009136

Type: Tool

Proper Citation

BPPH (RRID:SCR_009136)

Resource Information

URL: <http://www.csif.ucdavis.edu/~gusfield/bpph.html>

Proper Citation: BPPH (RRID:SCR_009136)

Description: Software application for inferring haplotypes from genotypes to determine if there are resulting haplotypes that fit a tree model (i.e. a perfect phylogeny, a coalescent). In population genetic terms, BPPH determines whether a set of SNP genotypes can be explained by haplotype pairs that could have evolved on a coalescent under the no-recombination, infinite sites model. (entry from Genetic Analysis Software)

Abbreviations: BPPH

Synonyms: Berkeley method for Perfect Phylogeny Haplotyping

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: BPPH

Resource ID: SCR_009136

Alternate IDs: nlx_154251

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260727T040444+0000

Ratings and Alerts

No rating or validation information has been found for BPPH.

No alerts have been found for BPPH.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.