Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

CHAPLIN

RRID:SCR_000833
Type: Tool

Proper Citation

CHAPLIN (RRID:SCR_000833)

Resource Information

URL: http://www.genetics.emory.edu/labs/epstein/software/chaplin/index.html

Proper Citation: CHAPLIN (RRID:SCR_000833)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022.Software application for identifying specific haplotypes or haplotype features that are associated with disease using genotype data from a case-control study. (entry from Genetic Analysis Software)

Abbreviations: CHAPLIN

Synonyms: Case-control HAPLotype INference package

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran90, (cvf 6.6) with imsl routines, ms-windows, (2000/xp)

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: CHAPLIN

Resource ID: SCR_000833

Alternate IDs: nlx_154266

Record Creation Time: 20220129T080203+0000

Record Last Update: 20260905T133222+0000

Ratings and Alerts

No rating or validation information has been found for CHAPLIN.

No alerts have been found for CHAPLIN.

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 PRECISE-TBI .

Bodoki L, et al. (2015) Vitamin D receptor gene polymorphisms and haplotypes in Hungarian patients with idiopathic inflammatory myopathy. BioMed research international, 2015, 809895.

Steinberg KK, et al. (2007) Genetic studies of a cluster of acute lymphoblastic leukemia cases in Churchill County, Nevada. Environmental health perspectives, 115(1), 158.