Resource Summary Report

Generated by RRID on Aug 5, 2026

VITESSE

RRID:SCR_009089
Type: Tool

Proper Citation

VITESSE (RRID:SCR_009089)

Resource Information

URL: http://watson.hgen.pitt.edu/register/soft_doc.html,

Proper Citation: VITESSE (RRID:SCR_009089)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16,2023. Software application that calculates likelihood on pedigrees (entry from Genetic Analysis Software)

Abbreviations: VITESSE

Synonyms: VITESSE (means speed in French)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, ansi c, unix, vms, ms-dos

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: VITESSE

Resource ID: SCR_009089

Alternate IDs: nlx_154106

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260805T044218+0000

Ratings and Alerts

No rating or validation information has been found for VITESSE.

No alerts have been found for VITESSE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 9 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Kennedy AC, et al. (2024) A robust evaluation of 49 high-dose-rate prostate brachytherapy treatment plans including all major uncertainties. Journal of applied clinical medical physics, 25(2), e14182.

Kennedy AC, et al. (2023) Being certain about uncertainties: a robust evaluation method for high-dose-rate prostate brachytherapy treatment plans including the combination of uncertainties. Physical and engineering sciences in medicine, 46(3), 1115.

Decker E, et al. (2008) PTHR1 loss-of-function mutations in familial, nonsyndromic primary failure of tooth eruption. American journal of human genetics, 83(6), 781.

Deng HW, et al. (2002) A genomewide linkage scan for quantitative-trait loci for obesity phenotypes. American journal of human genetics, 70(5), 1138.

Bartlett CW, et al. (2002) A major susceptibility locus for specific language impairment is located on 13q21. American journal of human genetics, 71(1), 45.

Huang QY, et al. (2002) Mutation patterns at dinucleotide microsatellite loci in humans. American journal of human genetics, 70(3), 625.

Gordon D, et al. (2001) A transmission/disequilibrium test that allows for genotyping errors in the analysis of single-nucleotide polymorphism data. American journal of human genetics, 69(2), 371.

Xu J, et al. (2001) Genomewide screen and identification of gene-gene interactions for asthma-susceptibility loci in three U.S. populations: collaborative study on the genetics of asthma. American journal of human genetics, 68(6), 1437.

Ewen KR, et al. (2000) Identification and analysis of error types in high-throughput genotyping. American journal of human genetics, 67(3), 727.