Resource Summary Report

Generated by NIF on May 17, 2025

ASP/ASPSHARE

RRID:SCR_009128
Type: Tool

Proper Citation

ASP/ASPSHARE (RRID:SCR_009128)

Resource Information

URL: http://www.uni-kiel.de/medinfo/mitarbeiter/krawczak/download/

Proper Citation: ASP/ASPSHARE (RRID:SCR_009128)

Description: Software applications including ASP, a power calculator for gene mapping using a sibpair design (concordant or discordant) and ASPSHARE, which complements ASP in that it allows rapid calculation of the expected ibd sharing at the trait locus, based upon the model parameters, and the incidence corresponding to the respective parameters. (entry from Genetic Analysis Software)

Abbreviations: ASP/ASPSHARE

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: ASP/ASPSHARE

Resource ID: SCR_009128

Alternate IDs: nlx_154233

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250513T061047+0000

Ratings and Alerts

No rating or validation information has been found for ASP/ASPSHARE.

No alerts have been found for ASP/ASPSHARE.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Pfefferle D, et al. (2016) Does the Structure of Female Rhesus Macaque Coo Calls Reflect Relatedness and/or Familiarity? PloS one, 11(8), e0161133.

Pfefferle D, et al. (2015) Male rhesus macaques use vocalizations to distinguish female maternal, but not paternal, kin from non-kin. Behavioral ecology and sociobiology, 69(10), 1677.

Semmler A, et al. (2011) Haplotype analysis of the 5,10-methylenetetrahydrofolate reductase (MTHFR) c.1298A>C (E429A) polymorphism. BMC research notes, 4, 439.