Resource Summary Report

Generated by dkNET on Apr 28, 2025

BAYESFST

RRID:SCR_013479
Type: Tool

Proper Citation

BAYESFST (RRID:SCR_013479)

Resource Information

URL: http://www.reading.ac.uk/Statistics/genetics/software.html

Proper Citation: BAYESFST (RRID:SCR_013479)

Description: Software application for Bayesian estimation of the coancestry coefficient FST (entry from Genetic Analysis Software)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c

Funding:

Resource Name: BAYESFST

Resource ID: SCR_013479

Alternate IDs: nlx_154236, biotools:bayesfst

Alternate URLs: https://bio.tools/bayesfst

Record Creation Time: 20220129T080316+0000

Record Last Update: 20250428T053753+0000

Ratings and Alerts

No rating or validation information has been found for BAYESFST.

No alerts have been found for BAYESFST.

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 dkNET.

Steele CD, et al. (2014) Worldwide F(ST) estimates relative to five continental-scale populations. Annals of human genetics, 78(6), 468.

Currat M, et al. (2002) Molecular analysis of the beta-globin gene cluster in the Niokholo Mandenka population reveals a recent origin of the beta(S) Senegal mutation. American journal of human genetics, 70(1), 207.