

Resource Summary Report

Generated by [dkNET](#) on Sep 5, 2026

MIXSCORE

RRID:SCR_009076

Type: Tool

Proper Citation

MIXSCORE (RRID:SCR_009076)

Resource Information

URL: <http://www.hsph.harvard.edu/faculty/alkes-price/software/>

Proper Citation: MIXSCORE (RRID:SCR_009076)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16,2023. Software application using a method for combining SNP association and admixture association statistics to increase power in GWAS in admixed populations. (entry from Genetic Analysis Software)

Abbreviations: MIXSCORE

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: MIXSCORE

Resource ID: SCR_009076

Alternate IDs: nlx_154074

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260904T000439+0000

Ratings and Alerts

No rating or validation information has been found for MIXSCORE.

No alerts have been found for MIXSCORE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Raj T, et al. (2017) Genetic architecture of age-related cognitive decline in African Americans. *Neurology. Genetics*, 3(1), e125.

Hinks A, et al. (2013) Dense genotyping of immune-related disease regions identifies 14 new susceptibility loci for juvenile idiopathic arthritis. *Nature genetics*, 45(6), 664.

Liu Y, et al. (2013) Softwares and methods for estimating genetic ancestry in human populations. *Human genomics*, 7(1), 1.

Zaitlen N, et al. (2012) Informed conditioning on clinical covariates increases power in case-control association studies. *PLoS genetics*, 8(11), e1003032.

Pasaniuc B, et al. (2011) Enhanced statistical tests for GWAS in admixed populations: assessment using African Americans from CARE and a Breast Cancer Consortium. *PLoS genetics*, 7(4), e1001371.

Price AL, et al. (2011) Single-tissue and cross-tissue heritability of gene expression via identity-by-descent in related or unrelated individuals. *PLoS genetics*, 7(2), e1001317.