

# Ancient admixture in human history genetics

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**What is the Patterson's D statistic?** Patterson's D, also known as the ABBA-BABA statistic, and related statistics such as the  $f_4$  -ratio, are commonly used to assess evidence of gene flow between populations or closely related species.

**What is the F4 ratio?** F4 ratio estimation allows inference of the mixing proportions of an admixture event, even without access to accurate surrogates for the ancestral populations. qpF4ratio requires that the input data is available in EIGENSTRAT format. To convert to the appropriate format, one can use CONVERTF program. See README.

**What is the F4 statistic in population genetics?** F4-ratio estimation (2009), is a method for estimating ancestry proportions in an admixed population, under the assumption that we have a correct historical model. Consider the phylogeny of Figure 4. The population X is an admixture of populations B? and C? (possibly with subsequent drift).

**What are D statistics?** A common tool for detecting ancient admixture events is the D-statistic. The D-statistic is based on the hypothesis of a genetic relationship that involves four populations, whose correctness is assessed by evaluating specific coincidences of alleles between the groups.

**What is the Durbin Watson D statistic?** The Durbin Watson statistic is a test for autocorrelation in a regression model's output. The DW statistic ranges from zero to four, with a value of 2.0 indicating zero autocorrelation. Values below 2.0 mean there is positive autocorrelation and above 2.0 indicates negative autocorrelation.

**What does the D statistic tell us?** A large Cohen's  $d$  indicates the mean difference (effect size = signal) is large compared to the variability (noise). For example, if Group A's Mean = 12 and Group B's Mean = 8, and the pooled standard deviation is 2, Cohen's  $d$  equals the following: The mean difference is twice the variability.

**What is the epistasis 13 3 ratio?** The phenotypic ratio is therefore 13:3. This type of epistasis is sometimes called dominant suppression, because the deviation from 9:3:3:1 is caused by a single allele that produces a dominant phenotype, and the action of this allele is to suppress the expression of some other gene.

**What is the CR4 ratio?** For example, the four-firm concentration ratio (CR4) refers to the market share of the four largest firms. The higher the concentration ratio, the more concentrated the industry. A market is generally considered highly concentrated if the CR4 is greater than 50 percent.

**What is the 4 0 ratio in genetics?** There is a 100% visibility rate in the single dominant phenotype. As there is no second phenotype, there is no phenotypic ratio. If we did put this result as a ratio, it would be 4:0. The genotypic ratio, however, does not look at the observable trait (the phenotype) but at potential allele combinations.

**What is the FIS in genetics?** FIS (inbreeding coefficient) is the proportion of the variance in the subpopulation contained in an individual. High FIS implies a considerable degree of inbreeding. Related measures:  $q$  (theta) of Weir and Cockerham (1984) and  $G_{ST}$  of Nei (1973, 1978). [See Weir, 1996; Avise, Box 6.3, p. 206].

**What is QQ in genetics?** • To search for evidence of systematic bias (from unrecognized population structure, analytical approach, genotyping artifacts, etc.), a quantile-quantile (QQ) plot can be used to characterize the extent to which the observed distribution of the test statistic follows the expected (null) distribution.

**What is the Wahlund effect?** In population genetics, the Wahlund effect is a reduction of heterozygosity (that is when an organism has two different alleles at a locus) in a population caused by subpopulation structure.

**What is an Abba Baba test?** The test considers ancestral ("A") and derived ("B") alleles and is based on the prediction that two particular single nucleotide

polymorphism (SNP) patterns, termed “ABBA” and “BABA” (see Materials and Methods), should be equally frequent under a scenario of incomplete lineage sorting without gene flow.

**What does UD mean in statistics?** In this example,  $\bar{d}$  is the mean value of the differences  $d$  for the population of all pairs of data, where each individual difference  $d$  is defined as the measurement from the right arm minus the measurement from the left arm.

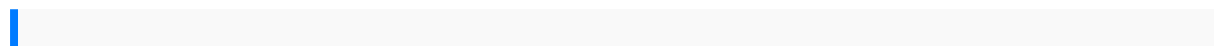
**What is drift in biology?** Genetic drift is the change in frequency of an existing gene variant in the population due to random chance. Genetic drift may cause gene variants to disappear completely and thereby reduce genetic variation. It could also cause initially rare alleles to become much more frequent, and even fixed.

**How do you interpret D statistic?** A positive D-statistic (i.e. an excess of ABBA) points to introgression between P2 and P3, whereas a negative D-statistic (i.e. an excess of BABA) points to introgression between P1 and P3. A Z-score can be calculated to assess the significance of the D-statistic.

**What is the meaning of Hopkins statistic?** Hopkins statistic (Hopkins and Skellam 1954) can be used to test for spatial randomness of data and for detecting clusters in data.

**What is a statistically significant D value?** Cohen suggested that  $d = 0.2$  be considered a “small” effect size, 0.5 represents a “medium” effect size and 0.8 a “large” effect size. This means that if the difference between two groups’ means is less than 0.2 standard deviations, the difference is negligible, even if it is statistically significant.

**What is  $\bar{d}$  statistics?**  $\bar{d}$  = mean difference in exhaustion times if everyone in the population were to run under both conditions.



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