

Genetic diversity of the Griffon vulture population in Serbia and its importance for conservation efforts in the Balkans

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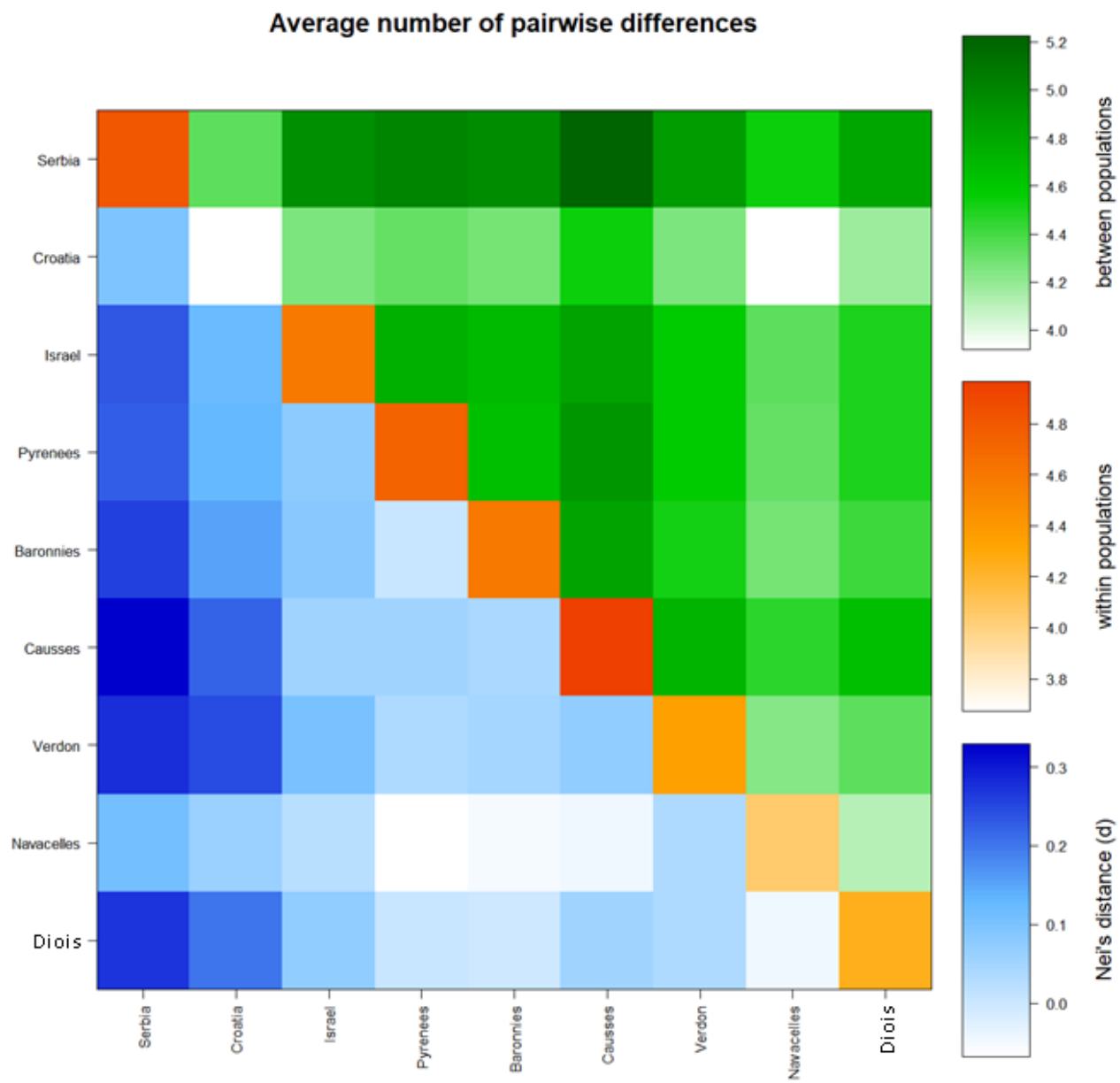
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Figure S1 Matrix of the average number of pairwise differences and Nei's distances detected in native and introduced *G. fulvus* populations based on the analysis of microsatellite loci. The average number of pairwise differences between populations is presented above diagonal, the average number of pairwise differences within the population is presented diagonal and Nei's distances are presented below diagonal. The average number of pairwise differences are presented in Table 1.

Matrix of pairwise F_{ST}

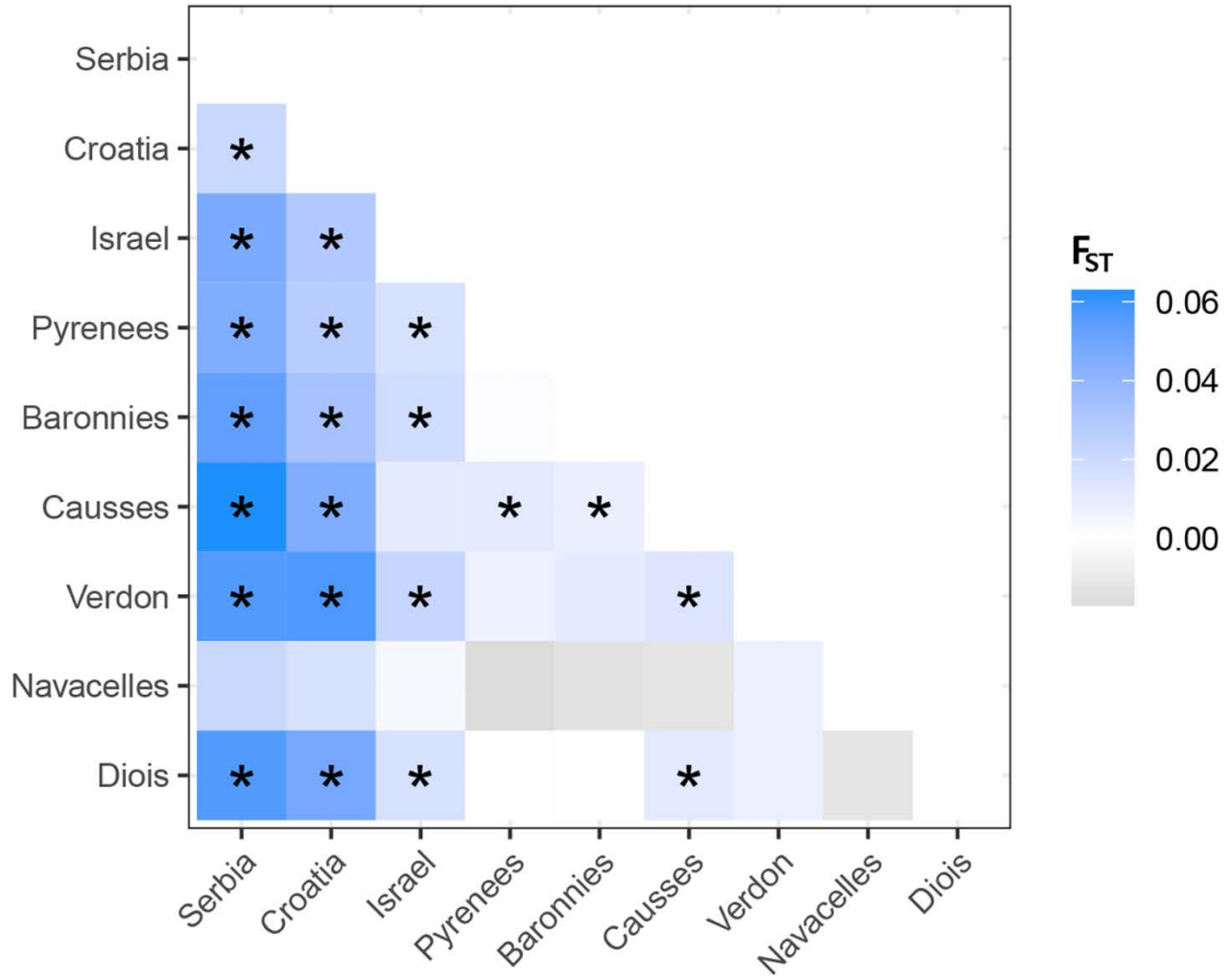


Figure S2 Matrix of pairwise F_{ST} distances between the *G. fulvus* population of Serbia and other native and introduced *G. fulvus* populations based on the analysis of 10 microsatellite loci. Statistically significant F_{ST} values are marked with an asterisk (*). Population pairwise F_{ST} values are presented in Table 3.

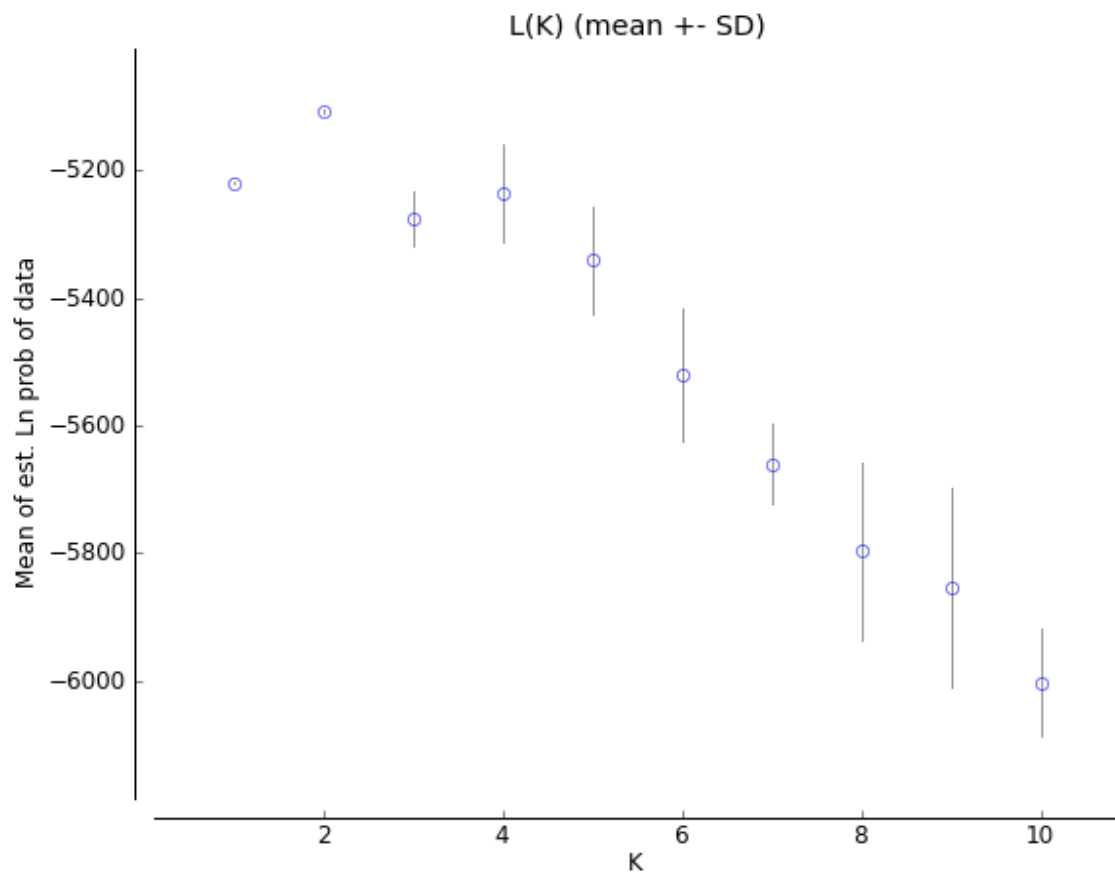


Figure S3 Ln values of probability for the assumed number of genetic clusters.

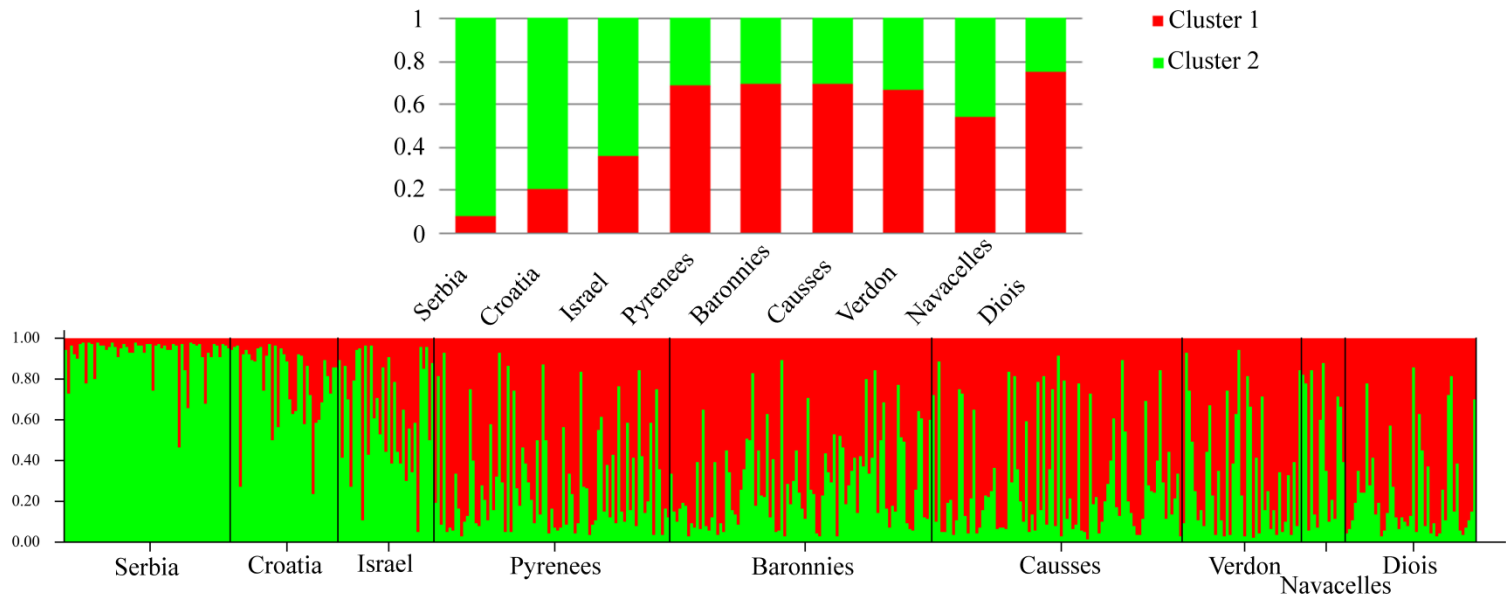


Figure S4 (a) Proportions of inferred STRUCLURE clusters ($K=2$) from the native and introduced populations.

(b) Proportions of the inferred STRUCLURE clusters ($K=2$) from the individuals.