SUPPLEMENTARY MATERIALS

Genetic position of Hungarian Grey among European cattle and identification of breed-specific markers.

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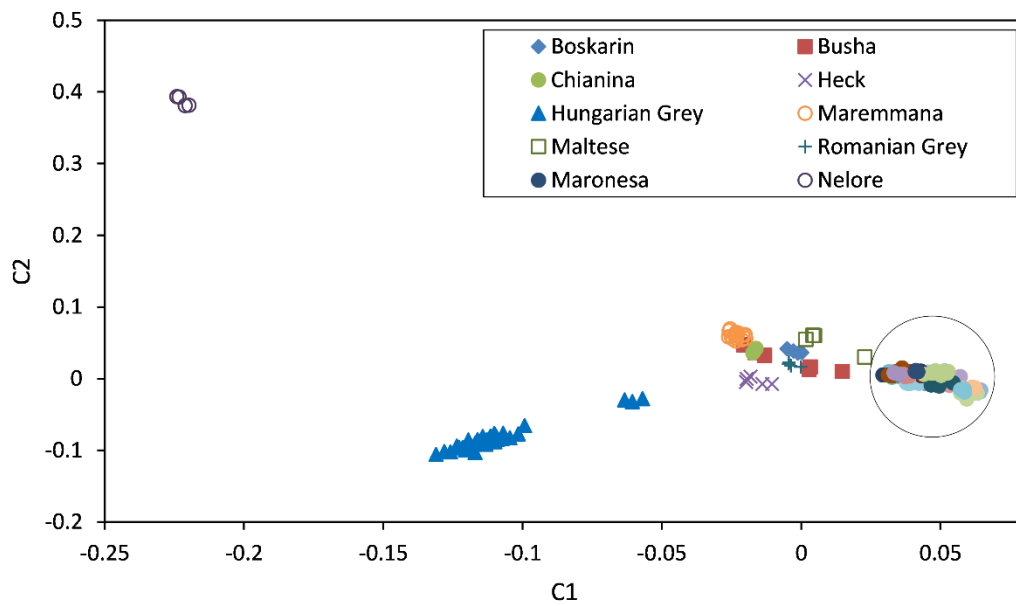
Supplementary Figure S1 Multidimensional scaling plot of Hungarian Grey and European cattle populations. Nelore was selected as an outgroup. Encircled breeds are from Upadhyay *et al.* (2017) including Fleckvieh and Holstein-Friesian sampled in Hungary. The following codes are used: C1=1st component (eigenvalue=7.681); C2=2nd component (eigenvalue=2.797)

Supplementary Figure S2 Graphical representation of pairwise coefficients of genetic differentiation (F_{ST}) values of Hungarian Grey cattle and the presented cattle breeds.
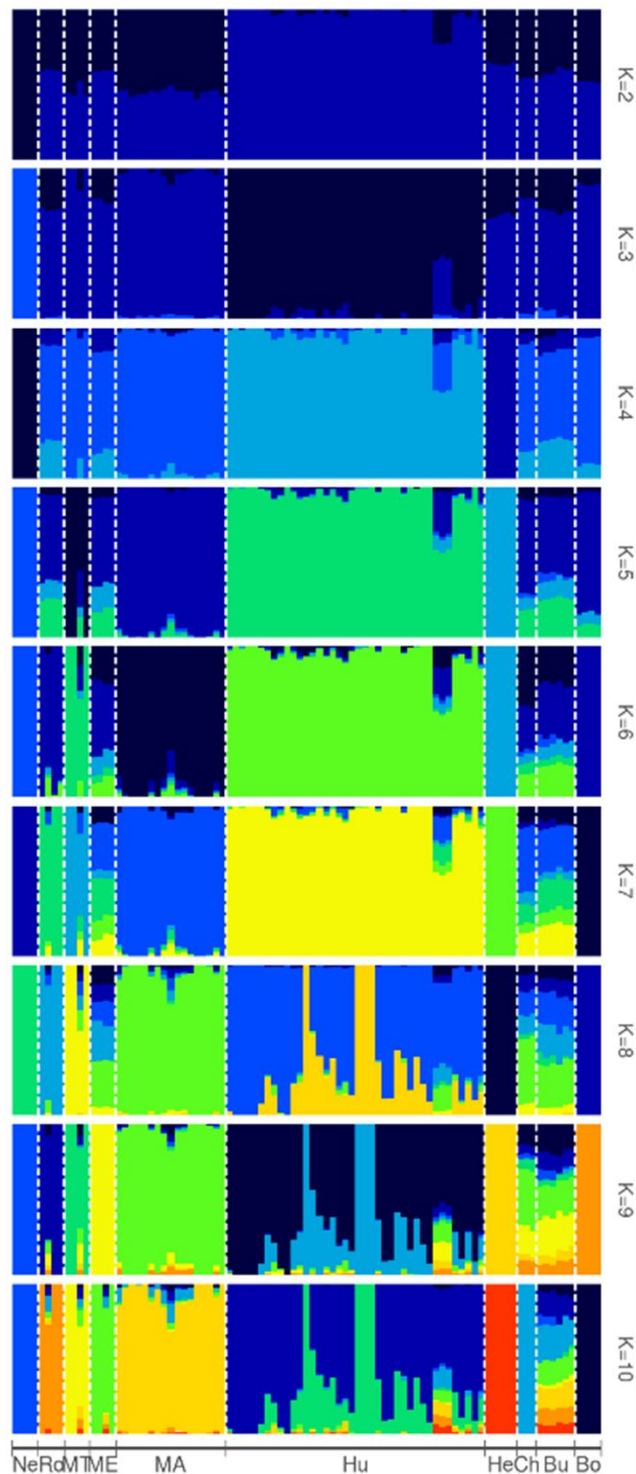
Supplementary Figure S3 Admixture analysis of Hungarian Grey and selected cattle populations for a range of K-values (K = 2-10). Each individual is represented by a single column divided into K colored segments, where K is the number of assumed clusters. Populations are separated by dashed lines. The following codes are used: Ne = Nelore; RO = Romanian Grey; MT = Maltese; ME = Maronese; MA = Maremmana; Hu = Hungarian Grey; He = Heck; Ch = Chianina; Bu = Busha; Bo = Boskarin