

Mattucci and Oliveira *et al.* Supplementary Figure S1.

Plot of delta K and mean likelihood L(K) as a function of K averaged over five independent runs of STRUCTURE run with the ‘admixture and the *F* model’. The Y-error bars are the standard deviation, and K is the assumed number of genetic clusters. Plot A has been obtained using all sampled European wild (*Fsi*) and domestic cats (*Fca*), including the admixed Hungarian and Scottish cats populations. Plot B has been obtained using only the European wildcat populations.

