

SUPPLEMENTARY MATERIALS FOR THE ARTICLE:

**Differentiation of the domestic pig and wild boar using genotyping-by-sequencing**

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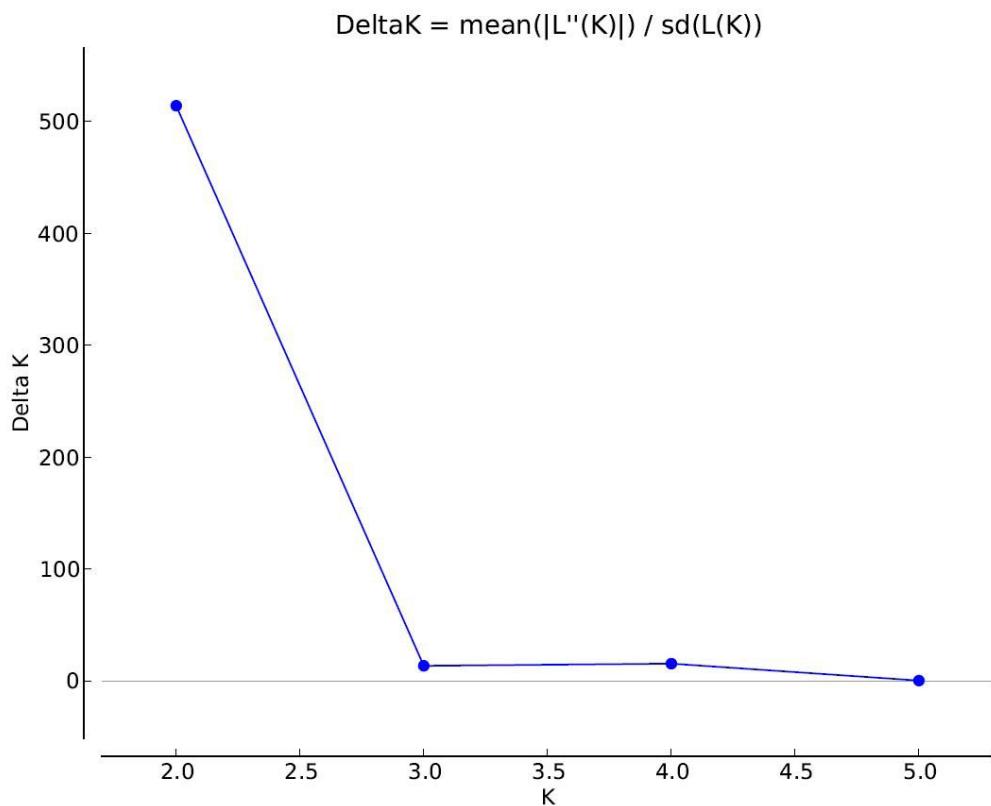
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**Supplementary Material 1.**

Suppl. Mat. 1 – Fig. 1. (below).

Delta K values for different numbers of populations assumed; K values ranging from one to six.



Suppl. Mat. 1 – Fig. 2. (next page).

Sliding-window averaged FST values between pigs and wild boars. A 5-SNP sliding window-averaged pairwise FST values were plotted against centered genomic positions of the windows. Threshold line marks top 1% of observations with FST above 0.426.

