

Supplementary Material

Setting up decision-making tools towards a quality-oriented participatory maize breeding program

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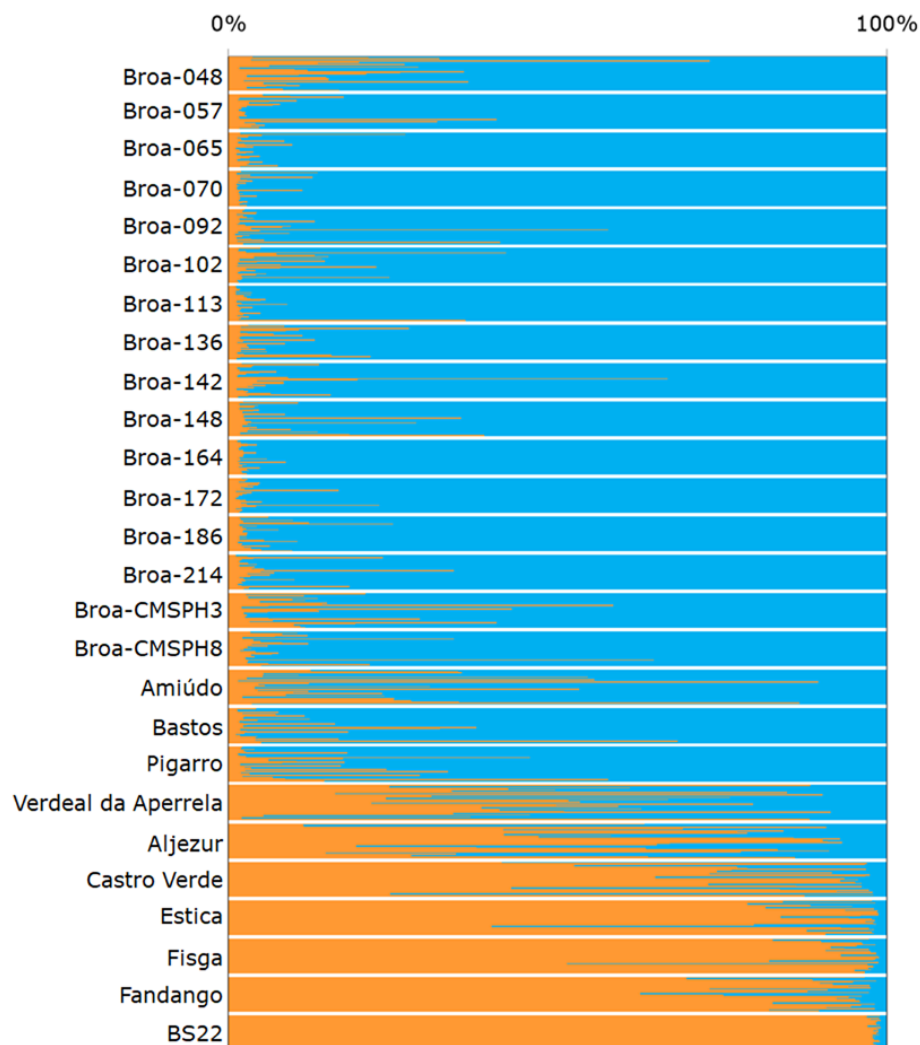


Figure S1. Proportion of membership of each maize population in each of the two gene pools inferred from multi-locus microsatellite data using a model-based clustering method. Each horizontal line within each population corresponds to an individual plant. Gene pool A is depicted in blue; gene pool B is depicted in orange. The *BS22(R)C6* synthetic population from the US, abbreviated for *BS22* in the figure.