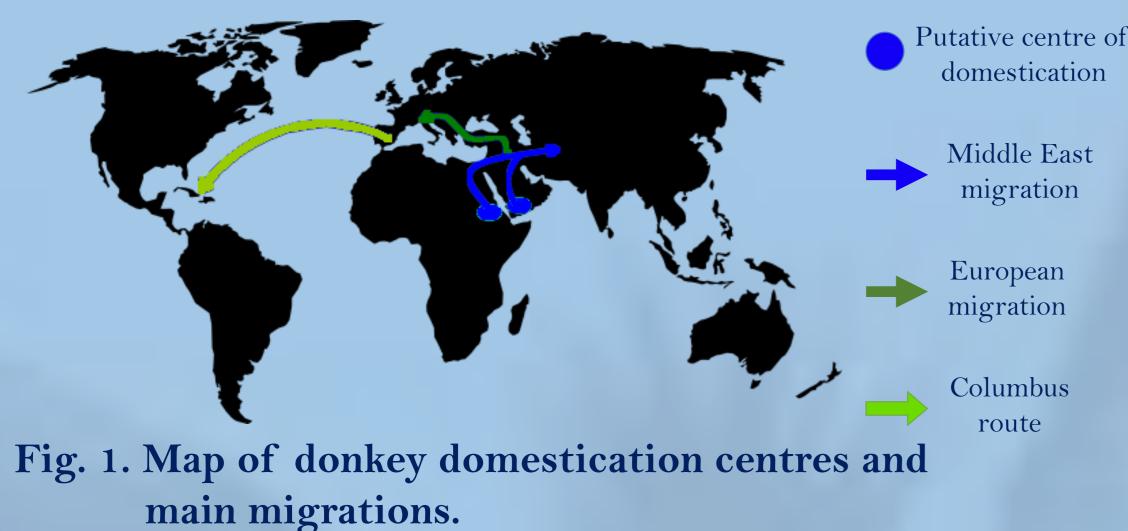
UAB **Universitat Autònoma** de Barcelona

GENETIC RELATIONSHIPS AMONG CUBAN DONKEY POPULATIONS



Valldeoriola Cardó, Anna – July 2016

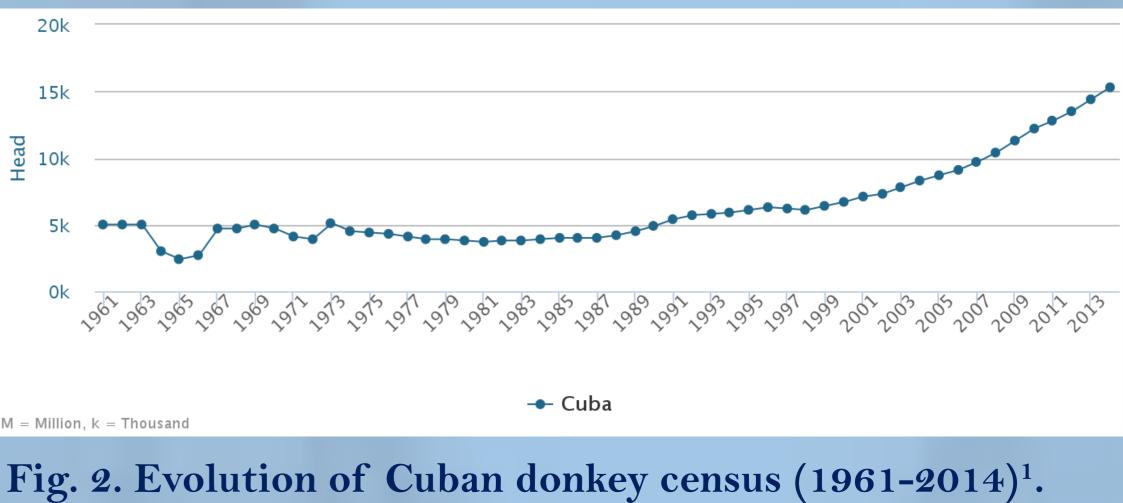




Cuba

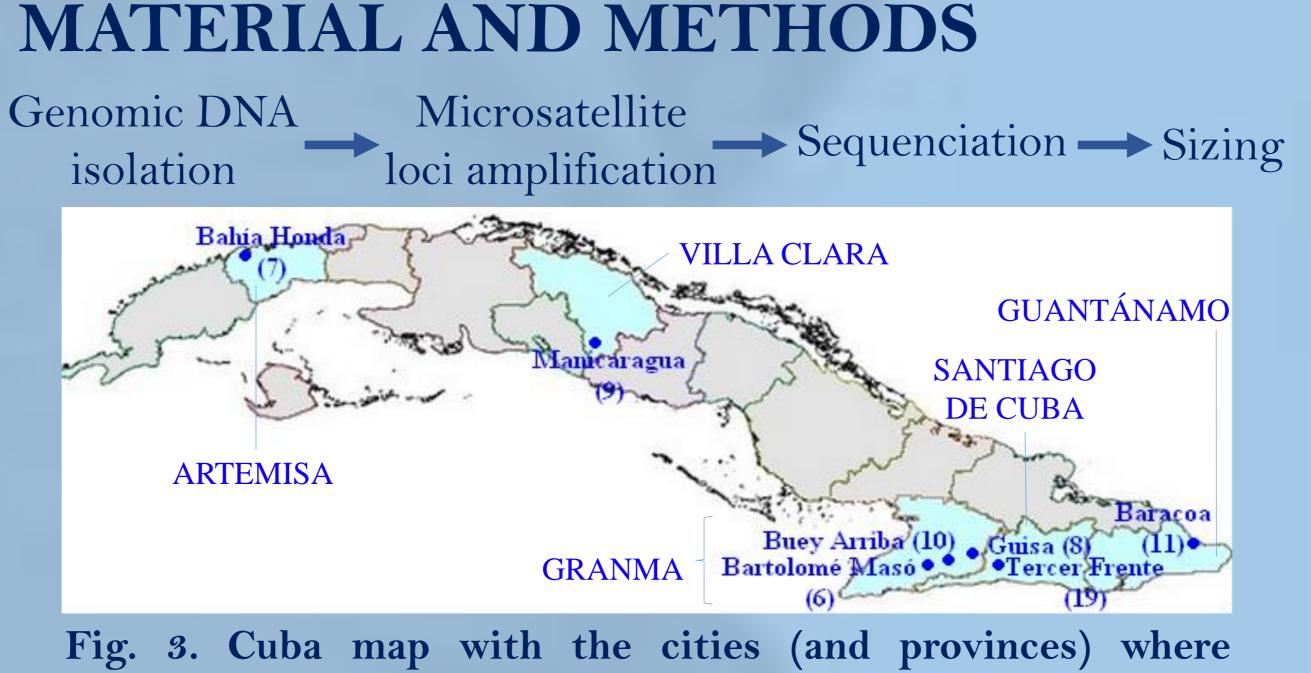
Donkeys were mainly used to breed mules in Cuba, which are used in mountainous regions. For this reason, donkeys are mostly found there, too.

However, it has changed in the last two decades (Fig. 2). A genetic improvement program in mule breeding, which affected donkeys and horses, has contributed.



OBJECTIVES

- Genetic characterization of Cuban donkey population and its subpopulations.
- Evaluation of population genetic structure, following geographic patterns.



samples were taken from and the number of it in brackets.

Genetic diversity analyses

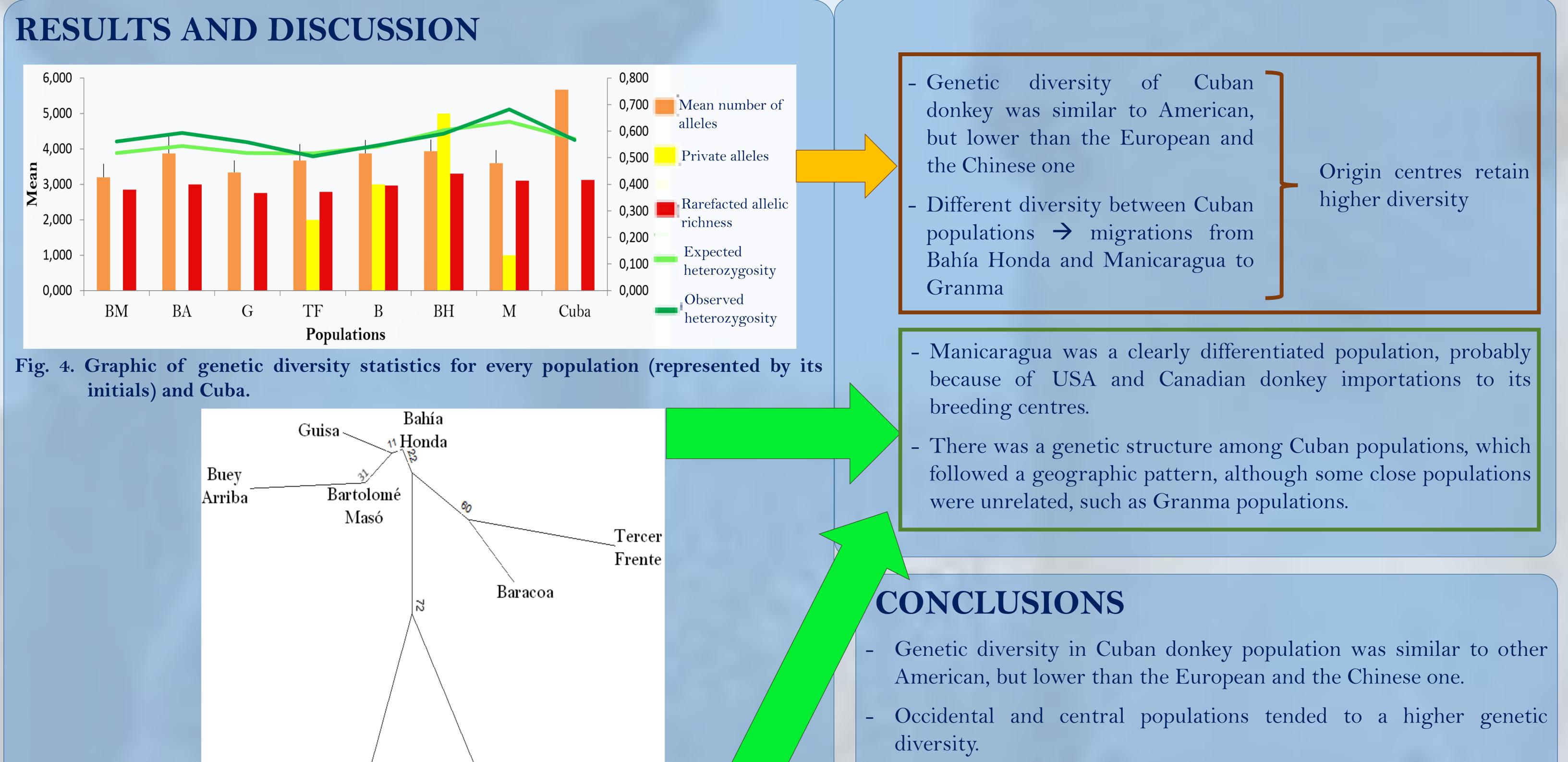
All diversity parameters, except from rarefacted allelic richness (rAR) - GENALEX v.6.502^{2,3}.

Genetic differentiation

Reynolds distance (D_R) matrix and the resulting genetic distance tree after 1000 bootstraps - POPULATIONS $V.1.2.32^4$.

Bayesian inference

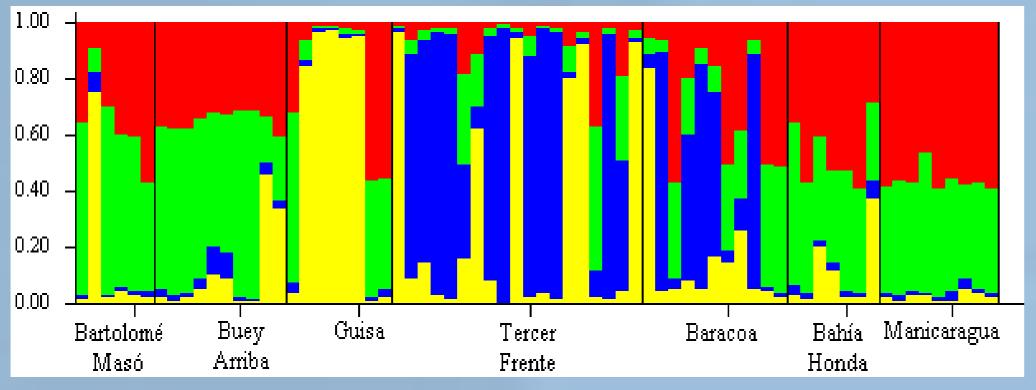
Population clusters - STRUCTURE v.2.3.4⁵.



0.002 Catalana

Raça

Fig. 5. Unrooted Neighbor-Joining distance tree with the weighted D_{R} after 1000 bootstraps using Catalan breed as an outgroup.



Manicaragua

Fig. 6. Summary of Q estimations for the most probable K (K=4). Every vertical fragmented line represents an individual. It is divided in K colours, according to its probability of belonging to that cluster.

Population genetic structure was demonstrated in Cuba, which mostly followed geographic pattern.

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