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To cite this article: M. Pellecchia, L. Colli, D. Bigi, P. Zambonelli, Verini Supplizi, L. Liotta, R. Negrini & Ajmone Marsan (2007) Mitochondrial DNA diversity of five Italian autochthonous donkey breeds, *Italian Journal of Animal Science*, 6:sup1, 185-185, DOI: [10.4081/ijas.2007.1s.185](https://doi.org/10.4081/ijas.2007.1s.185)

To link to this article: <https://doi.org/10.4081/ijas.2007.1s.185>



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Published online: 15 Mar 2016.



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# Mitochondrial DNA diversity of five Italian autochthonous donkey breeds

M. Pellecchia<sup>1</sup>, L. Colli<sup>1</sup>, D. Bigi<sup>2</sup>, P. Zambonelli<sup>2</sup>,  
A. Verini Supplizi<sup>3</sup>, L. Liotta<sup>4</sup>, R. Negrini<sup>1</sup>, P. Ajmone Marsan<sup>1</sup>

<sup>1</sup> Istituto di Zootecnica. Università Cattolica del Sacro Cuore, Piacenza, Italy

<sup>2</sup> Dipartimento di Produzione e valorizzazione Agroalimentare. Università di Bologna, Italy

<sup>3</sup> Dipartimento di Patologia, Diagnostica e Clinica Veterinaria. Università di Perugia, Italy

<sup>4</sup> Dipartimento di Morfologia Biochimica, Fisiologia e Produzioni Animali.  
Università di Messina, Italy

*Corresponding author:* Marco Pellecchia. Istituto di Zootecnica, Facoltà di Agraria, Università Cattolica del Sacro Cuore. Via Emilia Parmense 84, 29100 Piacenza, Italy - Tel. +39 0523 593204 - Fax: +39 0523 599276 - E-mail: marco.pellecchia@unicatt.it

## ABSTRACT

To investigate the mitochondrial DNA diversity of five Italian donkey breeds (Amiata, Martinafranca, Romagnolo, Asinara, and Ragusano), we sequenced the HVR I region (D-loop, 288 bp) and *cytochrome b* gene (274 bp) in 121 individuals. In the D-loop we found nineteen mutations corresponding to fourteen different haplotypes, while in *cyt b* coding gene only six mutations were found, originating five different haplotypes. In particular, three mutations out of six were non-synonymous, causing an aminoacidic substitution. About the D-loop region, the value of nucleotide diversity ( $\pi$ ) observed within breeds was relatively low, but not far from values detected in other European breeds. Phylogenetic and network analyses disclosed the presence of two divergent maternal lineages within Italian donkeys. These haplogroups correspond to the well known lineages of ancestors (*Equus asinus somaliensis* and *E. a. africanus*), as donkeys were domesticated from distinct wild subspecies living in Eastern Africa regions. In four of the investigated breeds we detected the presence of both mtDNA lineages, while Amiata donkeys were characterized by mitochondrial haplotypes belonging to the *E. a. somaliensis* lineage only. The genetic relationships between the Italian populations are discussed and interpreted according to the most recent bibliographic data.