

SUPPLEMENTARY MATERIALS FOR THE ARTICLE:

Skrzetuska W., Mackowski M., Borowska A., Kalski R., Musiał A., Bieniek A., Ropka-Molik K., Cieslak J.

**Studies of mitochondrial DNA D-loop sequence variation may support the Polish Primitive Horse (Konik) conservation programme**

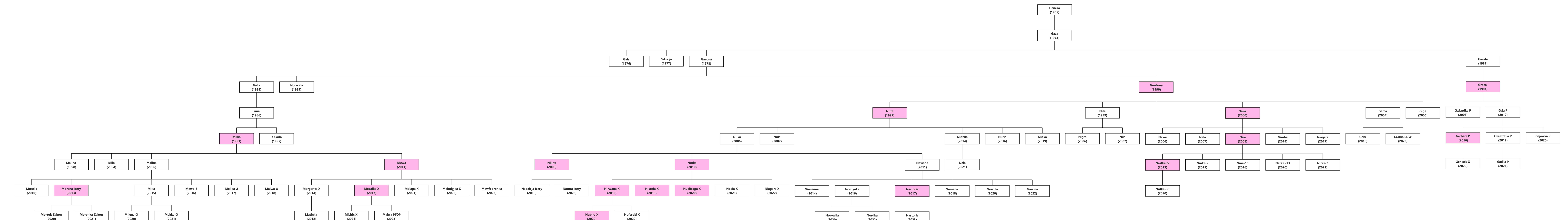
Folia Biologica (Kraków), vol. 73 (2025), No 1.

DOI: [https://doi.org/10.3409/fb\\_73-1.02](https://doi.org/10.3409/fb_73-1.02)

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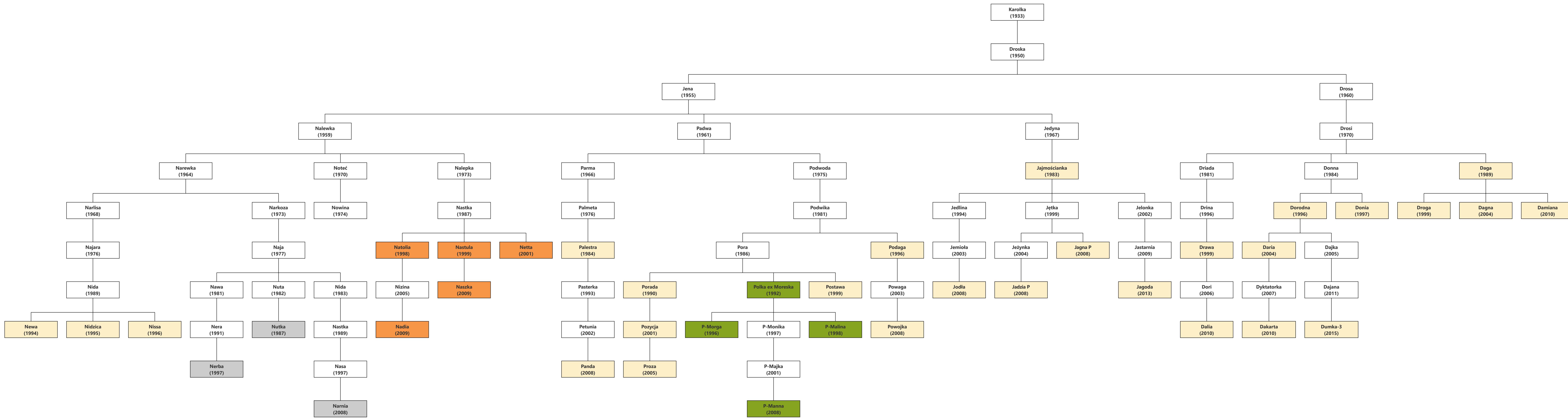
Figures S1-S4: Geneza, Karolka, Tunguska and Tarpanka lines pedigree charts.

**Fig. S1. Geneza line pedigree chart (simplified). Horses marked with the same color represent identical (molecularly confirmed) mtDNA haplotype.**



■ sample molecularly tested  
■ horses carried PPH6 mtDNA haplotype

**Fig. S2. Karolka line pedigree chart (simplified). Horses marked with the same color represent identical (molecularly confirmed) mtDNA haplotype.**



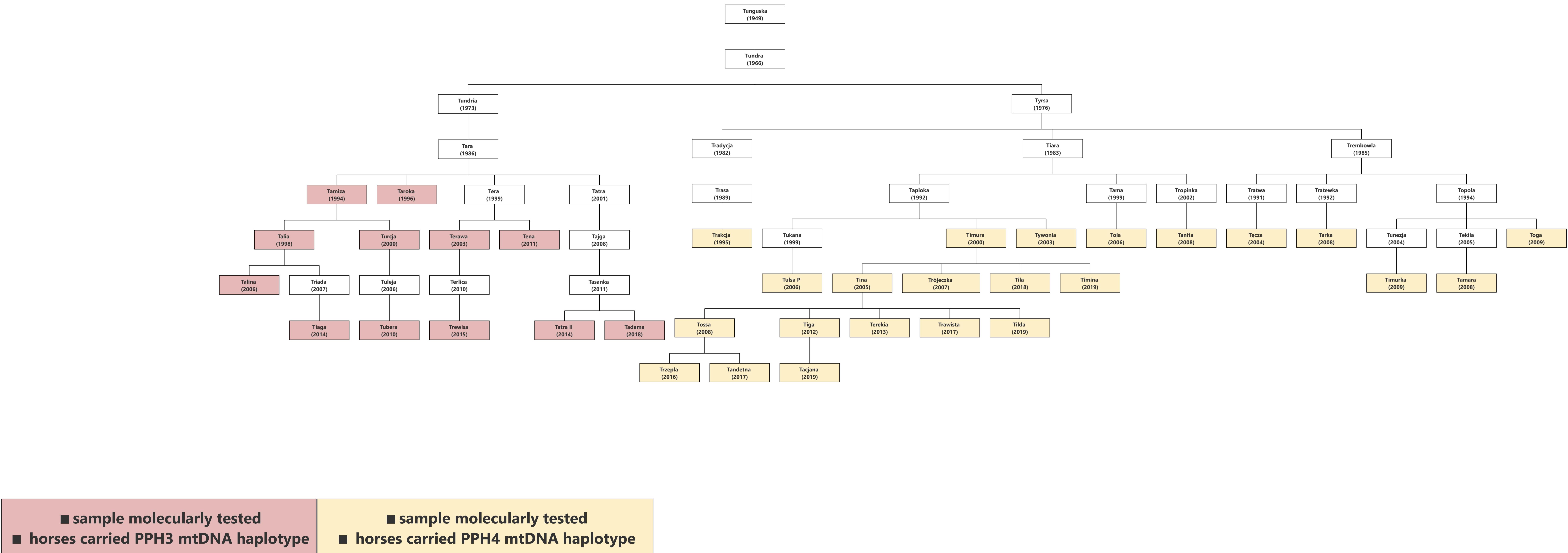
■ sample molecularly tested  
■ horses carried PPH4 mtDNA haplotype

■ sample molecularly tested  
■ horses carried PPH13 mtDNA haplotype

■ sample molecularly tested  
■ horses carried PPH14 mtDNA haplotype

■ sample molecularly tested  
■ horses carried PPH19 mtDNA haplotype

**Fig. S3. Tunguska line pedigree chart (simplified). Horses marked with the same color represent identical (molecularly confirmed) mtDNA haplotype.**



**Fig. S4. Tarpanka line pedigree chart (simplified). Horses marked with the same color represent identical (molecularly confirmed) mtDNA haplotype.**

