## HUMAN GENETICS =

## Gene Pool of Ethnic Groups of the Caucasus: Results of Integrated Study of the Y Chromosome and Mitochondrial DNA and Genome\_Wide Data

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Abstract—Genetic diversity has been analyzed in 22 ethnic groups of the Caucasus on the basis of data on Y\_chromosome and mitochondrial DNA (mtDNA) markers, as well as genome\_wide data on autosomal sin\_ gle\_nucleotide polymorphisms (SNPs). It has been found that the West Asian component is prevailing in all ethnic groups studied except for Nogays. This Near Eastern ancestral component has proved to be character\_ istic of Caucasian populations and almost entirely absent in their northern neighbors inhabiting the Eastern European Plain. Turkic\_speaking populations, except Nogays, did not exhibit an increased proportion of Eastern Eurasian mtDNA or Y\_chromosome haplogroups compared to some Abkhaz–Adyghe populations (Adygs and Kabardians). Genome\_wide SNP analysis has also shown substantial differences of Nogays from all other Caucasian populations studied. However, the characteristic difference of Nogays from other popu\_ lations of the Caucasus seems somewhat ambiguous in terms of the R1a1a\_M17(M198) and R1b1b1\_M73 haplogroups of the Y chromosome. The state of these haplogroups in Turkic\_speaking populations of the Caucasus requires further study.

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