

## ПРЕДВАРИТЕЛЬНЫЕ ДАННЫЕ О МОЛЕКУЛЯРНО-ГЕНЕТИЧЕСКОЙ СТРУКТУРЕ

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## PRELIMINARY DATA ON THE MOLECULAR GENETIC STRUCTURE OF *PELOPHYLAX RIDIBUNDUS* (AMPHIBIA: ANURA: RANIDAE) FROM THE SOUTHERN PART OF THE CRIMEAN PENINSULA, BASED ON MITOCHONDRIAL AND NUCLEAR DNA ANALYSIS

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Molecular genetic analysis of 28 marsh frogs *Pelophylax ridibundus* complex from 12 localities of the Crimean peninsula was conducted by two molecular markers, namely, *COI*, the gene of the first subunit of cytochrome oxidase mtDNA, and *SAL-1*, the first intron of the serum albumin gene nDNA. It has been found that the mtDNA type specific for the “eastern” form (the Anatolian *P. cf. bedriagae*) prevails for the Crimean marsh frogs, while the mtDNA type of the “western” form (the Central-European *P. ridibundus*) has been observed in a single case in the Chernaya River basin (the southwestern Crimea). At the same time, our nuclear marker analysis has revealed the presence of the “western” form alleles within the studied area which occurs there more rarely than the “eastern” one (in a 2:5 frequency ratio). The alleles of the “western” form were identified in contrasting landscape zones and various altitude-climatic belts of the peninsula which might give evidence of the occurrence of multidirectional colonization waves of *P. ridibundus* in the Pleistocene-Holocene epoch.

**Key words:** *Pelophylax ridibundus*, *Pelophylax cf. bedriagae*, cytochrome oxidase, serum albumin, Chernaya River basin, Crimea.

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