

Genetic homogeneity in a Pontocaspian crested newt species (*Triturus karelinii*) suggests recent isolation of its three allopatric range sections

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Supplementary material

Table S1. Sampling details for mitochondrial DNA for *Triturus karelinii*.

Table S2. Sampling details nuclear DNA and output Bayesian clustering programs for *Triturus karelinii*.

Table S3. Pairwise Fst distance matrix for the eight *Triturus karelinii* populations for which nuclear DNA was sequenced. Population numbers correspond to fig. 1 and table 1.

Table S4. Gene diversity and number of alleles per marker and population for *Triturus karelinii*.

Data Availability Statement

Sequence data and files associated with analyses are available from the Dryad Digital Repository: XXX.