

The distributions of the six species constituting the smooth newt species complex (*Lissotriton vulgaris sensu lato* and *L. montandoni*) – an addition to the New Atlas of Amphibians and Reptiles of Europe

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Figure S1. Results of the ADMIXTURE analysis for the smooth newt species complex (*Lissotriton vulgaris sensu lato* and *L. montandoni*), plotted on a map. This analysis is based on 1138 SNPs genotyped for 379 individuals from 329 populations. Pie slices reflect ancestry (the fraction of alleles derived from each parental species at the level of the population) and are colour coded. Raw scores are in Supplementary Table S1. The boxed area in panel A is shown in more detail in panel B. Grey shading in panel B reflects elevation.

