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Isolation and Gene Flow in a Speciation Continuum in Newts

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Highlights:

- Nine evolutionary lineages robustly delimited using sequences of 74 nuclear loci.
- Multiple phylogenetic approaches partially resolve the *Lissotriton* species tree.
- ABC modeling reveals gene flow of various extent among central European taxa.
- Four southern smooth newt lineages are distinct species.
- Strong statistical support for widespread mtDNA introgression among lineages.

Abstract

Because reproductive isolation often evolves gradually, differentiating lineages may retain the potential for genetic exchange for prolonged periods, providing an opportunity to quantify

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