

The range expansion of the pine processionary moth in a changing world: How a well understood multifactorial process could tangle up

Mathieu Laparie, Christelle Robinet, Jérôme Rousselet, Alain Roques

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Introduction: While distributional changes associated with climate change have been widely acknowledged in ectotherms in the light of their ecophysiological limits, causal relationships with range shifts have still rarely been demonstrated. In the pine processionary moth, a winter-developing insect, combined experimental and modelling evidence have elucidated the mechanisms in which average winter warming has lowered barriers to expansion. Nonetheless, warming may also alter life cycles, or vulnerable instars may be subjected to new stressors associated with climate fluctuations. Comprehensive responses to such varied conditions are not fully untangled yet, and the spread may be more complex than initially thought due to mixed effects of future climate change, its spatial dependency, and population differentiations.

Methods: This presentation will be based upon published work on this study system, spanning a number of techniques ranging from thermal biology under controlled conditions and semi-natural experiments to field surveys and *in silico* approaches.

Results/Conclusion: Tolerance to acute cold is considered high enough to withstand usual minimal temperatures near front edges, but larval susceptibility to cumulative chill injury and thermal induction of foraging were evidenced. Both processes impair colony survival of this gregarious species under natural conditions. Spread simulations from climate records and feeding thresholds showed that warmer winters have opened previously unsuitable areas that match those actually neo-colonized. However, thermal constraints depend on genotypes and life stages, which have yet to be considered. Demographic outcomes of climate variations are thus contingent on geography and phenology, the latter being known to vary among populations.

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Authors

Mathieu Laparie

INRA

Christelle Robinet

INRA

Jérôme Rousselet

INRA

Alain Roques

INRA

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