

How is dispersal integrated in life histories: A quantitative analysis using butterflies

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As dispersal plays a key role in gene flow among populations, its evolutionary dynamics under environmental changes is particularly important. The inter-dependency of dispersal with other life history traits may constraindispersal evolution, and lead to the indirect selection of other traits as a by-product of this inter-dependency.

Identifying the dispersal's relationships to other life-history traits will help to better understand the evolutionary dynamics of dispersal, and the consequences for species persistence and ecosystem functioning under global changes. Dispersal may be linked to other life-history traits as their respective evolutionary dynamics may be be beinter-dependent, or, because they are mechanistically related to each other.

We identify traits that are predicted to co-vary with dispersal, and investigated the correlations that may constrain dispersal using published information on butterflies. Our quantitative analysis revealed that:

- 1/ Dispersal directly correlated with demographic traits, mostly fecundity, whereas phylogenetic relationships among species had a negligible influence on this pattern,
- 2/ gene flow and individual movements are correlated with ecological specialisation and body size, respectively and
 - 3/ routine movements only affected short-distance dispersal.

Together, these results provide important insights into evolutionary dynamics under global environmental changes, and are directly applicable to biodiversity conservation.