

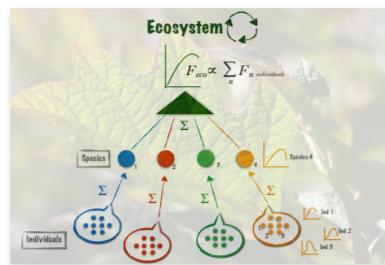
# SCALING UP INDIVIDUAL METABOLISM TO ECOSYSTEM FLUXES

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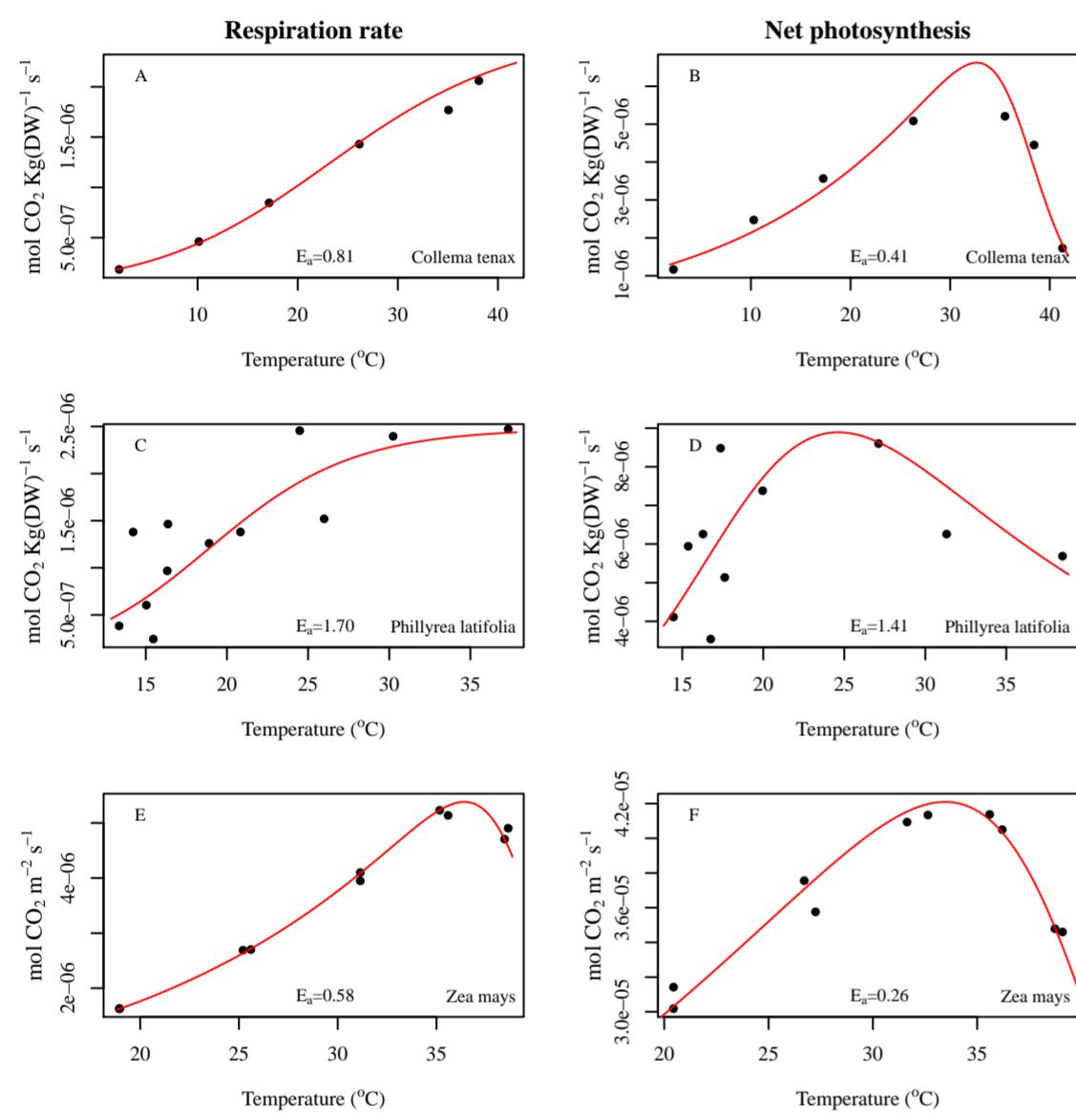
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Do species-level thermal performance curves (TPCs) scale up to ecosystem fluxes?

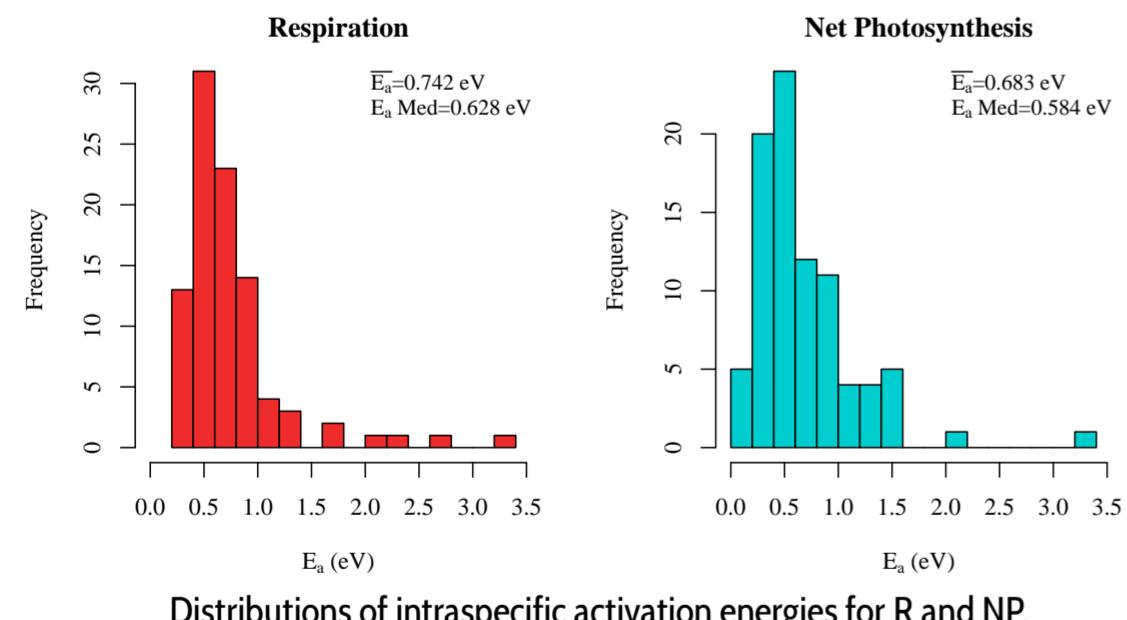


Previous studies assume that the temperature dependence of ecosystem function is a simple scaling up of all the component species' thermal responses. In this case, predicting the effects of climatic warming or cooling on ecosystem function would be a relatively straightforward task.

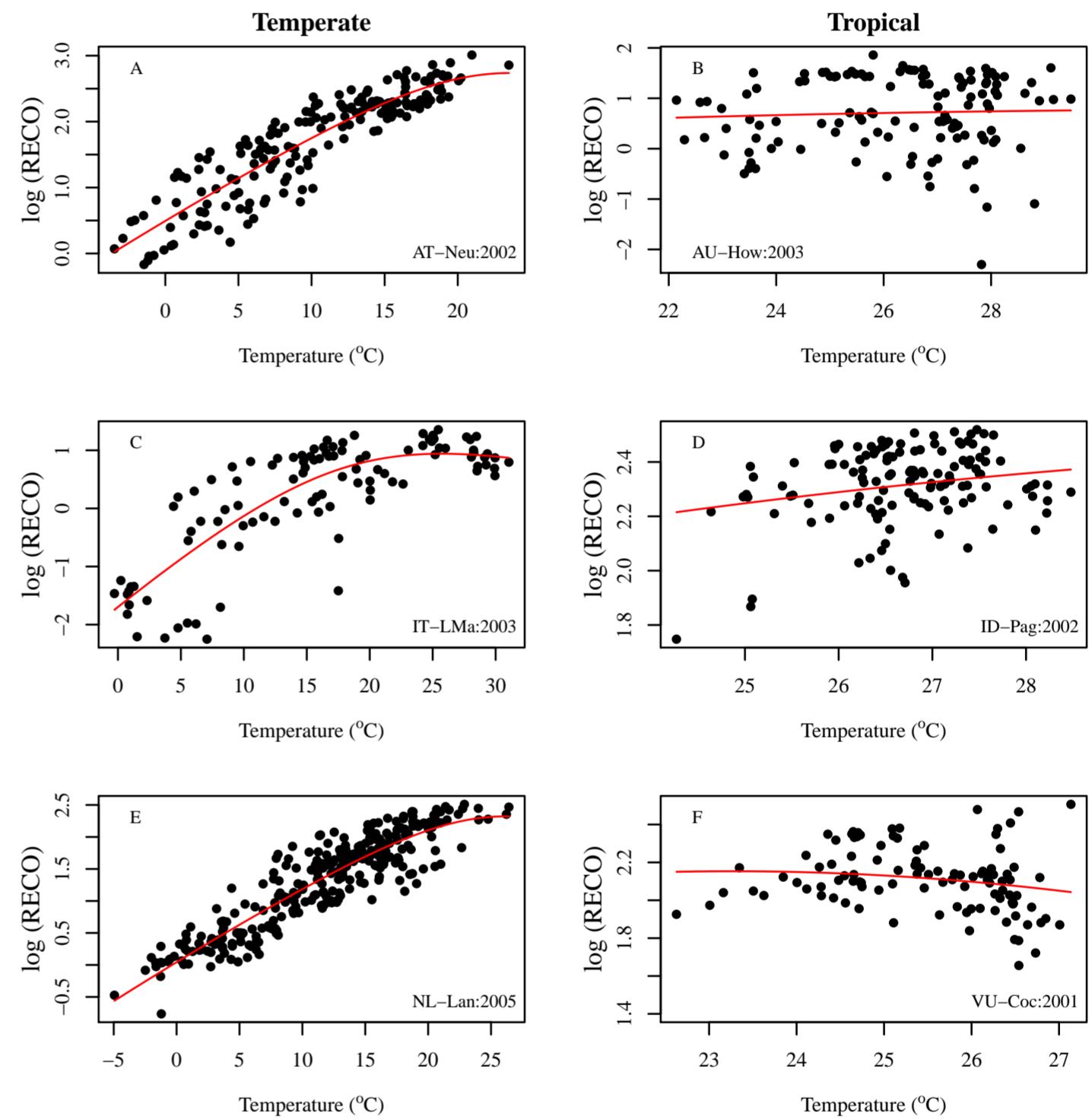
## Parametrization



Representative set of individual-level plot pairs for R (left) and NP (right). Each pair of R-NP corresponds to the same experiment and species. These species are associated to temperate regions.



## Validation



Representative set of ecosystem flux responses to temperature for different sites from tropical and temperate regions. Y axis show respiration in log scale.

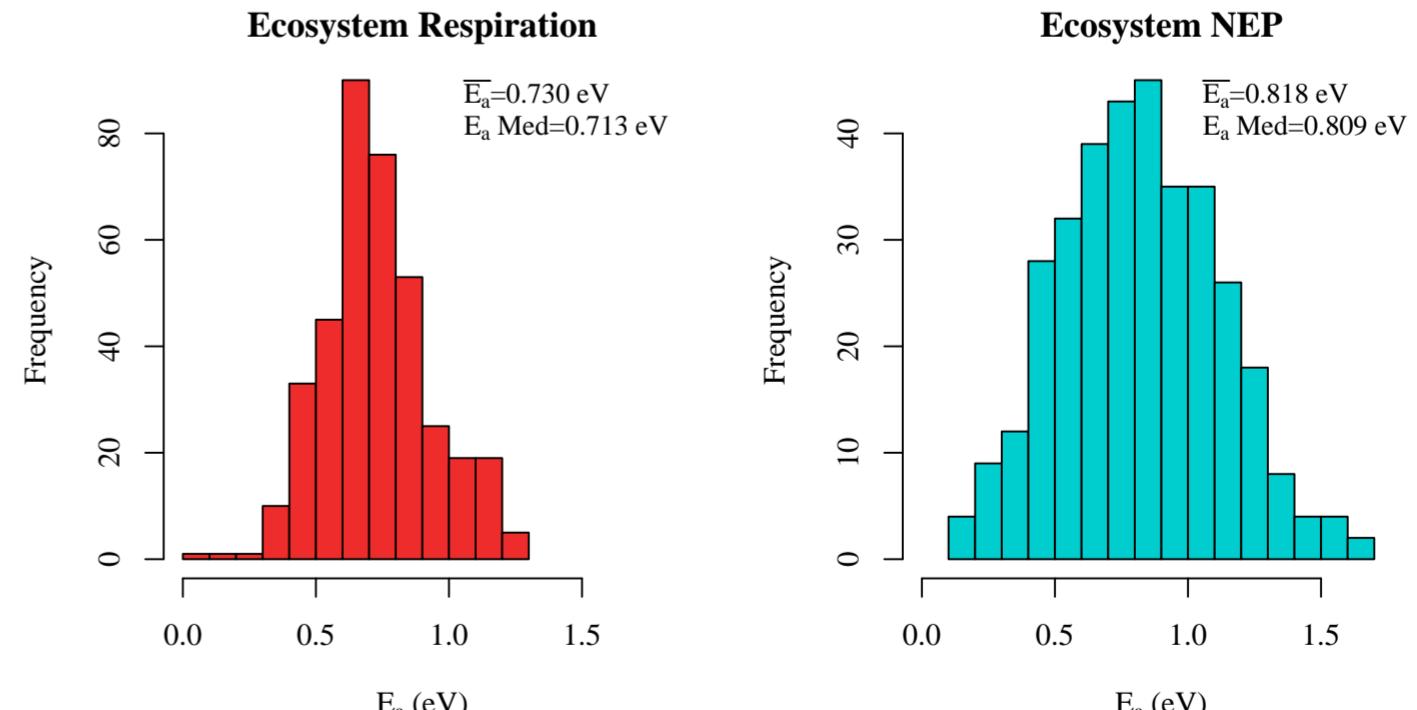


Figure 1: Distributions of activation energies for Reco and NEP at ecosystem level.

## Summary

- At the intra-specific level  $E_a$  and  $T_{max}$  for R are usually higher than for NP

## Ongoing

- At the intra-specific level  $E_a$  and  $T_{max}$  for R are usually higher than for NP