# Characters changing speed: the Covariomorph model and its impact on phylogenetic trees

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Evolution Meetings - June 23, 2025







# Phylogenetic inference using morphology



- Morphological datasets are analyzed using the Mk model [Lewis, 2001].
- Extensions like ACRV (e.g.,  $+\Gamma$ ) allow for among-character rate variation.
- But they assume constant rates for characters across all lineages.
- ▶ Do all characters evolve at the same speed in all lineages?

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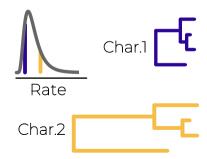


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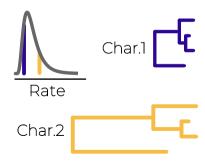


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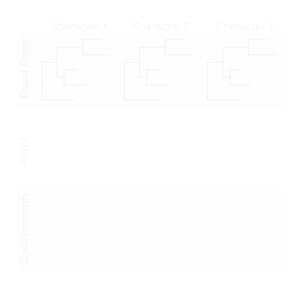


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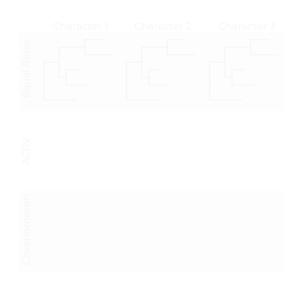






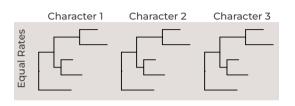
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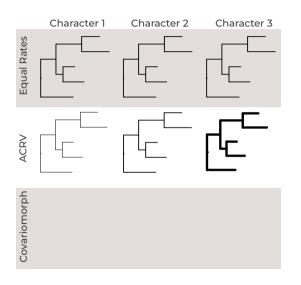


ACR\

Covariomorph

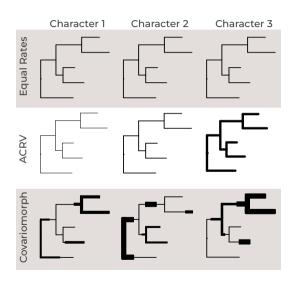
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- Generalizes the covarion idea for morphology.
- Characters can switch between rate regimes (e.g., slow, medium, and fast).
- Unifies character-level and lineage-level rate variation in one framework.



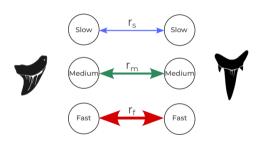




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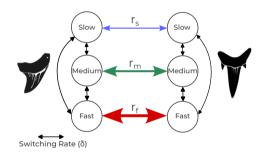
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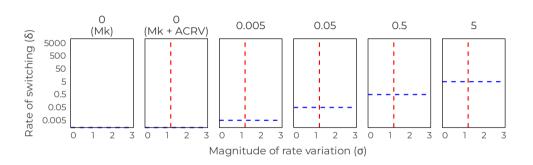
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### Simulation test



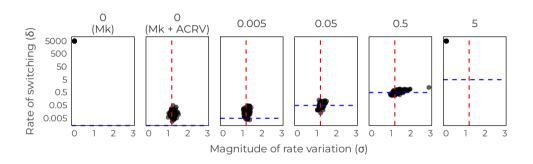


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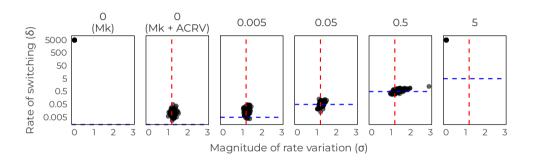




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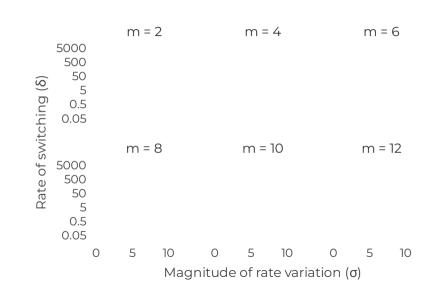




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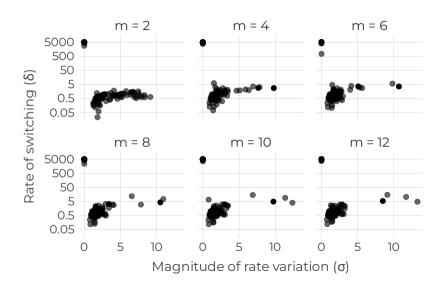
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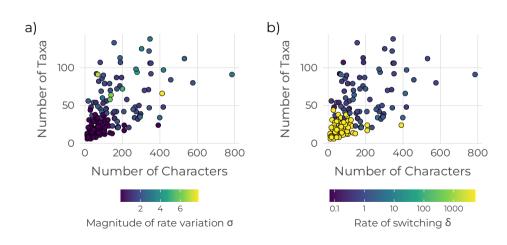




# Association between dataset size and heterotachy



#### Larger datasets show more rate switching.



### Focal data set of Rays



- ► Rays: 52 taxa, 124 characters [Marramà et al., 2023]
- Additional rate categories  $m = \{14, 16, 18\}$
- Explored posterior distributions of  $\delta$ ,  $\sigma$ , and the tree length and topology.
- Also, model selection.

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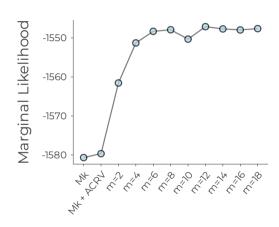


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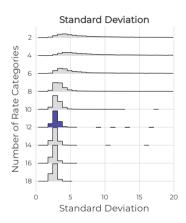


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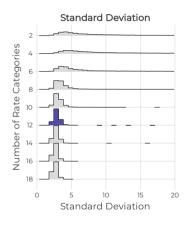
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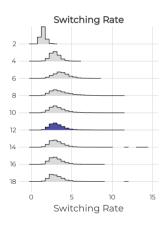




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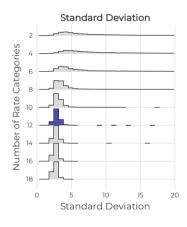


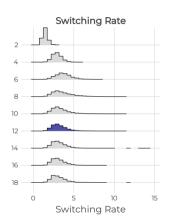


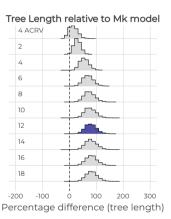


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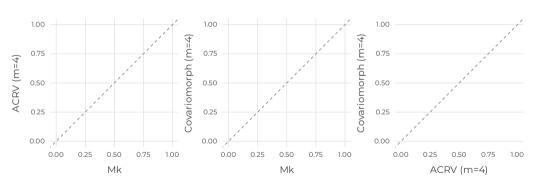




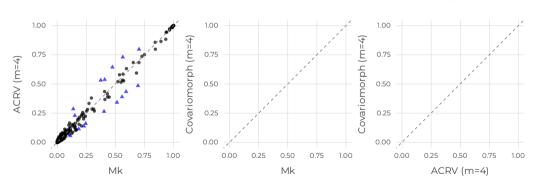




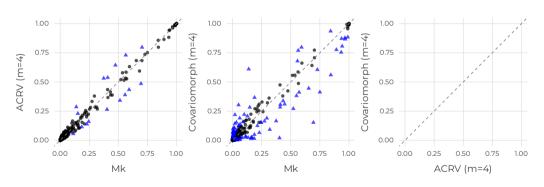




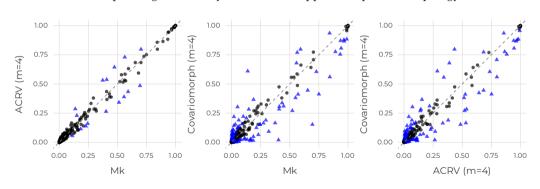












### **Take-home Messages**



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- Implications for downstream studies like divergence time estimation, and diversification studies.

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### Thank you!





Preprint coming soon!

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