Preprint template

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Abstract: This template is used by the Poisot lab at Université de Montréal to write manuscripts using github. It uses github actions as a way to generate a website that can be annotated using hypothes.is, a PDF document for copy-editing and submission to journals, and a PDF document for submission to preprint servers. At every push on the master branch, the whole series of documents will be updated automatically.

Keywords: pandoc pandoc-crossref github actions

1 Outline

Introduction: - Predictive ecology and predicting interactions - Data is scarce - Idea you can train a predictive model on simulated data and apply it to real data and it makes good predictions - It might seem wild that this can work, but here we show

Methods

- concept figure with (niche model, mpn) instead of (hypoth 1, 2)
- Empirical data from Streams in NZ
- Julia 1.6, EN, Flux, etc.

For each empirical network, we predict the interactions by training a neural network — structure of neural net todo. For each of these neural nets, we do not train them on any real data, but instead only on simulated networks from each of the candidate models with S_i species and betabinom connectance.

Summary stats to describe network properties are inevitably necessary.

- abc discussion and history
- generative learning discussion and history

explain why you can't predict interaction between two species based on anything but summary stats.

We generate networks from various candidate models: cascade, niche, mpn, etc.

Results

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Discussion