



Workflow for the calculation of phylogenetic indicators

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What is Phylogenetic Diversity?

Phylogenetic diversity is a measure of biodiversity which takes evolution into account. It is calculated as the sum of the lengths of the phylogenetic tree branches among a group of species.

Why Phylogenetic Diversity?

Phylogenetic diversity can be used in conservation planning to maximise a variety of features, which could be particularly useful in light of the changing environmental conditions.

Objectives

Objective 1:

Design an indicator

A metric that gives information about how well phylogenetic diversity is safeguarded by current protected areas for a certain higher taxonomic group

Objective 2:

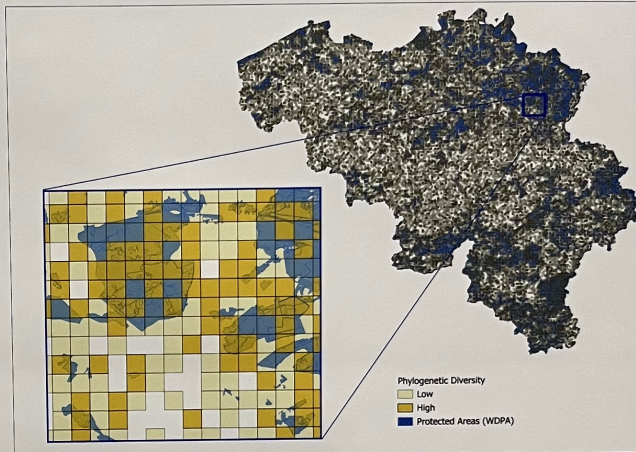
Create a map

A map that shows an overlay of high phylogenetically diverse areas with protected areas

Objective 3:

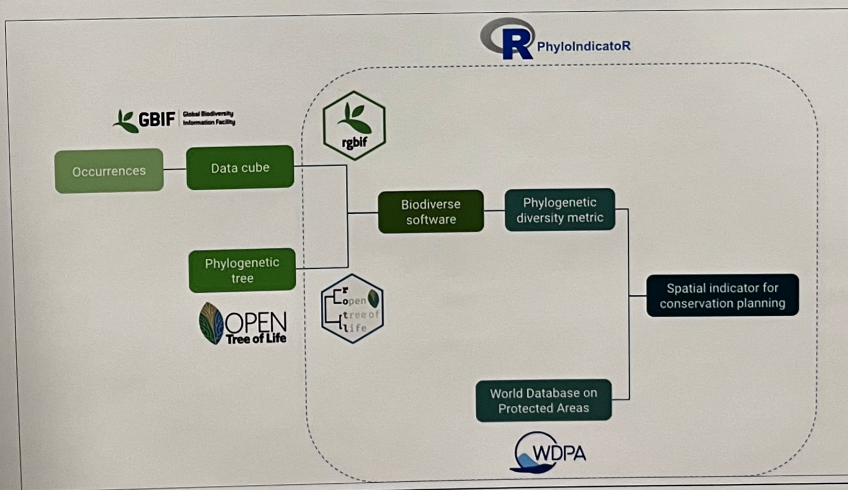
Design a workflow

A workflow to rapidly calculate this overlap to identify potential directions for future expansion of protected areas



Simulation of overlap between protected areas and areas of low/high phylogenetic diversity in Belgium with a randomly generated dataset

The components of PhyloIndicatoR, an R notebook to produce a phylogenetic indicator for conservation planning using species occurrence data cubes and a published phylogenetic tree as input.



Questions - Give your opinion by putting colored sticky dots in the boxes

- Have you ever considered using phylogenetic diversity?

- Where do you think phylogenetic diversity indicators are most useful?
 - CONSERVATION POLICY

- RESEARCH

- For which of the following higher taxa would you want to know the PD score?

- MAMMALS

- BIRDS

- AMPHIBIANS

- REPTILES

- FISH

Phylogenetic diversity as an indicator within the B-Cubed project



- Standardising access to biodiversity data, empowering policymakers to address the impacts of biodiversity change.
- Improve the existing policy evidence base and contribute to better alert systems by providing fast access to pre-aggregated and modelled biodiversity data and standardised biodiversity indicators responsive to the addition of new data.

