Evolutionary approach for mapping biodiversity priorities in Brazil

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Abstract

We have built a model to map biodiversity priorities across Brazil that integrates evolutionary approach with a set of biodiversity variables. The model uses a comprehensive dataset (circa of 500 thousand geo-records) on vertebrates, arthropods and angiosperms from which we derived indices of beta-diversity, species richness, area of endemism, endemicity, phylogenetic endemism and phylo-beta-diversity. Results indicate that 10% of the Brazil has high biological relevance for conservation with relatively good sampling effort. These areas encompass 87% of the species and 92% of the evolutionary lineages contained in our dataset. Sensitivity analyses show that the use of few biodiversity variables or only one group of organisms (vertebrates for example) is not sufficient for identifying biodiversity conservation priorities. Yet, vast areas of Brazil, especially in the Amazon, still lack biological inventories, making it impossible to compare their relevance in relation to other areas where biological knowledge is high. Some regions with high biological relevance are located in areas of deforestation pressure or of highly fragmented forest remnants, making them either a conservation or a restoration priority.

Keywords: Biodiversity conservation priorities, evolutionary conservation

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