

Evolution of Amazonian biodiversity: A review

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ABSTRACT

Amazonia (defined herein as the Amazon basin) is home to the greatest concentration of biodiversity on Earth, providing unique genetic resources and ecological functions that contribute to ecosystem services globally. The lengthy and complex evolutionary history of this region has produced heterogeneous landscapes and riverscapes at multiple scales, altered the geographic and genetic connections among populations, and impacted rates of adaptation, speciation, and extinction. In turn, ecologically diverse Amazonian biotas promoted further diversification, species coexistence, and coevolution, with biodiversity accumulating over tens of millions of years. Important events in Amazonian history included: (i) late Cretaceous and early Paleogene origin of major rainforest plant and animal groups; (ii) Eocene-Oligocene global cooling with rainforests contracting to tropical latitudes separating Atlantic coastal and Amazonian rainforests; (iii) Miocene uplift of central and northern Andes that separated Pacific coastal and Amazonian rainforests, spurred formation of mega-wetlands in the western Amazon, and contributed to the origin of the modern transcontinental Amazon River; (iv) late Neogene formation of the Panamanian Isthmus that facilitated the Great American Biotic Interchange; (v) Pleistocene climate oscillations followed by late Pleistocene-Holocene human colonization and megafaunal extinctions; and (vi) modern era of widespread anthropogenic deforestation, defaunation, and ecological transformations of regional landscapes and global climates. Amazonian conservation requires decade-scale investments into biodiversity documentation and monitoring to leverage existing scientific capacity, and strategic habitat planning to allow continuity of evolutionary and ecological processes now and into the future.

KEYWORDS: biogeography, conservation, extinction, Neotropics, speciation

Evolución de la biodiversidad amazónica: una revisión

RESUMEN

La Amazonía (definida como la cuenca amazónica) concentra la mayor biodiversidad de la Tierra, proporcionando recursos genéticos y funciones ecológicas únicas que contribuyen a los servicios ecosistémicos a nivel mundial. La compleja historia evolutiva de esta región produjo paisajes heterogéneos a múltiples escalas geográficas, alteró las conexiones geográficas y genéticas entre las poblaciones e influyó en las tasas de adaptación, especiación y extinción. Las biotas amazónicas, ecológicamente diversas, promovieron una mayor diversificación, coexistencia de especies y coevolución, acumulando biodiversidad a lo largo de decenas de millones de años. Acontecimientos importantes en la historia de la Amazonía incluyeron: (i) orígenes durante el Cretácico tardío y el Paleógeno temprano de los principales grupos de plantas y animales; (ii) enfriamiento global del Eoceno-Oligoceno, contrayendo los bosques a latitudes tropicales y separando los de la costa Atlántica de los amazónicos; (iii) levantamiento de los Andes centrales y del norte en el Mioceno, separando las selvas tropicales de la costa del Pacífico

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y de la Amazonía, estimulando la formación de megahumedales en la Amazonía occidental y contribuyendo al origen del moderno Río Amazonas transcontinental; (iv) formación del istmo de Panamá durante el Neógeno tardío, facilitando el Gran Intercambio Biótico Americano; (v) oscilaciones climáticas del Pleistoceno seguidas por la colonización humana y las extinciones de megafauna; (vi) era moderna de deforestación antropogénica generalizada, defaunación y transformaciones ecológicas de paisajes regionales y climas globales. La conservación de la Amazonía requiere inversiones por décadas en la documentación y el seguimiento de la biodiversidad para impulsar la capacidad científica existente, así como la planificación estratégica del hábitat para permitir la continuidad de los actuales y futuros procesos evolutivos y ecológicos.

PALABRAS CLAVE: biogeografía, conservación, extinción, Neotrópico, especiación

INTRODUCTION

This review was originally developed as Chapter 2 of the Science Panel for the Amazon Assessment Report (<https://www.theamazonwewant.org/>) (Guayasamin *et al.* 2021). The report aimed to scientifically assess the current state of the Amazon, and to identify and explore opportunities for relevant policy actions. We focused our review on the following main themes: (i) Amazonian biodiversity, (ii) evolution of Amazonian forests, (iii) assembling of Amazonian biota, (iv) species loss and turnover, and (v) conservation of ecological and evolutionary processes. Broad accessibility to this information is at the core of disseminating the knowledge

and understanding the complexity of the Amazon basin and the urgency for conservation actions.

AMAZONIAN BIODIVERSITY IS IMMENSE AND VASTLY UNDERESTIMATED

The Amazon basin (Figure 1) houses one of the highest organismal and ecosystemic diversity on Earth (Bass *et al.* 2010; Levêque *et al.* 2007). Approximately 10% of the world's vertebrate and plant species are contained in an area that corresponds to approximately 0.5% of the Earth's total surface (Jetz *et al.* 2012; Tedesco *et al.* 2017; ter Steege *et al.* 2020; Figure 2). Amazonian diversity also represents a bewildering

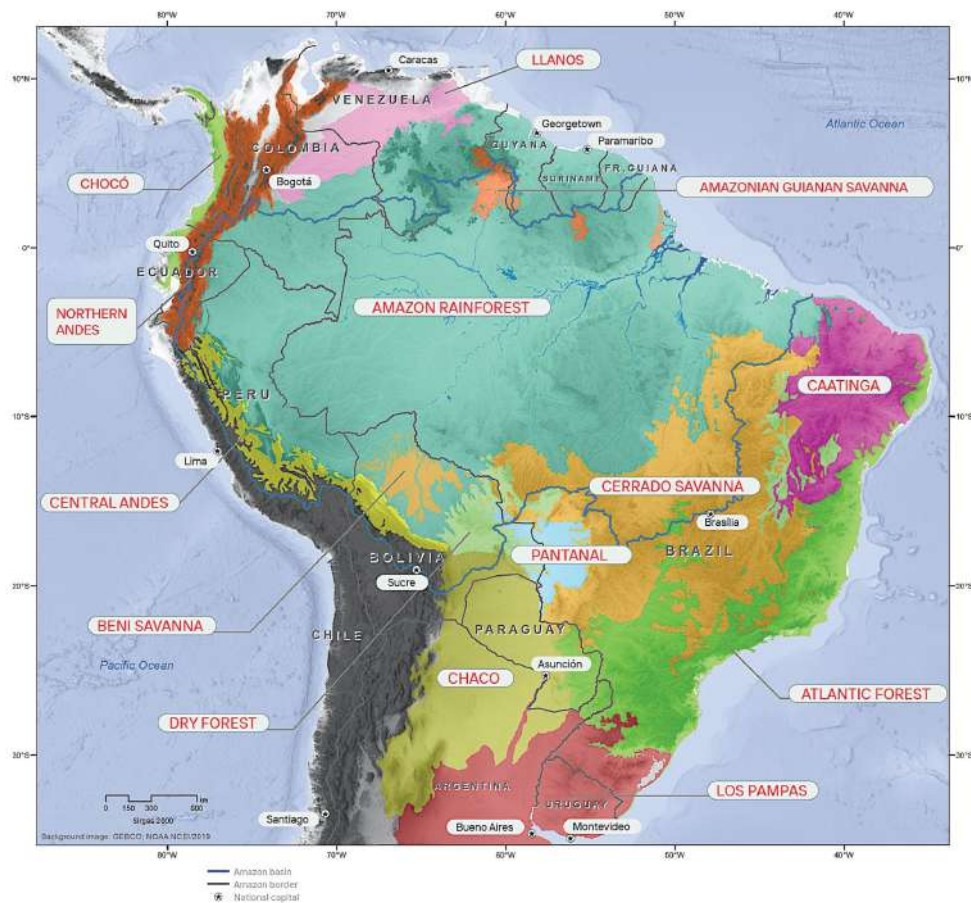


Figure 1. Main Neotropical biogeographical regions and the Amazon River drainage basin (blue polygon). Note that the Seasonally Dry Diagonal region (composed of the Caatinga, Cerrado, and Gran Chaco biomes) separates the Amazon and the Atlantic rainforests, while the northern Andes separate the Amazon and the Chocó rainforests. Reproduced with permission from Guayasamin *et al.* (2021).

range of life forms, ecological functions, chemical compounds, and genetic resources (Darst *et al.* 2006; Asner *et al.* 2014; Albert *et al.* 2020a; Figure 3). The highly diverse Amazonian ecosystems constitute the core of the Neotropical realm, which harbors about 30% of all species of vascular plants (Raven *et al.* 2020), vertebrates (Jenkins *et al.* 2013; Reis *et al.* 2016), and arthropods (Stork 2018) on Earth.

Despite decades of intensive study, the full dimensions of Amazonian diversity still remain vastly underestimated (da Silva *et al.* 2005; Barrowclough *et al.* 2016; García-Robledo *et al.* 2020). Critical information is lacking in all main biodiversity levels (see Hortal *et al.* 2015): taxonomic diversity (Linnaean shortfall), biogeographic distributions (Wallacean shortfall), species abundances (Prestonian shortfall), phylogenetic diversity (Darwinian shortfall), species traits (Raunkjæran shortfall), and species interactions (Eltonian shortfall). This knowledge shortfall arises from the extremely high number of species found in the region (Magurran and McGill 2011; Raven *et al.* 2020), the numerous species yet unrecognized due to the so called cryptic biodiversity (i.e., several species undescribed because they share a similar morphology) (Angulo and Icochea 2010; Benzaquem *et al.* 2015; Draper *et al.* 2020; Jaramillo *et al.* 2020), the logistical difficulties with sampling in remote and inaccessible regions (Cardoso *et al.* 2017; ter Steege *et al.* 2020), collection efforts that are biased towards accessible localities (Nelson *et al.* 1990;

Hopkins 2007; Loiselle *et al.* 2008), and a disproportionate number of studies applied to conspicuous organisms (Ritter *et al.* 2020) and broadly distributed species (Ruokolainen *et al.* 2002). As a result, many Amazonian species have never been collected, named, or studied; and often an entire group of closely related species (i.e., clade) is mistakenly treated as a single species (Albert *et al.* 2020b).

To fill this gap, integrated studies of Amazonian taxa conducted over the past two decades have employed a combination of molecular and morphological tools that allow scientists to recognize cryptic species of plants (Damasco *et al.* 2019; Francisco and Lohmann 2020), birds (Ribas *et al.* 2012; Whitney and Cohn-Haft 2013; Thom and Aleixo 2015; Schultz *et al.* 2017, 2019), amphibians (Gehara *et al.* 2014; Jaramillo *et al.* 2020; Vacher *et al.* 2020), fishes (Melo *et al.* 2016; Craig *et al.* 2017; García-Melo *et al.* 2019), and primates (Lynch Alfaro *et al.* 2015). Between 1999 and 2015, many new species of plants (1,155 species), fishes (468), amphibians (321), reptiles (112), birds (79), and mammals (65) were described from the Amazon Basin (WWF 2016).

Spectacular Amazonian species continue to be described. They include, for instance, a new critically endangered titi monkey (*Plecturocebus grovesi* Boubli *et al.* 2019; Byrne *et al.* 2016), 15 new species of Amazonian birds described in a single publication (Whitney and Cohn-Haft 2013), 44 new species of lungless *Bolitoglossa* salamanders that await

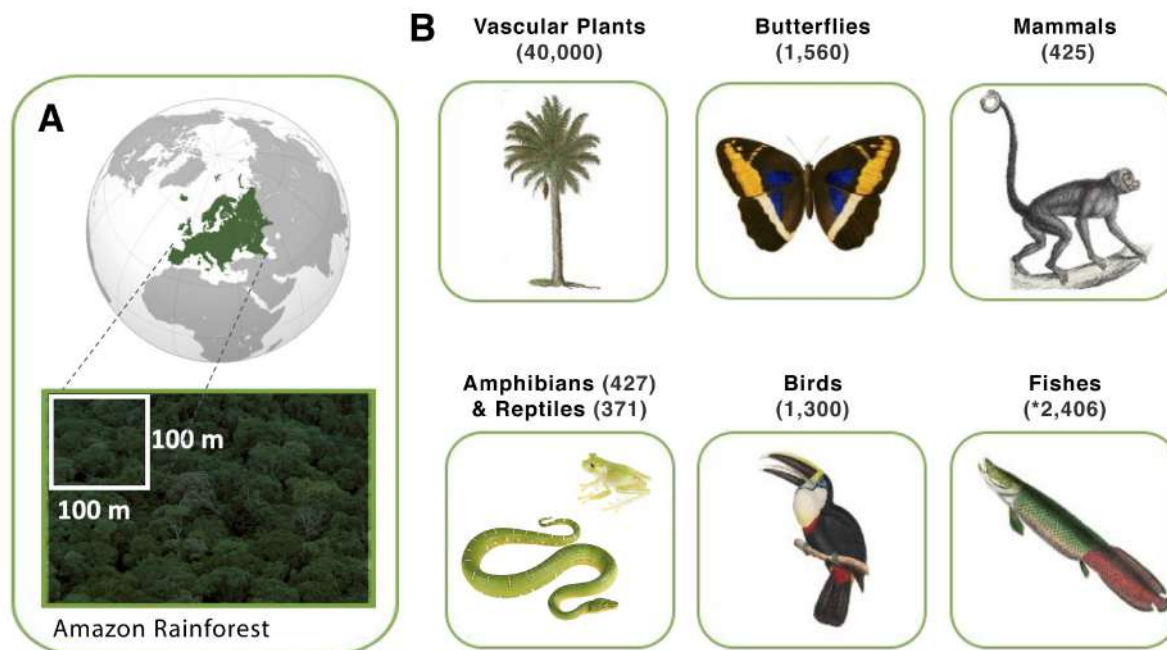


Figure 2. The Amazonian lowlands in numbers (minimum estimates based on current knowledge). **A** – More tree species are found in a 10,000 m² area of Amazon rainforest than in the whole of Europe (ter Steege *et al.* 2006); **B** – Estimated numbers of species of selected Amazonian lineages, including vascular plants (Hubbell *et al.* 2008; Mittermeier *et al.* 2003; image by Roberts 1839), butterflies (Vieira and Höfer 2021; image by Hewitson 1856), mammals (Mittermeier *et al.* 2003; image by Jardine and Courier 1840), amphibians and reptiles (Mittermeier *et al.* 2003; image by Jose Vieira), birds (Mittermeier *et al.* 2003; image by Gould *et al.* 1852), and fishes (Oberdorff *et al.* 2019, Jézéquel *et al.* 2020; image by Castelnau 1855). Note that the number of fish species corresponds to the whole basin, but most of them (> 95%) are from elevations below 250 m (Albert *et al.* 2011, 2020; Dagosta and de Pinna 2020). Reproduced with permission from Guayasamin *et al.* (2021).

formal description (Jaramillo *et al.* 2020), a distinctive new and critically endangered vanilla orchid (*Vanilla denshikoira* Flanagan and Ospina-Calderón, 2018), and a new worm-like fish species (*Tarumania walkerae* de Pinna *et al.*, 2017) that inhabits moist leaf litter deep within the rainforest, and which represents an entirely new family, the Tarumaniidae (de Pinna *et al.* 2018).

Comprehensive knowledge of the species that inhabit hyperdiverse Amazonian ecosystems is central to better understanding their ecosystem functions (Malhi *et al.* 2008) and the emergent properties that arise from interactions among Amazonian species and their abiotic environments. For example, while it is clear that the Amazonian hydrological cycles depend on forest transpiration, and that they impact climate at a continental scale (Costa *et al.* 2021), the influence of local species and their traits on precipitation patterns and climate remains to be understood (Chambers *et al.* 2007). Large-scale approaches aiming at quantifying unknown biodiversity, such as metagenomics, are also contributing to a deeper understanding of poorly studied life forms (e.g., bacteria, fungi, microorganisms) and ecosystem-level biochemical processes in Amazonian soils (Ritter *et al.* 2020) and rivers (Ghai *et al.* 2011; Santos *et al.* 2019). While still under-utilized, these approaches are revolutionizing our understanding of Amazonian biodiversity patterns and

their inherent processes, guiding conservation prioritization approaches and management plans for the basin.

Understanding the evolutionary history of Amazonian biodiversity is crucial to managing its exceptional biodiversity and ecosystem functions (Rull 2011; Figure 3). This knowledge, in turn, holds key information for guiding conservation of endemic species and ecosystem services in times of climatic change. Until recently, fragmentary comprehension of Amazonian biodiversity at finer taxonomic levels led scientists to use more inclusive taxonomic categories (e.g., genera, families) in studying diversification patterns in this region (Antonelli *et al.* 2009). While these higher taxonomic categories have provided important insights into biodiversity patterns (Terborgh and Andresen 1998), they cannot be objectively defined nor compared across taxa, rendering generalizations difficult (Cracraft *et al.* 2020). Integrative approaches that combine standardized field sampling, DNA barcoding (García-Melo *et al.* 2019; Vacher *et al.* 2020), comparative phylogenomics (Alda *et al.* 2019; Santos *et al.* 2019), and artificial intelligence (Draper *et al.* 2020) have accelerated the fine-scale documentation of Amazonian biodiversity (Ritter *et al.* 2020; Vacher *et al.* 2020). These approaches involve new sampling efforts while also relying on museum specimens, which significantly leverage taxonomic work (e.g., Thom *et al.* 2020; Vacher *et*



Figure 3. Examples of the Amazonian biota. First column: wire-tailed manakin (*Pipra filicauda* Spix, 1825), Zimmer's woodcreeper (*Dendroplex kienerii* [DesMurs, 1856]), Amazonian royal flycatcher (*Onychorhynchus coronatus* [Müller, 1776]). Second column: Amazon flying fish (*Thoracocharax stellatus* [Kner, 1858]), red bellied piranha (*Pygocentrus cariba* [Humboldt, 1821]), redspot killifish (*Trigonectes rubromarginatus* Costa, 1990). Third column: Andean glassfrog (*Hyalinobatrachium pellucidum* [Lynch and Duellman, 1973]), red howler monkey (*Alouatta seniculus* [Linnaeus, 1766]), La Salle's shadow-snake (*Synopsis lasallei* Nicéforo-María, 1950). Fourth column: columelia (*Columellia oblonga* Ruiz y Pavón, 1798), quinine (*Cinchona officinalis* Linnaeus, 1753), red passion flower (*Passiflora manicata* [Juss] Persoon, 1806). Photos by Camila Ribas and Tomaz Melo (first column), James Albert (second column), Tropical Herping (third column), and Carmen Ulloa Ulloa (fourth column). Reproduced with permission from Guayasamin *et al.* (2021).

al. 2020). Unfortunately, Amazonian museum collections are still undervalued despite offering a rich source of information (Escobar 2018). Local institutions need support to hire experts in the field, and to maintain and expand their biological collections (Fontaine *et al.* 2012; Funk 2018). Human resources and infrastructure support are also crucial for the maintenance of the large databases of Amazonian species compiled to date. While important and useful, they should be constantly vetted and updated to address knowledge gaps and misidentifications.

EVOLUTION OF AMAZONIAN FORESTS

Flowering plants constitute the main physical structure of Amazonian rainforests. They exhibit a wide variety of growth forms, including woody trees, shrubs, and lianas, as well as epiphytes, herbaceous sedges, grasses, and colonial bamboos (Rowe and Speck 2005). DNA studies suggest that flowering plants first diversified in the Early Cretaceous or ca. 145–100 million years ago (Ma) (Magallón *et al.* 2015), and fossil data suggest that they did not dominate Neotropical ecosystems until the Late Cretaceous (ca. 100–66 Ma; Burnham & Johnson 2004; Dino *et al.* 1999; Mejia-Velasquez *et al.* 2012; Carvalho *et al.* 2021).

While some Amazonian lineages have ancient origins dating back to the Early Cenozoic or Cretaceous (Cracraft *et al.* 2020), most species that currently inhabit the Amazon originated within the past few million years (Da Silva *et al.* 2005; Rull 2008, 2011, 2020; Santos *et al.* 2019). The wide distribution of evolutionary ages of Amazonian species suggests that the formation of its modern-day biodiversity took place over an immense time span (Cracraft *et al.* 2020), being influenced by the many changes in the physical landscape during the Cenozoic (Antonelli *et al.* 2009; Hoorn *et al.* 2010), producing extinctions and turnovers of several lineages (Jaramillo *et al.* 2010a, 2010b).

The Amazon was substantially modified by a sudden mass extinction triggered by the Chicxulub asteroid impact about 66 million years ago at the Cretaceous–Paleogene [K–Pg] boundary (De La Parra *et al.* 2008; Carvalho *et al.* 2021; Jacobs and Currano 2021). Many groups of Neotropical birds (Claramunt and Cracraft 2015; Oliveros *et al.* 2019), butterflies (Espeland *et al.* 2015, 2018; Seraphim *et al.* 2018), and fishes (Friedman 2010; Hughes *et al.* 2018) diversified rapidly following this event. Plant communities similar to those seen in today's Neotropical rainforests, although with fewer species, evolved in the Paleocene (ca. 66–56 Ma) (Wing *et al.* 2009; Jaramillo *et al.* 2010a), with many plant lineages diversifying in the Eocene (ca. 56–34 Ma) (Lohmann *et al.* 2013). Indeed, Neotropical rainforest plants seem to have reached a pinnacle of diversity during the Eocene (ca. 56–34 Ma), when warm and moist climates still predominated (Jaramillo *et al.* 2006). Eocene forests are thought to have

been highly rich in species (Burnham and Graham 1999; Jaramillo *et al.* 2006, 2010a, 2010b). Conspicuous elements of Paleocene Neotropical rainforests include members of key plant groups such as palms and herbs (e.g., families Araceae and Zingiberaceae), shrubs (e.g., Malvaceae), lianas (e.g., Menispermaceae), and trees (e.g., Lauraceae) (Burnham and Johnson 2004; Wing *et al.* 2009; Carvalho *et al.* 2011).

The drier seasons and cooler climates of the Early Oligocene (ca. 30 Ma) contributed to extensive vegetational changes throughout South America. Namely, the once continuous and broadly distributed wet South American rainforests were divided in two, the Amazon and Atlantic rainforests, due to expansion of open subtropical woodland forests in central South America and the establishment of the Seasonally Dry Diagonal (Bigarella 1975; Costa 2003; Orme 2007; Fouquet *et al.* 2012; Sobral-Souza *et al.* 2015; Thode *et al.* 2019). These vegetational changes coincided with the beginning of the uplift of the Mantiqueira Mountains of eastern Brazil, and of the northern Andes in Colombia, causing substantial changes in South American air currents. Increasingly drier climates and the expansion of open savannah vegetation types were accompanied by substantial changes in species composition (e.g., more palms), the origin of C4 grasses (Vicentini *et al.* 2008; Urban *et al.* 2010; Bouchenak-Khelladi *et al.* 2014), and the expansion of grasslands and open woodlands at the expense of closed-canopy forested habitats (Edwards and Smith 2010; Edwards *et al.* 2010; Kirschner and Hoorn 2020).

In the Miocene, uplift of the northern and central Andes led to a profound reorganization of the river network of the whole of northern South America, including the formation of the Pebas mega-wetland (Hoorn *et al.* 1995, 2010, 2017; Albert *et al.* 2018), a vast region (ca. 1 million km²) of lacustrine and swampy environments located in the area of the modern western Amazon (Hoorn 1993; Wesselingh and Salo 2006; Bicudo *et al.* 2019). Progressive uplift of the northern Andes also affected the regional climate, leading to increased precipitation due to the orography (Poulsen *et al.* 2010). Vast areas of flooded forests were then established, composed of palms (i.e., *Grimsdalea*), ferns, and grasses (Poaceae), among others (Hoorn 1994; Jaramillo *et al.* 2017; Hoorn *et al.* 2017; Kirschner and Hoorn 2020; Hoorn *et al.* 2023). In addition, marine incursions into the western Amazon from the Caribbean Sea allowed estuarine taxa to colonize the northern Pebas shorelines (Hoorn 1993; Boonstra *et al.* 2015; Jaramillo *et al.* 2017; Hoorn *et al.* 2023).

In the Late Miocene and Pliocene, a major landscape reshaping took place, caused in part by overfilling of sedimentary basins in the western Amazon with Andean-derived sediments; this led to a renewed drainage reorganization and the onset of the modern transcontinental Amazon River (Hoorn *et al.* 2010, 2017, 2023). The former Pebas wetland

surfaces were colonized by many different lineages (Antonelli *et al.* 2009; Roncal *et al.* 2013), in a process of upland forest expansion that is suggested to have continued until the Late Pleistocene (Pupim *et al.* 2019). Landscape changes also led to increased diversification of numerous plant lineages, such as the flowering plant genera *Inga* (Legumes; Richardson *et al.* 2001) and *Guatteria* (Annonaceae; Erkens *et al.* 2007). At around the same time, the Andean slopes were colonized by many plant lineages, including species of the Malvaceae (Hoorn *et al.* 2019), Arecaceae (i.e., palms; Bacon *et al.* 2018), and Chloranthaceae families (i.e., *Hedyosmum*; Martínez *et al.* 2013). From the Late Miocene to the Pliocene (ca. 11–4 Ma), the rise of the Eastern Cordillera of the Colombian Andes completed the isolation of the *cis*-Andean (Orinoco-Amazon) from the *trans*-Andean (Pacific slope, Magdalena, and Maracaibo) basins, resulting in the isolation of their resident aquatic biotas (Albert *et al.* 2006). Evidence suggests that high levels of plant species diversity existed in Amazonia during the Miocene thanks to a combination of low seasonality, high precipitation, and edaphic heterogeneous substrate (Jaramillo *et al.* 2010a).

The Neogene uplift of the Northern Andes (ca. 23–2.6 Ma) had profound effects on Amazonian landscapes, impacting the diversification of both lowland and highland lineages (Hoorn *et al.* 2010; Albert *et al.* 2011b; Givnish *et al.* 2016; Rahbek *et al.* 2019; Montes *et al.* 2021). Yet, despite its importance for biogeography, the specific role of mountain ranges as a dispersal barrier between South and Central American lowland plant lineages is still poorly understood (Pérez-Escobar *et al.* 2017). Different diversification patterns have been detected within and between upland and lowland groups, with higher species richness in lowlands and higher species endemism in uplands. The uplift of the northern Andes and its associated dynamic climate history were key drivers of the rapid radiation of Andean-centered plants (Gentry 1982; Jost 2004; Madriñán *et al.* 2013; Luebert and Weigend 2014; Lagomarsino *et al.* 2016; Vargas *et al.* 2017) and animals (Albert *et al.* 2018; Rahbek *et al.* 2019; Perrigo *et al.* 2020). Near mountain tops, plants of the páramo ecosystem underwent one of the highest speciation rates ever recorded (Madriñán *et al.* 2013; Padilla-González *et al.* 2017; Pouchon *et al.* 2018).

During the Quaternary (last ca. 2.6 Ma), global climate cooling in combination with geomorphological processes strongly altered the western Amazonian landscape. Alluvial megafans (large sediment aprons >10,000 km²) extended from the Andes into the Amazon (e.g., Räsänen *et al.* 1990, 1992; Wilkinson *et al.* 2010), and floodplains varied in size according to changes in precipitation patterns (Pupim *et al.* 2019). The effect of these cyclic climatic changes on landscape and vegetation composition is yet to be fully understood. Direct studies of the sedimentary and fossil records (Jaramillo *et al.* 2017; Hoorn *et al.* 2017; Mason *et al.* 2019), as well as climatic models (Arruda *et al.* 2017; Costa

et al. 2017; Häggi *et al.* 2017), suggest that general patterns of regional vegetation cover (i.e. forest, savannah) did not change drastically in tropical South America in comparison with other regions of the world over the past 100,000 years, but did vary spatially and over time under the influence of both geological and climatic changes (Hoorn *et al.* 2010; Antoine *et al.* 2016; Wang *et al.* 2017). The dynamic nature of Amazonian vegetation cover during the Quaternary may not have been extremely drastic (e.g., rapidly replacing closed canopy forest by savanna), but sufficient to change the forest cover and to affect the distribution of specialized species (Arruda *et al.* 2017; Wang *et al.* 2017; Silva *et al.* 2019; but see Sato *et al.* 2021). Current evidence fails to support one of the better-known hypotheses for Amazonian diversification, the Pleistocene Refugia hypothesis as originally proposed by Haffer (1969). The Refugia hypothesis proposed that Pleistocene climatic oscillations led to the cyclic replacement of forest- and savanna-covered landscapes, resulting in recurrent isolation and merging of populations, and leading to an increased rate of formation of new species. Although available data from multiple sources now indicate that savannah and open grassland ecosystems have never been widespread in the Amazon (Liu and Colinvaux 1985; Colinvaux *et al.* 2000; Bush and Oliveira 2006), the eastern Amazon probably experienced substantial changes in vegetation structure, with possible episodes of open vegetation expansion (Cowling *et al.* 2001; Arruda *et al.* 2017, Sato *et al.* 2021) that may have affected species distributions and diversification. Nevertheless, it is important to stress that the effects of Pleistocene climate oscillations on the diversification of Amazonian biotas are incomplete and generalizations should be taken cautiously.

ASSEMBLING THE MEGADIVERSE AMAZONIAN BIOTA

Diversification dynamics

Amazonian biodiversity was assembled through a unique and unrepeatable combination of processes that intermingle geological, climatic, and biological factors across broad spatial and temporal scales, involving taxa distributed across the whole of the South American continent and evolving over a period of tens of millions of years. From a macroevolutionary perspective, the number of species in a geographic region may be modeled as a balance between rates of speciation and immigration that increase overall species numbers, and extinction that decreases species richness (Voelker *et al.* 2013; Castroviejo-Fisher *et al.* 2014; Roxo *et al.* 2014). A region that accrues high species richness due to elevated speciation rates has been referred to as an “evolutionary cradle” of diversity, i.e., a place of high species origination (Gross 2019). By contrast, a region where species tend to accumulate through low rates of extinction may be called an “evolutionary museum” of diversity (Stebbins 1974; Stenseth

1984). Although a useful heuristic in some contexts, this model is a poor fit to Amazonian biodiversity. Amazonian species and higher taxa exhibit a broad range of evolutionary ages, such that the Amazon serves simultaneously as both an evolutionary cradle and museum. Still, groups with different average phylogenetic ages tend to inhabit different geographic portions of the Amazon basin. Species assemblages in the upland Guianas and Brazilian Shields (>250 – 300 m elevation) often include a mix of both older and younger lineages, while the lowland sedimentary basins often harbor younger lineages. This pattern is observed in many taxonomic groups (e.g., plants, Ulloa Ulloa and Neill 2006; Amazonian rocket frogs *Allobates*, Réjaud *et al.* 2020; fishes, Albert *et al.* 2020a), although exceptions also exist (Castroviejo-Fisher *et al.* 2014; Bonaccorso and Guayasamin 2013). Similar contrasting core-periphery patterns are observed in many Neotropical taxa, including birds, mammals, snakes, frogs, and plants (Antonelli *et al.* 2018; Azevedo *et al.* 2020; Vasconcelos *et al.* 2020). Diversification in response to geographic barriers is one of the most widespread processes that facilitates speciation. In the Amazon, this process is thought to have played an important role in the evolution of the local biota (e.g., Crouch *et al.* 2018). Geographic barriers can isolate individuals that once belonged to a continuous population of a given species into two (or more) non-overlapping sets of populations (Coyne and Orr 2004). When this geographic separation is maintained for long periods of time, new species may be formed through a process called allopatric speciation (Figure 4). For instance, the uplift of the Andes separated previously connected lowland taxa, preventing dispersal, and establishing new habitats that have fostered the evolution of novel, independent lineages (Albert *et al.* 2006; Hutter *et al.* 2013; Canal *et al.* 2019; Figure 5). This event fragmented the aquatic fauna of northwestern South America, leaving a clear signal on all major taxa (Albert *et al.* 2006). Among families of freshwater fishes, species diversity is significantly correlated with a minimum number of cis-/trans-Andean clades, which indicates that the relative species diversity and biogeographic distributions of Amazonian fishes were effectively modern by the Late Miocene (Albert *et al.* 2006).

Changes in river drainage networks have also strongly affected dispersal, gene flow, and biotic diversification within the Amazon. Large lowland Amazonian rivers represent important geographic barriers for groups of primates (e.g., Wallace 1852; Ayres and Clutton-Brock 1992), birds (Ribas *et al.* 2012; Silva *et al.* 2019), fishes (Albert *et al.* 2011a), butterflies (Brower 1996; Rosser *et al.* 2021), wasps (Menezes *et al.* 2020), and plants (Nazareno *et al.* 2017, 2019a, b, 2021). Similarly, past climatic change is believed to have cyclically changed the distribution of Amazonian habitats such as closed-canopy forests, open forests, non-forest vegetation, and cold-adapted forests, often causing population fragmentation

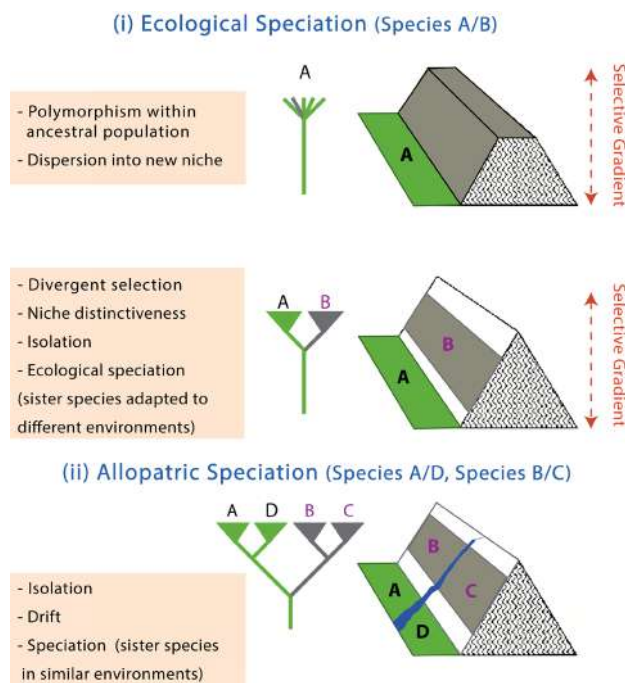


Figure 4. Common speciation mechanisms in the Amazon basin. Ecological speciation: the process by which new species form as a consequence of selection along climatic or ecological gradients, such as those encountered in the Andes. Note that the resulting species occupy distinct environments. Allopatric speciation: when populations of the same species become isolated because of geographical barriers, such as rivers or mountain ranges. Note that the resulting sister species occupy the same environment. Modified from Guayasamin *et al.* (2020).

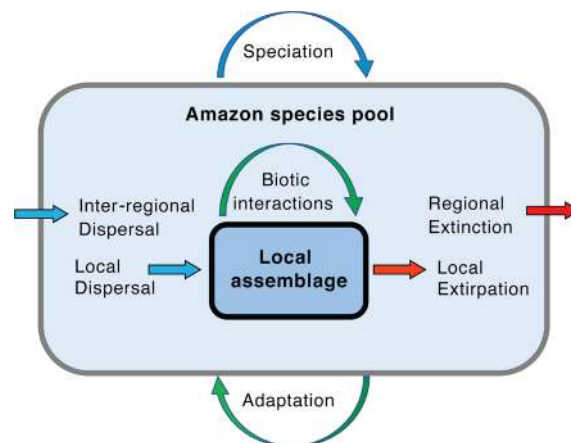


Figure 5. Regional and local processes underlying the assembly of the Amazonian biota. The regional species pool (outer light-blue box) is defined as the sum of all the local species assemblages (inner dark-blue box). Blue arrows indicate processes that increase species richness, red arrows highlight those that reduce species richness. Green arrows represent processes that modify or filter species traits. Speciation and dispersal contribute new species to the regional pool, while extinction removes species. Habitat filtering, dispersal ability, and facilitation affect the richness of local assemblages by limiting or enhancing the establishment of species pre-adapted to local conditions. Local extinction may arise from biotic interactions (such as predation and competition), or abiotic factors (e.g., tectonics or climate change). Adapted from Ricklefs and Schluter (1993), Vellend and Orrock (2009) and Antonelli *et al.* (2018). Reproduced with permission from Guayasamin *et al.* (2021).

and speciation (Cheng *et al.* 2013; Arruda *et al.* 2017; Wang *et al.* 2017; Silva *et al.* 2019).

Apart from the importance of past geographic isolation and speciation due to habitat discontinuity, adaptation to specific habitats has also contributed significantly to species diversification in this region (Figure 5). The large geographical extension of the Amazon, tied to its diverse soil types, provided multiple opportunities for ecological specialization (Fine *et al.* 2005; Tuomisto *et al.* 2019). This soil heterogeneity reflects the complex geological history of northern South America.

While the erosion of the Guiana and Brazilian shields produced the soils of the eastern Amazon, younger sediments that are products of Andean orogeny have developed soils in the western Amazon that tend to be more fertile (Tuomisto *et al.* 2014). This east-to-west gradient in soil fertility is paralleled by a gradient in species composition, wood density, seed mass, and wood productivity (but not forest biomass, see Ter Steege *et al.* 2006; Tuomisto *et al.* 2014). Likewise, different levels of forest inundation during the annual flooding cycle

have contributed to the formation of diverse habitat types and specializations in groups of birds and fishes (Albert *et al.* 2011a; Wittmann *et al.* 2013; Luize *et al.* 2018; Thom *et al.* 2020).

Habitat heterogeneity has played an important role in the formation of Amazonian biodiversity, with geological changes also impacting the ecological conditions available to the Amazonian biota. Andean uplift, for instance, has had a major effect on the Neotropical climate, as it created both habitat and climate heterogeneity while leading to the humidification of Amazonian lowlands and the aridification of Patagonia (Blisniuk *et al.* 2005; Rohrmann *et al.* 2016). The Andes, with an average elevation of 4,000 m, exhibit an immense gradient of humidity and temperature. This has provided numerous opportunities for colonization, adaptation, and speciation events in lowland species, such as frogs, birds, and plants, at different times (Ribas *et al.* 2007; Hutter *et al.* 2013; Hoorn *et al.* 2019; Cadena *et al.* 2020a; Réjaud *et al.* 2020; Figure 6). As a consequence, the Andes are disproportionately

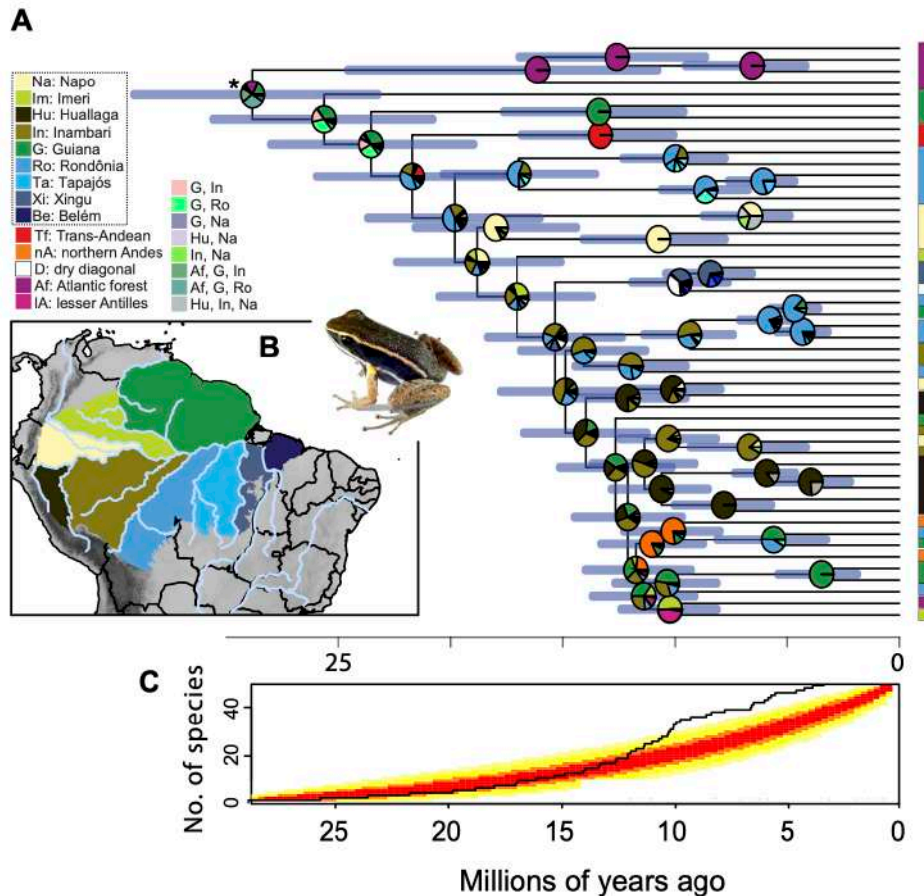


Figure 6. Diversification and endemism in Amazonian rocket frogs (*Allobates* spp.). Closely related species display an allopatric pattern of distribution, matching interfluves delimited by modern Amazonian rivers. **A** – Evolutionary relationships, represented as a phylogenetic tree. Time is provided along the horizontal axis. Blue bars denote the confidence intervals around the inferred time of speciation; pie charts indicate how probable are the estimated ancestral areas of each clade; and colored squares represent the current distribution of each species. **B** – Amazonian areas of endemism. **C** – Inferred number of lineages accumulated through time. Modified from Réjaud *et al.* (2020).

more biodiverse relative to their surface area (e.g., Testo *et al.* 2019). This dynamic interaction between lowlands and adjacent mountains are known to generate diversity worldwide (Quintero and Jetz 2018; Rahbek *et al.* 2019). Repeated cycles of ecological connectivity and spatial isolation in the high Andes (as observed in today's páramos) may have acted as a "species pump" and significantly increased speciation rates in high-elevation Andean taxa due to the joint action of allopatry, natural selection, and adaptation (Madriñán *et al.* 2013; Rangel *et al.* 2018; Pouchon *et al.* 2018).

The contributing roles of abiotic and biotic processes in biodiversification have been neatly summarized as the so-called Court Jester and Red Queen perspectives, respectively (Benton 2009). The Court Jester hypothesis emphasizes the role of abiotic physical and chemical factors as major drivers of speciation (emphasizing, for example, the role of adaptation to climate, substrate, or water condition; Barnosky 2001). Abiotic factors deriving directly from geographic space, climatic and elevation gradients, topographic relief, hydrology, and sediment and water chemistry all serve to facilitate organismal diversification into major habitat types. Intertwined with these landscape processes are innumerable biotic processes that create new species and prevent extinction (e.g., competition, predation, parasitism, mutualism, and cooperation). These biotic interactions can lead to the co-evolution of new traits, increase the structural heterogeneity and functional dimensions of habitats, and enhance the genetic and phenotypic diversity of Amazonian ecosystems. Together with the evolutionary processes that emerge from them, these biological interactions are emphasized in the Red Queen hypothesis. As we discuss below, the immense biodiversity of the Amazon results from both abiotic (see Geographical connectivity through time) and biotic (see How biodiversity generates and maintains biodiversity) factors.

Geographical connectivity through time

The Amazon basin is a highly heterogeneous set of landscapes and riverscapes that form a mosaic of habitat types, often characterized by distinct floras and faunas (e.g., Duellman 1999; Cardoso *et al.* 2017; Tuomisto *et al.* 2019; Albert *et al.* 2020a; Figure 7). Abiotic changes and shifts in the distribution and connection among these different habitats across space and through time drove the accumulation of the impressive number of Amazonian species (Dambros *et al.* 2020). Because organisms differ so widely in their functional traits (such as their dispersal abilities and physiological tolerances), the same landscape conditions that allow for demographic and genetic connections in some groups can have no effects or even reduce connections in others. For example, while large lowland rivers such as the Amazon and the Negro constitute effective barriers to dispersal in upland species of monkeys and birds (representing boundaries between closely related species of those groups; Cracraft 1985), these very

same waterways serve as dispersal corridors for riverine and floodplain species of fishes, birds, mammals, and plants with seeds dispersed by fishes or turtles (e.g., Albert *et al.* 2011b; Parolin *et al.* 2013).

This habitat heterogeneity may be one of the reasons why past landscape changes that promoted the diversification of co-existing lineages in the Amazon resulted in different geographical patterns of species distributions among groups, and different times of speciation (Da Silva *et al.* 2005; Naka and Brumfield 2018; Silva *et al.* 2019). In this heterogeneous and dynamic landscape, the effectiveness of an isolating barrier depends on the biological characteristics of individual species, such as their habitat affinity, their ability to move through the landscape, their tolerance to temperature and precipitation extremes, their generation time, clutch size, and abundance patterns, among other factors (Paz *et al.* 2015; Papadopoulou and Knowles 2016; Capurucho *et al.* 2020). Low dispersal ability, for example, facilitates geographic isolation and genetic differentiation that tend to increase speciation rates (e.g., tropical insects, Polato *et al.* 2018), but also increase the risk of local extinction (Cooper *et al.* 2008). Thermal tolerances, on the other hand, mediate the impact of climate on diversity maintenance and speciation rates (Janzen 1967). Because tropical species experience relatively stable environmental temperatures across their annual cycle, they have evolved more narrow thermal tolerances and reduced dispersal capacities relative to temperate species (Janzen 1967; Shah *et al.* 2017), which promotes speciation, especially in mountain gradients (Polato *et al.* 2018). Lowland tropical species also live under temperature conditions close to their thermal maximum, which places them at risk in the face of increased global warming (Colwell *et al.* 2008; Deutsch *et al.* 2008; Campos *et al.* 2018; Diele-Viegas *et al.* 2018, 2019). This is especially true for species that inhabit areas that are either distant from mountain ranges or fragmented, impeding migration to higher (and cooler) environments.

Because Amazonian species have unique evolutionary trajectories and variable environmental requirements, they have been differentially affected by past geological and climatic events. Patterns of historical connectivity among populations that inhabit upland rainforest habitats have been profoundly influenced by the changing courses of major lowland rivers and their associated floodplains over millions of years, and also by prominent topographic and habitat discontinuities, such as patches of rugged terrain, open savannah vegetation, and sandy soils (Capurucho *et al.* 2020; Cracraft *et al.* 2020). As an example, while the relatively narrow and young Branco River delimits the distribution of some primate species (Boubli *et al.* 2015), this river has had a different role in the evolution of some birds (Naka and Brumfield 2018), plants (Nazareno *et al.* 2019a, b, 2021), and some small-bodied fishes (Dagosta and Pinna 2017), serving as an effective barrier for some species but not for others. Traits are hence important not only to

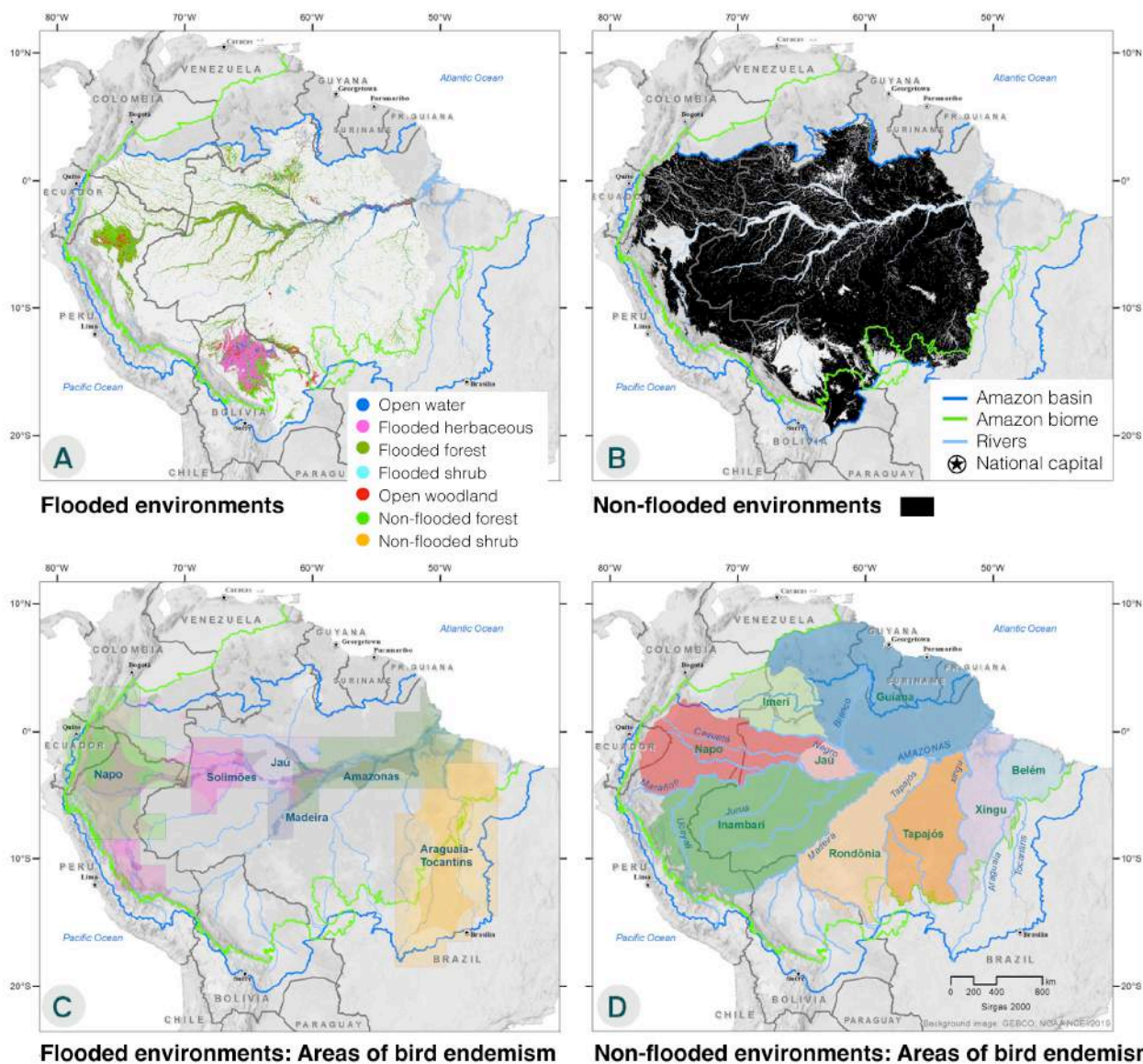


Figure 7. Habitat heterogeneity and bird distribution and endemism in the Amazonian lowland floodplain areas. Distribution of flooded (~14% of the total area) (A) and non-flooded environments (B) (modified from Hess *et al.* 2015). Areas of endemism for birds associated with flooded (C) (Cohn-Halt *et al.*, 2007) and non-flooded (D) (Silva *et al.* 2019) environments. Reproduced with permission from Guayasamin *et al.* (2021).

define the distribution and degree of connectivity of extant populations, but they have also influenced their evolutionary history over time (see Tamme *et al.* 2014; Weeks *et al.* 2022).

Both terrestrial and aquatic Amazonian habitats have been profoundly affected by climate change, especially changing precipitation patterns and sea levels, over millions of years (Vonhof and Kaandorp 2010). Many studies have discussed the influence of past climates on Amazonian landscapes while focusing on changes of the relative cover of forest and savanna (e.g., Bush and Oliveira 2006). However, more subtle changes in forest structure may also affect species distributions and landscape connectivity (Cowling *et al.* 2001; Arruda *et al.* 2017). Understanding how to maintain population connectivity is key

to protecting Amazonian biodiversity. For instance, it is believed that the resilience of upland Amazonian forest taxa in face of environmental changes that occurred through time could be explained by the large dimensions of suitable habitat that allowed them to track appropriate climatic conditions, possibly explaining why so many upland forest species exhibit signs of changes in population size during the Pleistocene (Silva *et al.* 2019). These historical dynamics lay the foundation for predictions of how future climate change will affect patches of humid forests, which are becoming increasingly fragmented due to deforestation and other human land-use activities (Albert *et al.* 2023).

Trait mediated diversification in a heterogeneous Amazon

Studies that consider the habitat affinities of Amazonian species show that the history of each taxon, and its resilience through time, is deeply linked to the kinds of environments it occupies. This view is transforming the way scientists and the general public view the Amazon. Because the heterogeneity of lowland Amazonian habitats has been underappreciated, and because the region has been (wrongly) perceived as a large and homogeneous ecosystem, many taxa have been mistakenly considered widespread and generalist, and, consequently, resilient to landscape change (Bates and Demos 2001). Among birds, one of the best studied groups in Amazonia, it has been demonstrated that species from upland non-flooded forest have different ecological associations and evolutionary histories relative to the species that inhabit the floodplains and to those in open vegetation areas (Figures 7–9). Consequently, the geographical distribution of biological diversity differs among those three groups, and likely also their resilience to future environmental shifts (Capurucho *et al.* 2020; Cracraft *et al.* 2020; Thom *et al.* 2020).

Birds associated with upland non-flooded forest are the most diverse (currently comprising about 1,000 species; Billerman *et al.* 2020). Within these groups, distinct species, although closely related, are found in each main Amazonian interfluvium (Silva *et al.* 2019). Similar patterns have also been described for other groups of Amazonian lineages, mostly distributed in upland forests (e.g., Craig *et al.* 2017; Godinho and da Silva 2018; Figure 9). In contrast, populations associated

with seasonally flooded environments, whose available habitats are currently distributed along the main Amazonian rivers, have been impacted by drastic habitat change due to shifts in the drainage system during the last 5 Ma (Bicudo *et al.* 2019), including significant changes even within the last 45 thousand years ago (Ka) (Pupim *et al.* 2019). While large rivers are barriers for the dispersal of small-bodied understory birds in humid non-flooded forests, the seasonally flooded vegetation that grows along these rivers promotes connections across populations of floodplain-associated species adapted to the annual flooding cycle of river floodplains. Differently from the upland non-flooded forest birds, floodplain species have little intraspecific diversity, but they represent older lineages that originated during the Middle to Late Miocene (5–11 Ma; Thom *et al.* 2020). The largest genetic differences within these widespread floodplain species are observed between populations from the western sedimentary basins and populations from the eastern shields (Thom *et al.* 2018, 2020). These distinct evolutionary trajectories have helped to shape the history of Amazonian floodplains (Bicudo *et al.* 2019). Data from floodplain-adapted birds and fishes, for instance, indicate historically larger and more connected populations in the western Amazon (Santos *et al.* 2007; Thom *et al.* 2020), and cycles of connectivity and isolation between species that occupy seasonally flooded habitats in the eastern vs. western Amazon. Organisms adapted to seasonally flooded landscapes are particularly vulnerable to disruptions of connectivity caused either by historical landscape change or by anthropogenic impacts such as dams and waterways (Latrubesse *et al.* 2017; Anderson *et al.* 2018).

Species associated with open vegetation growing on sandy soils have yet a third pattern of diversity distribution in the Amazon. Among plants and birds, for instance, populations of the same species are distributed in patches of open habitat separated by upland and flooded forests and located thousands of kilometers apart, spanning all the main interfluvia (Capurucho *et al.* 2020). Despite having a naturally fragmented distribution today, these species were less isolated in the past, suggesting that, although present in the Amazon for millions of years, the distribution of open vegetation has varied through time (Adeney *et al.* 2016).

Together, these contrasting patterns indicate that the Amazonian landscape and its different habitats have been spatially dynamic during millions of years, and that the current distribution of habitats and species represents a snapshot in time.

How biodiversity generates and maintains biodiversity

There is little doubt that diverse biotas with many functionally distinct organisms, complex biotic environments, and multiple ecological interactions facilitate species coexistence and elevate regional species richness and density



Figure 8. Amazonian habitat heterogeneity: Plant and bird species adapt to habitats with different soils. Species of the plant family Burseraceae (in the genera *Protium*, *Crepidospermum*, and *Tetragastris*) are examples of specialization to the different types of soil that occur throughout *terra firme* habitats: **A** – clay-soil forest; **B** – terrace-soil forest; **C** – white-sand vegetation (Fine *et al.* 2005). Amazonian bird species that occur exclusively in patches of white sand vegetation are often related to species from open habitats outside Amazonia, like the Cerrado and Tepuis (Capurucho *et al.* 2020; Ritter *et al.* 2020), and do not have close relatives occupying the adjacent humid forest. This result suggests that the adaptations necessary to occupy these open vegetation habitats may not be common within forest specialized groups. Photos by Camila Ribas; reproduced with permission from Guayasamin *et al.* (2021).

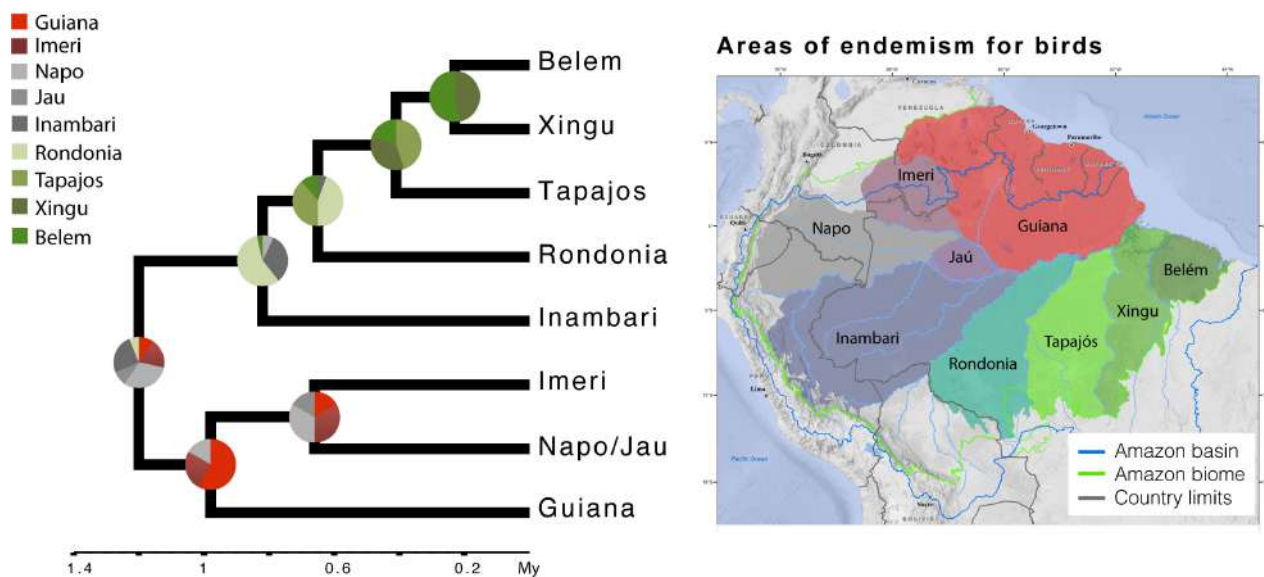


Figure 9. Areas of endemism and diversification patterns for 21 taxonomic clades of Amazonian birds restricted to the upland forest (*terra firme*) understorey. **A** – Relationship among nine areas of endemism, inferred from genetic data; pie charts denote ancestral area probabilities. **B** – Areas of endemism currently recognized for upland forest birds in the Amazon region. Notice how the diversification history of this group matches the location of Amazonian rivers that delimit areas of endemism (e.g., the Tocantins River between the Belém and Xingu endemism areas). Also evident is an initial differentiation between clades north of the Amazonas River (represented by the areas Guiana, Imeri, and Napo/Jaú) from those south of the Amazonas River (Inambari, Rondonia, Tapajós, Belém, Xingu). Modified from Silva *et al.* (2019).

values. In this regard, biological diversity may be understood to be autocatalytic: species richness itself is a key feature in the origin of hyperdiverse Amazonian ecosystems (Sombroek 2000; Albert *et al.* 2011b; Dáttilo and Dyer 2014).

The notion that biotic interactions help drive organismal diversification is not new. In a famous article, the paleontologist Leigh Van Valen observed that the life span of species as shown by the fossil record was roughly constant (van Valen 1973). Borrowing from a line in *Through the Looking Glass* by Lewis Carroll, where the Red Queen tells Alice “*It takes all the running you can do, to keep in the same place*”, he proposed the Red Queen hypothesis as a metaphor to express the idea that lineages do not increase their ability to survive through geological time (van Valen 1973). In modern evolutionary theory, Red Queen dynamics refers to phenotypic evolution in response to biotic interactions, such as the coevolution of parasites and their hosts, chemically defended prey and their predators, and interactions between pollinators and the plant species they visit. In all these biotic interactions, adaptive changes in one species may be followed by adaptations in another species, spurring an evolutionary arms race that may result in co-evolution or extinction, or both (Strotz *et al.* 2018).

Other examples of potential Red Queen dynamics include organisms that affect the physical environment experienced by other species, such as plants that constitute structural habitat (e.g., tank bromeliads, which provide breeding habitat for frog species and invertebrates), or organisms that modify the physical and chemical environments utilized by several other

taxa (e.g., fungi and earthworms that change soil and water chemistry). Organismal interactions such as those, which benefit at least one member of a local species assemblage, are referred to as biotic facilitation (Zélé *et al.* 2018). Below, we provide several examples of how biotic interactions have facilitated the evolution of Amazonian diversity.

Host-parasite interactions: Because the species composition of many parasite groups often tracks that of their hosts, it is possible to estimate a minimum number of parasite species by comparison with the diversity of their host taxa (e.g., McNew *et al.* 2021; but see Weckstein 2004). Given that many fish parasites exhibit strong host-specificity, it is believed that the actual diversity of the parasites could rival the immense diversity of their fish hosts (Salgado-Maldonado *et al.* 2016). At present, only about 300 species of Neotropical monogenoid flatworms are described, all ectoparasites of fish gills and the external body surface; however, these numbers are rising rapidly due to ongoing taxonomic research (see Vianna and Boeger 2019). Moreover, tight associations between helminth (flatworm and roundworm) and haemosporidian (*Plasmodium*) parasites and host species have been reported in many groups of Amazonian vertebrates, including fishes (Thatcher 2006), amphibians and reptiles (McAllister *et al.* 2010), and birds (Fecchio *et al.* 2018). The diversity of protozoan parasites of vertebrate hosts in the Amazon is presumably much greater still, based on what is known from better-studied faunas (Dobson *et al.* 2008). Even less is known about the diversity of Amazonian insect and plant parasites, but glimpses provided by recent studies using environmental genomics indicate the existence of extraordinary genetic and

functional diversity of metazoan and protozoan parasites in the Amazon (Mahé *et al.* 2017; Puckett 2018).

Niche construction: Biological diversity also contributes to the evolution of more diversity through the many ways by which organisms modify their external environments. The process by which organismal behaviors alter their local environments is called niche construction, which also affects the ecological conditions for all organisms in a local assemblage (Odling-Smee *et al.* 2013). Organismal behaviors strongly affect and even create many important habitats in the Amazon. These activities include nest-burrow construction and fruit-seed-pollen dispersal by animals, the formation of vegetation structure and shade by plants, and the roles of plants, fungi, and soil or water microbes in nutrient and energy cycling, soil and water chemistry, and fire regimes (Mueller *et al.* 2016; Santos-Júnior *et al.* 2017). Earthworms (*Clitellata*, Annelida) represent a classic example of how niche construction elevates habitat heterogeneity and biodiversity in the Amazon. Earthworms are important ecosystem engineers, whose activity helps to mineralize soil organic matter, construct and maintain soil structure, stimulate plant growth, and protect plants from pests (Marichal *et al.* 2017). Several other Amazonian taxa are also important engineers of terrestrial ecosystems, including fungi (Palin *et al.* 2011), termites (Duran-Bautista *et al.* 2020), and ants (Folgarait 1998).

Keystone species: The high number of fish species in aquatic Amazonian ecosystems can strongly affect nutrient and energy cycling (Winemiller and Jepsen 1998; Arruda *et al.* 2017). A striking example is the ecological role of the “coporo” or “sábalo” (*Prochilodus mariae* Eigenmann, 1922), a detritivorous and migratory characiform fish that

is functionally important in Andean foothill streams of the western Amazon and Orinoco basins. Selective exclusion of this single species qualitatively changes the structure of local aquatic communities, as measured by sediment accrual and the composition of algal and invertebrate assemblages (Flecker 1996). Another example is provided by planktivorous electric fishes (Gymnotiformes) that constitute the base of aquatic food webs in the Amazon and Orinoco basins (Lundberg *et al.* 1987; Fernandes *et al.* 2004). Because these food webs are essential to support the regional fisheries on which millions of Amazonian people depend as a primary source of animal protein (Goulding *et al.* 2019), planktivorous fishes are keystone species to human-dominated Amazonian landscapes.

Predator-prey interactions and the evolution of chemical diversity (Figure 10): Predator-prey dynamics is one of the most powerful evolutionary forces in nature, resulting in a myriad of strategies and weaponry to prey or avoid predation. Some long-evolved interactions between Amazonian species are responsible for the generation and accumulation of natural products amenable to bioprospection. Amazonian poison frogs (family Dendrobatidae), for instance, are known to sequester chemical defenses from the arthropod prey that they feed upon (Saporito *et al.* 2011 and references therein). These alkaloids are used by indigenous people and are explored by the medical community and the pharmaceutical industry (Daly *et al.* 2000; Cordell *et al.* 2001; Philippe and Angenot 2005). Mites, ants, beetles, and millipedes have all been flagged as sources of alkaloids for poison frogs worldwide (Saporito *et al.* 2009; McGugan *et al.* 2016), and several species of frogs are able to further modify them chemically, leading to other alkaloids (Daly *et al.* 2003, 2009). Moreover, although more research is pending, some poison frog alkaloids appear to be derived from

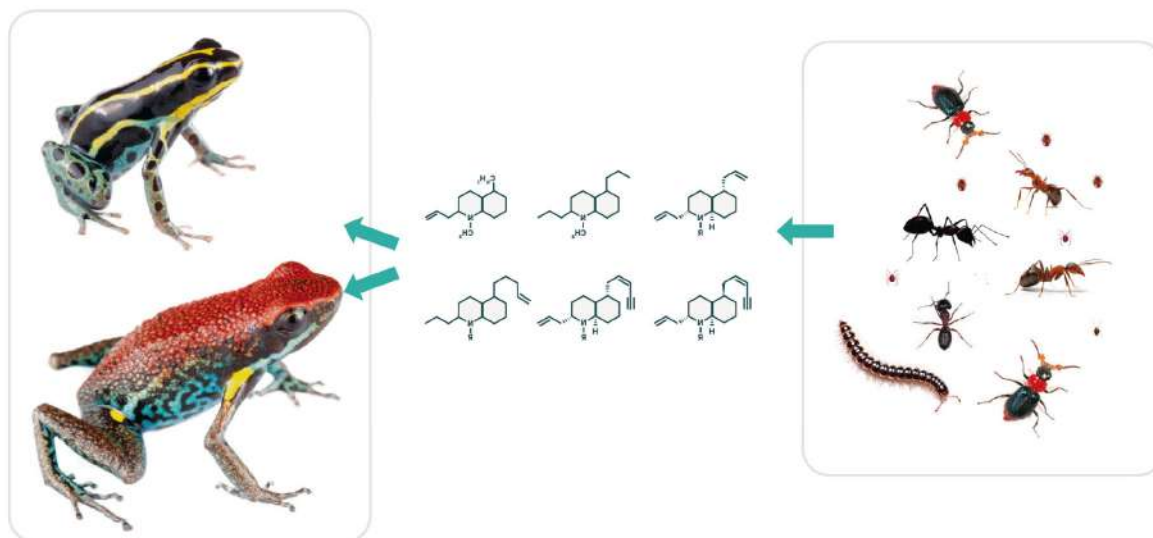


Figure 10. The evolution of jumping pharmacies. Poison dart frogs (family Dendrobatidae) are protected by alkaloids that they sequester from their prey, including ants, mites, millipedes, and melyrid beetles (see Saporito *et al.* 2011 and references therein). Alkaloid sequestration and modification is both an outcome of biotic interactions between Amazonian frogs and their invertebrate prey, and a mediator of interactions between those same frogs and their predators (e.g., Brodie and Tumbarello 1978; Fritz *et al.* 1981; Macfoy *et al.* 2005; Darst *et al.* 2006; Weldon *et al.* 2006). Reproduced with permission from Guayasamin *et al.* (2021).

plants. This reflects the complex trophic interactions between plants, the arthropods that feed on them, and the frogs that prey on those arthropods (Tokuyama and Daly 1983).

The potential of plants for the Amazonian bioeconomy is enormous. For instance, Amazonian people have known the effects of plant alkaloids as medicine for centuries. Plant alkaloids evolved as a defense mechanism against herbivory (Gauld *et al.* 1992) and are synthesized in the roots, stems (e.g., banisterine), leaves (e.g., caffeine), flowers, fruits, seeds (e.g., strychnine), and bark (e.g., quinine). Some of the known plant alkaloids include the antimalarial quinine, hunting poisons (barbasco, curare), stimulants (guayusa, nicotine, coca), and ritualistic herbs (ayahuasca, scopolamine) (Heinrich *et al.* 2021; Uzor 2020). Many of these compounds are precursors for modern medicine; however, due to their complex chemical structures, only a fraction go into commercial production (Reis *et al.* 2019). Moreover, allochemicals from some Amazonian plants might prove useful as sources of biodegradable pesticides; the “piquiá” (*Caryocar*), for instance, produces a compound that seems to be toxic to the leaf-cutter ant (*Atta*), which causes large financial losses to South American agriculture each year (Plotkin 1988). Today, entire companies are dedicated to screening chemical compounds in plants, insects, and frogs, in search for potential drugs. Natural products and their derivatives have been, and continue to be, a primary source in the drug discovery domain (Lopes *et al.* 2019).

SPECIES LOSS AND TURNOVER IN AMAZONIA: LESSONS FROM THE FOSSIL RECORD

Extinction rates vary throughout evolutionary time. It has been estimated that more than 99% of all species that have ever lived on Earth are now extinct (Raup 1986). The fossil record offers unique evidence to study extinctions. Paleontologists have identified 18 time intervals with elevated extinction rates over the past 540 million years, five of which are classified as mass extinction events (Bambach 2006). Models based on DNA analyses and the fossil record, especially of marine invertebrates and mammals, show that background extinction rates over geological time have ranged from 0.02 to 0.14 extinctions per million species per year. In turn, speciation rates are estimated to be about twice this value, ranging from 0.05 to 0.20 speciation events per million species per year (Jablonski 2005; De Vos *et al.* 2015). The fossil record also shows changes in biodiversity over geological time with occasional catastrophic mass extinction events, when extinction rates increased by thousands of times, eliminating large clades with distinctive genes and body plans (Bambach 2006; Ceballos *et al.* 2015).

This understanding of the past allows us to put in perspective the wave of extinctions faced by the modern biota, which is estimated to be 1,000 to 10,000 times larger than the background rate, and therefore similar in scope to that of past mass extinction events (Barnosky *et al.* 2011; Ceballos *et al.* 2015). While its causes are multiple, the increase in the concentration of carbon dioxide in the atmosphere, and the acidification of the oceans caused by human action, match the great natural environmental changes that triggered mass extinction events in the deep past.

Throughout its lengthy geological history, the Amazonian region has undergone extensive environmental changes, driven primarily by regional tectonic and global climatic forces. It once extended over most of northern South America, with lowlands characterized by alternating fluvial and lacustrine conditions and marginal marine embayments. Modern lineages of Amazonian organisms have survived and adapted to five major rearrangements of landforms and habitats during the Cenozoic (66–0 Ma), as follows (summarized from Val *et al.* 2021, and references therein):

The Paleogene uplift of the central Andes, caused by plate subduction along the Pacific margin and the breakup of the Pacific plate (ca. 23 Ma), resulted in the establishment of a sub-Andean river basin draining north towards a large embayment of the Caribbean Sea. The basin extended over the area that is now occupied by the Colombian and Venezuelan Llanos.

Mountain building in the central and northern Andes narrowed the Caribbean influence and led to the origin and movement of mega-wetlands in the western Amazon ca. 22–10 Ma. The Pebas mega-wetland system resulted from this expansion, reaching more than 1 million km².

Intense Andean mountain building since the late-middle Miocene (last 10 Ma), which coincided with global fluctuations in sea level, prevented further marine influences in the western Amazon and along the northern Andean foreland basin. This retained much of the drainages that flowed into the Pacific and the Caribbean, and formed the wide floodplain named the Acre System.

From the end of the Miocene (ca. 7 Ma) on, further Andean uplift forced the mega-wetland to be completely drained and the onset of the modern Amazon River system. This led to the development of widespread river terrace systems with expanded *terra firme* rainforests.

The closure of the Central American Seaway and the emergence of the Panama Isthmus (ca. 13–3.5 Ma) provided opportunities for extensive migrations of North American lineages to both the Amazon and new montane habitats in the Andes.

The biotic responses to these immense environmental changes included dispersal and habitat shifts at the organismal level, adaptation and geographic range shifts at the population

level, and speciation and extinction at the species level. While the geological record does not provide evidence of sudden mass extinction events during the Cenozoic in the Amazon, some groups of animals once abundant in both terrestrial and aquatic environments were extirpated by one or more of the aforementioned events, including species expected to provide a variety of ecological functions (Scheyer *et al.* 2013).

Extinctions and Amazonian mega-wetlands (Figure 11)

The fossil record evidences pulses of extinctions in Amazonia that occurred in the transition from the lacustrine-fluvial Pebas to the fluvio-lacustrine Acre mega-wetland systems, in association with the origin of the modern transcontinental Amazon River, ca. 9–4.5 Ma (Albert *et al.* 2018). The most significant extinctions were those affecting the rich and endemic lacustrine fauna, notably bivalve mollusks (Wesselingh and Ramos 2010) and crocodylian reptiles (Riff *et al.* 2010; Scheyer *et al.* 2013; Salas-Gismondi *et al.* 2015).

Mollusks and crocodylians are among the best represented clades in the fossil record of the Amazon; they exemplify the diversification and subsequent extinction of aquatic fauna in association with the evolution of mega-wetlands during the Neogene. About 85 species of mollusks were documented from the last stages of the Pebas System (Middle to Late Miocene). This fauna was dominated by Pachydontinae bivalves, which originated in coastal Pacific and Caribbean marine waters. Marine mollusks colonized the western Amazon during pulses of marine incursions ca. 23–15 Ma, together with other aquatic animal groups such as freshwater stingrays, anchovies, needlefishes, dolphins, manatees, and various parasitic lineages (Lovejoy *et al.* 1998). Small, blunt-snouted crocodylians evolved crushing dentitions that allowed them to feed on hard-shelled organisms and prey on the Pebasian malacofauna (Salas-Gismondi *et al.* 2015). The crocodylian fauna of the Pebas system also included species specialized in eating fish (long-snouted gharials), large to giant prey (*Purussaurus*), “gulp-feeding” of small prey (*Mourasuchus*), and generalized small prey (*Caiman* and *Paleosuchus*). On land, the last representatives of an extinct group of terrestrial crocodyliforms, the Sebecidae, competed with mammals as top-predators. This group included the largest terrestrial predator of the Amazon during the Middle Miocene, *Barinasuchus arveloi* Paolillo and Linares, 2007, from the Parangula Formation in Venezuela, which reached up to 6 meters in length (Paolillo and Linares 2007). Because top predators are very susceptible to drastic environmental changes, it is possible that the changes in the mega-wetland impacted the survivorship of these organisms (Salas-Gismondi *et al.* 2015).

With the end of the Pebas System, most of the associated molluscan fauna became extinct. Consequently, modern Amazonian mollusk diversity is remarkably poor and dominated by cosmopolitan freshwater groups, such as freshwater mussels, clams, and snails (Wesselingh and Ramos 2010). The disappearance of the Pebasian endemic mollusks adversely

affected crocodylians, who then suffered their first large-scale extinction event (Salas-Gismondi *et al.* 2015; Souza-Filho *et al.* 2019).

Still, most of the crocodylian lineages survived to the formation of the Acre System ca. 10–7 Ma. In the extensive wetlands of the Acre System flourished a notable diversity of around 30 species showing morphological variation greater than any other crocodylian fauna, extant or extinct (Riff *et al.* 2010; Cidade *et al.* 2019). Similarly, the period witnessed a large diversity of turtles, including one of the largest turtles that ever lived on Earth, more than 2.5 m in length and with an estimated body mass of about 1,000 kg (Cadena *et al.* 2020b). Beyond some generalist taxa that have been present in the Amazon since the Middle Miocene through to today (e.g., *Caiman*, *Melanosuchus*, and *Paleosuchus*), the availability of large-bodied prey and competition with other aquatic predators likely triggered the evolution of giant top predators. Examples include *Purussaurus brasiliensis* Barbosa-Rodrigues, 1892, with its 12-meter long body (Aureliano *et al.* 2015), highly specialized forms such as the bizarre species in the genus *Mourasuchus*, known for their long, wide, dorsoventrally flat skull, and tiny dentition (Cidade *et al.* 2019), and the long-snouted gharials, some also giant in size (Riff *et al.* 2010).

The transition from the Acre System to the modern fluvial and *terra firme* Amazonian environments, starting at around 7 Ma, led to a large extinction event affecting crocodylian fauna. All specialized forms, from small to giant, vanished. The extant South American crocodylians are now a small fraction of their former diversity. Entire body types and ecological roles among aquatic fauna disappeared after the demise of the Amazonian Miocene mega-wetlands.

In stark contrast to the turnover of mollusks and crocodylians, modern Amazonian fish fauna has remained largely unchanged at the genus level and above. Direct evidence from the fossil record indicates that all but one fossil genus known from the Miocene is still living (Lundberg *et al.* 1998). Further, molecular phylogenies of most Amazonian fish genera are now available, including more than 1,000 of the 3,000 known species (van der Sleen and Albert 2017). In combination, these datasets indicate that most genera that compose today’s rich Amazonian fish fauna were present by the middle Miocene (ca. 15–10 Ma). The evolutionary origins of most Amazonian fish forms and their ecological roles predate the geological assembly of the modern Amazon and Orinoco basins during the Late Miocene and Pliocene (ca. 9–4.5 Ma; Albert *et al.* 2011b).

The American Biotic Interchange and the influence of humans on Amazonian biota

The tectonics that elevated the Andes and caused the great environmental changes also elevated the terrestrial route that ended a long-lasting isolation of South America from other

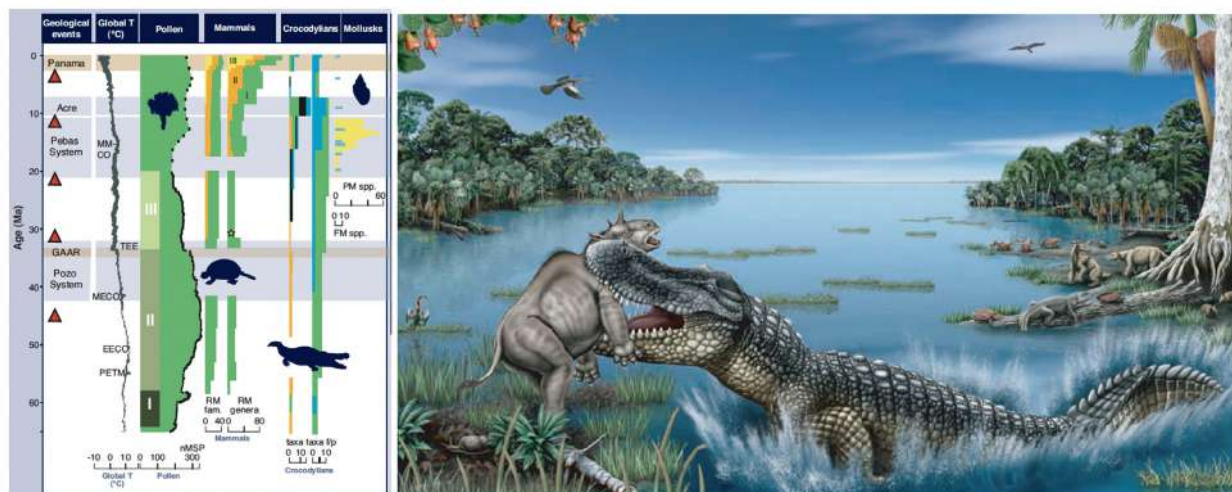


Figure 11. Past diversity in Amazonia and the mega-wetland landscape. Left: Diversity changes through time, as shown by the fossil record. Notice that floral diversity has remained high since the Paleogene (ca. 60 Ma), and crocodiles and mollusks diversified with the onset of the megawetlands and declined with their demise (modified from Hoorn *et al.* 2010). Right: Reconstruction of an Amazonian mega-wetland landscape during the middle to Late Miocene (16–7 Ma) highlighting the giant caiman *Purussaurus brasiliensis* Barbosa-Rodrigues, 1892, preying on a *Trigonodon* toxodont. Illustration by Orlando Grillo, in Hoorn *et al.* (2010).

continents during most of the Cenozoic (Croft 2016). This isolation, which led South America to harbor a peculiar and endemic mammalian megafauna (Defler 2019), ceased when the formation of the Isthmus of Panama facilitated the biotic interchange between North and South America, through the event known as the Great American Biotic Interchange (GABI; Stehli and Webb 1985). This connection had great implications for the historical assembly of the Amazonian fauna and flora. Plants, which have a greater dispersal ability, dispersed before animals did, even before a land bridge was fully established between the continents (ca. 50–20 Ma; Cody *et al.* 2010). The fossil record of terrestrial mammals, which is abundant in both continents and therefore illustrates dispersal dynamics, shows that the interchange was initially symmetrical, but followed by an increasing dominance of mammals of North American origin in South America (Marshall *et al.* 1982), caused by a higher extinction of South American mammals (Carrillo *et al.* 2020). Because the fossil record mostly reflects patterns of the temperate regions (Carrillo *et al.* 2015), molecular phylogenies have also been employed to understand the GABI; they show that dispersal from South to North America occurred most likely between the tropical regions of the two continents (Bacon *et al.* 2015). Indeed, many groups of mammals that are found today in tropical forests from Central America originated in the Amazon, and many of the Neotropical placental mammals, such as felids, canids, peccaries, deer, otters, tree squirrels, camelids, as well as the extinct proboscideans and horses, are descendants of North American migrants (Webb 1991; Antonelli *et al.* 2018).

Global-scale extinction of megafauna impacted the Amazon at the end of the Pleistocene. It reduced megafauna

diversity worldwide by two thirds approximately 50,000–10,000 years ago (Barnosky *et al.* 2004). Hunting by humans was an important cause of extinctions, in some regions in synergy with climate change (Barnosky *et al.* 2004; Barnosky and Lindsey, 2010). South America lost about 83% of its megafauna (adult body weight > 44 kg sensu Martin 1973) during this extinction event, more than any other continent (Barnosky and Lindsey 2010; Prado *et al.* 2015). This loss affected some important ecosystem processes. Because large animals play an important role in the spatial movement of nutrients from areas of high to low nutrient concentration, megafauna extinctions resulted in reduced nutrient flows (Doughty *et al.* 2016a). Extinctions likely reduced the population size of large-seeded tree species that depended on large herbivores for dispersal. In the Amazon basin, the size range of large seeded trees decreased by about 26-31% (Doughty *et al.* 2016b). Furthermore, because fruit size correlates with wood density, the reduction of large-seeded trees dispersed by animals is thought to have reduced the carbon content in the Amazon by about 1.5% after megafauna extinction (Doughty *et al.* 2016b).

The global fossil record shows us that species with specialized diet, larger body size, broader geographic distribution, longer life span, slower reproduction rate, and fewer offspring, are more susceptible to change and in greater risk of extinction (McKinney 1997; Purvis *et al.* 2000). On the other hand, short-lived species with rapid population growth, more generalist diet, and with high phenotypic plasticity are better suited to adapt and cope with environmental change (Chichorro *et al.* 2019). The Amazonian fossil record of Cenozoic crocodylians illustrates this pattern, with large and dietarily-specialized forms occupying large areas that were

heavily impacted by environmental change (Scheyer *et al.* 2013; Cidade *et al.* 2019). In the face of environmental pressures currently faced by the Amazon, such as deforestation, fires, hydroelectric dams, and other anthropogenic disturbances (Escobar 2019; Albert *et al.* 2023), it is possible that species with more specialized diet might face greater extinction risk (Bodmer *et al.* 1997; Shahabuddin and Ponte 2005; Benchimol and Peres 2015).

Humans may have occupied the Americas much earlier than previously thought, with records dating back to 33,000–31,000 years ago in Mexico (Ardelean *et al.* 2020) and 13,000 years ago in lower latitudes (Roosevelt *et al.* 2013). As such, human impact on local ecosystems, including the Amazon, has a lengthy history (Levis *et al.* 2017; Watling *et al.* 2017). Studies from multiple disciplines suggest that pre-Columbian human settlements in the Amazon basin were complex and culturally diverse, and that they influenced current patterns of Amazonian biodiversity (Heckenberger and Neves 2009; Shepard and Ramirez 2011).

Although human influence in the Amazon basin has changed through time, one of the most outstanding legacies of these interactions over many millennia is the abundance and widespread distribution of plant species commonly used by indigenous peoples. These trees, now identified as hyperdominant, include the Brazil nut (*Bertholletia excelsa* Humboldt and Bonpland, 1807), several species of palms (e.g., *Astrocaryum murumuru* Martius, 1824, *Oenocarpus bacaba*), cacao (*Theobroma cacao*, Linnaeus, 1753), and the caimito (*Pouteria caimito* [Ruiz & Pavón, 1802]) (Shepard and Ramirez 2011; Levis *et al.* 2017). These domesticated/managed species have been vital to the livelihood of Amazonian peoples, who have interacted with the forest for many centuries (Levis *et al.* 2017; Montoya *et al.* 2020).

Accumulating evidence demonstrates that the socially and culturally complex pre-Columbian Amerindians modified the riverine, *terra firme*, and wetland areas of the Amazon, directly impacting the distribution of local species assemblages (Heckenberger 2005; Montoya *et al.* 2020). Examples include anthropogenic soils (*terra preta*) and artificial earthworks such as fish ponds, ring ditches, habitation mounds, and raised fields (Heckenberger and Neves 2009; Prestes-Carneiro *et al.* 2016). The magnitude of these changes varied considerably. In areas such as the Llano de Moxos (Bolivia), natives created a landscape that comprised approximately 4,700 artificial forest islands within a seasonally flooded savannah (Lombardo *et al.* 2020). This region has been confirmed as a hotspot for early plant cultivation, including squash (*Cucurbita* sp.), at about 10,250 calibrated years before present (cal. yr bp), manioc (*Manihot* sp.) at about 10,350 cal. yr bp, and a secondary improvement center for the partially domesticated maize (*Zea mays* Linnaeus, 1753), at about 6,850 cal. yr bp (Kistler *et al.* 2018; Lombardo *et al.* 2020).

Changes across the Amazon basin accelerated with Portuguese and Spanish colonization in the past 500 years, and accelerated again during with the transition to modern socio-economic activities (reviewed by Albert *et al.* 2023). The modern Amazon basin is now home to about 35 million people, including about 400 indigenous and traditional communities, but also a large mestizo population concentrated in urban and rural áreas (see Nobre *et al.* 2021). In the last 40 years, the Amazon has undergone unprecedented demographic and ecological transformations, in which the original indigenous populations suffered population crashes because of new diseases, displacement, and hostility (Walker *et al.* 2015; Cuví *et al.* 2021), and ecosystems have been degraded by industrial and agricultural activities (Albert *et al.* 2023). The magnitude of past and current human interlinks and impacts on Amazonian biodiversity was recently reviewed by the Science Panel for the Amazon (Nobre *et al.* 2021).

CONSERVATION OF ECOLOGICAL AND EVOLUTIONARY PROCESSES

One key goal of conservation biology is to provide effective principles and tools for preserving biodiversity (Soulé 1985), especially in complex and threatened ecosystems. Critical information for conservation planning in the Amazon is lacking in all major biodiversity dimensions, including taxonomic diversity, geographic distributions, species abundances, phylogenetic relationships, species traits, and species interactions.

The main threats to Amazonian diversity, just like its ecosystems and landscapes, are heterogeneously distributed (RAISG 2020; Figure 12). As such, a “one-plan-fits-all” strategy will not work in the region. Effective conservation strategies must consider the evolutionary and ecological processes that generate and maintain local species diversity in the many unique biological communities present in this large and ecologically relevant area. However, the legal structure for biodiversity conservation in the Amazon (and globally) is based primarily on individual species. Both governmental initiatives (e.g., US Endangered Species Act: <https://www.fws.gov/media/endangered-species-act>) and non-governmental initiatives (e.g., IUCN Red List: <https://www.iucnredlist.org/>) are organized around the ideas and actions of species conservation status and threat categories. In a similar manner, deforestation processes and impacts of infrastructure development, like roads, dams, and waterways, often ignore the compartmentalization of Amazonian diversity, and the unique characteristics of each region and habitat type (Da Silva *et al.* 2005; Latrubesse *et al.* 2017).

While current initiatives focused on endangered species are crucial, it is important not to lose sight of the processes that keep these species alive and those that generate new diversity. For instance, when conservation priorities are viewed from



Figure 12. Relation of forests, anthropized areas and bird endemism in the Amazon: deforested (red), forested (green), dams (black diamonds), and avian areas of endemism (yellow polygons). Note that the greatest immediate threats to Amazonian biodiversity are located along the agricultural frontier in the southeastern Amazon, especially impacting southeastern areas of endemism. Note the large number of dams in the Andes and on the Brazilian Shield. Modified from Guayasamin *et al.* (2021).

an evolutionary standpoint, areas that hold the same number of species may not share the same conservation relevance. Instead, the preservation of areas holding distinct, unique, and/or higher amounts of evolutionary lineages should be given higher conservation priority (Forest *et al.* 2007; Castro *et al.* 2020; Jézéquel *et al.* 2020; Figure 13). By prioritizing connected regions that host widely-divergent lineages, higher levels of phylogenetic uniqueness, and a broader spectrum of the genealogy of life (Meffe and Carroll 1994), scientists can maximize future options, both for the continuing evolution of life on Earth and for the benefit of society (Forest *et al.*

2007). Maximum levels of global phylogenetic diversity lead to higher ecosystem services globally and higher plant services in general for humankind (Molina-Venegas *et al.* 2021).

Conservation priorities based on a deep understanding of how biodiversity patterns have emerged allow us to preserve a potential for future evolution and adaptation (Erwin 1991; Brooks *et al.* 1992). By prioritizing lineages that are rapidly speciating and adapting we might, for instance, be able to preserve higher potential to resist future climatic and ecological change (Gavin *et al.* 2018). Likewise, by increasing evolutionary diversity, we are likely to increase trait diversity

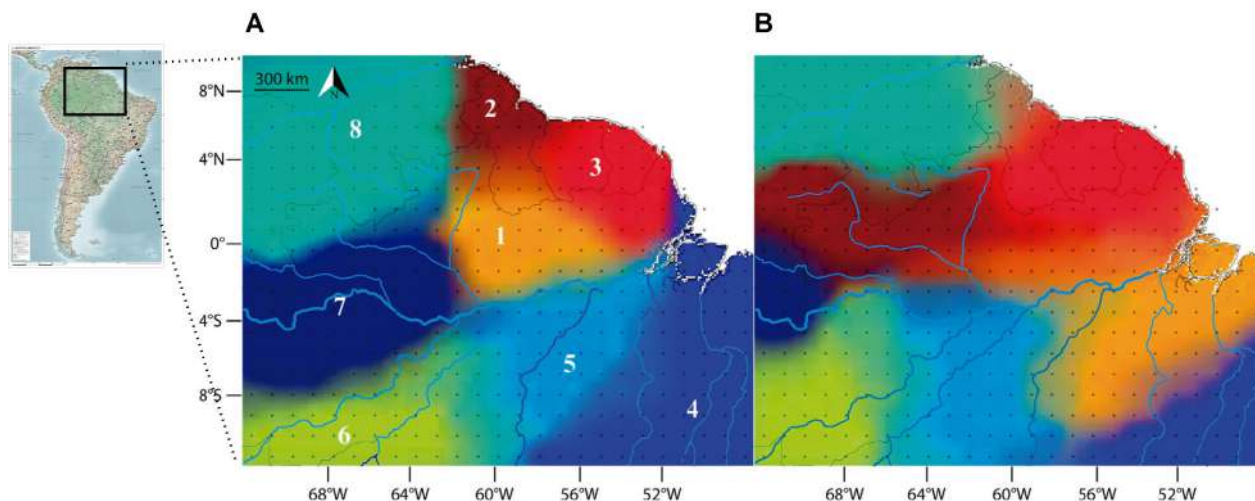


Figure 13. Bioregionalization based on species occurrence data for frogs in the eastern Amazon. **A** – Data from DNA-based species delimitation. **B** – Data from morphology-based taxonomy. Colors represent affiliations of cells to bioregions. Modified from Vacher *et al.* (2020).

and provide increased resilience for Amazon rainforests (Sakschewski *et al.* 2016).

Another way to incorporate evolutionary thinking into conservation is to focus on landscape attributes that generate unique variation or maintain connectivity among populations. Geographic barriers, for instance, restrict species ranges and lead to allopatric diversification (Figure 4). In the Amazon, rivers have imposed limits to the distribution of closely related species (Ribas *et al.* 2012). On the other hand, rivers may also be corridors of connectivity for species associated with floodplain habitats.

Free flowing rivers are hence fundamental not only for the species they support, but also for the evolutionary processes that they drive (Barbarossa *et al.* 2020; Bem *et al.* 2021; Vasconcelos *et al.* 2021). Similarly, the conservation of regions of steep environmental gradients, which are expected to promote ecological speciation (Figure 4), is relevant from an evolutionary standpoint. In the Amazon, for instance, adjacent yet distinct soil types are intimately associated with plant specialization and differentiation (Fine *et al.* 2005; Tuomisto *et al.* 2014). Promoting conservation of these gradients and diverse habitats associated with distinct soil types is therefore important in the short and long term.

The singular diversity of Amazonian organisms was generated over a period of millions of years and represents a large portion of Earth's known and unknown diversity. Because the Amazon has been functioning as a primary source of biodiversity to all other Neotropical biomes (Antonelli *et al.* 2018), forest destruction and species loss in the Amazon (WWF 2016) likely has a direct impact on biodiversity and ecosystem function, threatening evolutionary processes governing the origin and maintenance of species diversity in all other South and Central American regions. A strong regional network of biological collections combined

with long term monitoring of Amazonian populations, such as those conducted by the RAINFOR network (<https://rainfor.org/en/>), ForestGeo (<https://forestgeo.si.edu/>), LTER (<https://lternet.edu/>), and PPBio programs (PPBio 2005), are urgently needed to improve our understanding of Amazonian biodiversity, ecology, evolution, biogeography, and demography (Stouffer *et al.* 2021).

Conservation efforts in the Amazon must take into account the unique ecological properties and evolutionary processes of its constituent biotas. Organismal habits and behaviors are one important example. The annual migrations of fishes (*piracema*), birds, and insects, as well as tree fruiting blooms, all constitute important biotic resources for human agroecosystems and other natural Amazonian ecosystems (Goulding *et al.* 2019). These behaviors are the basis for important ecological phenomena and annual life cycles, including mast flowering, phenological patterns, reproductive booms, and natural flood regimes (Alho 2020; Cunha-Machado *et al.* 2021). Such phenomena need to be considered in regional planning and during rainforest conservation efforts. The establishment of river impoundments, for instance, interrupt natural flood regimes and disrupt migration corridors that are critical for the survival of Amazonian freshwater organisms (Winemiller *et al.* 2016; Latrubesse *et al.* 2017; Barthem *et al.* 2017; Albert *et al.* 2020b).

CONCLUSIONS

Amazonian biodiversity is immense and vastly underestimated

Amazonian biodiversity is among the highest on Earth and constitutes the core of the Neotropical realm. The Amazon basin encompasses a vast range of life forms, genetic resources, and ecological functions, including hydrological cycles

that impact climate at continental and global scales. This bewildering biodiversity arose from evolutionary diversification over highly heterogeneous landscapes and lengthy time periods in which rates of speciation exceeded those of extinction. A comprehensive understanding of Amazonian biodiversity is fundamental for data-driven conservation and management plans for these crucial ecosystems. Yet Amazonian biodiversity remains poorly known, due to high numbers of cryptic species, logistic difficulties of samplings, and insufficient human and infrastructure resources for assessments. Critical information is lacking in all major biodiversity dimensions: taxonomic diversity (Linnaean shortfall), biogeographic distributions (Wallacean shortfall), species abundances (Prestonian shortfall), phylogenetic diversity (Darwinian shortfall), species traits (Raunkiaeran shortfall), and species interactions (Eltonian shortfall). Resolutions to all these information shortfalls will require increased investments in both governmental and non-governmental organizations, in particular in universities, research institutes, and natural history museums, within Amazonia and elsewhere, to leverage existing information and resources to fill existing gaps. Resources must also be invested in large-scale metagenomic screenings to quantify the known and document the as-yet unknown aspects of biodiversity, especially targeting poorly-studied but ecologically important life forms (e.g. bacteria, microorganisms, fungi, meiofauna) that drive ecosystem-level biochemical processes in Amazonian soils and waters.

Macroevolutionary history of Amazonian terrestrial and aquatic ecosystems

Tropical rainforests composed primarily of flowering plants originated on the supercontinent of Western Gondwana during the Early Cretaceous (ca. 145 – 100 Ma), and most modern tropical plant and animal groups diversified in the super-greenhouse world of the Late Cretaceous (ca. 100 – 66) and Early Paleogene (ca. 66 – 30 Ma). Two landmark events during this time were the final separation of South America from Africa (ca. 100 Ma) that isolated the biotas of these continental blocks, and the Chicxulub asteroid impact (ca. 66 Ma) that caused the End-Cretaceous global mass extinction, and which spurred rapid Early Paleogene (ca. 66–55 Ma) diversification of most modern terrestrial and aquatic taxa.

Global cooling starting in the Early Oligocene (ca. 30 Ma) resulted in extensive vegetational changes throughout South America, with tropical rainforests contracting to lower latitudes and grasslands expanding at temperate latitudes. These vegetation changes divided the formerly continuous moist Neotropical rainforest into disjunct Amazon and Atlantic rainforests. Climatic warming and tectonic processes during the Miocene (ca. 23 – 4.5 Ma) resulted in the formation of enormous mega-wetland systems in western Amazonia. Late Miocene (ca. 10 Ma) uplift of the northern Andes reorganized river drainage patterns across northern South America, driving

the formation of the modern trans-continental Amazon river, as well as the courses of the modern Orinoco and Magdalena rivers, among many others.

Late Miocene to Pliocene (ca. 12–3.5 Ma) uplift of the Panama Isthmus allowed reciprocal dispersal between the biotas of North and South America, an event known as the Great American Biotic Interchange (GABI). The interchange was approximately symmetrical at first, with about equal numbers of taxa moving north and south, however the subsequent survival was highly asymmetrical, with mammals of North American origin surviving better than those from South America. Global climate oscillations during the Pleistocene (ca. 2.5 – 0.1), combined with human hunting, are associated with the extinctions of megafaunas worldwide. The extinction of large-bodied mammals in South America was more severe than elsewhere, where 83% of the megafauna (adult body weight > 44 kg) genera were lost.

Assembly of the megadiverse Amazonian biota

Amazonia's outstanding biodiversity was assembled over a period of tens of millions of years, through a unique history of geological, climatic, and biological factors, all operating over partially-overlapping time scales. Geological and climatic factors operating over evolutionary time scales (thousands to millions of years) constrained the landscape and riverscape processes that generated heterogeneous soil and water chemistry profiles and other factors, which in turn affected the geographic, demographic, and genetic connections among populations. Through their controls on organismal dispersal, these abiotic factors strongly affected rates of adaptation, speciation and extinction. Lowland Amazonian landscapes and riverscapes are highly dynamic over time periods ranging from tens to hundreds of thousands of years, under the perennial influence of river capture over the broad low-relief topography, and of climate fluctuations over the course of the Plio-Pleistocene (ca. 5.3–0.01 Ma).

Biodiversity itself also contributes to elevated Amazonian species richness, through autocatalytic feedback mechanisms within hyperdiverse Amazonian ecosystems. Functionally and structurally diverse biotas provide more complex and multifarious environmental substrates that facilitate the evolution of physiological and behavioral specializations that may promote ecological coexistence and, in some cases, genetic isolation and speciation. Abiotic factors deriving directly from geographic space, climatic and elevation gradients, topographic relief, hydrology, and sediment and water chemistry, all serve to facilitate organismal diversification into major habitat types. Intertwined with these landscape processes are biotic processes that allow species to coexist within the same habitats and thereby lower their extinction risks. These ecological interactions include competition, predation, parasitism, mutualism, and cooperation, and the many ways in which organisms modify their environment.

Such activities, known as niche construction, include the building of nests and burrows by animals, the formation of vegetation structure and shade by plants, and nutrient and energy cycling in soils and waterways by plants, fungi, and microbes. These biotic interactions lead to the evolution of new traits, increase the structural heterogeneity and functional dimensions of habitats, and enhance the genetic and phenotypic diversity of Amazonian ecosystems.

The human footprint on Amazonia

Human activities have greatly impacted Amazonian biodiversity for at least 20 Ka. The main effects by indigenous peoples were in plant domestication and agricultural practices that altered local vegetation structure and species abundance, and hunting practices that, combined with climatic changes, produced the extinction of pre-Pleistocene megafauna. Changes to the Amazonian ecosystems accelerated with Portuguese and Spanish colonization in the past 500 years, and accelerated again with the transition to modern socio-economic activities during the past 40 years. During this time, the Amazon has undergone profound demographic and ecological transformations, in which the original indigenous tribal populations suffered tremendous population crashes because of new diseases, displacement, and hostility. The modern Amazon basin is now home to about 35 million people, including about 400 indigenous and traditional communities, but also a large mestizo population concentrated in urban and rural areas. Rapid changes in land-use and other human activities (logging, mining, hunting, fishing, dams, roads) are profoundly affecting the species richness and evolutionary processes of the Amazon basin, by altering the distribution, abundance, connectivity, and ecology of species.

Conservation of ecological and evolutionary processes

The exceptional Amazonian biota accumulated over the course of millions of years by the action of numerous ecological and evolutionary processes that promoted ecological coexistence, facilitated both dispersal and genetic isolation, and ultimately resulted in a biota in which rates of speciation exceeded those of extinction. However, population sizes of many Amazonian species have been falling rapidly in recent years, due to human activities, imperiling many species and degrading the forest biome as a whole. The most effective conservation efforts prioritize regions characterized by: (1) high lineage and functional trait diversity (e.g., mature or “old growth” rainforests); (2) a high proportion of geographically restricted species (e.g. tepuí table-top mountains and other uplands areas of the Guiana and Brazilian Shields); (3) environmentally extreme habitats (e.g., white sand forests, acidic blackwater rivers) where species exhibit distinctive and specialized physiological and genetic traits; and (4), ongoing and rapid environmental change and lineage diversification

(e.g. cloud forests of the Andean foothills, Páramos at higher elevations).

The most effective conservation strategies are both dynamic and pluralistic, balancing the irreplaceability, representativeness, and vulnerability of species and ecosystems (Jézéquel *et al.* 2020). These strategies prioritize species with ecological and evolutionary resilience, with the goal to preserve lineages with a greater potential to resist and respond to ongoing and future climatic and ecological changes. Effective conservation planning seeks to maintain population connectivity, dispersal and gene flow, which facilitate ongoing evolutionary and ecological processes. Special attention and resources are required in areas of rapid economic and infrastructure development (e.g. road and dam construction), or where major anthropogenic habitat changes have fragmented natural populations via deforestation and defaunation for agriculture, hunting and mining.

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