bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

TITLS: The flickering connectivity system of the north Andean páramos

SHORT RUNNING TITLE: Andean flickering connectivity since 1 Ma

AUTHORS

Suzette G.A. FLANTUA[1,2*], Aaron O'DEA[3], Renske E. ONSTEIN[4], Henry HOOGHIEMSTRA[1]

[1] Institute for Biodiversity and Ecosystem Dynamics (IBED), University of Amsterdam, Amsterdam, The Netherlands.

[2] Department of Biological Sciences, University of Bergen, Bergen, Norway

[3] Smithsonian Tropical Research Institute, Box 0843-03092, Balboa, Republic of Panama

[4] German Centre for Integrative Biodiversity Research (iDiv), Halle-Jena-Leipzich, Deutscher Platz 5e, 04103 Leipzig, Germany.

* Corresponding author: s.g.a.flantua@gmail.com. ORCID: https://orcid.org/0000-0001-6526-3037

ACKNOWLEDGEMENTS

This work was part of SGAF's doctoral thesis funded by Netherlands Organization for Scientific Research (NWO, grant 2012/13248/ALW to HH.). The Hugo de Vries foundation is acknowledged for financially supporting multiple grant proposals during the project including the development of the visualization accompanying this paper. The Sistema Nacional de Investigadores (SNI) de SENACYT supported AO. REO acknowledges the support of the German Centre for Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig funded by the Deutsche Forschungsgemeinschaft (DFG, German Research Foundation)—FZT 118. Carina Hoorn and Daniel Kissling are thanked for the educational environment shaped by the parallel paper on mountain diversity (Antonelli et al., 2018). Colin Hughes is thanked for comments on previous versions of this paper. We thank Mauricio Bermúdez for help with the geological delimitation of the Northern Andes and Francisco Cuesta for facilitating the map by Josse et al. (2009).

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

ABSTRACT AND KEYWORDS

Aim To quantify the effect of Pleistocene climate fluctuations on habitat connectivity across páramos in the Neotropics.

Location The Northern Andes

Methods The unique páramos habitat underwent dynamic shifts in elevation in response to changing climate conditions during the Pleistocene. The lower boundary of the páramos is defined by the upper forest line, which is known to be highly responsive to temperature. Here we reconstruct the extent and connectivity of páramos over the last 1 million years (Myr) by reconstructing the UFL from the long fossil pollen record of Funza09, Colombia, and applying it to spatial mapping on modern topographies across the Northern Andes for 752 time slices. Data provide an estimate of how often and for how long different elevations were occupied by páramos and estimates their connectivity to provide insights into the role of topography in biogeographic patterns of páramos.

Results Our findings show that connectivity amongst páramos of the Northern Andes was highly dynamic, both within and across mountain ranges. Connectivity amongst páramos peaked during extreme glacial periods but intermediate cool stadials and mild interstadials dominated the climate system. These variable degrees of connectivity through time result in what we term the 'flickering connectivity system'. We provide a visualization (video) to showcase this phenomenon. Patterns of connectivity in the Northern Andes contradict patterns observed in other mountain ranges of differing topographies.

Main conclusions Pleistocene climate change was the driver of significant elevational and spatial shifts in páramos causing dynamic changes in habitat connectivity across and within all mountain ranges. Some generalities emerge, including the fact that connectivity was greatest during the most ephemeral of times. However, the timing, duration and degree of connectivity varied substantially among mountain ranges depending on their topographic configuration. The flickering connectivity system of the páramos uncovers the dynamic settings in which evolutionary radiations shaped the most diverse alpine biome on Earth.

Keywords (6-10): alpine biome, evolutionary radiations, flickering connectivity system, fossil pollen, mountain fingerprint, Neotropical biodiversity, páramos, paleotemperature, past habitat connectivity, species pump

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

MAIN TEXT

1. INTRODUCTION

Mountains are regarded as powerhouses of biodiversity in the world (Barthlott, Rafiqpoor, Kier, & Kreft, 2005; Kreft & Jetz, 2007; Antonelli et al., 2018) and harbour numerous examples of very rapid and recent species diversifications ('radiations'; Hughes & Atchison, 2015). It is thought that a large part of this diversity arose geologically recently, during the Plio-Pleistocene (last 5.3 million years, [Ma]), but there is no consensus on the drivers of these radiations. One favoured hypothesis is that the combination of high topographic relief and Plio-Pleistocene climatic oscillations led to rapidly changing distributions of montane species, which generated new lineages (e.g. Qian & Ricklefs, 2000; Graham et al., 2014; Mutke, Jacobs, Meyers, Henning, & Weigend, 2014). However, the relative contributions of isolation (e.g. Schönswetter, Stehlik, Holderegger, & Tribsch, 2005; Wallis, Waters, Upton, & Craw, 2016; Weir, Haddrath, Robertson, Colbourne, & Baker, 2016) vs. gene flow and dispersal (e.g. Smith et al., 2014; Cadena, Pedraza, & Brumfield, 2016; Kolář, Dušková, & Sklenář, 2016; Knowles & Massatti, 2017) in driving fast diversification rates (i.e. the 'species-pump' effect, Rull, 2005; Rull & Nogué, 2007; Winkworth, Wagstaff, Glenny, & Lockhart, 2005; Ramírez-Barahona & Eguiarte, 2013; Steinbauer et al., 2016; Flantua & Hooghiemstra, 2018) are still debated. It is likely that these radiations have been the results of the interchange between phases of isolation, causing allopatric, *in situ* speciation, and connectivity, triggering diversification through dispersal and settlement in new areas and hybridization of differentiated taxa from previously isolated populations (Flantua & Hooghiemstra, 2018). The fastest and most spectacular radiations may therefore occur in mountain regions with variable degrees of past connectivity and isolation during climate fluctuations, which, complex in space and time, are inherently related to the mountain topography (Flantua & Hooghiemstra, 2018). It is therefore critical to quantify connectivity of montane habitats using our understanding of topography and past climate fluctuations (**Fig. 1**).

The Northern Andes is an ideal model system to quantify connectivity, due to the large variation in topography and the advanced paleoecological knowledge on Plio-Pleistocene climate fluctuations derived during the last five decades (Hooghiemstra & Flantua, 2019). The Northern Andes is topographically-rich with high elevations, steep ridges and valleys (see illustrations by Von Humboldt during his trips in Latin America, 1773-1858), composed of several mountain ranges, some of which are parallel running from North to South. The area hosts the treeless tundra-like alpine biome, the páramos, regarded the richest alpine flora in the world in terms of endemism and species richness (Sklenář, Hedberg, & Cleef, 2014) and is known for its bursts of Plio-Pleistocene species diversification amongst plants (see overview in

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Hughes & Atchison, 2015). In terms of quantifying Plio-Pleistocene temperature fluctuations, the palaeoecological history of the páramos has been studied extensively (e.g. Van der Hammen, 1974; Cleef, 1979; Hooghiemstra, 1984; Hooghiemstra & Van der Hammen, 2004) because of the unique high elevation fossil pollen records that cover most of the Pleistocene (Groot et al., 2011; Groot, Hooghiemstra, Berrio, & Giraldo, 2013; Bogotá-Angel et al., 2011; Bogotá-A., Hooghiemstra, & Berrio, 2016; Torres, Hooghiemstra, Lourens, & Tzedakis, 2013). Under current conditions, the páramos form isolated archipelagos of alpine (sky) islands (McCormack, Huang, & Knowles, 2009) but the rich collection of fossil pollen sequences throughout this region (Flantua et al., 2015) show that the páramos underwent substantial elevational shifts during the Pleistocene, resulting in extensive changes in surface area and connectivity (Van der Hammen, 1974; Hooghiemstra & Van der Hammen, 2004; Flantua et al., 2014; Sklenář et al., 2014). Thus, the topographic diversity and the robust catalogue of palaeoecological reconstructions make the Northern Andes a highly suitable model region to explore patterns of connectivity in mountain biomes in response to Pleistocene climate fluctuations.

In this study, we aim to quantify the biogeographic changes of the páramos in terms of spatial scale and connectivity based on modern topography and pollen-based records of past climate change. Specifically, we developed a novel tool to explore the complex temporal and spatial patterns of páramo connectivity. We constrain our model by using the last 1 Myr of the high-resolution fossil pollen record of Funza09, a 586 m deep core taken from the Bogotá basin of Colombia (Torres et al., 2013). Available surface area (Elsen & Tingley, 2015) and connectivity (Flantua et al., 2014; Bertuzzo et al., 2016) is variable along elevational gradients of mountains. We therefore hypothesize that the different mountain ranges that compose the Northern Andes display variable patterns of past páramo connectivity dependent upon their topography (**Fig. 1**). We discuss the implications of our outcomes for evolutionary processes and how defining and quantifying past connectivity in mountain systems is essential to help reveal mechanisms of ecological, biogeographical and evolutionary change. Ultimately, our quantification of páramo connectivity through space and time provides a unique opportunity to disentangle some of the mechanistic drivers ('modulators') of radiations in this biome (Bouchenak-Khelladi, Onstein, Xing, Schwery, & Linder, 2015).

2. METHODS

2.1 Geographical features

The Northern Andes (ca. 448.000 km^2) covers parts of Venezuela, Colombia and Ecuador (**Fig. 2a**), and

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

can be partitioned into six principal mountain ranges or 'cordilleras' (**Fig. 2c**), namely the Sierra Nevada de Santa Marta (SNSM), Cordillera de Mérida, Eastern, Central and Western Cordillera and the Ecuadorian Cordilleras. Most of the Northern Andes is considered a highly to extremely high rugged landscape (**Fig. 2b**; See mountain illustrations by Von Humboldt (1845)where the high peaks and deep inter-Andean valleys cause strong contrasts in climate throughout the region (Flantua et al., 2016). Surface area in mountains does not decrease monotonically with elevation as has been shown previously in southern Colombia by Flantua et al. (2014) and on a global scale by Elsen & Tingley (2015). The Northern Andes shows a decrease of surface area going upslope where there is a slight peak around 900-1200 m asl but then continues to decrease up to 6260 m asl (**Fig. 2d**), following a typical 'pyramid shape'. However, the different cordilleras show different patterns of elevational surface area (**Fig.2d**) where the Eastern Cordillera shows a sharp peak around 2600 m asl and the Ecuadorian Cordillera shows high values of surface area at much higher elevations than the other cordilleras (**for more details see Table S1.1, Appendix S1 in Supporting Information**). Of all tropical alpine floras, such as in East Africa and New Guinea, the páramos are home to the highest species richness and endemism (Luteyn, 1999; Sklenář, Dušková, & Balslev, 2011), with low between-mountain similarity in species; (Sklenář et al., 2014). The páramos today are spread out over the Northern Andes as an archipelago of small and highly fragmented páramo complexes (**Figure S2.1**, **Appendix S2**).

2.2 Quantifying temperature and upper forest line based on fossil pollen data

To quantify temperature fluctuations during the Pleistocene (and consequently estimate páramo connectivity), we used fossil pollen data from the Northern Andes. The composite pollen record Funza09 (4.83°N, 75.2°W; 2550 m asl, **Fig. S2.1. Red star**) reveals vegetation and climate dynamics over the past 2.25 Myr (Torres et al., 2013). We reconstructed the páramos' elevational fluctuations, and consequently páramo connectivity, by estimating the upper forest line (the transition from the upper montane forest to the páramos; UFL) from the Funza09 record. Though this record covers the last 2.2 Myr, we only used the last 1 Myr as this interval reflects continuous lake conditions in comparison with variable hydrological conditions between 2.2-1 Ma which obscure the quantification of changes to the UFL. We follow the methodology described and implemented by Hooghiemstra (1984), Groot et al. (2011), and Hooghiemstra et al. (2012) to derive the Andean UFL and paleotemperature curve (for detailed methodology on the UFL reconstruction see **Appendix S3**)

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

2.3 Calculations of connectivity per páramo "island"

To calculate the degree of connectivity between páramos, we used a graph-based habitat availability index called probability of connectivity (PC) metric. This metric takes into account the area of the páramo "island" itself and the distances to other islands where a user-defined distance threshold defines the 'reachability' of other islands (Saura & Pascual-Hortal, 2007; Saura, Estreguil, Mouton, & Rodríguez-Freire, 2011), even if they are not physically connected (i.e. 'structural connectivity', Tischendorf & Fahrig, 2000). The metric assigns a value to each páramo island representing its contribution in maintaining the overall connectivity of the páramo biome (Saura & Pascual-Hortal, 2007; Saura et al., 2011). The total PC is built up in three 'fractions', namely the 'intrapatch', the 'flux', and the 'connector' fractions (Saura & Rubio, 2010). The first fraction focusses on the available surface area and habitat quality (if applicable) within the individual island. The second fraction assesses how well the individual island is connected to other islands given additional importance to the other islands' attributes (surface and quality) and its strategic position to other páramo islands. The third fraction quantifies the contribution of the island to maintain connectivity between the rest of the islands, in other words its role as an intermediate stepping stone between non-adjacent islands. Additionally, we calculated the equivalent connected area (ECA), which is derived directly from the PC, as a measure of the overall connectivity of a region (Saura et al., 2011). Conefor Sensinode 2.2 software and ESRI ArcGIS 10.3 were used to calculate the straight-line distances between islands, the PC and ECA (Saura & Pascual-Hortal, 2007; Saura & Torné, 2009). We calculated connectivity for the entire Northern Andes and for each mountain range separately.

2.4 Calculations of corridors between páramo islands

We identified corridors between páramo islands within and between cordilleras under different climatic conditions. We used the Gnarly Landscape Utilities (V0.1.3; McRae, Shirk, & Platt, 2013) with ESRI ArcGIS 10.3 to create a raster grid of 'landscape resistance' based on ruggedness (**Fig. 2b**) and habitat suitability. We assumed an increased landscape resistance with increased ruggedness, assigning values between 0 (no resistance) to 100 (maximum resistance) using an equal interval classification. For the habitat suitability map, we started by assigning a "perfectly suitable" score of 100 to each páramo island, while outside the island the score of 0 reflects maximum unsuitability. To soften this boundary, an exponential decay function was then used by increasing resistance in 5 elevational steps of 100 m where we assigned a suitability score of 40 to the boundary of the páramo. As a result of the decay function the highest suitability of páramo - its core area - was restrained 200 m above the UFL and 200 m below the

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

snowline.

We used Linkage mapper to calculate the least-cost pathways, or corridors, based on the produced raster grid of landscape resistance (McRae & Kavanagh, 2011). These corridors are expressed as 'conductance maps' that represent gradients of cumulative corridors. Where the densities of corridors is highest, it is assumed that there is a high probability of dispersal and migration possible between islands (McRae, Dickson, Keitt, & Shah, 2008). The full landscape of the Northern Andes is considered an area where corridors could exist, with exception of the region between SNSM and the Sierra de Perijá (**Fig. S2.1**).

We resampled the 30 m Digital Elevation Model (DEM, **Fig. 2**) to a 1 km resolution to reduce computing time for each Linkage mapper down to on average 2 hours. We allowed Linkage mapper to create corridors through (instead of only between) core areas to represent the full arsenal of connectivity through the landscape. Only corridors between páramo islands larger than 1 km^2 were considered at any given moment in time. From the final output maps, only values lower than 200k conductance (default threshold) are selected to highlight the strongest corridors. The outputs were weighted according to the percentage of time they occurred during the last 1 Myr.

3. RESULTS

3.1 A million years of temperature fluctuations

Temperatures at Funza (2550 masl) are estimated to have fluctuated between ca. 15 and 6°C causing an estimated maximum 1600 m elevational shift of the UFL between ca. 3500 and ca. 1900 m asl (**Fig. 3**). The Pleistocene glacial-interglacial dynamics were not replicated cycles of temperature change showing repeated patterns of high and lows, but display a high temporal variability between each glacial-interglacial cycle. Conditions similar to the current warm, interglacial conditions occurred several times during the last 1 Myr and accounted for around a quarter of the time. Extreme cool glacial conditions, ~ 6 - 8°C cooler than today, were relatively rare, occurring less than 10 percent of the time. On the whole, intermediate cool stadials and mild interstadials dominated the last 1 Myr, occurring over two thirds of the time.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

3.2 Calculations of páramo connectivity

Our estimations on the spatial and elevational extent of ancient páramos and their connectedness at different times in the past reveals that páramos underwent frequent spatial alterations between fragmented and connected spatial configurations, but the exact patterns were highly dependent on mountain chain topography (**Fig. 4a,b. See Appendices 4 and 5**). The páramos in the Ecuadorian Cordillera generally maintained a high degree of connectivity over the last 1 Myr, rarely enduring severe fragmentation. Fragmentation did however occur when the snowline plunged significantly during colder and wetter glacial periods, causing a break up of páramo areas on lateral flanks of the mountains. Likewise, the level of connectivity between páramos on the Central Cordillera fragmented substantially through a descending snowline, breaking the upper elevation limit of páramo connectivity. In contrast, the Eastern Cordillera shifted substantially between periods of connectivity and fragmentation, always, however, maintaining two large páramo islands surrounded by smaller 'satellite islands'. Páramos in the Cordillera de Mérida seem to have been restricted during interglacials to one core area only, while during colder periods a relatively high fragmentation is observed possibly due to glaciers pushing páramos to lateral distributions. Here connectivity increased mainly towards the southwest and during colder periods (UFL $\leq$ 2300 m asl). The páramos of the SNSM and the Western Cordillera endured the highest degree of rates of change in fragmentation of all ranges. In the latter, páramo habitats are estimated to have often completely disappeared. In contrast, páramos of the Central Cordillera maintained a long latitudinal distribution, forming a chain of isolated populations in small patches that on the whole remained connected. Even in very cold conditions, no continuous connectivity of core areas seems to have been possible between the Eastern Cordillera and Cordillera de Mérida, or the region of Sierra de Perijá. Towards the south of the Eastern Cordillera a low-elevation barrier was possibly crossed at 1900 m asl forming a brief bridge suitable for páramo habitat into the Macizo Colombiano of the Central Cordillera.

The reconstruction of putative corridors shows a complex spatial pattern through the mountainous landscapes of the Northern Andes (**Fig. 4c,d**). The long ridge of the Central Cordillera forms the starting point of numerous corridors to the páramos in the Western Cordillera. The Eastern Cordillera shows a complex internal pattern of corridors, where there are neither strong corridors towards Sierra de Perijá in the North, nor towards the Cordillera de Mérida, while a high concentration of corridors is found between the large páramos complexes in the Eastern Cordillera (Páramos of Boyacá and Cundinamarca, **Fig. 1**). In the Ecuadorian Cordillera a more lateral pattern of high/low potential corridors is observed following the intra-Andean valleys and peaks within this mountain range. Corridors to the southernmost páramos of Ecuador as also the northernmost páramos of the Western Cordillera are weak and occurred infrequent during the last million years, shown by the thin lines.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

3.3 Flickering connectivity systems

Páramo connectivity through time shows a highly variable pattern (**Fig. 5.a**) introduced by Flantua & Hooghiemstra (2018) as a flickering connectivity system (**see visualization in Appendix S6**). We find support for the hypothesis that this system with fluctuating, highly variable connectivity in spatial and temporal dimension is unique for each mountain range of the Northern Andes (**Fig. 1**). For example, changes in connectivity within the Ecuador Cordillera are substantial but the system 'flickers' around a high average when compared to other mountain ranges. The flickering connectivity systems within the Eastern and Central Cordillera are surprisingly similar, though the peaks of connectivity during glacial periods and the dips of connectivity during interglacials are more extreme in the former (**Fig. 5a**). The Western Cordillera is a larger mountain range than the Cordillera of Mérida and the SNSM (**Table S.1**), and its variation of connectivity has been correspondingly larger (**Fig. 5b**) but with the lowest occurrence of connectivity compared to the other mountain ranges (**Fig. 5a**). Considering only the frequency in the distribution of data (**Fig. 5b**), the Ecuadorian Cordillera and the SNSM stand out for their relatively small within-mountain range variation in connectivity, compared to the Eastern and Central Cordillera (similar patterns) and the Western Cordillera.

When frequencies of connectivity are weighted by the amount of time that connectivity lasted two main patterns emerge (**Fig. 5c**). The first is shared by the Western, Central, and Eastern Cordilleras, which all display an elongated pattern where the highest values are around a centroid, resembling a 'humming top' or, as Elsen & Tingley (2015) recognized in mountain hypsographies, a 'diamond' shape. Ecuadorian Cordilleras, Cordillera de Mérida and SNSM instead reveal a different pattern with a narrower centroid that widens towards the upper and lower section, resembling an 'hourglass' shape. Here, the Ecuadorian Cordillera and SNSM show a surprising similarity though at different connectivity ranges. The Central and Eastern Cordilleras are strikingly similar overall.

4. DISCUSSION

4.1 Variable degrees of past connectivity of different mountain ranges

Although currently isolated, evolutionary radiations and the assembly of the páramo ecosystem formed during times when the páramos were flickering in and out of connectivity (**Fig. 5b**.) The concept of 'mountain fingerprints' (Flantua & Hooghiemstra 2018) proposes that the region's complex topography would have meant that páramos in different mountain regions would have fragmented and connected at

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

different periods of time and with different rates and frequencies (as summarized in **Fig. 1**). This means that in some mountain ranges the páramos are a mix of somewhat even occurrence of connectivity and fragmentation events through time (**Fig.1b**, here representative of the Eastern Cordillera), or could have been dominantly fragmented (**Fig.1a**, e.g. Western Cordillera), or more connected (**Fig. 1c**, e.g. Ecuadorian Cordilleras). These regional differences in the temporal and spatial variation in past páramo connectivity (**Figs 4** and **6**) are likely to have resulted not only in regional differences in biogeographical patterns through time, but also varying ecological and evolutionary processes. We therefore propose that the data we present can be used to test hypotheses of the drivers of species richness, endemism and degrees of Pleistocene diversification in the Northern Andes, and likewise are applicable to other mountain regions around the world.

4.2 Evolutionary implications of the flickering connectivity system

The dynamic history of the páramos elucidated by the flickering connectivity system can provide three important insights in terms of evolutionary processes. First of all, the regional differences in past páramo connectivity - the mountain fingerprint – support temporally and spatially discordant phylogeographic patterns (Pennington et al., 2010; Massatti & Knowles, 2014; Papadopoulou & Knowles, 2015; 2016). This means that the timing of diversification in the different mountain regions would not be expected to have occurred synchronously, even if all phylogenetic studies on páramo species could overcome current issues in techniques, spatial resolution and time-calibration points (Rull, 2011). Secondly, diversification rates might differ along the elevational gradient and this might be the rule rather than the exception. Elevational differences in surface availability and connectivity are likely to influence at what elevation the strongest phylogeographic processes will occur, and these processes are thus expected to differ between mountain systems resulting in elevational differences of diversification (see for instance Kropf, Kadereit, & Comes, 2003; Lagomarsino, Condamine, Antonelli, Mulch, & Davis, 2016). And thirdly, the flickering connectivity system, which is expected to cause isolation followed by connectivity of populations, is expected to cause pulses of diversification (Knowles, 2000), possibly resulting in series of sub-radiations in the páramos. Where isolation resulted in allopatric, *in situ* speciation, connectivity triggered diversification through dispersal and settlement in new areas ("dispersification", Moore & Donoghue, 2007), and hybridization of differentiated taxa from previously isolated populations (Petit et al., 2003; Grant, 2014). Phylogenetic studies are increasingly supportive of the important role of gene flow, dispersification and hybridization, alongside periods of isolation, in driving (explosive) diversification in mountains (e.g. Knowles & Massatti, 2017; Hazzi, Moreno, Ortiz-Movliav, & Palacio, 2018), as well as in

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

other systems such as tropical rainforests (e.g. Onstein et al., 2017) and islands (e.g. Ali & Aitchison (2014). In the páramos, examples originate from studies on birds (Quintero & Jetz, 2018; Cadena et al., 2016) and plants, such as *Neobartsia* (Uribe-Convers & Tank, 2015), *Lupinus* (Hughes & Eastwood, 2006; Nevado, Contreras-Ortiz, Hughes, & Filatov, 2018; Contreras-Ortiz, Atchison, Hughes, & Madriňán, 2018), *Loricaria* (Kolář et al., 2016), *Espeletia* (Diazgranados, 2012; Diazgranados & Barber, 2017; Pouchon et al., 2018) and *Hypericum* (Nürk, Scheriau, & Madriñán, 2013), supporting the strong relationship between changing degrees of connectivity and radiations (Flantua & Hooghiemstra, 2018). Interestingly, the Funza09 pollen record shows a clear shift in the rhythm of climate change around the mid-Pleistocene transition (ca. 0.9 Ma) after which 100 kyr cycles with high amplitudes started to dominate the climate, overruling the lower-amplitude 41-kyr cycles that continued in the background. Strikingly, changes in speciation rates of high elevation birds (Weir, 2006) and the Espeletiinae in the Cordillera de Mérida (Pouchon et al., 2018), echo the mid-Pleistocene transition by an acceleration of diversification during the last 1 Myr suggesting a close link between the intensity of the flickering connectivity system and radiations.

4.3 Limitations and model assumptions

Inherent to any model in mountains and concerning connectivity are simplifications of the temporal and spatial complexity of the real world. For instance, the UFL is asymmetric on wet and dry mountain slopes (e.g. Cleef, 1981), the current elevation of the UFL shows a range of variation of ca. 200 m (incidentally to 300 m), surface processes have changed topography on a million years time scale (Herman et al., 2013; Antonelli et al., 2018), the elevational temperature gradient (lapse rate) seems higher during glacial conditions than at present (Wille, Hooghiemstra, Behling, van der Borg, & Negret, 2001; Loomis et al., 2017), and the current subdivision of páramo vegetation into a 300 m : 600 m : 200 m interval for shrubpáramo, grasspáramo, and superpáramo, respectively, is subject to change (Van der Hammen, 1981; Hooghiemstra, 1984), potentially related to changing atmospheric pCO_2 levels (Boom, Mora, Cleef, & Hooghiemstra, 2001; Boom, Marchant, Hooghiemstra, & Sinninghe Damsté, 2002). We estimate the potential impact of these factors on the estimated connectivity of little significance in determining the overall patterns observed in the flickering connectivity systems.

Any study concerning connectivity also uses a number of assumptions on the probability of dispersal through the landscape. Here, we used a generalized PC value of 0.5 at 10 km to estimate the degree of connectivity. However, species traits, life histories and dispersal capacities may strongly influence dispersal distances (Onstein et al., 2017), and thus influence the probability of connectivity between

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

populations. Implementing taxon-specific traits when calculating the landscape resistance grid (see Methodology) may thus improve the accuracy of the connectivity estimates. Also, family or taxa specific connectivity maps could uncover why certain plant genera do not show any evolutionary diversification during the Pleistocene, e.g. Distichia (Juncaceae; Colin Hughes, personal comm.) and Arcytophyllum (Rubiaceae; Madriñán, Cortés, & Richardson, 2013). Additionally, *a priori* "hard" barriers can be imposed to emphasize areas where habitat connectivity is unlikely to have occurred (see for instance how we maintained SNSM isolated from the other mountain ranges). Defining these barriers *a priori* is not indispensable, though, as the connectivity analysis hints at strong dispersal restrictions when resistance values of corridors are high and indicative of highly constrained dispersal. In the Northern Andes, this is shown by the multiple single line corridors between the Central and Eastern Cordillera, confirmed by the lack of gene flow between plant populations of these regions (Jabaily & Sytsma, 2013; Diazgranados & Barber, 2017; Contreras-Ortiz et al., 2018). This example illustrates the added value of integrating different lines of evidence (e.g. genetic, fossil, paleoclimate) in a spatial and temporal context to understand the biogeographical patterns observed in phylogeographic studies.

4.4 Future research

Our spatio-temporal estimates of past connectivity lay a foundation for further research on elucidating the causal mechanisms of mountain diversifications. Models of past connectivity (**Figs. 4 and 5**), when combined with phylogeographic data, could help reveal the role of interspecific gene flow and allopatric speciation in driving radiations in the high Andes (Nevado et al., 2018) and contribute to a better understanding of the relative importance of geography vs adaptive radiation that underpin Andean diversifications (Contreras-Ortiz et al., 2018). In such a complex system it may also be useful to pay attention to commonalities. For example, when considering both frequency and duration, our data show that two connectivity patterns emerge (i.e. hourglass vs. non hourglass; **Fig.5.c**). Research could explore if cordilleras with shared connectivity patterns also share phylogenetic histories and contemporary (endemic) species' biogeographies to test for universal mechanisms that have shaped present day alpine biomes. This would be especially useful if used in conjunction with information on the reproductive life histories, growth and dispersal capacities of specific taxa.

Finally, past patterns of connectivity are critical to interpret biogeographical patterns of currently isolated or fragmented systems in a wide variety of terrestrial ecosystems including mountains (Flantua & Hooghiemstra, 2018), islands (e.g. Simpson, 1974; Weigelt, Steinbauer, Cabral, & Kreft, 2016; Norder et al., 2018), fresh water systems (e.g. Dias et al., 2014), rainforests (e.g. Graham, Moritz, & Williams,

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

2006), and grasslands (e.g. Lindborg & Eriksson, 2004; Münzbergová et al., 2013), and marine coastal ecosystems (Hoeksema, 2007) that similarly experienced major spatial changes during rapid sea-level fluctuations over the Pleistocene. The approach developed here, to quantify historical connectivity, can therefore provide a platform for interpreting contemporary biogeographies and past drivers of diversification in a wide array of both marine and terrestrial ecosystems where available space has been altered by climatic fluctuations. We postulate that quantifying flickering connectivity systems will facilitate a much more detailed and much needed quantitative basis to compare diversity patterns across the mountain regions of the world.

5. CONCLUSIONS

We present a pollen-based biogeographical model for the páramos biome spanning the northern Andes (Venezuela, Colombia and Ecuador) over the last 1 Myr. Our models suggest substantial temperature oscillations where extreme temperature lows were ca. 8°C cooler than today, causing a lowering of the UFL of ca. 1600 vertical meters. These extreme cool events were however rare and during glacial periods most of the time cool stadials and interstadials prevailed. Our analysis shows that páramos on all mountain ranges underwent frequent alterations between fragmented and connected configurations, but the estimated degrees and amount of connectivity varied among mountain ranges. Most páramos expanded during glacial periods even though extensive glaciers were present. To a large extent the current páramo distribution was replaced by glaciers, implicating a substantial range size change of populations and a highly dynamic system during Pleistocene times.

In light of Von Humboldt's work of relevance of different topographies for mountain biota, we show that topography and climate change together dictated páramo connectivity through time with high spatial variability. The interplay of the topographic and paleoclimatic conditions created a unique pattern of connecting and fragmenting páramo patches through time, here described as the flickering connectivity system. Our spatially explicit model quantifies the complexity of mountain biome dynamics during climate oscillations, in terms of the degree, frequency and duration of past connectivity of alpine biome mountains (**Figs 4 and 5**) and can be applied to other mountain regions. Our connectivity estimates can contribute to answering long-standing questions on the drivers of evolutionary diversification in phylogenetic and phylogeographic studies, and enrich our understanding of the biogeographic history of mountain ecosystems more generally.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

"There the different climates are ranged the one above the other, stage by stage, like the vegetable zones, whose succession they limit; and there the observer may readily trace the laws that regulate the diminution of heat, as they stand indelibly inscribed on the rocky walls and abrupt declivities of the Cordilleras" (Von Humboldt, 1877 (1845), I, p 46).

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

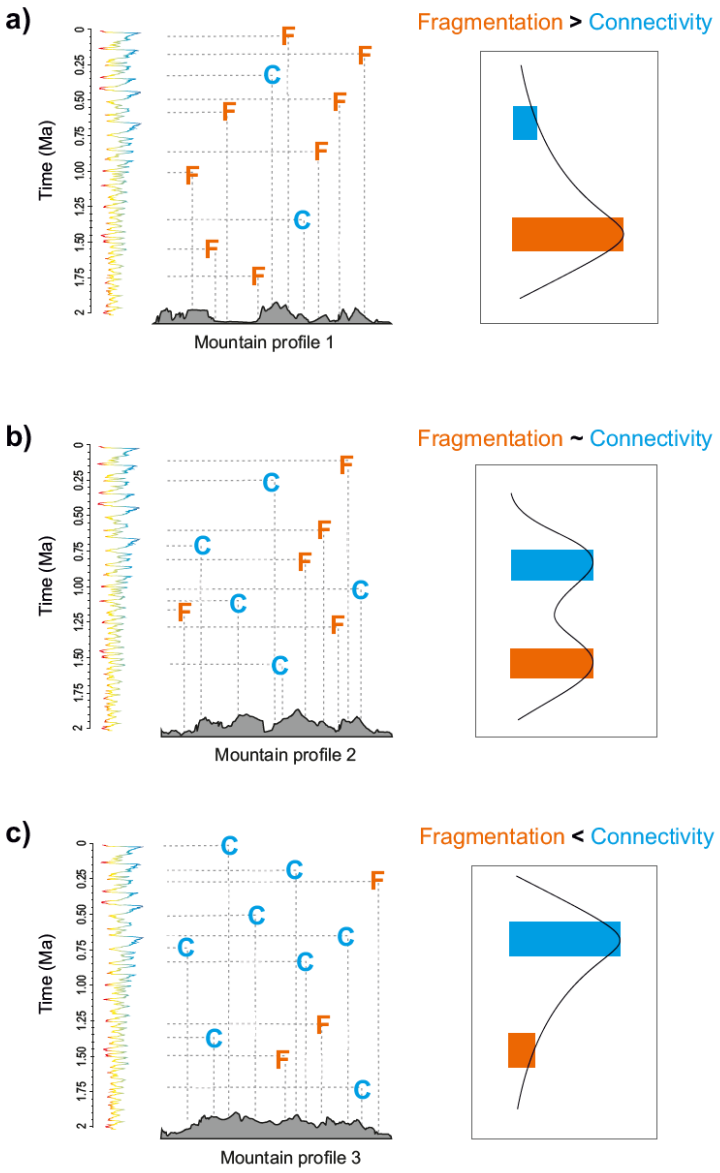

Figure 1 | Connectivity and fragmentation in a mountain landscape. Connectivity (blue) and fragmentation (orange) events ocurred in a spatially and temporally variable manner. This complex pattern in space (latitude, longitude, elevation) and time resemblances a multi-dimensional "mountain fingerprint" which is unique for each mountain range (Flantua & Hooghiemstra, 2018). Three hypothetical mountain profiles are shown with elevational shifts in vegetation distribution driven by climate change (pollen-based record at the left indicating temperature). We recognise mountains with where a) only few events of connectivity occurred during the Pleistocene compared to fragmentation events ('fragmentation-prone mountain fingerprint'), connectivity events interchanged with isolation events in an evenly manner ('mixed connectivity-fragmentation mountain fingerprint'), c) connectivity is facilitated and occured more often than fragmentation events ('connectivity-prone mountain fingerprint'). The right panel is only based on frequency, not the duration of each event.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

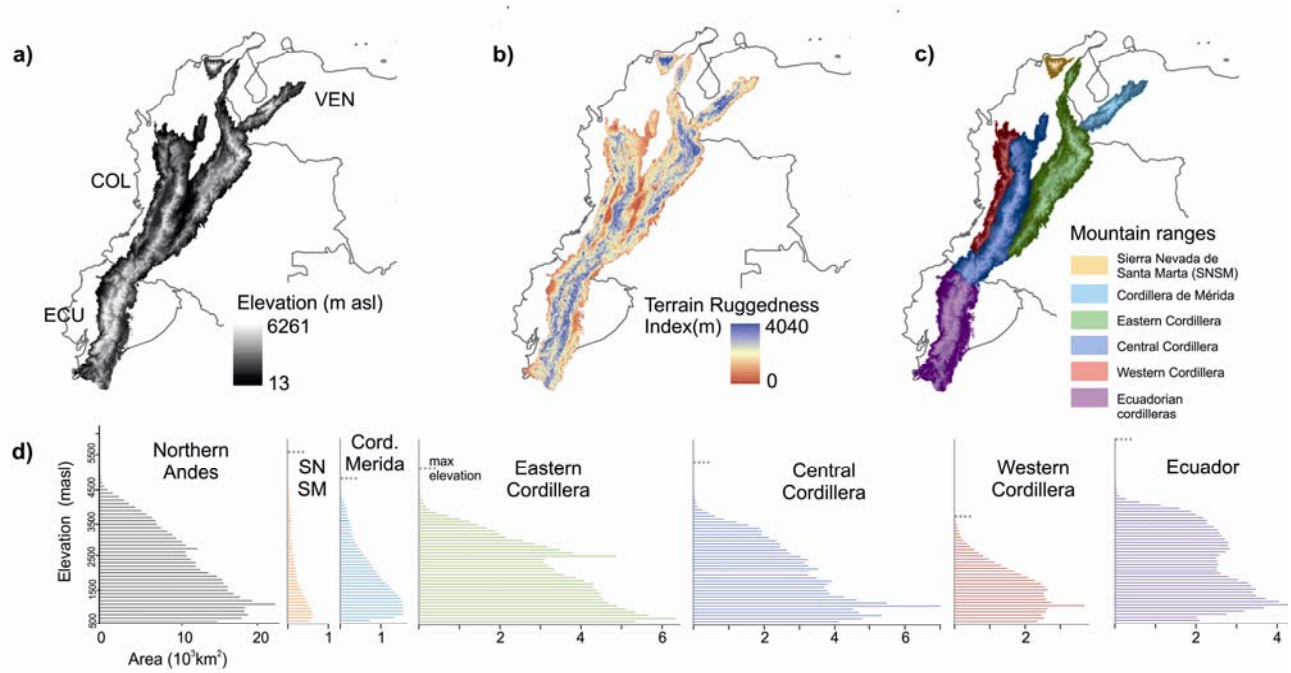

Figure 2 | Hypsographic curves of the Northern Andes. a) Elevation (m asl). b) Terrain ruggedness index calculates the sum change in elevation between a grid cell and its eight neighbor grid cells (Riley et al. (1999) using a ca. 30m DEM (NASA STRM Global 1arc second V003). c) Delimitation of mountain ranges. d) Elevational availability of surface area for the Northern Andes and each mountain range separately shown for 100 m bins. Hypsographic curves based on the Shuttle Radar Topography Mission 1-arc second Digital Terrain Elevation Data (~ 30 m resolution; USGS), taking an elevational threshold of 500 m asl as the horizontal reference plane. Maximum elevation per cordillera is indicated. VEN: Venezuela; COL: Colombia; ECU: Ecuador.

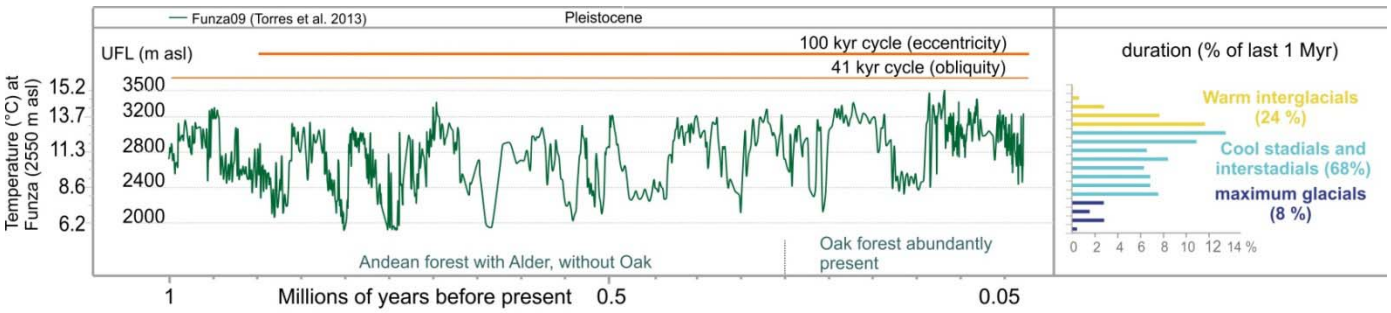

Figure 3 | Upper forest line (UFL) curve of Funza09 (Torres et al. 2013) and reconstructed temperature record covering the last 1 Myr (last ca. 30 kyr BP not included).

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

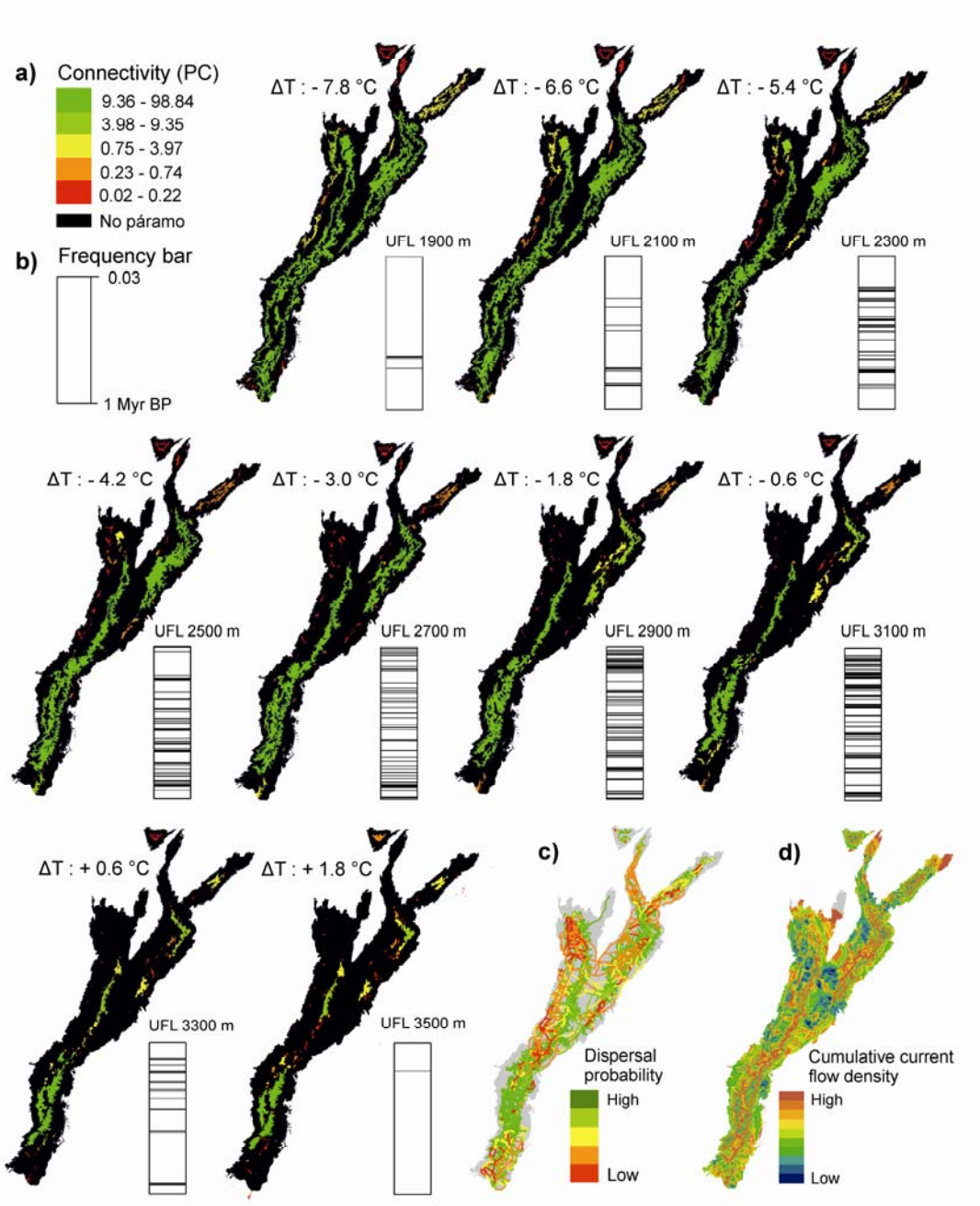

Figure 4 | Páramo connectivity at different upper forest line (UFL) elevations. a) Probability of connectivity metric (PC; distance = 10 km, p = 0.5) (Saura & Rubio, 2010) calculated for all páramos larger than 1 km^2. Maps are plotted with natural-breaks classification. Temperature at 2550 m elevation are relative to the present. b) Frequency bar indicates when the corresponding UFL elevation occurred during the last 1 Myr. Dispersal pathway analysis among páramos using c) Least cost pathways and d) Circuit model expressed in cumulative current flow density (McRae et al., 2008). Areas with high dispersal probability (c) and high current flow (d) indicate frequent and highly probable corridors during the last 1 Myr (weighted by frequency and duration). See Appendices 4 and 5 for all maps and frequencies.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

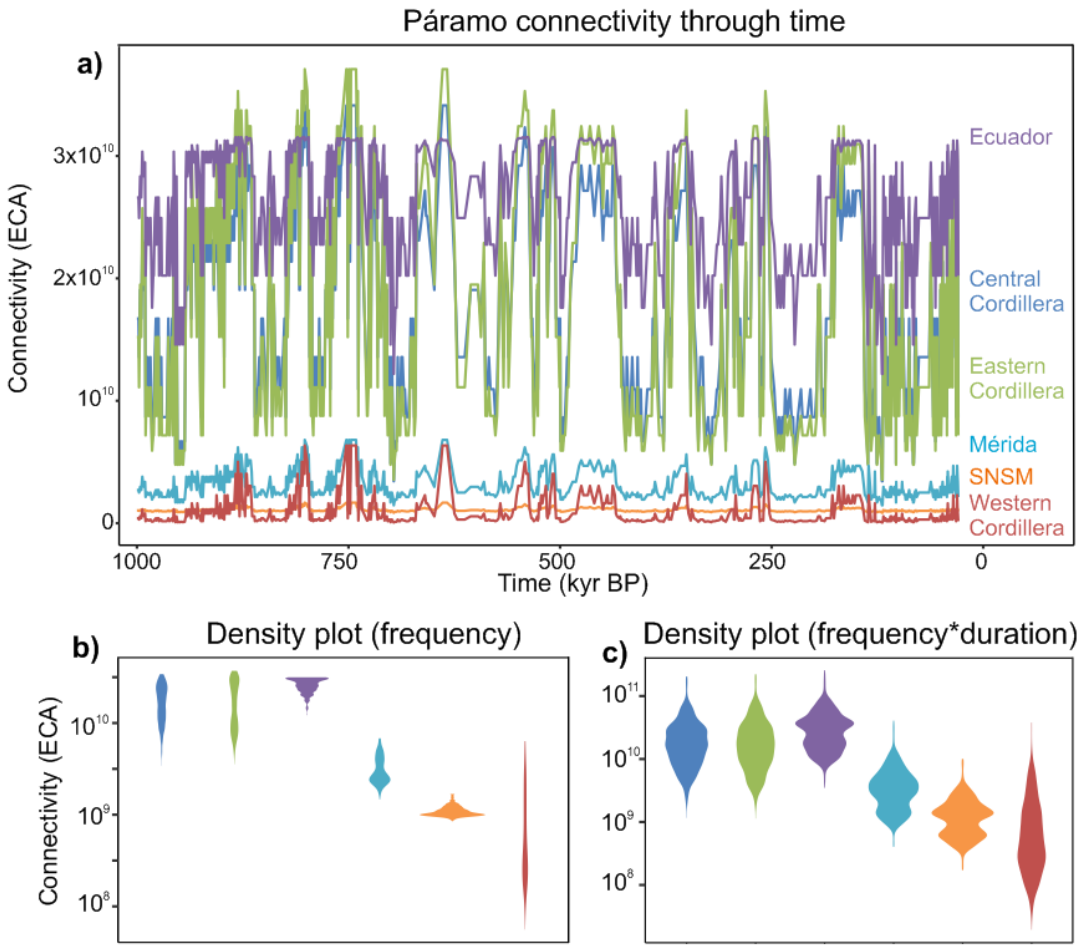

Figure 5 | The 'Flickering Connectivity System' of the Northern Andes. a) Páramo connectivity (ECA) through time (3000-30 kyr BP) for each cordillera. b) Violin plot showing the distribution of the data and its probability density. Plot only considers how often certain connectivity occurred, not how long it lasted. c) Violin plot showing the distribution of the data and its probability density multiplied by how long connectivity persisted.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

REFERENCES

Ali, J. R., & Aitchison, J. C. (2014). Exploring the combined role of eustasy and oceanic island thermal subsidence in shaping biodiversity on the Galápagos. *Journal of Biogeography*, *41*(7), 1227–1241. https://doi.org/10.1111/jbi.12313

Antonelli, A., Kissling, W. D., Flantua, S. G. A., Bermúdez, M. A., Mulch, A., Muellner-Riehl, A. N., … Hoorn, C. (2018). Geological and climatic influences on mountain biodiversity. *Nature Geoscience*, *11*(10), 718–725. https://doi.org/10.1038/s41561-018-0236-z

Barthlott, W., Rafiqpoor, D., Kier, G., & Kreft, H. (2005). Global centers of vascular plant diversity. *Nova Acta Leopoldina NF*, *92*(342), 61–83.

Bertuzzo, E., Carrara, F., Mari, L., Altermatt, F., Rodriguez-Iturbe, I., & Rinaldo, A. (2016). Geomorphic controls on elevational gradients of species richness. *Proceedings of the National Academy of Sciences*, *113*(7), 1737–1742.

Bogotá-A., R. G., Hooghiemstra, H., & Berrio, J. C. (2016). North Andean environmental and climatic change at orbital to submillennial time-scales: Vegetation, water-levels and sedimentary regimes from Lake Fúquene between 284 and 130ka. *Review of Palaeobotany and Palynology*, *226*, 91–107. https://doi.org/10.1016/j.revpalbo.2015.09.007

Bogotá-Angel, R. G., Groot, M. H. M., Hooghiemstra, H., Lourens, L. J., Van der Linden, M., & Berrio, J. C. (2011). Rapid climate change from north Andean Lake Fúquene pollen records driven by obliquity: implications for a basin-wide biostratigraphic zonation for the last 284 ka. *Quaternary Science Reviews*, *30*(23–24), 3321–3337. https://doi.org/10.1016/j.quascirev.2011.08.003

Boom, A., Marchant, R., Hooghiemstra, H., & Sinninghe Damsté, J. S. (2002). CO2- and temperature-controlled altitudinal shifts of C4- and C3-dominated grasslands allow reconstruction of palaeoatmospheric pCO2. *Palaeogeography, Palaeoclimatology, Palaeoecology*, *177*, 151–168.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Boom, A., Mora, G., Cleef, A. M., & Hooghiemstra, H. (2001). High Altitude C4 grasslands in the northern Andes: Relicts from glacial conditions? *Review of Palaeobotany and Palynology*, *115*, 147–160.

Bouchenak-Khelladi, Y., Onstein, R. E., Xing, Y., Schwery, O., & Linder, H. P. (2015). On the complexity of triggering evolutionary radiations. *New Phytologist*, *207*(2), 313–326. https://doi.org/10.1111/nph.13331

Cadena, C. D., Pedraza, C. A., & Brumfield, R. T. (2016). Climate, habitat associations and the potential distributions of Neotropical birds: Implications for diversification across the Andes. *Revista de La Academia Colombiana de Ciencias Exactas, Físicas y Naturales*, *40*(155), 275–287.

Cleef, A.M. (1981). *The vegetation of the páramos of the Colombian Cordillera Oriental*. J Cramer, Vaduz. Retrieved from http://repository.naturalis.nl/record/534752

Cleef, Antoine M. (1979). The phyogeographical position in the Neotropical vascular páramo flora with special reference to the Colombian Cordillera Oriental. In K. Larsen & L. B. Holm-Nielsen (Eds.), *Tropical Botany* (pp. 175–184). London: Academic Press.

Contreras-Ortiz, N., Atchison, G. W., Hughes, C. E., & Madriňán, S. (2018). Convergent evolution of high elevation plant growth forms and geographically structured variation in Andean Lupinus (Fabaceae). *Botanical Journal of the Linnean Society*. https://doi.org/10.1093/botlinnean/box095

Dias, M. S., Oberdorff, T., Hugueny, B., Leprieur, F., Jézéquel, C., Cornu, J.-F., … Tedesco, P. A. (2014). Global imprint of historical connectivity on freshwater fish biodiversity. *Ecology Letters*, *17*(9), 1130–1140. https://doi.org/10.1111/ele.12319

Diazgranados, M. (2012). *Phylogenetic and biogeographic relationships of Frailejones (Espeletiinae, Compositae): An ongoing radiation in the Tropical Andes* (Ph.D. Dissertation). Saint Louis University, St. Louis.

Diazgranados, M., & Barber, J. C. (2017). Geography shapes the phylogeny of frailejones (Espeletiinae Cuatrec., Asteraceae): a remarkable example of recent rapid radiation in sky islands. *PeerJ*, *5*, e2968. https://doi.org/10.7717/peerj.2968

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Elsen, P. R., & Tingley, M. W. (2015). Global mountain topography and the fate of montane species under climate change. *Nature Climate Change*, *5*(8), 772–776. https://doi.org/10.1038/nclimate2656

Flantua, S. G. A., Hooghiemstra, H., Vuille, M., Behling, H., Carson, J. F., Gosling, W. D., … González-Arango, C. (2016). Climate variability and human impact in South America during the last 2000 years: synthesis and perspectives from pollen records. *Climate of the Past*, *12*(2), 483–523. https://doi.org/10.5194/cp-12-483-2016

Flantua, S.G.A., & Hooghiemstra, H. (2018). Historical connectivity and mountain biodiversity. In C. Hoorn, A. Perrigo, & A. Antonelli (Eds.), *Mountains, climate and biodiversity* (1st ed., pp. 171–185). Wiley-Blackwell.

Flantua, Suzette G. A., Hooghiemstra, H., Grimm, E. C., Behling, H., Bush, M. B., González-Arango, C., … Van Boxel, J. H. (2015). Updated site compilation of the Latin American Pollen Database. *Review of Palaeobotany and Palynology*, *223*, 104–115. https://doi.org/10.1016/j.revpalbo.2015.09.008

Flantua, Suzette G. A., Hooghiemstra, H., Van Boxel, J. H., Cabrera, M., González-Carranza, Z., & González-Arango, C. (2014). Connectivity dynamics since the Last Glacial Maximum in the northern Andes; a pollen-driven framework to assess potential migration. In W. D. Stevens, O. M. Montiel, & P. H. Raven (Eds.), *Paleobotany and Biogeography: A Festschrift for Alan Graham in His 80th Year* (1st ed., Vol. 2014, pp. 98–123). St. Louis: Missouri Botanical Garden.

Graham, C. H., Carnaval, A. C., Cadena, C. D., Zamudio, K. R., Roberts, T. E., Parra, J. L., … Sanders, N. J. (2014). The origin and maintenance of montane diversity: integrating evolutionary and ecological processes. *Ecography*, *37*(8), 711–719. https://doi.org/10.1111/ecog.00578

Graham, C. H., Moritz, C., & Williams, S. E. (2006). Habitat history improves prediction of biodiversity in rainforest fauna. *Proceedings of the National Academy of Sciences of the United States of America*, *103*(3), 632–636. https://doi.org/10.1073/pnas.0505754103

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Grant, P. R. (2014). Adaptive radiation. In J. B. Losos, D. A. Baum, D. J. Futuyma, H. E. Hoekstra, R. E. Lenski, A. J. Moore, … M. C. Whitlock (Eds.), *The Princeton guide to Evolution*. Princeton University Press.

Groot, M. H. M., Bogotá, R. G., Lourens, L. J., Hooghiemstra, H., Vriend, M., Berrio, J. C., … Westerhoff, W. (2011). Ultra-high resolution pollen record from the northern Andes reveals rapid shifts in montane climates within the last two glacial cycles. *Clim. Past*, *7*(1), 299–316. https://doi.org/10.5194/cp-7-299-2011

Groot, Mirella H. M., Hooghiemstra, H., Berrio, J. C., & Giraldo, C. (2013). North Andean environmental and climatic change at orbital to submillennial time-scales: Vegetation, water levels and sedimentary regimes from Lake Fúquene 130–27ka. *Review of Palaeobotany and Palynology*, *197*(Supplement C), 186–204. https://doi.org/10.1016/j.revpalbo.2013.04.005

Hazzi, N. A., Moreno, J. S., Ortiz-Movliav, C., & Palacio, R. D. (2018). Biogeographic regions and events of isolation and diversification of the endemic biota of the tropical Andes. *Proceedings of the National Academy of Sciences*, *115*(31), 7985–7990. https://doi.org/10.1073/pnas.1803908115

Herman, F., Seward, D., Valla, P. G., Carter, A., Kohn, B., Willett, S. D., & Ehlers, T. A. (2013). Worldwide acceleration of mountain erosion under a cooling climate. *Nature*, *504*(7480), 423–426. https://doi.org/10.1038/nature12877

Hoeksema, B. W. (2007). Delineation of the Indo-Malayan centre of maximum marine biodiversity: The coral triangle. In W. Renema (Ed.), *Biogeography, Time, and Place: Distributions, Barriers, and Islands* (pp. 117–178). Dordrecht: Springer Netherlands. https://doi.org/10.1007/978-1-4020-6374-9_5

Hooghiemstra, H. (1984). *Vegetational and climatic history of the high plain of Bogotá, Colombia*. (Dissertationes Botanicae, Vol. 79).

Hooghiemstra, H., & Flantua, S. G. A. (2019). Colombia in the Quaternary: an overview of environmental and climatic change. In *The Geology of Colombia* (1st ed.). Bogotá, Colombia: INGEOMINAS.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Hooghiemstra, Henry, Berrio, J. C., Groot, M. H. M., Bogotá-A, R. G., Olivera, M. M., & González-Carranza, Z. (2012). The dynamic history of the upper forest line ecotone in the Northern Andes. In R. W. Myster (Ed.), *Ecotones Between Forest and Grassland* (pp. 229–246). Springer New York. Retrieved from http://link.springer.com/chapter/10.1007/978-1-4614-3797-0_10

Hooghiemstra, Henry, & Van der Hammen, T. (2004). Quaternary Ice-Age Dynamics in the Colombian Andes: Developing an Understanding of Our Legacy. *Philosophical Transactions: Biological Sciences*, *359*(1442), 173–181.

Hughes, C. E., & Atchison, G. W. (2015). The ubiquity of alpine plant radiations: from the Andes to the Hengduan Mountains. *New Phytologist*, *207*(2), 275–282. https://doi.org/10.1111/nph.13230

Hughes, C., & Eastwood, R. (2006). Island radiation on a continental scale: Exceptional rates of plant diversification after uplift of the Andes. *Proceedings of the National Academy of Sciences*, *103*(27), 10334–10339. https://doi.org/10.1073/pnas.0601928103

Jabaily, R. S., & Sytsma, K. J. (2013). Historical biogeography and life-history evolution of Andean Puya (Bromeliaceae). *Botanical Journal of the Linnean Society*, *171*(1), 201–224. https://doi.org/10.1111/j.1095-8339.2012.01307.x

Knowles, L. L. (2000). Tests of Pleistocene Speciation in Montane Grasshoppers (genus Melanoplus) from the Sky Islands of Western North America. *Evolution*, *54*(4), 1337–1348. https://doi.org/10.1111/j.0014-3820.2000.tb00566.x

Knowles, L. L., & Massatti, R. (2017). Distributional shifts – not geographic isolation – as a probable driver of montane species divergence. *Ecography*, *40*(12), 1475–1485. https://doi.org/10.1111/ecog.02893

Kolář, F., Dušková, E., & Sklenář, P. (2016). Niche shifts and range expansions along cordilleras drove diversification in a high-elevation endemic plant genus in the tropical Andes. *Molecular Ecology*, *25*(18), 4593–4610. https://doi.org/10.1111/mec.13788

Kreft, H., & Jetz, W. (2007). Global patterns and determinants of vascular plant diversity. *Proceedings of the National Academy of Sciences*, *104*(14), 5925–5930. https://doi.org/10.1073/pnas.0608361104

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Kropf, M., Kadereit, J. W., & Comes, H. P. (2003). Differential cycles of range contraction and expansion in European high mountain plants during the Late Quaternary: insights from Pritzelago alpina (L.) O. Kuntze (Brassicaceae). *Molecular Ecology*, *12*(4), 931–949. https://doi.org/10.1046/j.1365-294X.2003.01781.x

Lagomarsino, L. P., Condamine, F. L., Antonelli, A., Mulch, A., & Davis, C. C. (2016). The abiotic and biotic drivers of rapid diversification in Andean bellflowers (Campanulaceae). *New Phytologist*, *210*(4), 1430–1442. https://doi.org/10.1111/nph.13920

Lindborg, R., & Eriksson, O. (2004). Historical landscape connectivity affects present plant species diversity. *Ecology*, *85*(7), 1840–1845.

Loomis, S. E., Russell, J. M., Verschuren, D., Morrill, C., Cort, G. D., Damsté, J. S. S., … Kelly, M. A. (2017). The tropical lapse rate steepened during the Last Glacial Maximum. *Science Advances*, *3*(1), e1600815. https://doi.org/10.1126/sciadv.1600815

Madriñán, S., Cortés, A. J., & Richardson, J. E. (2013). Páramo is the world's fastest evolving and coolest biodiversity hotspot. *Frontiers in Genetics*, *4*. https://doi.org/10.3389/fgene.2013.00192

Massatti, R., & Knowles, L. L. (2014). Microhabitat Differences Impact Phylogeographic Concordance of Codistributed Species: Genomic Evidence in Montane Sedges (carex L.) from the Rocky Mountains. *Evolution*, *68*(10), 2833–2846. https://doi.org/10.1111/evo.12491

McCormack, J. E., Huang, H., & Knowles, L. L. (2009). Sky islands. In R. G. Gillespie & D. Clague (Eds.), *Encyclopedia of Islands* (pp. 839–843). Berkeley: University of Chicago Press. Retrieved from http://sites.oxy.edu/mccormack/McCormack/picks/McCormack_2009_sky_islands_REVIEW.pdf

McRae, B. H., & Kavanagh, D. M. (2011). *Linkage Mapper Connectivity Analysis Software*. Seattle WA: The Nature Conservancy. Retrieved from http://www.circuitscape.org/linkagemapper

McRae, B. H., Shirk, A. J., & Platt, J. T. (2013). *Gnarly Landscape Utilities: Resistance and Habitat Calculator User Guide*. Fort Collins, CO: The Nature Conservancy. Retrieved from http://www.circuitscape.org/gnarly-landscape-utilities.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

McRae, Brad H., Dickson, B. G., Keitt, T. H., & Shah, V. B. (2008). Using circuit theory to model connectivity in ecology, evolution, and conservation. *Ecology*, *89*(10), 2712–2724. https://doi.org/10.1890/07-1861.1

Moore, B. R., & Donoghue, M. J. (2007). Correlates of Diversification in the Plant Clade Dipsacales: Geographic Movement and Evolutionary Innovations. *The American Naturalist*, *170*(S2), S28–S55. https://doi.org/10.1086/519460

Münzbergová, Z., Cousins, S. A. O., Herben, T., Plačková, I., Mildén, M., & Ehrlén, J. (2013). Historical habitat connectivity affects current genetic structure in a grassland species. *Plant Biology*, *15*(1), 195–202. https://doi.org/10.1111/j.1438-8677.2012.00601.x

Mutke, J., Jacobs, R., Meyers, K., Henning, T., & Weigend, M. (2014). Diversity patterns of selected Andean plant groups correspond to topography and habitat dynamics, not orogeny. *Frontiers in Genetics*, *5*. https://doi.org/10.3389/fgene.2014.00351

Nevado, B., Contreras-Ortiz, N., Hughes, C., & Filatov, D. A. (2018). Pleistocene glacial cycles drive isolation, gene flow and speciation in the high elevation Andes. *New Phytologist*, *219*(2), 779–793.

Norder, S. J., Proios, K., Whittaker, R. J., Alonso, M. R., Borges, P. A. V., Borregaard, M. K., … Rijsdijk, K. F. (2018). Beyond the Last Glacial Maximum: Island endemism is best explained by long-lasting archipelago configurations. *Global Ecology and Biogeography*, *0*(0). https://doi.org/10.1111/geb.12835

Nürk, N. M., Scheriau, C., & Madriñán, S. (2013). Explosive radiation in high Andean Hypericum—rates of diversification among New World lineages. *Evolutionary and Population Genetics*, *4*, 175. https://doi.org/10.3389/fgene.2013.00175

Onstein, R. E., Baker, W. J., Couvreur, T. L. P., Faurby, S., Svenning, J.-C., & Kissling, W. D. (2017). Frugivory-related traits promote speciation of tropical palms. *Nature Ecology & Evolution*, *1*(12), 1903. https://doi.org/10.1038/s41559-017-0348-7

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Papadopoulou, A., & Knowles, L. L. (2015). Species-specific responses to island connectivity cycles: refined models for testing phylogeographic concordance across a Mediterranean Pleistocene Aggregate Island Complex. *Molecular Ecology*, *24*(16), 4252–4268. https://doi.org/10.1111/mec.13305

Papadopoulou, A., & Knowles, L. L. (2016). Toward a paradigm shift in comparative phylogeography driven by trait-based hypotheses. *Proceedings of the National Academy of Sciences*, *113*(29), 8018–8024. https://doi.org/10.1073/pnas.1601069113

Pennington, R. T., Lavin, M., Särkinen, T., Lewis, G. P., Klitgaard, B. B., & Hughes, C. E. (2010). Contrasting plant diversification histories within the Andean biodiversity hotspot. *Proceedings of the National Academy of Sciences*, *107*(31), 13783–13787. https://doi.org/10.1073/pnas.1001317107

Petit, R. J., Aguinagalde, I., Beaulieu, J.-L. de, Bittkau, C., Brewer, S., Cheddadi, R., … Vendramin, G. G. (2003). Glacial Refugia: Hotspots but not melting pots of genetic diversity. *Science*, *300*(5625), 1563–1565. https://doi.org/10.1126/science.1083264

Pouchon, C., Fernández, A., Nassar, J. M., Boyer, F., Aubert, S., Lavergne, S., … Antonelli, A. (2018). Phylogenomic Analysis of the Explosive Adaptive Radiation of the Espeletia Complex (Asteraceae) in the Tropical Andes. *Systematic Biology*, *67*(6), 1041–1060. https://doi.org/10.1093/sysbio/syy022

Qian, H., & Ricklefs, R. E. (2000). Large-scale processes and the Asian bias in species diversity of temperate plants. *Nature*, *407*(6801), 180–182. https://doi.org/10.1038/35025052

Quintero, I., & Jetz, W. (2018). Global elevational diversity and diversification of birds. *Nature*, *555*(7695), 246–250. https://doi.org/10.1038/nature25794

Ramírez-Barahona, S., & Eguiarte, L. E. (2013). The role of glacial cycles in promoting genetic diversity in the Neotropics: the case of cloud forests during the Last Glacial Maximum. *Ecology and Evolution*, *3*(3), 725–738. https://doi.org/10.1002/ece3.483

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Rull, V. (2005). Biotic diversification in the Guayana Highlands: a proposal. *Journal of Biogeography*, *32*(6), 921–927. https://doi.org/10.1111/j.1365-2699.2005.01252.x

Rull, Valentí. (2011). Neotropical biodiversity: timing and potential drivers. *Trends in Ecology & Evolution*, *26*(10), 508–513. https://doi.org/10.1016/j.tree.2011.05.011

Rull, Valentí, & Nogué, S. (2007). Potential migration routes and barriers for vascular plants of the Neotropical Guyana highlands during the Quaternary. *Journal of Biogeography*, *34*(8), 1327–1341.

Saura, S., Estreguil, C., Mouton, C., & Rodríguez-Freire, M. (2011). Network analysis to assess landscape connectivity trends: Application to European forests (1990–2000). *Ecological Indicators*, *11*(2), 407–416. https://doi.org/10.1016/j.ecolind.2010.06.011

Saura, S., & Pascual-Hortal, L. (2007). A new habitat availability index to integrate connectivity in landscape conservation planning: Comparison with existing indices and application to a case study. *Landscape and Urban Planning*, *83*(2–3), 91–103. https://doi.org/10.1016/j.landurbplan.2007.03.005

Saura, S., & Rubio, L. (2010). A common currency for the different ways in which patches and links can contribute to habitat availability and connectivity in the landscape. *Ecography*, *33*, 523–537. https://doi.org/10.1111/j.1600-0587.2009.05760.x

Saura, S., & Torné, J. (2009). Conefor Sensinode 2.2: A software package for quantifying the importance of habitat patches for landscape connectivity. *Environmental Modelling & Software*, *24*(1), 135–139. https://doi.org/10.1016/j.envsoft.2008.05.005

Schönswetter, P., Stehlik, I., Holderegger, R., & Tribsch, A. (2005). Molecular evidence for glacial refugia of mountain plants in the European Alps. *Molecular Ecology*, *14*(11), 3547–3555. https://doi.org/10.1111/j.1365-294X.2005.02683.x

Simpson, B. B. (1974). Glacial migrations of plants: Island biogeographical evidence. *Science*, *185*(4152), 698–700. https://doi.org/10.1126/science.185.4152.698

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Sklenář, P., Hedberg, I., & Cleef, A. M. (2014). Island biogeography of tropical alpine floras. *Journal of Biogeography*, *41*(2), 287–297. https://doi.org/10.1111/jbi.12212

Smith, B. T., McCormack, J. E., Cuervo, A. M., Hickerson, M. J., Aleixo, A., Cadena, C. D., … Brumfield, R. T. (2014). The drivers of tropical speciation. *Nature*, *515*(7527), 406–409. https://doi.org/10.1038/nature13687

Steinbauer, M. J., Field, R., Grytnes, J.-A., Trigas, P., Ah-Peng, C., Attorre, F., … Beierkuhnlein, C. (2016). Topography-driven isolation, speciation and a global increase of endemism with elevation. *Global Ecology and Biogeography*, *25*(9), 1097–1107. https://doi.org/10.1111/geb.12469

Tischendorf, L., & Fahrig, L. (2000). On the usage and measurement of landscape connectivity. *Oikos*, *90*(1), 7–19. https://doi.org/10.1034/j.1600-0706.2000.900102.x

Torres, V., Hooghiemstra, H., Lourens, L., & Tzedakis, P. C. (2013). Astronomical tuning of long pollen records reveals the dynamic history of montane biomes and lake levels in the tropical high Andes during the Quaternary. *Quaternary Science Reviews*, *63*, 59–72. https://doi.org/10.1016/j.quascirev.2012.11.004

Uribe-Convers, S., & Tank, D. C. (2015). Shifts in diversification rates linked to biogeographic movement into new areas: An example of a recent radiation in the Andes. *American Journal of Botany*, *102*(11), 1854–1869. https://doi.org/10.3732/ajb.1500229

Van der Hammen, T. (1974). The Pleistocene changes of vegetation and climate in tropical South America. *Journal of Biogeography*, *1*, 3–26. https://doi.org/10.2307/3038066

Van der Hammen, Thomas. (1981). Glaciales y glaciaciones en el cuaternario de Colombia: paleoecología y estratigrafía. *Revista CIAF*, *6*(1–3), 635–638.

Von Humboldt, A. (1845). *Kosmos: Entwurf einer physischen Weltbeschreibung* (Vol. 1). Eichborn Verlag.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Wallis, G. P., Waters, J. M., Upton, P., & Craw, D. (2016). Transverse Alpine Speciation Driven by Glaciation. *Trends in Ecology & Evolution*, *31*(12), 916–926. https://doi.org/10.1016/j.tree.2016.08.009

Weigelt, P., Steinbauer, M. J., Cabral, J. S., & Kreft, H. (2016). Late Quaternary climate change shapes island biodiversity. *Nature*, *532*(7597), 99–102. https://doi.org/10.1038/nature17443

Weir, J. T. (2006). Divergent timing and patterns of species accumulation in lowland and highland neotropical birds. *Evolution*, *60*(4), 842–855. https://doi.org/10.1554/05-272.1

Weir, J. T., Haddrath, O., Robertson, H. A., Colbourne, R. M., & Baker, A. J. (2016). Explosive ice age diversification of kiwi. *Proceedings of the National Academy of Sciences*, 201603795. https://doi.org/10.1073/pnas.1603795113

Wille, M., Hooghiemstra, H., Behling, H., van der Borg, K., & Negret, A. J. (2001). Environmental change in the Colombian subandean forest belt from 8 pollen records: the last 50 kyr. *Vegetation History and Archaeobotany*, *10*(2), 61–77.

Winkworth, R. C., Wagstaff, S. J., Glenny, D., & Lockhart, P. J. (2005). Evolution of the New Zealand mountain flora: Origins, diversification and dispersal. *Organisms Diversity & Evolution*, *5*(3), 237–247. https://doi.org/10.1016/j.ode.2004.12.001

BIOSKETCHES

Suzette Flantua has a background in paleoecology, biogeography, landscape ecology and spatial analyses, and enjoys integrating them all. She is interested in a wide range of topics from the Miocene to the present, from islands to mountains, to understand contemporary patterns of species richness and endemism.

Henry Hooghiemstra is a terrestrial and marine tropical palynologist working on time-scales from the full Quaternary to the Anthropocene. His research focuses on a wide variety of biomes in Central and South America, Saharan and East Africa and in Mauritius.

bioRxiv preprint doi: https://doi.org/10.1101/569681; this version posted March 7, 2019. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. It is made available under aCC-BY 4.0 International license.

Aaron O'Dea is a marine paleobiologist and uses the marine fossil record in Tropical America to explore drivers of macroevolution in the seas, and takes cores on coral reefs from French Polynesia to the Dominican Republic to reconstruct how reefs changed over millennia with the aim of improving their future resilience.

Renske E. Onstein is an evolutionary ecologist who enjoys collecting (and eating) tropical megafaunal fruits, e.g. on Borneo and Madagascar, while studying how fruit functional traits interact with frugivores to affect diversification dynamics. She is generally interested in the broad-scale distribution and diversification of functional and taxonomic diversity of flowering plants.

APPENDICES

Additional Supporting Information may be found in the online version of this article:

Appendix 1 | Surface areas, elevational ranges and hypsographies of the Northern Andes

Appendix 2 | Background on the páramo alpine biome

Appendix 3 | Methodology underlying the use of fossil pollen data to reconstruct the upper forest line changes

Appendix 4 | Degree of connectivity of páramos at all UFL elevations during the last 1 Myr.

Appendix 5 | Frequency analysis of all UFL elevations during the last 1 Myr.

Appendix 6 | Visualization of the flickering connectivity system in the Northern Andes. Artwork by Catalina Giraldo Pastrana in collaboration with Suzette G.A. Flantua and Henry Hooghiemstra.

Appendix 7 | Further suggestions for future work.

AUTHOR CONTRIBUTIONS

S.G.A.F. and H.H. conceived the ideas. H.H. provided the AP% of the Funza09 dataset. S.G.A.F. performed the spatial analyses. S.G.A.F and H.H. led the writing and figure design with critical contributions by A.O. and R.E.O. All authors contributed to versions of the manuscript and revisions.