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**IMPORTANCIA DE LA CONFIGURACIÓN DEL PAISAJE SOBRE LAS
COMUNIDADES DE PLÁNTULAS Y MAMÍFEROS DE UN BOSQUE TROPICAL
HIPER FRAGMENTADO**

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RESUMEN

La deforestación y cambio de uso de suelo en los bosques tropicales húmedos son la principal amenaza para la biodiversidad a nivel global al transformar coberturas continuas de bosque en paisajes tropicales modificados por el hombre. Estos paisajes consisten en fragmentos de bosque inmersos en una matriz heterogénea de usos de suelo y vegetación en distintos estados de sucesión. Esto genera una compleja estructura en estos paisajes, tanto en su composición de coberturas, como en la configuración espacial de los remanentes de bosque, los cuales ejercen una fuerte influencia sobre las especies remanentes dentro de los fragmentos. A su vez, esta flora y fauna remanentes dependen de la calidad de hábitat y disponibilidad de recursos dentro de los fragmentos, los cuales están asociados con la estructura y diversidad de la comunidad de árboles. Por lo tanto, preservar la regeneración del bosque dentro de fragmentos es crucial para la conservación de la biodiversidad en estos paisajes en el mediano y largo plazo. La regeneración de los bosques tropicales húmedos está directamente regulada por los patrones de diversidad de la comunidad de plántulas que se desarrollan dentro de ellos. Sin embargo, esta comunidad es influenciada por factores a escala local y de paisaje que modifican la llegada de especies nuevas, las fluctuaciones en las abundancias de especies con distintos atributos funcionales y la llegada de los mamíferos terrestres que promueven su diversidad. No obstante, se conoce muy poco sobre como estos factores influyen directa o indirectamente en la abundancia de plántulas demandantes de luz/tolerantes a la sombra, y en la diversidad alfa y beta, así como en la presencia de mamíferos terrestres. En este trabajo analizamos los efectos directos y en cascada de factores a escalas de fragmento y de paisaje sobre la abundancia y diversidad de plántulas, así como el papel de la estructura del paisaje y las funciones de la matriz sobre la ocupación de mamíferos terrestres en un bosque tropical hiper fragmentado. Encontramos que el disturbio por fuego tiene un efecto en cascada positivo sobre las abundancias de plántulas demandantes de luz al reducir el área basal de la vegetación arbórea. A su vez, el incremento en la abundancia de estas plántulas redujo la diversidad beta de especies típicas y dominantes. Por otro lado, la agregación de fragmentos tuvo un efecto en cascada positivo sobre la riqueza de plántulas al incrementar la abundancia de especies tolerantes a la sombra. También encontramos efectos directos del paisaje sobre la diversidad de plántulas, siendo la configuración del paisaje la que influyó de manera significativa en la diversidad beta, particularmente en paisajes con una alta densidad de fragmentos y un bajo contraste de borde. Finalmente, paisajes con una alta densidad de fragmentos y una matriz que proporciona un mayor porcentaje de hábitat complementario a pequeñas escalas y una menor resistencia al movimiento a grandes escalas incrementó la ocupación de mamíferos terrestres. Estos resultados nos indican que la configuración del paisaje juega un papel muy importante en la diversidad de plántulas y en facilitar la llegada de mamíferos en paisajes hiper fragmentados. Es urgente promover patrones configuracionales que incrementen la diversidad de la comunidad de plántulas dentro de planes de manejo y restauración de estos paisajes modificados por el hombre.

ABSTRACT

Deforestation and land-use change in tropical rainforests are the major threat for global biodiversity by transforming continuous tracts of undisturbed forest into human-modified tropical landscapes. These landscapes consist in forest patches embedded in a heterogeneous matrix of a variety of land uses and secondary forests. This creates a complex landscape structure, whether in the composition of land covers or in the spatial configuration of forest patches, which strongly influence the remnant species within patches. At the same time, the remaining vegetation and fauna are dependent on the habitat quality and resources availability provided by the structure and diversity of the tree forest community. Thus, sustaining forest regeneration within patches is critical for biodiversity conservation at long term. Tropical forest regeneration is mostly driven by the diversity patterns of tree seedling community. However, seedlings are also affected by local- and landscape-scale factors that alter species arrival, abundance fluctuations of species with contrasting functional traits and the arrival of terrestrial mammals that shape seedling diversity. Nonetheless, the factors that have direct or cascading effects on seedling abundance and diversity, and mammal persistence are poorly understood. We assessed the direct and cascading effects of patch- and landscape-scale factors on seedling abundance and diversity, as well as the role of landscape structure and matrix functionality on mammal occupancy in a hyper-fragmented tropical rainforest. We found that fire disturbance had positive cascading effects on the abundance of light-demanding seedlings, which in turn reduced beta diversity of typical and dominant seedling species. These effects were mediated by forest basal area, which was reduced in patches with previous fire events. Contrarily, patch aggregation had positive cascading effects on seedling richness by increasing the abundance of shade-tolerant species. We also found direct effects of landscape structure on seedling diversity, particularly by landscape configuration on beta diversity in landscapes with a high patch density and a low edge contrast. Finally, a high patch density and matrices that increased habitat availability at smaller scales while reduced resistance to movement at larger scales promoted patch occupancy of terrestrial mammals. These findings indicated that landscape configuration played a key role on seedling diversity and mammal occupancy within patches. Promoting configurational patterns that sustain seedling diversity is crucial in management and restoration actions in these human-modified tropical landscapes.

CAPÍTULO I. INTRODUCCIÓN GENERAL

Fragmentación de los bosques tropicales húmedos y sus efectos sobre la comunidad de plántulas

Paisajes modificados por el hombre

La extensión, biodiversidad e integridad ecosistémica de los bosques a nivel mundial están fuertemente amenazadas por la deforestación y cambio de uso de suelo (Grantham et al., 2020; Song et al., 2018). Actualmente solo el 40% de los bosques remanentes presentan una alta integridad ecosistémica (Grantham et al., 2020), mientras que el resto ha sido transformado en paisajes modificados por el hombre (PMH). Los PMH están constituidos por fragmentos de bosque inmersos dentro de un mosaico de coberturas de uso de suelo de origen antropogénico (DeClerck et al., 2010; Perring & Ellis, 2013) y se caracterizan por presentar un ensamble de especies y funciones ecológicas alteradas (DeClerck et al., 2010; Richard J. Hobbs et al., 2013; Western, 2001). Esto limita tanto sus servicios ecosistémicos como su resiliencia a eventos de perturbación a múltiples escalas espaciales y temporales (Watson et al., 2018; Wurtzebach & Schultz, 2016). No obstante, los PMH tienen un alto valor de conservación y sus servicios ecosistémicos son fundamentales para la provisión de alimento, regulación climática y mitigación del disturbio antropogénico al ser una fuente de propágulos para la regeneración del bosque (Cumming & Peterson, 2005; Kennedy et al., 2013; Perring & Ellis, 2013). Mantener estas capacidades requiere de un manejo adaptativo del paisaje que permitan conservar la biodiversidad remanente o manejar los nuevos ensambles de especies y sus funciones asociadas (Hulvey et al., 2013).

Estos PMH son más frecuentes en los bosques tropicales húmedos, donde el 10% de los bosques remanentes persisten como fragmentos menores a las 10,000 ha, inmersos dentro de coberturas agrícolas, pastizales ganaderos, plantaciones agroindustriales (ej., cultivos extensivos de soya y palma de aceite) y bosques secundarios en distintos estadios

sucesionales (Arroyo-Rodríguez et al., 2017, 2020; DeClerck et al., 2010). Solamente en América, se estima que existen 55.5 millones de fragmentos con un tamaño promedio de 17 ha y cuyo número se espera que aumente en un factor de 20-33 durante los próximos 30 años, trayendo consigo efectos negativos sobre la biodiversidad que habita en ellos, sus funciones ecológicas y la integridad ecosistémica de los PMH tropicales (Grantham et al., 2020; Haddad et al., 2015; Taubert et al., 2018). Conservar las especies y funciones de los PMH tropicales es fundamental debido a que estos bosques albergan más de la mitad de las especies conocidas en el planeta y juegan un papel importante en procesos como el ciclo de carbono y la regulación del clima a múltiples escalas (Le Quéré et al., 2018; Wright, 2005).

Los fragmentos de bosque tropical son críticos para la preservación de especies de plantas y animales en PMH tropicales (Bueno & Peres, 2019; Laurance et al., 2011; Pfeifer et al., 2014), mantienen la conectividad del paisaje (Jesus et al., 2012) y representan la principal fuente de propágulos para la regeneración de bosques secundarios (Chazdon, 2003; Martínez-Ramos et al., 2016). Estas funciones están fuertemente vinculadas a la vegetación arbórea de los fragmentos, siendo indicadores de calidad de hábitat, disponibilidad de recursos y funciones ecológicas (Cudney-Valenzuela et al., 2021; San-José et al., 2014; Wies et al., 2021). Por lo tanto, para preservar la integridad ecosistémica de los PMH en el trópico es fundamental mantener la regeneración de la comunidad arbórea dentro de los fragmentos.

Comunidades de plántulas dentro de fragmentos

La regeneración de los bosques tropicales húmedos es principalmente a través de la comunidad de plántulas de árboles, la cual representa la principal fuente de propágulos y su diversidad y composición determinan la estructura y diversidad de los bosques futuros (Comita et al., 2007; Nicotra et al., 1999; Wills et al., 2006). Entonces, comprender los factores y

procesos que determinan y moldean la comunidad de plántulas de árboles dentro de los fragmentos nos permitirá comprender como preservar la regeneración de fragmentos dentro de PHM tropicales (Benítez-Malvido et al., 2018; Santo-Silva et al., 2013). Estudios a escala de fragmento han encontrado que la comunidad de plántulas es influenciada por el tamaño y la forma del fragmento (Benitez-Malvido, 1998; Benítez-Malvido et al., 2018), los cuales afectan la abundancia y riqueza de plántulas (Benítez-Malvido et al., 2018; Benitez-Malvido & Martinez-Ramos, 2003; Santo-Silva et al., 2013). En PMH tropicales hiper fragmentados, los efectos de borde promueven la abundancia de plántulas pertenecientes a especies generalistas y demandantes de luz (especies dominantes en bosques secundarios) que limitan el establecimiento de plántulas pertenecientes a especies especialistas tolerantes a la sombra (especies dominantes en bosques no perturbados), trayendo consigo una reducción en la riqueza de especies (diversidad α) y una homogenización biótica que consiste en la pérdida de la heterogeneidad espacial de las especies de plántulas (diversidad β) dentro y entre los fragmentos (Tabarelli et al., 2010, 2012).

Asimismo, la diversidad α y β de las plántulas de árboles tropicales también es modificada por la depredación de semillas y plántulas por parte de la comunidad de mamíferos terrestres (Dirzo & Miranda, 1990; Villar et al., 2019; Young et al., 2016). La pérdida de estos mamíferos trae consigo modificaciones tanto en la depredación y dispersión secundaria de semillas de árboles (Galetti et al., 2015; Orrock et al., 2006; Wright et al., 2000), como en la herbivoría de plántulas (Villar et al., 2019), resultando en patrones de regeneración alterados debido a la formación de comunidades de plántulas con una alta abundancia de individuos, con una menor riqueza, dominadas por pocas especies y una baja diversidad β (Camargo-Sanabria et al., 2014; Galetti & Dirzo, 2013; Villar et al., 2019). La deforestación y fragmentación afecta severamente la persistencia de mamíferos terrestres, particularmente aquellos cuyo tamaño, dieta y patrones de forrajeo proveen las funciones ecológicas que promueven la diversidad de

plántulas (Magioli et al., 2021; Young et al., 2016). En el caso de los fragmentos de selva, el tamaño tiene un papel importante en explicar la presencia de mamíferos terrestres de talla mediana y grande, mientras que aquellos fragmentos de menor tamaño pueden ser usados de manera temporal como sitios de paso (Magioli et al., 2021; Prugh et al., 2008; Thornton et al., 2011).

Como consecuencia, los bosques resultantes de estos fragmentos presentan una estructura y diversidad semejante a la de bosques secundarios, comprometiendo la integridad ecosistémica de los PMH tropicales al reducirse la calidad del hábitat para plantas y animales de hábitos especialistas y pauperizando sus funciones ecológicas y servicios ecosistémicos (Laurance, Nascimento, Laurance, Andrade, Fearnside, et al., 2006; Laurance, Nascimento, Laurance, Andrade, Ribeiro, et al., 2006; Tabarelli et al., 2012). El resultado de los trabajos a escala de fragmento sugiere que las actividades de conservación y manejo deben orientarse a detener la deforestación de estos fragmentos y promover su restauración activa a lo largo de sus bordes para aminorar los efectos que tienen sobre la comunidad de plántulas, promoviendo así su diversidad a largo plazo.

No obstante, existen otros factores que afectan a la comunidad de plántulas que van más allá del tamaño y forma de los fragmentos. La composición de especies de plántulas y árboles está fuertemente influenciada por la estructura de la vegetación arbórea (Nicotra et al., 1999; Rose, 2000), el tipo de matriz adyacente (Benítez-Malvido & Martínez-Ramos, 2006; Laurance et al., 2007), la heterogeneidad de hábitats entre distintos fragmentos (Arroyo-Rodríguez et al., 2013; Sánchez-Gallen et al., 2010) y la persistencia de mamíferos a través de la conectividad entre fragmentos (Cassano et al., 2012; Magioli et al., 2016; Umetsu & Pardini, 2007).

Efectos de la estructura del paisaje y la escala sobre la biodiversidad

La biodiversidad es significativamente influenciada por la estructura del paisaje (Fahrig, 2003), la cual está constituida por la composición (número y extensión) y la configuración (formas, distribución y arreglo espacial) de las coberturas de vegetación y uso de suelo (Fletcher & Fortin, 2018; McGarigal et al., 2012). En hábitats fragmentados, los efectos de la composición del paisaje están asociados a la pérdida del hábitat (ej., reducción del porcentaje de hábitat original), mientras que los efectos de la configuración se encuentran relacionados con los efectos de la fragmentación *per se* (Fahrig, 2003). Estos factores tienen un poder explicativo más alto que el tamaño y forma del fragmento sobre la abundancia y diversidad de especies, particularmente la pérdida de hábitat, que suele tener un efecto significativamente mayor (Fahrig, 2003). No obstante, en regiones hiper fragmentadas (cobertura de hábitat < 30%), los efectos de la fragmentación *per se* pueden ser mayores (Andren, 1994), teniendo efectos positivos o negativos sobre las especies remanentes, dependiendo de la interacción que cada una tenga con el paisaje (Fahrig, 2017, 2019).

Además, la escala del efecto que tiene la estructura del paisaje sobre las especies varía entre respuestas biológicas y niveles de disturbio (Jackson & Fahrig, 2012, 2015; Miguet et al., 2016). Por un lado, se espera que respuestas biológicas afectadas por procesos de natalidad/mortalidad, tales como la abundancia, serán afectadas por factores locales o a escalas menores que aquellas respuestas controladas por eventos de colonización/extinción, como la riqueza (Miguet et al., 2016; Moraga et al., 2019). Por otro lado, se ha hipotetizado una escala de efecto menor en paisajes altamente deforestados, donde la dispersión es baja y los efectos son determinados por factores locales (Andren, 1994; Miguet et al., 2016). Determinar las escalas del efecto entre la mirada de escalas, respuestas biológicas y métricas de composición y configuración del paisaje requieren de análisis multiescalares que determinen la

escala del efecto de cada métrica para cada respuesta y posteriormente la importancia relativa que tendrá respecto a otras métricas (Jackson & Fahrig, 2012, 2015; Martin, 2018; Miguet et al., 2016).

Posibles efectos de la estructura del paisaje sobre la comunidad de plántulas

Abundancia

La comunidad regenerativa de los bosques tropicales húmedos no está exenta de la influencia de la estructura del paisaje y la escala a la que son afectados los distintos atributos de la comunidad. Se presume que la abundancia de plántulas podría estar afectada por la estructura del paisaje a través de efectos en cascada. En los Tutlxas, Veracruz, se encontró que la densidad de tallos de plantas del sotobosque se incrementa al reducirse la abundancia de una palma altamente competitiva, la cual es negativamente afectada por la proporción de matrices abiertas y el grado de fragmentación de la cobertura de bosque primario (Hernández-Ruedas et al., 2018). Las abundancias de plántulas están influenciadas por la abundancia y diversidad de árboles reproductivos (Comita et al., 2007; Wandrag et al., 2017), la disponibilidad de luz regulada por la estructura de la comunidad arbórea (J S Denslow, 1987; Julie S. Denslow & Guzman G., 2000; Nicotra et al., 1999) y por factores densodependientes entre especies conespecíficas (Comita et al., 2010; Harms et al., 2000). A su vez, los PMH tropicales la estructura y diversidad de árboles es afectada por el porcentaje de bosque remanente, la exposición a bordes y el tipo y manejo de las matrices adyacentes (Laurance et al., 2007; Oliveira et al., 2008; Tenius Ribeiro et al., 2019). Además, las abundancias de plántulas también están sujetas a la extensión, duración y frecuencia del disturbio asociados al manejo agrícola y ganadero de la región, así como a la historia de uso de suelo (Zermeño-Hernández et al., 2015,

2016). Por lo tanto, es posible que la abundancia de plántulas este influenciada por factores locales asociados con atributos estructurales del fragmento, tales como el área basal y la diversidad de los árboles que a su vez son influenciados por la estructura del paisaje, en particular la exposición a bordes y la composición de la matriz adyacente.

Diversidad

Por otro lado, estudios de paisaje han señalado que la diversidad α de plántulas es directamente afectada por la estructura del paisaje. Tanto la diversidad α , como la proporción de especies tolerantes a la sombra de semillas y brinzales están asociadas a la composición del paisaje, en particular a los porcentajes de bosque primario a escalas grandes y de matrices desprovistas de vegetación arbórea a escalas pequeñas (Arasa-Gisbert et al., 2021; San-José et al., 2019; San-José et al., 2019). Estas variables también afectan la abundancia y diversidad de aves frugívoras y primates arborícolas (Carrara et al., 2015; Galán-Acedo et al., 2019; Kormann et al., 2018), los cuales son los principales dispersores de semillas de árboles en los trópicos (Jordano, 2000). Lo anterior sugiere que la reducción de la cobertura de bosque primario y la dominancia de coberturas uso de suelo abiertas pueden limitar la diversidad α de plántulas al restringir la llegada de nuevas especies a través de la dispersión de semillas provenientes de otros fragmentos.

En el caso de la diversidad β , se desconoce si la estructura del paisaje influye o no en su variación. No obstante, evidencia de trabajos a escala de fragmento indican que esta diversidad se encuentra fuertemente influenciada por los efectos de borde y las fluctuaciones de abundancias entre plántulas demandantes de luz y tolerantes a la sombra por los factores locales mencionados anteriormente (Krishnadas et al., 2019, 2020; Tabarelli et al., 2012). Lo anterior a través de procesos de exclusión competitiva y una mayor heterogeneidad ambiental

de los bordes que promueve el recambio de especies (Krishnadas et al., 2019; Laurance et al., 2007; McKinney & Lockwood, 1999). Asimismo, trabajos con comunidades arbóreas indican que la diversidad β es promovida en paisajes fragmentados, donde la heterogeneidad espacial entre fragmentos promueve el establecimiento de comunidades arbóreas distintas entre ellos (Arroyo-Rodríguez et al., 2013). Por lo tanto, se sugiere que factores locales como la estructura de la vegetación y la configuración del paisaje relativa al contraste de borde y la fragmentación de los bosques primarios pueden jugar un papel importante sobre la diversidad β de las plántulas. Sin embargo, aún no existen estudios que nos permitan responder estas preguntas.

Finalmente, la estructura del paisaje también tiene una fuerte influencia sobre la persistencia de fauna dentro de fragmentos. Una mayor cobertura de matrices arboladas y la presencia de una mayor cantidad de fragmentos a lo largo del paisaje promueven la riqueza de mamíferos terrestres, siendo la complementación de recursos o la conectividad que provee el paisaje las posibles explicaciones (Arce-Peña et al., 2022; Garmendia et al., 2013; Thornton et al., 2011). Lo anterior indica que la matriz juega un papel importante en la persistencia de mamíferos terrestres en los PMH tropicales. Sin embargo, se carece de estudios que identifiquen el papel de la matriz para facilitar la presencia de mamíferos dentro de fragmentos (complementación/conectividad) y a que escalas están operando. Comprender las funciones que desempeña la matriz permitirá establecer que criterios de manejo se deben implementar ya sea para aumentar su capacidad de complementación de recursos o para aumentar su permeabilidad para el movimiento entre fragmentos (Blaum & Wichmann, 2007; Fischer & Lindenmayer, 2006; Franklin & Lindenmayer, 2009).

Esclarecer lo anterior es primordial para elaborar estrategias que mantengan la diversidad de plántulas en selvas hiper fragmentadas, ya sea a través de mejorar la calidad de hábitat dentro de los fragmentos y/o promover patrones de composición y configuración en el paisaje que aminoren los efectos de la pérdida de hábitat, reduzcan los efectos de borde y

faciliten la presencia de mamíferos terrestres. Por lo tanto, es necesario elaborar estudios que evalúen la contribución que tiene la estructura del paisaje sobre los factores que afectan la abundancia y diversidad de plántulas de árboles dentro de fragmentos e identifiquen las coberturas de matriz y los patrones configuracionales prioritarios de conservación y/o manejo en PTH tropicales.

Objetivos

General

Esta tesis tiene como objetivo evaluar la respuesta de la comunidad de plántulas de árboles tropicales a la estructura del paisaje en una selva hiper fragmentada.

Particulares

1. Identificar los efectos directos e indirectos de los factores a escala de fragmento que afectan la abundancia y diversidad α y β de las plántulas de árboles tropicales (capítulo II)
2. Evaluar la el efecto de la estructura del paisaje y sus escalas de efecto sobre la diversidad α y β de las plántulas de árboles tropicales (capítulo III)
3. Analizar la contribución que los efectos de la matriz (complementación y conectividad) y la fragmentación *per se* tienen sobre la presencia de mamíferos terrestres dentro de fragmentos a distintas escalas espaciales (capítulo IV)
4. Discutir las implicaciones de los resultados obtenidos en la regeneración de fragmentos (capítulo V)

La información generada por este trabajo permitirá identificar aquellos factores a escala de fragmento y de paisaje que moldean la comunidad de plántulas dentro de PMH tropicales, los cuáles serán la cobertura dominante de los bosques tropicales húmedos en el mediano y largo plazo. Favorecer y preservar la regeneración de fragmentos a través del manejo integral de estos paisajes será fundamental tanto para la conservación de la biodiversidad de los bosques tropicales húmedos y sus servicios ecosistémicos, como para aumentar la resiliencia de estos nuevos ecosistemas a los efectos del cambio global.

Hipótesis

A partir de la evidencia revisada anteriormente y de acuerdo con la hipótesis del umbral de fragmentación de Andren (1994), se plantea que la comunidad de plántulas y los mamíferos terrestres que la afectan, responderán a tanto a factores a escala de fragmento, como a elementos de la estructura del paisaje relacionados con la composición y efectos de la matriz circundante y con los patrones configuracionales asociados a la densidad, aislamiento y agregación de fragmentos.

En el caso de la comunidad de plántulas, la abundancia de especies demandantes de luz será limitada en fragmentos con una mayor área basal y diversidad de la vegetación arbórea, y un menor disturbio asociado al manejo ganadero, que es la actividad productiva dominante en la región (Berget et al., 2021; Lohbeck et al., 2022). Para la diversidad α , se espera que aumente dentro de paisajes dominados por matrices arbóreas (i.e., porcentaje de bosques secundarios) y donde los patrones de configuración reduzcan el aislamiento entre fragmentos, ya sea una mayor densidad, una menor distancia o una mayor agregación de

fragmentos en el paisaje. Por su parte, la diversidad β será afectada por factores directos e indirectos, siendo directamente promovida en paisajes con un bajo contraste de borde con la matriz adyacente y una mayor densidad de fragmentos e indirectamente beneficiada tanto por el área basal del fragmento (a través de la fluctuación de abundancias de plántulas). A su vez, la escala de los efectos de la estructura del paisaje sobre la diversidad α y β de las plántulas variará dependiendo de la abundancia de las especies evaluadas, siendo mayor para especies con bajas abundancias y menor para especies dominantes.

Por último, esperamos que la presencia de los mamíferos terrestres que influyen a la comunidad de plántulas sea favorecida por la matriz circundante, la cual tendrá funciones de complementación del paisaje a escalas pequeñas y de conectividad funcional a escalas grandes.

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**CAPÍTULO II. CASCADING EFFECTS OF FIRE DISTURBANCE AND
LANDSCAPE CONFIGURATION ON SEEDLING DIVERSITY WITHIN
RAINFOREST PATCHES**

Cascading effects of fire disturbance and landscape configuration on seedling diversity within rainforest patches

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Abstract

Seedling community drives forest regeneration in fragmented rainforests. In these novel ecosystems, seedlings are influenced by the structure of patches (i.e., size, shape, vegetation structure and disturbance) and the surrounding landscape (i.e., composition and configuration). However, the attributes of seedling community respond differently to patch and landscape structure, whose direct or cascading effects on seedling composition and diversity are unknown. Given the hierarchical relationship among landscapes, patches and seedlings, we assessed the direct and cascading effects of patch and landscape structure on seedling community within 16 forest patches of a hyper-fragmented rainforest in southeastern Mexico. We used a multi-scale approach to determine the effects of landscape composition/configuration on patch structure. Then, we used piecewise structural equation models to assess the direct and cascading effects of patch and landscape structure on seedling abundance, composition and α - and β -diversity. Our results showed that seedling composition and β -diversity were affected by fire disturbance, whereas seedling richness was influenced by landscape configuration. On one hand, fire disturbance had cascading negative effects on seedling composition and β -diversity through positive effects on the abundance of light-demanding seedlings (i.e., promoting early-successional species). On the other hand, the aggregation of patches had cascading positive effects on seedling richness through positive effects on the abundance of shade-tolerant seedlings (i.e., sustaining seed dispersal). These findings support that patch-scale and landscape configuration effects are greater in hyper-fragmented landscapes, as well as the positive effects of fragmentation on biodiversity.

Keywords. abundance, alpha-diversity, beta-diversity, structural equation models, fragmentation, landscape structure

Introduction

Deforestation has reduced 10% of the remaining rainforests in small-sized patches (< 20 ha), which will dominate tropical landscapes in the next decades (Taubert et al., 2018). Nonetheless, these patches preserve forest species, ecological processes (i.e., pollination and seed dispersal/predation), and landscape connectivity (Laurance et al., 2011; Hernández-Ruedas et al., 2014; Cudney-Valenzuela et al., 2021), urging to promote their natural regeneration to sustain their biodiversity (Crouzeilles et al., 2016; Brancalion et al., 2019). In this sense, it is decisive to understand the abundance and diversity patterns of tree seedlings within patches, since the regeneration, structure and diversity of rainforests is largely determined by the tree seedling community (Brokaw & Scheiner, 1989; Whitmore, 1989; Dalling, Hubbell & Silvera, 1998; Brokaw & Busing, 2000).

The abundance, richness (α -diversity) and spatial heterogeneity (β -diversity) of forest seedling species are shaped by seed production and dispersal, as well as seedling mortality by density-dependence and predation/trampling of herbivorous mammals (Nathan & Muller-Landau, 2000; Harms et al., 2000; Comita et al., 2007; Young et al., 2016). Following forest fragmentation, the size and shape of patches drive the extent and magnitude of edge effects (Haddad et al., 2015), favoring the abundance of light-demanding seedlings – tree species with acquisitive traits commonly found in forest gaps and early-successional stages –, reduce the abundance of shade-tolerant seedlings – tree species with conservative traits dominant in undisturbed forests and late-successional stages – (Benitez-Malvido, 1998; Santo-Silva et al., 2013; Benítez-Malvido, Lázaro & Ferraz, 2018) and impoverish α - and β -diversity (Tabarelli, Peres & Melo, 2012; Krishnadas, Kumar & Comita, 2019; Liu et al., 2019; Krishnadas, Agarwal & Comita, 2020). Nonetheless, seedlings can also be affected by components of patch structure aside from its size and shape.

Tree basal area plays a key role on light availability, microclimate regulation, nutrient cycling and seed production (Hardwick et al., 2015; Lohbeck et al., 2015; Minor & Kobe, 2019; Wies, Nicasio Arzeta & Martínez Ramos, 2021), conditions that regulate the abundance of light-demanding seedlings and shade-tolerant seedlings (Whitmore, 1989; Denslow & Guzman G., 2000). Also, patches with a high basal area harbor a greater diversity of forest-dependent birds and mammals that favor seedling α - and β -diversity (Zurita & Bellocq, 2010; Wearn et al., 2017; Cudney-Valenzuela et al., 2021). Additionally, seedling abundance and composition can be influenced by tree diversity (Terborgh et al., 2001; Comita et al., 2007), particularly where seed input is limited to the surrounding adult trees (Wandrag et al., 2017). Furthermore, disturbance regimes by agricultural land uses affect forest regeneration in the tropics (Guariguata & Ostertag, 2001; Fontúrbel et al., 2015). The size, duration and severity of disturbance by agricultural and cattle management can limit seed rain, reduce soil fertility, and increase temperature and water deficit, leading to a reduced abundance and diversity of seedlings (Zermeño-Hernández et al., 2015). Such effects are higher in landscapes dominated by pastures (Zermeño-Hernández, Pingarroni & Martínez-Ramos, 2016), where cattle grazing and trampling limit seedling establishment. Moreover, these areas are subject to recurrent fires, which not only decrease soil fertility, but also reduce the diversity and basal area of the remaining forest (Barlow, Haugaasen & Peres, 2002).

Additionally, it is well documented that composition (i.e., proportion of primary forest and matrix covers in the landscape) and configuration (i.e., number/aggregation of patches and edge contrast between patches and matrix) of landscape structure have significant effects on seedling communities. Landscape structure can influence seedling abundance, composition and diversity whether by sustaining seed dispersal in landscapes where forest cover and patch density/aggregation are high (Jesus et al., 2012; San-José, Arroyo-Rodríguez & Meave, 2019), or by alleviating edge effects and facilitating the arrival of seed/seedling predators in landscapes

with low edge contrast (Arasa-Gisbert et al., 2021; Nicasio-Arzeta et al., 2021). It is noteworthy that landscape fragmentation (modulated by landscape configuration) have stronger effects on understory vegetation, and tree seedlings/saplings than habitat loss (modulated by landscape composition) in highly deforested rainforests (Nicasio-Arzeta et al., 2021; Arasa-Gisbert et al., 2022). Their findings support the “fragmentation threshold hypothesis”, which stipulates that patch-scale and landscape configuration have stronger effects on biodiversity within landscapes with < 30% of forest cover (Andren, 1994). Thus, the composition and diversity of tree seedlings from hyper-fragmented rainforests might be driven by the synergistic effect of patch structure and landscape configuration.

However, seedling community and patch structure are expected to be influenced by landscape structure at multiple spatial scales, since the scale of effect vary among biological variables (i.e., structure, abundance, composition, richness), habitat specialization (i.e., light-demanding/shade-tolerant) and disturbance extent (Miguet et al., 2016; Martin, 2018; Suárez-Castro et al., 2018). The latter suggests that landscape structure might have cascading effects on seedling communities, a process observed elsewhere (Hernández-Ruedas et al., 2018). These cascading effects are possible operating through patch-structural attributes. The proportion of forest cover in the landscape is directly related to patch size (Fahrig, 2013; Hernando et al., 2017) and tree basal area/diversity (Laurance et al., 2000; Wies, Nicasio Arzeta & Martinez Ramos, 2021; Brindis-Badillo et al., 2022). Alternatively, tree diversity is affected by the surrounding secondary forests and the aggregation of forest patches (Nascimento et al., 2006; Laurance et al., 2007; Jesus et al., 2012). Forest disturbance by fires and management intensity can also be modulated by edge contrast, forest cover and patch density. On one hand, edge exposure increases fuel availability and fire ignition (Román-Cuesta & Martínez-Vilalta, 2006; Avila-Flores et al., 2010; Farfán Gutiérrez et al., 2018). On the other hand, cattle presence is limited where forest cover or patch number are high, since they increase the occurrence of

large predators (Trolle, 2003; Desbiez, Bodmer & Aparecida, 2009; de Souza et al., 2018).

Although patch and landscape structure can potentially have direct and/or cascading effects on seedling community, their evaluation remains unexplored.

We aimed to identify the direct and cascading effects of patch and landscape structure on seedling abundance, composition and diversity in a fragmented rainforest. We hypothesized that patch structure and landscape configuration are exerting direct or cascading effects on seedling composition and diversity. We expect that seedling composition and β -diversity are more influenced by patch structure, since these community attributes are sensitive to abundance fluctuations of species with contrasting life-history traits (Comita et al., 2007; Santo-Silva et al., 2015). Hence, we expect a similar composition of a set of species and greater β -diversity within patches with lower abundances of light-demanding seedlings and higher abundances of shade-tolerant seedlings, fluctuations favored in large-sized, highly-diverse and low-disturbed patches. Additionally, we expect that seedling α -diversity will be favored in landscapes with high forest cover or high density/aggregation of patches, since species richness is strongly associated to large-scale processes that sustain seed dispersal (San-José et al., 2019; Arasa-Gisbert et al., 2022). These effects can occur whether by cascading effects of landscape structure by influencing patch structure (Fig 1, hypothetical pathway 1), or by cascading effects of patch structure by mediating abundance fluctuations (Fig 1, hypothetical pathway 2). We employed structural equation models to assess the direct and indirect effects of patch components (Fig S1).

Material and methods

Study site.

We conducted the study at the Marqués de Comillas region in the Lacandona rainforest in southeastern Mexico (Fig 2a). The majority of cleared areas in the study site are employed to raise cattle (Carabias, De la Maza & Cadena, 2015; Berget et al., 2021), and use of forest patches to provide shade for cattle is a common practice. The latter implies that some landowners slash the understory vegetation in forest fragments to facilitate the movement of cows and people along patches. Land clearing by fire is another practice in the area, and sometimes, mismanaged fires expand into patches, partially burning tree vegetation. Another disturbance factor in the study area is the periodic flooding of patches nearby the Lacantún river during rainy season.

We selected 16 forest patches (Fig 2b). Land tenure of Marqués de Comillas is private, and land property is distributed among the inhabitants (Carabias, De la Maza & Cadena, 2015; Kolb & Galicia, 2018; Berget et al., 2021). Therefore, we conducted our study only in patches where permission was granted beforehand by the owners. Patches were distributed over alluvial terraces, so diversity and composition of tree assemblages, soil type and abiotic factors remain relatively constant (Navarrete-Segueda et al., 2018; Lohbeck et al., 2022).

Seedling sampling.

Patches were sampled in 1-ha blocks placed at the center of each forest patch from February to June of 2014. Each block contained ten 1-m² plots randomly arranged in groups of two or three plots along five equidistant transects (20 m apart; Fig 2d). For 1-ha forest patches, we never positioned plots at the edge in order to have a 20-m buffer zone of vegetation that

protect them from strong edge effects as much as possible (Fig 2d). Within each 1-m² plot we counted and identified all tree seedlings (10-100 cm height) to the lowest possible taxonomic level with the help of a local parataxonomist and field guides (Martínez, Ramos-A. & Chiang, 1994; Sousa, 2009). When field identification was not possible, we took samples for identification at several herbariums (MEXU, ECO-SC-H). This study did not involve the extraction or damage of endangered species. Plant nomenclature followed the Missouri Botanical Garden database Tropicos (Missouri Botanical Garden, 2019). Once identified, seedlings species were classified by dispersal syndrome (Ibarra-Manríquez, Martínez Ramos & Oyama, 2001; Ibarra-Manríquez & Cornejo-Tenorio, 2010) and light requirements for establishment and growth: light-demanding species, shade-tolerant species and intermediate species (Nicotra, Chazdon & Iriarte, 1999; Rose, 2000; Comita et al., 2007; Kitajima, Cordero & Wright, 2013).

Seedling community.

We assessed the sampling completeness with the sample coverage estimator of Chao and Shen (Chao et al., 2016). We combined the data of the 10 sampling plots within each patch and then we estimated the proportion of the total number of individuals that belong to the species represented in the sample (Chao & Lee, 1992). The sample coverage among patches was of $91.07 \pm 7.47\%$, indicating that our sampling effort was adequate to estimate diversity (Chao & Jost, 2012).

We summed the seedlings of the 10 1-m² plots to obtain the abundance of light-demanding and shade-tolerant seedlings for each patch (seedlings · 10 m⁻²). Additionally, we employed a Nonmetric Multidimensional Scaling (NMDS) to estimate species compositional

similarity among patches (Laurance et al., 2007; Santo-Silva et al., 2015; Liu et al., 2019). We used the *metaMDS* function based on the Bray-Curtis dissimilarity index of the *vegan* package (Oksanen et al., 2019) and the whole seedling dataset. We used the NMDS scores of the first two axes as response variables.

We used the diversity decomposition of effective numbers of species or the Hill numbers (Jost, 2006, 2007). Hill numbers (qD) are expressed in units of species, which permits to characterize the distribution of species abundances and provide complete information about the community diversity (Jost, 2006; Chao, Chiu & Hsieh, 2012). We calculated Hill numbers of all species (species richness or 0D), typical species (the exponential of Shannon's entropy index or 1D), and dominant species (the inverse of Simpson's index or 2D). The 0D is not sensitive to individual abundances (Jost, 2007; Tuomisto, 2010), permitting to include rare species in assessments, while 1D and 2D give a high weight to the equally abundant and the most abundant species, respectively (Jost, 2007, 2010; Chao, Chiu & Hsieh, 2012).

We followed Jost (2007) and Tuomisto (2010) procedure to calculate the gamma (γ), alpha (α) and beta (β) diversity (Fig S2c and Table S2). The α -diversity values are expressed in species \cdot m⁻², whereas β -diversity is the effective number of completely distinct assemblages, ranging between one (when the assemblages of all 1-m² plots are identical) and 10 (when the assemblages of the 10 1-m² plots are completely different from each other; Table 1). We used the package *vegan* in R for the entire procedure (Oksanen et al., 2019).

Patch structure.

We employed a multispectral SPOT-5 satellite image of 10 × 10 m pixel resolution recorded in March 2013 to carry out a supervised classification using the GRASS GIS software

(GRASS Development Team, 2017). We elaborated a forest/non-forest classification and used sampling points to assess classification accuracy. The overall classification accuracy was 79%. We exported the resulting raster file into a shapefile to calculate patch perimeter in meters (m) and patch area in m^2 using the *rgeos* package (Bivand & Rundel, 2019). We employed these data to measure patch size in hectares and patch shape as a limitless index that measures patch complexity compared to a square of the same size (Patton, 1975). For patch shape, we controlled for patch size effects by using the residuals of a linear model between patch shape and size ($R^2 = 0.63$; $F_{1,14} = 26.32$; $P = 0.0001$).

Within each patch, we sampled tree community by recording the diameter at breast height (dbh, 130 cm above the ground) of all trees with $\text{dbh} \geq 10$ cm within ten 50×2 m transects (Gentry, 1982). When possible, we minimized the forest diversity and structural biases derived from edge effects by locating all transects at least 20 m from patch edges (Laurance et al., 2007). Within each transect, we counted, measured the diameter and identified each tree to species level (Martínez, Ramos-A. & Chiang, 1994; Missouri Botanical Garden, 2019). Then, we calculated tree abundance ($\text{trees} \cdot 0.1 \text{ ha}^{-1}$), basal area ($\text{m}^2 \cdot \text{ha}^{-1}$), and richness, exponential of Shannon's entropy index and the inverse of Simpson's index ($\text{species} \cdot 0.1 \text{ ha}^{-1}$).

Additionally, we gathered information regarding disturbance factors within patches through a questionnaire applied to landowners and field observations throughout the study period. We used the questionnaire results to elaborate a binary database of the following disturbance factors of patches (Table S3): flooding during rainy season, fire events during the last 10 years, presence of cattle, slashing of understory vegetation and the predominant land use in the adjacent matrix. Then, we employed a logistic PCA (Landgraf & Lee, 2020) to construct disturbance indices from binary data. We previously estimated the number of dimensions (k) and the tuning parameter (m) through a cross-validation of the negative Log-likelihood, and obtained $k = 3$ and $m = 3$ to perform the analysis (Fig S2). We final model contained the 77.8%

of deviance explained. We used the *logisticPCA* package for the entire procedure (Landgraf & Lee, 2015). According to Spearman correlation between disturbance factors and PCA components, the slash of understory vegetation, the adjacent land use and the presence of cattle were strongly associated to the first component, whereas flooding and fire events were associated to the second component (Table S4). Accordingly, we used the first component as the management disturbance index (D_M) and the second component as the fire disturbance index (D_F).

Landscape structure

We calculated five landscape metrics (Table S5) known to influence patch size, vegetation, and disturbance in the tropics, as well as seedling community. The composition metrics were the proportion of forest cover (FC) and secondary forest (SF), whereas the configuration metrics were patch density (PD), patch aggregation AI and edge contrast (EC). We calculated the EC by assigning quality values to each matrix cover. These values were indicators of matrix cover capacity to alleviate edge effects facilitate the movement of terrestrial mammals. The quality values were the following: 1 (water bodies, with the lowest suitability); 2 (anthropogenic cover); 3 (cattle pasture); 4 (arboreal crops); 5 (floodplains); 6 (secondary forest); and 7 (old-growth forest, representing the highest suitability). To obtain a more robust and realistic representation of landscape structure effects, we estimated the area-weighted mean of the EC index (Li & Archer, 1997).

We measured these landscape metrics within 13 circular buffers (300 to 1500 m radius, at 100 m intervals) from the center of each focal patch (Fig 2c). Radius were based in the dispersal distances of seed dispersers and terrestrial mammals, landscape size to assess seedling

community and disturbances, and spatial scales at which landscape structure affects forest diversity and structure in the study region (Zermeño-Hernández, Pingarroni & Martínez-Ramos, 2016; San-José, Arroyo-Rodríguez & Meave, 2019; Wies, Nicasio Arzeta & Martinez Ramos, 2021; Nicasio-Arzeta et al., 2021). Independence among sampling sites, landscape metrics and buffer sizes can be found in Nicasio-Arzeta et al. (2021).

Statistical analysis

We estimated the scale of effect of each landscape metric on patch structure using linear models. We fitted a patch size, shape, basal area, diversity and disturbance by fire and management with a single landscape metric for each buffer and obtained its coefficient of determination (R^2). Then, we plotted the resulting 13 R^2 values and selected the one with the strongest response as the scale of effect for that landscape metric. We repeated this procedure for each patch variable and landscape metric. Thereafter, we evaluated the effects of landscape metrics and fire disturbance on each patch structure variable through multiple linear models. We only used the landscape metrics at the scale of effect identified in the previous step (Table S6). All models were additive due to our limited sample size of 16 forest patches.

We employed the dredge function of the *MuMIn* package (Barton, 2018) to create all possible combinations of explanatory variables plus the null model (only the intercept) to assess the relative importance and effects of landscape metrics and fire disturbance using an information theory approach and multimodel inference (Burnham & Anderson, 2002). We used the accumulated sum of Akaike weights (w_i) of each model to select the subset of models that $\sum w_i \leq 0.95$. Then, we employed the w_i of the model's subset to calculate the relative importance and the model-averaged parameter estimates of each explanatory variable. We considered as

influential variable those for which the confidence interval did not include zero in the averaged parameters. We carried out all statistical analysis in the R 3.5.2 statistical computing environment (R Development Core Team, 2018). We selected the significant landscape metrics and disturbance factor for each patch structural variable for further analysis (Table S7).

We analyzed the direct and indirect effects of patch and landscape structure on seedling abundance, composition and diversity through structural equation models (SEM). SEM is a statistical method that test for causal relationships (Fan et al., 2016; Tarka, 2018). We proposed a conceptual model of multiple casualties to explain seedling abundance, diversity and composition in relation to the patch components (size, shape, basal area, tree diversity, and disturbance by fire and management), which served as the exogenous predictors (Fig S1). Indirect paths included the abundance of light-demanding and shade-tolerant seedlings, representing the hypothesis regarding cascading effects mediated by abundance fluctuations. We previously assessed the association among explanatory variables to avoid variance inflation. We found that patch structural variables were strongly correlated (Table S8), and only tree basal area and tree diversity were employed for further analysis.

We fitted the SEM using the *piecewiseSEM* package (Lefcheck, 2016). Multivariate normality of seedling ($\gamma_{1,p} = 217.39$; $P = 0.54$ and $\gamma_{2,p} = -1.95$; $P = 0.052$) and patch data ($\gamma_{1,p} = 67.41$; $P = 0.14$ and $\gamma_{2,p} = -1.03$; $P = 0.30$) was validated. We applied a logarithmic transformation to patch size and abundance of light-demanding seedlings, and a square root transformation to abundance of shade-tolerant seedlings to meet normality assumptions. We performed simple model structures (few variables) due to sample size limitations. Hence, we tested different combinations of patch structure (D_M , T_{BA} , T_D , P_{SZ} and P_{SI}) in alternative models instead of all together. We employed the fire disturbance index and the selected landscape metrics (D_F , FC_{1200} , PD_{600} , PD_{900} , AI_{600}) to explain variation of exogenous variables. Alternative models consisted in combinations of one and two exogenous predictors. This produced 15

alternative models for each diversity order (all species: ${}^0\alpha$, ${}^0\beta$; typical species: ${}^1\alpha$, ${}^1\beta$; dominant species: ${}^2\alpha$, ${}^2\beta$) and composition index (MDS1, MDS2). Model selection procedure consisted in rejecting all models with lack-of-fit ($P < 0.05$), then, we excluded the models that had no significant links to seedling abundance, composition or diversity variables we attempted to explain. Finally, we selected the best-fitting models based on the lowest AICc values.

Results

We recorded 1378 seedlings belonging to 27 families, 38 genera and 67 species. We only considered animal-dispersed species for further analysis because they comprise up to 90% of the seed rain in rainforests (Jordano, 2000; San-José et al., 2019). The 94.7% of seedlings (1305) were animal-dispersed, represented in 24 families, 30 genera and 56 species. From this subset, we found 668 seedlings (52.72%) from 27 shade-tolerant tree species, followed by 527 seedlings (40.38%) from 18 light-demanding species and 90 seedlings (6.9%) from 11 intermediate or indetermined species (Table S1).

Our candidate models were good-fitted, indicating that the conceptual model adequately described the data (Fig 1). The best-fitted models (those with the lowest AICc; Table S9) included patch size and aggregation for diversity of all species (Fig 3b), whereas tree basal area explained seedling composition (Fig 3a) and diversity of typical and dominant species, and (Fig 3c and d). For the latter models, tree basal area decreased in patches affected by previous fire events ($B = -0.63$), and light-demanding seedling abundance was reduced by tree basal area ($B = -0.54$). The latter indicated that fire disturbance had indirect effects in the increase of light-demanding seedlings ($B = -0.63 \times 0.54 = 0.34$).

Seedling composition model explained 39% of basal area, 30% of light-demanding seedling abundance, 12% of shade-tolerant seedling abundance, and 64% and 56% of the first and second NMDS axis, respectively (Fig 3a). The first axis was negatively influenced by light-demanding seedlings ($B = -0.55$), whereas the second NMDS axis was negatively affected by the abundance of shade-tolerant-seedlings ($B = -0.76$). The indirect effect of fire disturbance on the first composition axis was lower (i.e., $-0.63 \times -0.55 \times 0.54 = -0.18$) than the indirect effect of basal area (i.e., $-0.55 \times 0.54 = 0.29$). Seedling composition differed between fire-disturbed and non-disturbed patches ($F_{1,14} = 2.64$; $R^2 = 0.16$; $P = 0.01$), where the abundance of light-demanding seedlings was higher in fire-disturbed patches than in their non-disturbed counterparts (Fig 4).

The typical species model (Fig 3c) explained 40% of α -diversity, and was positively influenced only by the abundance of shade-tolerant seedlings ($B = 0.54$), whereas the 68% explained of β -diversity variation was strongly attributed to the negative effect of light-demanding seedlings ($B = -1$), and indirectly promoted by tree basal area (i.e., $-1 \times 0.54 = 0.54$). This also held true for the dominant species model (Fig 3d), which explained 69% of β -diversity through the negative effects of light-demanding ($B = -0.94$) and shade-tolerant ($B = -0.43$) seedling abundances. Tree basal area also had a positive indirect effect on β -diversity of dominant species (i.e., $-0.94 \times 0.54 = 0.50$). Finally, fire disturbance had overall indirect effects on the decline of β -diversity of typical ($B = -1 \times -0.54 \times -0.63 = 0.34$) and dominant species ($B = -0.94 \times -0.54 \times -0.63 = 0.32$).

Finally, the model for all species explained 40% of patch size, 10% of light-demanding seedlings, 51% of shade-tolerant seedlings, 50% of α -diversity and 17% of β -diversity (Fig 3b). Patch aggregation had direct effects on patch size ($B = 0.64$) and shade-tolerant seedlings ($B = 0.63$), which was also affected by light-demanding seedlings ($B = -0.58$). In turn, shade tolerant seedlings had positive effects on α -diversity ($B = 0.56$), which was indirectly reduced by the

abundance of light-demanding seedlings ($B = -0.58 \times 0.56 = -0.32$). Patch aggregation had cascading effects on the increase of seedling α -diversity ($B = -0.63 \times 0.54 = 0.34$).

Discussion

Our findings confirm that landscape configuration and disturbance factors within patches, such as fire incidence, defined species composition and α - and β -diversity of typical and dominant species of seedling community. As expected, patches with low disturbance regimes and higher basal area reduced the abundance of light-demanding seedlings, which in turn led to a more diversified and complex regenerative community. Also, landscapes with more aggregated patches increased the abundance of shade-tolerant seedlings with cascading effect that favored α - diversity of species in the regenerative community. Thus, cascading effects of patch disturbance and landscape configuration shaped species composition and α - and β -diversity, respectively.

Disturbance by fire was the main local factor having major cascading effects that modulated the abundance, composition and diversity of the regenerative community. Fire is an atypical natural disturbance in tropical rainforests where most of their tree species are considered fire-sensitive (Rodriguez Trejo 2008). Nevertheless, land-use changes to pastures and farmlands, forest fragmentation and extreme weather events are increasing fire frequency, severity, and extent (Cochrane & Laurance, 2008). In this study, fire occurrence, which was present in almost 50% of forest patches, significantly reduced the basal area of adult trees leading to an increased abundance of light-demanding species in the understory. This is consistent with other studies where surface fires reduced basal by the loss of medium- and large-sized trees and lianas (Nepstad et al., 1999; Barlow et al., 2003; Cochrane & Laurance,

2008). The consequent canopy openness -which can be four times greater than in unburned forests-, lead to an increased light incidence in the forest floor (Barlow et al., 2003). These microclimate alterations favor the dominance of disturbance-loving tree species in the understory vegetation (Cochrane & Laurance, 2002), as was found here. In turn, the abundance of light-demanding seedlings had direct effects on seedling composition and the turnover of equally common and abundant species. In this sense, patches that had fires during the last 10 years, had a different seedling community from that found in patches without fires, which was dominated mainly by light-demanding species such as *Inga punctata*. Some tropical rain forest tree species are reported to respond well to fire, for example *Swietenia macrophylla* which is promoted by fire species reported for tropical rain forests of southern Mexico. However, the vast majority of tree species in tropical forests are fire-sensitive. Then, the presence of species like *Brosimum alicastrum* and *Ampelocera hottlei*, found exclusively in unburned patches dominated by shade tolerant, could be threatened by the occurrence of fires.

In our study, basal area only affected the abundance of light-demanding seedlings, suggesting that disturbed fragments (lower abundance of large trees) promote the establishment and survival of these species by increasing light availability (Denslow, 1987, 1996). These conditions promote the establishment of a subset of species that are ecologically redundant, that lack phylogenetic relationship and that give rise to impoverished species assemblages dominated by few species with high abundances (Oliveira, Santos & Tabarelli, 2008; Tabarelli, Peres & Melo, 2012). These assemblages indicate a process of biotic homogenization, characterized by low community composition (i.e., less dissimilarity between communities), as well as a reduction in β -diversity (i.e., less species turnover) at local and landscape scales (Olden & Rooney, 2006; Tabarelli, Peres & Melo, 2012; Solar et al., 2015). Basal area influences the abundance and composition of seedlings through fruit production (Chapman et

al., 1992), the availability of light, water and nutrients in the soil (Denslow, 1987; Martinez-Ramos & Soto-Castro, 1993; Turner, 2004) and conspecific seedling mortality (Wright, 2002).

This is consistent with our observations, where a greater abundance of light-demanding seedlings reduced both the composition of seedlings and the β -diversity of typical and dominant species, variables that are sensitive to abundance fluctuations (Jost, 2007; Roden et al., 2018). In addition, the absence of direct effects by the landscape structure supports the hypotheses that establish that generalist species and populations with a high density of individuals are mainly affected by factors at small scales (Miguet et al., 2016). This supports the cascading effects of landscape structure on the abundance of a light-competing palm, which represented the local factor that had the greatest impact on biological responses affected at small scales, such as the density of species and individuals (Hernández-Ruedas et al., 2018).

On the contrary, the abundance of shade-tolerant seedlings was mainly explained by the aggregation of fragments in the landscape, which supports the hypotheses of Miguet et al. (2016) mentioned above. Shade-tolerant seedlings have higher habitat specialization than light-demanding species, as well as lower population densities (Denslow, 1996; Turner, 2004). In addition, their abundances are regulated by the conspecific mortality of seedlings and the foraging of fruits, seeds, and seedlings by terrestrial mammals (Dirzo & Miranda, 1990; Comita et al., 2007, 2010; Camargo-Sanabria et al., 2014). This suggests that landscape structure affects the abundance of shade-tolerant seedlings by facilitating or limiting the arrival of seed dispersers and seed/seedling predators.

This assumption is supported by studies that found a significant contribution of the percentage of primary forest in the landscape on the richness and abundance of seed rain and saplings of specialist tree species of biotic dispersal (San-José, Arroyo-Rodríguez & Meave, 2019; Arasa-Gisbert et al., 2021). These studies concluded that the loss of shade-tolerant species is the result of a lack of seed dispersal in landscapes where low forest cover limits the

movement of fauna, which disperses 90% of tropical forest tree species (Jordano, 2000). However, the size of the home range of fauna can vary depending on the aggregation of fragments in landscapes with a low percentage of habitat (Radford, Bennett & Cheers, 2005; Hodgson et al., 2012; Gardiner et al., 2019).

In this work, we found that the abundance of shade-tolerant seedlings increased in landscapes with high fragment aggregation, suggesting that this configurational pattern promotes the arrival of seed dispersers and predators of seeds and seedlings. The functions of dispersal and predation, together with the conspecific mortality of seedlings that promote their diversity (Harms et al., 2000; Wright, 2002; Comita et al., 2010), explain the positive effect that the abundance of seedlings tolerant to the shadow has on the α diversity of all seedling species. Interestingly, the abundance of shade-tolerant seedlings was affected by the abundance of light-demanding seedlings, suggesting that the latter exert competitive pressure for resources. Competitive exclusion may play an important role in reducing α diversity within forests with high light availability and low herbivore pressure (Wright, 2002). This supports the indirect effect that the abundance of light-demanding seedlings had on species richness.

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Table 1. Summary of response and explanatory variables of the study

Variable	Component	Name	Range	Label
Seedling	Abundance	Light-demanding seedlings	0 – 109	S _{LD}
		Shade-tolerant seedlings	4 – 130	S _{ST}
	Diversity	α -diversity of all species	1.60 – 4.11	${}^0\alpha$
		β -diversity of all species	3.33 – 6.12	${}^0\beta$
		α -diversity of typical species	1.45 – 3.60	${}^1\alpha$
		β -diversity of typical species	1.25 – 3.90	${}^1\beta$
		α -diversity of dominant species	1.34 – 2.98	${}^2\alpha$
		β -diversity of dominant species	0.94 – 3.11	${}^2\beta$
Patch	Spatial	Size	1.96 – 71.13	P _{SZ}
		Shape	1.40 – 5.30	P _{SI}
	Vegetation	Tree basal area	20.04 – 186	T _{BA}
		Tree abundance	33 – 74	T _A
		Number of adult tree species	10 – 27	T _{0D}
		Number of typical adult tree species	6.12 – 20.1	T _{1D}
		Number of dominant adult tree species	4.3 – 16.3	T _{2D}
	Disturbance	Fire index	-5.28 – 5.57	D _F
		Management index	-5.99 – 5.9	D _M
Landscape	Composition	Forest cover (1200 m radius)	1.83 – 3.49	FC ₁₂₀₀
	Configuration	Patch density (600 m radius)	0.89 – 6.2	PD ₆₀₀
		Patch density (900 m radius)	1.97 – 5.9	PD ₉₀₀
		Patch aggregation (600 m radius)	89.56 – 97.56	AI ₆₀₀

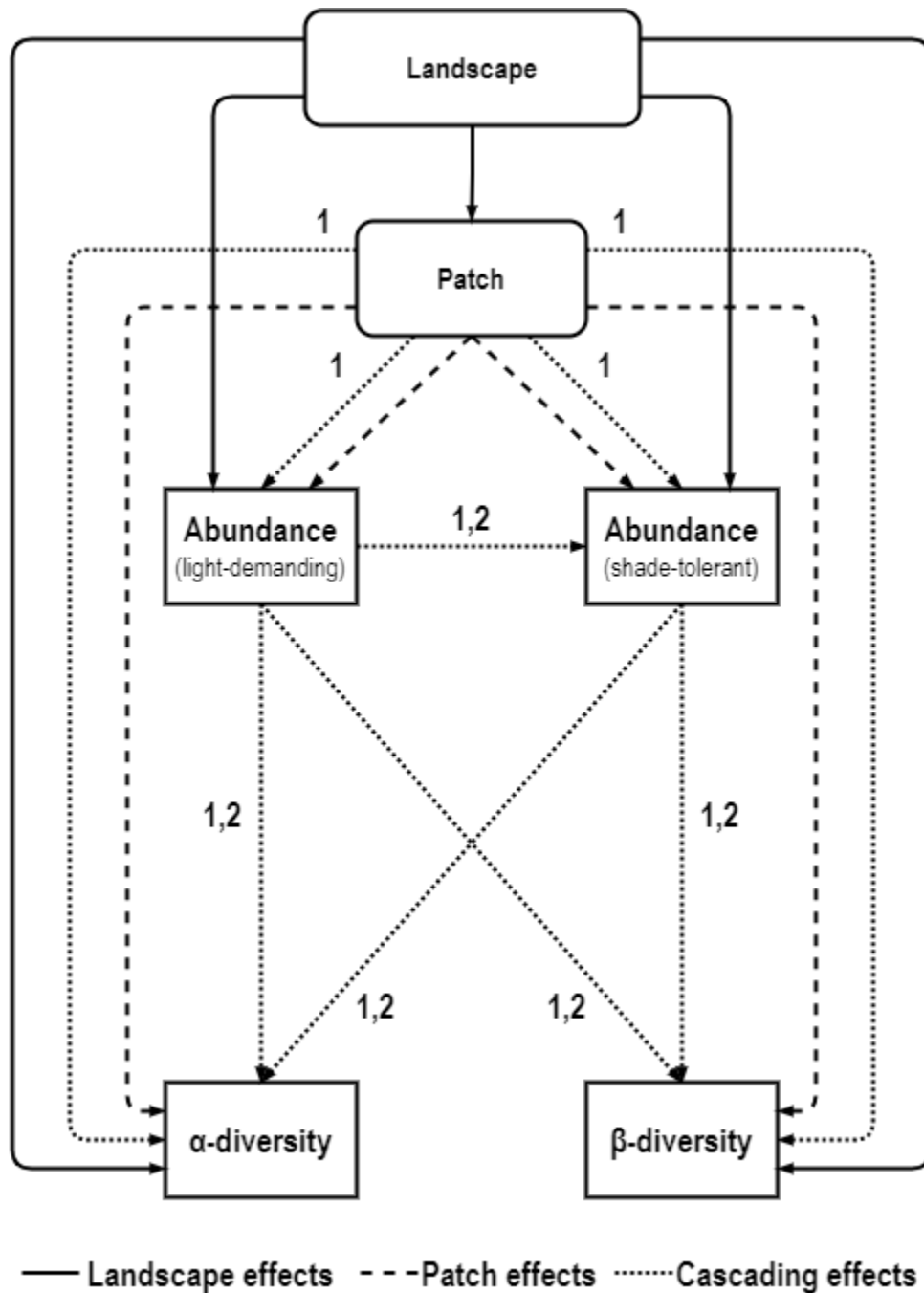


Fig 1. Conceptual model employed to assess the association among landscape (i.e., composition and configuration) and patch structure (i.e., size, shape, tree basal area, tree diversity, and disturbance), and seedling community (i.e., abundance and diversity). The hypothesis evaluated in this study are presented in two hypothetical pathways: direct effects of landscape/patch and cascading effects whether by landscape structure through patches (1) or by patch structure through seedling abundances (2).

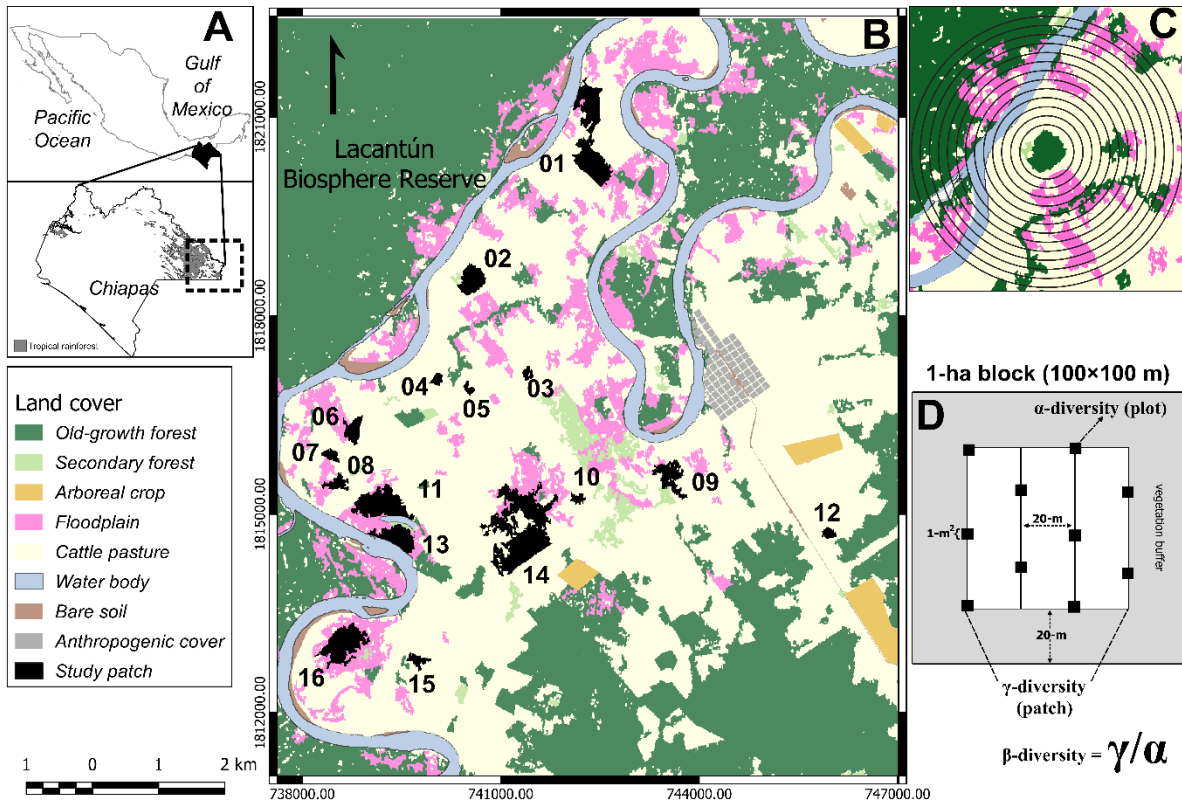
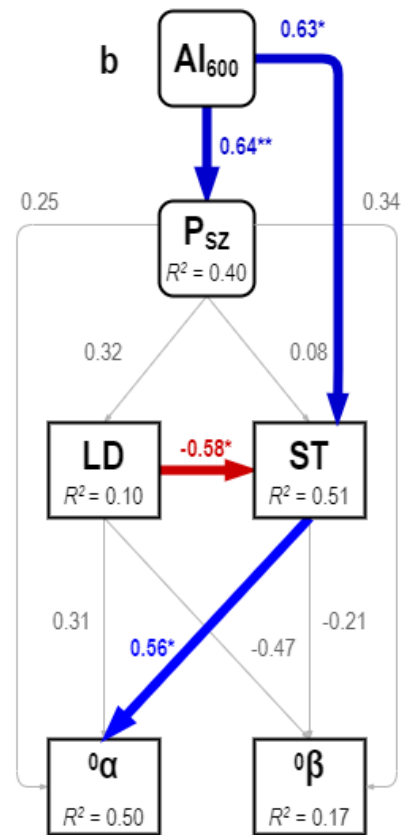
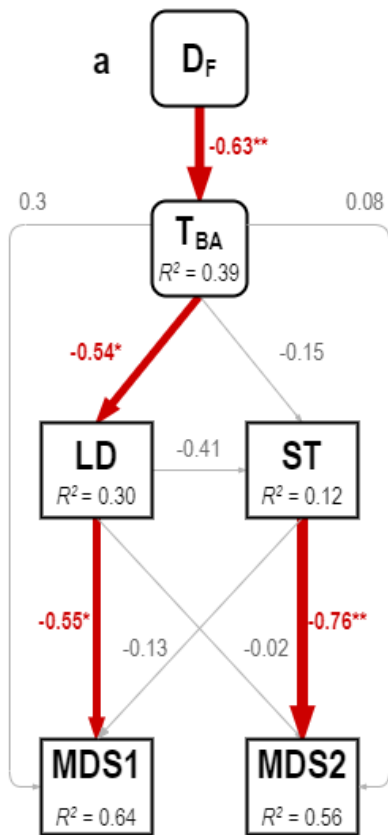


Fig 2. Location of the study area in the Lacandona rain forest in Chiapas, southeastern Mexico (a). We show the location of the 16 study forest fragments (black) in the Marqués de Comillas Region (b). The 13 buffer sizes (300-1500ha) around the geographic center of the focal patch (c) and the seedling sampling procedure (d) are also indicated.



█ Positive
█ Negative
█ Non-significant

* $P < 0.05$
 ** $P < 0.01$
 *** $P < 0.001$

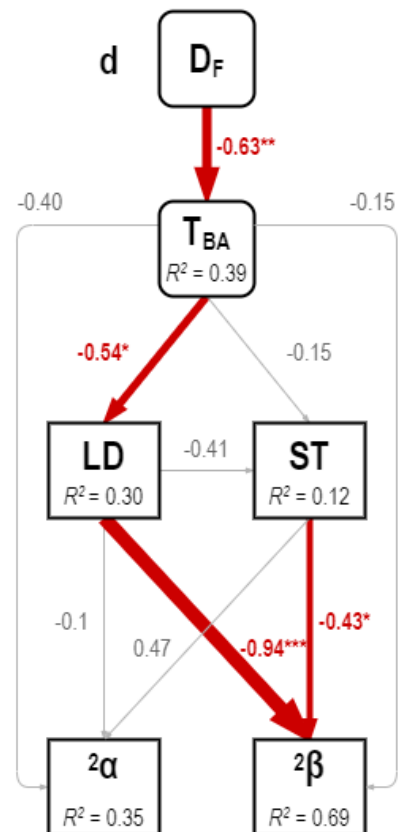
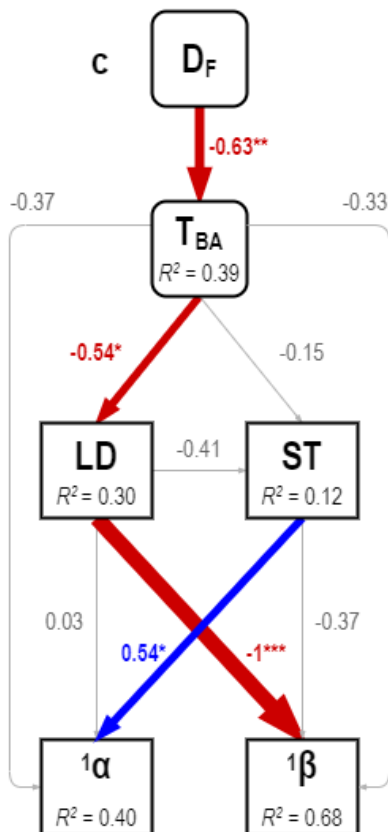


Fig 3. Best-fitted structural equation models for seedling (a) composition ($df = 12$), and α - and β -diversity of (b) all species ($df = 10$), (c) typical species ($df = 10$) and (d) dominant species ($df = 10$) in 16 forest patches in the Lacandona rainforest, southeast Mexico. Arrow thickness are proportional to the standardized path-coefficient value. The coefficients of determination of linear models (R^2) are reported. Patch size (P_{SZ}), tree basal area (T_{BA}), disturbance by fire (D_F), patch aggregation (AI), as well as the abundances of light-demanding (LD), and shade-tolerant seedlings (ST), compositional axes from the NMDS analysis ($MDS1$ and $MDS2$), and α - and β -diversity of all species (${}^0\alpha$, ${}^0\beta$) typical species (${}^1\alpha$, ${}^1\beta$) and dominant species (${}^2\alpha$, ${}^2\beta$) are shown.

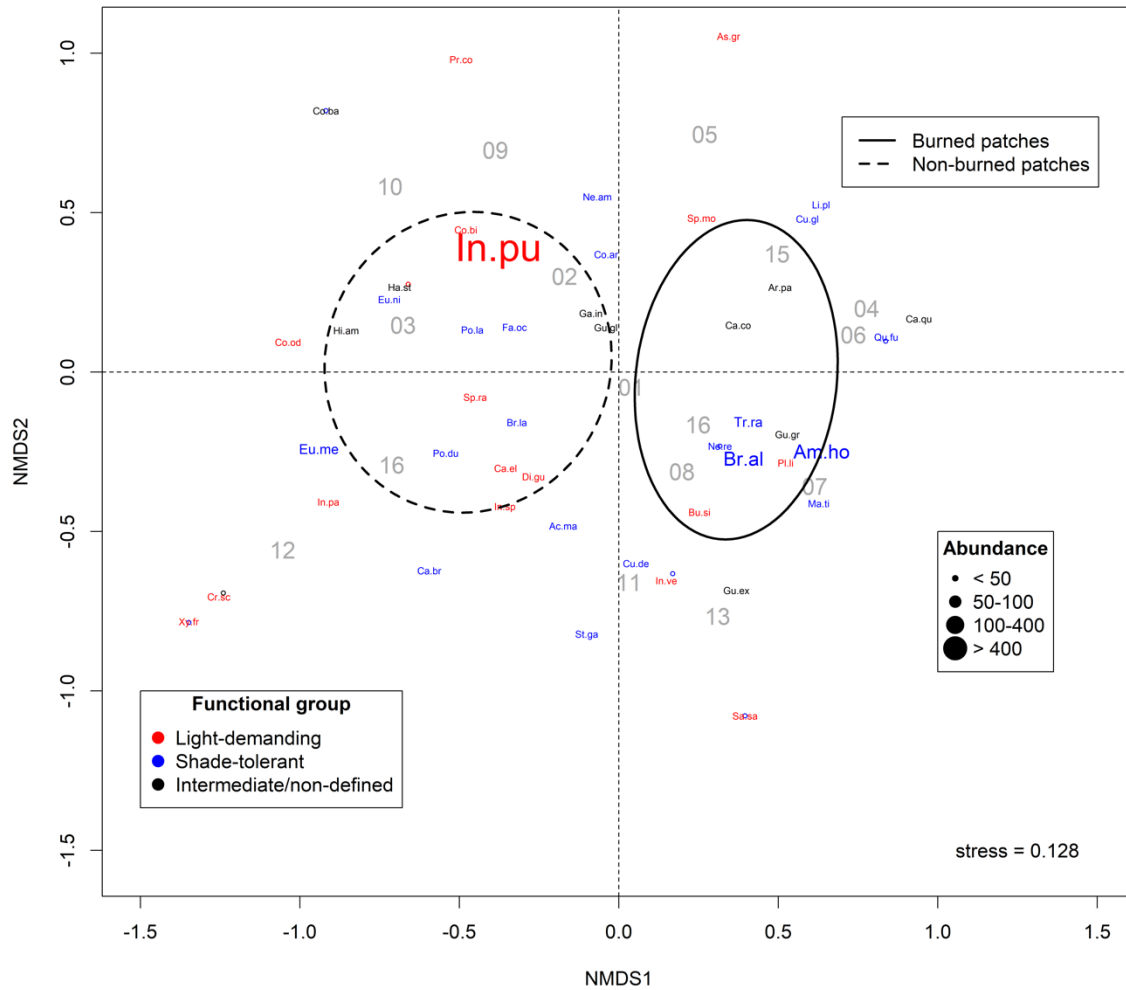


Fig 4. NMDS ordination of seedling assemblages of 16 forest patches (grey numbers) in the Lacandona rainforest, southeast Mexico. Abbreviations are seedling species (Table S1).

Supporting information

Table S1. Seedling species recorded in 16 forest patches in the Lacandona rainforest, southeast Mexico.

Family	Specie	Abbreviation	Light	Dispersion	Abundance
ACANTHACEAE	<i>Bravaisia integririma</i>	Br.in	L	A	29
	<i>Astronium graveolens</i>	As.gr	L	B	1
ANACARDIACEAE	<i>Spondias mombin</i>	Sp.mo	L	B	7
	<i>Spondias radlkoferi</i>	Sp.ra	L	B	4
ANNONACEAE	<i>Annona sp</i>	An.sp	S	B	4
	<i>Xylopia frutescens</i>	Xy.fr	L	B	1
APOCYNACEAE	<i>Stemmadenia galeottiana</i>	St.ga	S	B	2
BIGNONIACEAE	<i>Tabebuia rosea</i>	Ta.ro	S	A	1
BORAGINACEAE	<i>Cordia bicolor</i>	Co.bi	L	B	2
	<i>Cordia odorata</i>	Co.od	L	B	4
BURSERACEAE	<i>Bursera simaruba</i>	Bu.si	L	B	1
	<i>Protium copal</i>	Pr.co	L	B	7
CAPPARACEAE	<i>Capparis quiringuensis</i>	Ca.qu	I	B	1
	<i>Hirtella americana</i>	Hi.am	I	B	15
CHRYSOBALANACEAE	<i>Licania hypoleuca</i>	Li.hy	I	B	8
	<i>Licania platypus</i>	Li.pl	S	B	1
CLUSIACEAE	<i>Calophyllum brasiliense</i>	Ca.br	S	B	10
	<i>Garcinia intermedia</i>	Ga.in	I	B	4
EUPHORBIACEAE	<i>Croton schiedeanus</i>	Cr.sc	L	B	11
	<i>Acacia mayana</i>	Ac.ma	S	B	12
FABACEAE	<i>Albizia leucocalyx</i>	Al.le	S	A	5
	<i>Andira inermis</i>	An.in	S	B	4

	<i>Cojoba arboreum</i>	Co.ar	S	B	4
	<i>Dialium guianense</i>	Di.gu	L	B	10
	<i>Inga dulis</i>	In.du	L	B	3
	<i>Inga pavoneana</i>	In.pa	L	B	1
	<i>Inga punctata</i>	In.pu	L	B	418
	<i>Inga sp</i>	In.sp	L	B	4
	<i>Inga vera</i>	In.ve	L	B	2
	<i>Lonchocarpus cruentus</i>	Lo.cr	L	A	3
	<i>Lonchocarpus pubescens</i>	Lo.pu	L	A	8
	<i>Platymiscium yucatanum</i>	Pl.yu	I	A	10
	<i>Pterocarpus hayesii</i>	Pt.ha	S	A	1
	<i>Pterocarpus yucatanum</i>	Pt.yu	S	A	1
ICACINACEAE	<i>Calatola costaricanum</i>	Ca.co		B	3
LASCISTEMATAACEAE	<i>Lacistema aggregatum</i>	La.ag	S	B	2
	<i>Licaria capitata</i>	Li.ca	S	B	8
LAURACEAE	<i>Nectandra ambigiens</i>	Ne.am	S	B	17
	<i>Nectandra reticulata</i>	Ne.re	S	B	5
	<i>Hampea stipitata</i>	Ha.st	I	B	4
MALVACEAE	<i>Luehea seemannii</i>	Lu.se	L	A	4
	<i>Quararibea funebris</i>	Qu.fu	S	B	5
	<i>Cederela odorata</i>	Ce.od	L	A	3
	<i>Guarea excelsa</i>	Gu.ex	I	B	9
MELIACEAE	<i>Guarea glabra</i>	Gu.gl	I	B	10
	<i>Guarea grandifolia</i>	Gu.gr	I	B	5
	<i>Trichilia havanensis</i>	Tr.ha	S	B	3
MORACEAE	<i>Brosimum alicastrum</i>	Br.al	S	B	177
	<i>Brosimum latescens</i>	Br.la	S	B	20

	<i>Castilla elastica</i>	Ca.el	L	B	32
	<i>Maclura tinctoria</i>	Ma.ti	S	B	3
	<i>Trophis racemosa</i>	Tr.ra	S	B	94
	<i>Eugenia edulis</i>	Eu.ed	S	B	4
MYRTACEAE	<i>Eugenia mexicana</i>	Eu.me	S	B	56
	<i>Eugenia nigrita</i>	Eu.ni	S	B	5
POLYGONACEAE	<i>Coccoloba barbadensis</i>	Co.ba	I	B	1
PRIMULACEAE	<i>Ardisia paschalis</i>	Ar.pa	I	B	30
	<i>Faramea occidentalis</i>	Fa.oc	S	B	20
RUBIACEAE	<i>Posoqueria latifolia</i>	Po.la	S	B	18
SALICACEAE	<i>Pleuranthodendron lindenii</i>	Pl.li	L	B	18
	<i>Cupania dentata</i>	Cu.de	S	B	8
SAPINDACEAE	<i>Cupania glabra</i>	Cu.gl	S	B	15
	<i>Sapindus saponaria</i>	Sa.sa	L	B	1
	<i>Chrysophyllum mexicanum</i>	Ch.me	S	B	10
SAPOTACEAE	<i>Pouteria durlandii</i>	Po.du	S	B	5
ULMACEAE	<i>Ampelocera hottlei</i>	Am.ho	S	B	176
VOCHYSIACEAE	<i>Vochysia guatemalensis</i>	Vo.gu	S	A	8

Table S2. Formulas employed to estimate Hill numbers of seedling α - and β -diversity in the Lacandona rainforest, southeast Mexico.

Diversity	Order	Formula
Gamma	${}^0D_\gamma$	${}^qD_\gamma = \left(\sum_{i=1}^S \bar{p}_i^q \right)^{\frac{1}{(1-q)}}$
	${}^2D_\gamma$	
	${}^1D_\gamma$	${}^1D_\gamma = \exp \left(- \sum_{i=1}^S \bar{p}_i \log \bar{p}_i \right)$
Alpha	${}^0D_\alpha$	${}^qD_\alpha = \left(\frac{1}{N} \sum_{i=1}^S p_{i1}^q + \frac{1}{N} \sum_{i=1}^S p_{i2}^q + \dots \right)^{\frac{1}{(1-q)}}$
	${}^2D_\alpha$	
	${}^1D_\alpha$	${}^1D_\alpha = \exp \left\{ - \frac{1}{N} \left(\sum_{i=1}^S (p_{i1}^q \ln p_{i1}^q) + \sum_{i=1}^S (p_{i2}^q \ln p_{i2}^q) + \dots \right) \right\}$
Beta	${}^0D_\beta$	${}^qD_\beta = \frac{{}^qD_\gamma}{{}^qD_\alpha}$
	${}^1D_\beta$	
	${}^2D_\beta$	

where p_i denotes the relative abundance of the i th species in each of the N 1-m² plots, \bar{p}_i denotes the mean relative abundance of the i th species in the N 1-m² plots and q is a parameter that determines the sensitivity of the measure to the relative abundances (Jost, 2007; Tuomisto, 2010).

References:

- Jost, L. (2007). Partitioning diversity into independent alpha and beta components. *Ecology*, 88(10), 2427–2439. <https://doi.org/10.1890/06-1736.1>
- Tuomisto, H. (2010). A diversity of beta diversities: Straightening up a concept gone awry. Part 1. Defining beta diversity as a function of alpha and gamma diversity. *Ecography*, 33(1), 2–22. <https://doi.org/10.1111/j.1600-0587.2009.05880.x>

Table S3. Binary data of disturbance factors of the 16 forest patches in the Lacandona rainforest, southeast Mexico.

Patch	Flood	Fire	Slash	Cow	Use
1	0	1	0	0	1
2	1	0	1	1	1
3	0	0	1	1	1
4	1	1	0	1	1
5	1	1	1	1	1
6	0	0	1	1	1
7	1	0	1	1	1
8	0	0	1	1	0
9	0	1	0	0	0
10	1	1	1	0	0
11	0	0	0	0	0
12	1	0	0	0	1
13	0	0	0	0	0
14	1	0	0	0	0
15	1	1	0	1	0
16	1	1	0	1	1

The disturbance factors are the following: flooding during rainy season (*Flood*: 0 = no; 1 = yes), fire events during the last 10 years (*Fire*: 0 = no; 1 = yes), presence of cattle (*Cow*: 0 = cattle do not enter to the patch; 1 = cattle roams within the patch), understory vegetation (*Slash*: 0 = non-slashed; 1 = slashed) and the predominant land use in the adjacent matrix (*Use*: 0 = temporal agriculture; 1 = cattle pasture)

Table S4. Spearman correlation matrix between disturbance factors and logistic PCA components.

	Flood	Fire	Slash	Use	Cow	PC1	PC2	PC3
Flood	1							
Fire	0.27	1						
Slash	0.016	-0.27	1					
Use	0.238	0.016	0.27	1				
Cow	0.238	0.016	0.524*	0.492	1			
PC1	0.397	-0.123	0.699**	0.726**	0.863***	1		
PC2	0.644**	0.863***	-0.425	0.151	0.014	-0.027	1	
PC3	-0.452	0.069	-0.206	0.617*	0.069	0.062	-0.024	1

* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$

Table S5. Abbreviation and description of the landscape metrics employed in the analysis

Landscape metric	Abbreviation	Formula	Description
Aggregation index	AI	$AI = 100 \times \left[\frac{g_{ii}}{\max \rightarrow g_{ii}} \right]$	Where g_{ii} is the number of joins between pixels of forest patch i and $\max \rightarrow g_{ii}$ is the maximum number of joins between pixels of forest patch i based on the single-count method
Edge contrast index	EC	$EC = 100 \times \left[\frac{\sum_{k=1}^m p_{ijk} \cdot d_{ik}}{p_{ij}} \right]$	Where p_{ijk} is the edge length (m) of patch ij adjacent to cover type k , d_{ik} is the edge contrast weight between patch i and cover k , and p_{ij} is the length of perimeter of patch ij
Forest cover	FC	$FC = 100 \times \left[\frac{\sum_{j=1}^n a_i}{A} \right]$	Where a is the area (m ²) of forest patch i and A the is the total landscape area
Secondary forest	SF	$SV = 100 \times \left[\frac{\sum_{j=1}^n a_i}{A} \right]$	Where a is the area (m ²) of cover patch i and A the is the total landscape area
Patch density	PD	$PD = 100 \times \left[\frac{n_i}{A} \times (10,000) \right]$	Where n_i is the total number of patches in the landscape of the land cover i and A is the total landscape area (m ²)

Table S6. Landscape metrics selected from the multi-scale assessment for each patch structural variable

Patch structure	Landscape metric	Radius (m)	R ²
D_F	PD	600	0.23
D_M	FC	1200	0.59
	PD	900	0.44
T_{BA}	EC	1100	0.21
T_{1D}	FC	700	0.27
P_{SZ}	PD	1100	0.22
	AI	600	0.36
P_{SI}	PD	900	0.23

Table S7. Relative importance and significance of disturbance and landscape variables for each patch structural variable

Patch	Variable	β (SE)	$\sum w_i$
DF	PD₆₀₀	-0.532 (0.528)	0.754
D_M	FC₁₂₀₀	-0.611 (0.424)	1
	PD₉₀₀	-0.41 (0.38)	0.827
	P _{SZ}	-0.138 (0.381)	0.127
T_{BA}	D_F	-0.575 (0.494)	0.919
	EC ₁₁₀₀	-0.394 (0.493)	0.552
	P _{SZ}	-0.032 (0.517)	0.114
T_{1D}	D _F	0.306 (0.551)	0.248
	FC ₇₀₀	0.486 (0.563)	0.596
	P _{SZ}	0.329 (0.661)	0.222

P_{SZ}	Al₆₀₀	0.611 (0.516)	0.834
	D _F	0.223 (0.507)	0.308
	PD ₁₁₀₀	0.396 (0.653)	0.202
P_{SI}	D_F	0.541 (0.452)	1
	PD₉₀₀	-0.473 (0.435)	0.808

Table S8. Pearson correlation matrix among explanatory variables

	D _F	D _M	T _{BA}	T _A	T _{0D}	T _{1D}	T _{2D}	P _{SZ}
D _M	0.001							
T _{BA}	-0.63**	0.35						
T _A	-0.25	0.27	0.73**					
T _{0D}	0.014	0.23	0.32	0.49				
T _{1D}	0.31	0.16	-0.09	0.14	0.88***			
T _{2D}	0.46	0.06	-0.31	-0.05	0.72**	0.95***		
P _{SZ}	0.23	-0.46	-0.14	-0.12	0.25	0.41	0.49	
P _{SI}	0.58*	-0.21	-0.13	0.31	0.28	0.35	0.40	0

* $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$.

Table S9. Best-fitted structural equation models for seedling composition and diversity in 16 forest patches in the Lacandona rainforest, southeast Mexico.

Model	Exogenous variables	χ^2	df	P	AICc	Δi
Seedling composition	T _{BA}	15.78	10	0.11	-178.5	0
	D _M	22.6	16	0.12	-152.24	26.26
	P _{SZ}	8.54	8	0.38	-134.78	43.72
	P _{SZ} , P _{SI}	37.27	32	0.24	-108.88	69.62
	T _{BA} , D _M	41.19	30	0.08	-107.94	70.56
	T _{BA} , P _{SZ}	25.7	22	0.26	-100.55	77.95
	D _M , P _{SZ}	32.35	30	0.35	-98.51	79.99
	T _{BA} , T _{1D}	19.37	12	0.08	-97.82	80.68
	D _M , P _{SI}	30.33	30	0.45	-96.35	82.14
T _{BA} , P _{SI}	20.44	20	0.43	-89.65	88.85	
All-species diversity	P _{SZ}	9.15	8	0.33	-136.39	0
	D _M	11.2	16	0.8	-126.17	10.23
	D _M , T _{1D}	38.9	32	0.19	-110.74	25.65
	D _M , P _{SI}	33.53	30	0.3	-99.77	36.62
	T _{BA} , D _M	29.68	30	0.48	-95.66	40.73
	D _M , P _{SZ}	25.15	30	0.72	-90.83	45.56
Typical-species diversity	T _{BA}	15.35	10	0.12	-177.11	0
	D _M	18.56	16	0.29	-143	34.12
	P _{SZ}	8.47	8	0.39	-134.59	42.53
	D _M , T _{1D}	45.67	32	0.06	-118.48	58.63

	D_M, P_{SI}	42.3	30	0.07	-109.12	67.99
	T_{BA}, D_M	34.37	30	0.27	-100.66	76.45
	D_M, P_{SZ}	29.39	30	0.5	-95.35	81.76
	T_{BA}, P_{SZ}	20.78	22	0.53	-94.5	82.62
	T_{BA}, T_{1D}	14.52	12	0.27	-91.36	85.76
	T_{BA}, P_{SI}	21.52	20	0.37	-90.89	86.23
Dominant-species diversity	T_{BA}	13.84	10	0.18	-172.29	0
	D_M	16.25	16	0.44	-137.71	34.59
	P_{SZ}	6.23	8	0.62	-128.62	43.67
	D_M, T_{1D}	38.14	32	0.21	-109.87	62.42
	P_{SZ}, P_{SI}	38.05	32	0.21	-109.77	62.52
	D_M, P_{SI}	37.03	30	0.18	-103.5	68.79
	T_{BA}, D_M	31.1	30	0.41	-97.17	75.12
	T_{BA}, P_{SZ}	18.75	22	0.66	-92	80.29
	D_M, P_{SZ}	24.83	30	0.73	-90.48	81.81
	T_{BA}, T_{1D}	12.7	12	0.39	-88.94	83.35

Results of the alternative models that fitted selection criteria for each index of seedling composition and diversity and the exogenous variables included in each model (patch size: P_{SZ} , patch shape: P_{SI} , tree basal area: T_{BA} , tree diversity: T_{1D} and patch disturbance by management: D_M). Given are the degrees of freedom (df), Fisher's C value statistic (χ^2) and p-value (P), Akaike Information Criterion corrected for small samples (AICc), and AICc differences compared to be best-supported model by data (Δi).

Table S10. Test statistics for the single best model for each index of seedling composition and diversity in 16 forest patches in the Lacandona rainforest, southeast Mexico.

Seedling variable	Response	Predictor	β_i	SE	df	t	P	β'_i	
Composition	T_{BA}	DF	-8.069	2.683	14	-3.007	0.009	-0.626	
	LD	T_{BA}	-0.007	0.003	14	-2.426	0.029	-0.544	
	ST	LD		-1.694	1.273	13	-1.331	0.206	-0.412
		T_{BA}		-0.009	0.017	13	-0.499	0.626	-0.154
	MDS1	LD		-0.504	0.201	12	-2.504	0.028	-0.547
		ST		0.028	0.041	12	0.685	0.507	0.126
		T_{BA}		0.004	0.002	12	1.487	0.163	0.308
	MDS2	LD		-0.02	0.186	12	-0.108	0.916	-0.026
		ST		-0.141	0.038	12	-3.713	0.003	-0.758
		T_{BA}		0.001	0.002	12	0.341	0.739	0.078
All-species diversity	P_{SZ}	AI₆₀₀	0.335	0.108	14	3.083	0.008	0.636	

	LD	P _{sz}	0.17	0.134	14	1.267	0.226	0.321
		LD	-2.384	0.893	12	-2.669	0.02	-0.579
	ST	P _{sz}	0.177	0.574	12	0.308	0.763	0.081
		Al₆₀₀	0.726	0.306	12	2.372	0.035	0.632
		LD	0.353	0.28	12	1.26	0.232	0.308
⁰ α		ST	0.157	0.068	12	2.326	0.038	0.564
		P _{sz}	0.154	0.147	12	1.046	0.316	0.253
		LD	-0.514	0.346	12	-1.484	0.164	-0.467
⁰ β		ST	-0.057	0.084	12	-0.688	0.505	-0.214
		P _{sz}	0.198	0.182	12	1.086	0.299	0.338
Typical-species diversity	T _{BA}	D_F	-8.069	2.683	14	-3.007	0.009	-0.626
	LD	T_{BA}	-0.007	0.003	14	-2.426	0.029	-0.544
		LD	-1.694	1.273	13	-1.331	0.206	-0.412
	ST	T _{BA}	-0.009	0.017	13	-0.499	0.626	-0.154
		LD	0.026	0.24	12	0.107	0.917	0.03
¹ α		ST	0.111	0.049	12	2.258	0.043	0.539
		T _{BA}	-0.004	0.003	12	-1.369	0.196	-0.368
		LD	-1.339	0.277	12	-4.84	0	-1.011
¹ β		ST	-0.117	0.056	12	-2.069	0.061	-0.364
		T _{BA}	-0.006	0.004	12	-1.681	0.119	-0.333
Dominant-species diversity	T _{BA}	D_F	-8.069	2.683	14	-3.007	0.009	-0.626
	LD	T_{BA}	-0.007	0.003	14	-2.426	0.029	-0.544
		LD	-1.694	1.273	13	-1.331	0.206	-0.412
	ST	T _{BA}	-0.009	0.017	13	-0.499	0.626	-0.154
		LD	-0.07	0.214	12	-0.325	0.751	-0.096
² α		ST	0.083	0.044	12	1.895	0.082	0.47
		T _{BA}	-0.004	0.003	12	-1.442	0.175	-0.403

	LD	-0.987	0.214	12	-4.613	0.001	-0.944
${}^2\beta$	ST	-0.108	0.044	12	-2.479	0.029	-0.427
	T_{BA}	-0.002	0.003	12	-0.759	0.463	-0.147

The response and predictor variables are patch size (P_{SZ}), patch shape (P_{SI}), tree basal area (T_{BA}), patch disturbance by fire (D_F), abundance of light-demanding seedlings (LD), abundance of shade-tolerant seedlings (ST), and alpha and beta diversity of all species (${}^0\alpha, {}^0\beta$), typical species (${}^1\alpha, {}^1\beta$) and dominant species (${}^2\alpha, {}^2\beta$). Given are unstandardized path coefficients (β_i) and associated standard error (SE), degrees of freedom (df), t-value (t), p-value (P) and standardized path coefficients (β'_i). The significant paths are highlighted in bold.

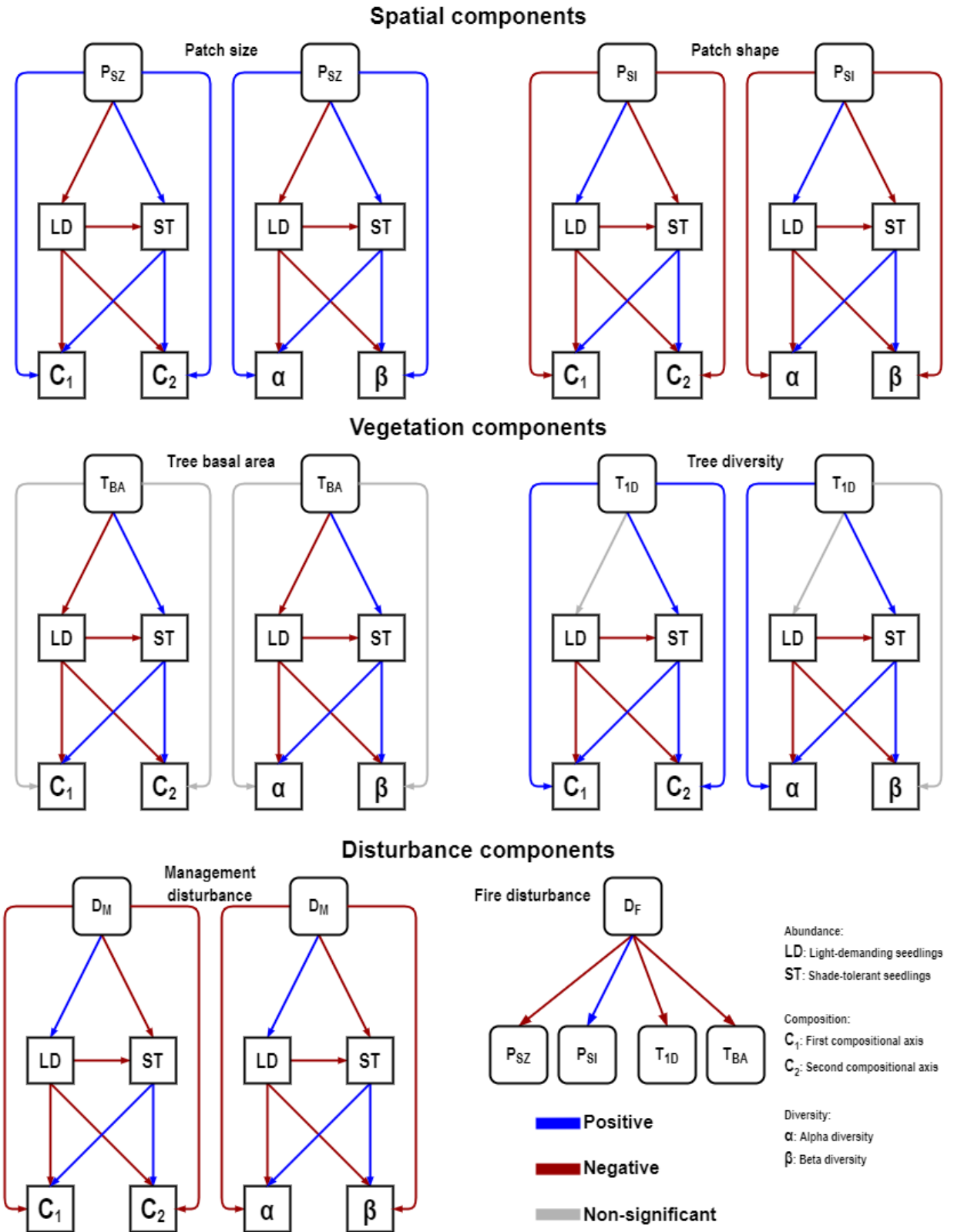


Figure S1. Expected relationships between patch components and seedling community

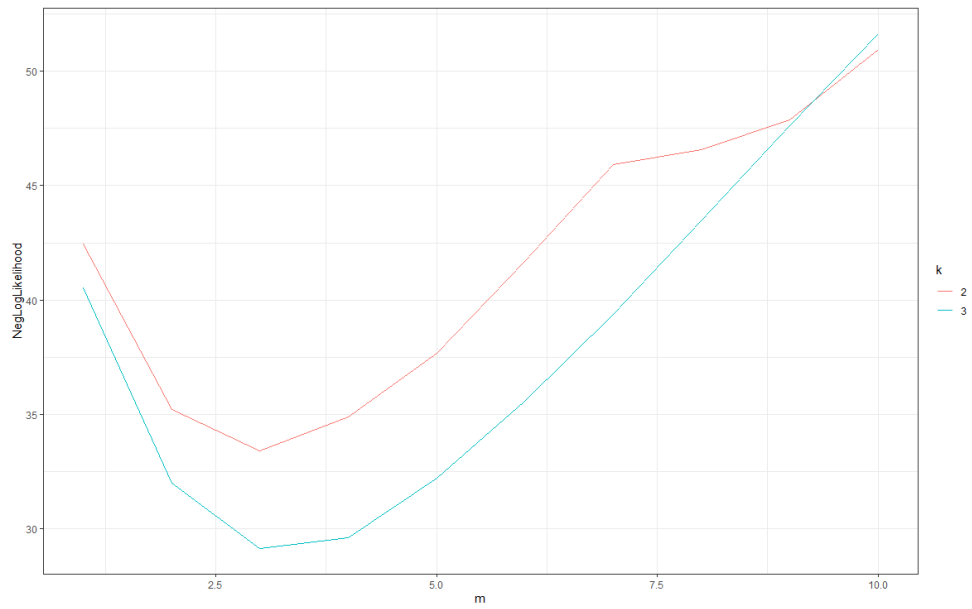


Figure S2. Association between the tuning parameter (m) and the negative Log-likelihood of models with varying m values. The red line represents a logistic PCA of two dimensions and the blue line a PCA of three dimensions.

**CAPÍTULO III. LANDSCAPE STRUCTURE SHAPES THE DIVERSITY OF
TREE SEEDLINGS AT MULTIPLE SPATIAL SCALES IN A
FRAGMENTED TROPICAL RAINFOREST**

RESEARCH ARTICLE

Landscape structure shapes the diversity of tree seedlings at multiple spatial scales in a fragmented tropical rainforest

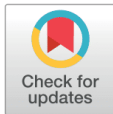
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Data Availability Statement: The data of seedling abundance and landscape metrics have been deposited with the Dryad Data Repository

Abstract

The maintenance of seedling diversity of animal-dispersed tree species is fundamental for the structure and function of forest patches in fragmented tropical rainforests. Nonetheless, the effects of landscape structure at different spatial scales on α - and β -diversity of tree seedling communities are recently explored. Using a multi-scale approach, we assessed the relative effect of landscape composition and configuration on α - and β -diversity of animal-dispersed seedlings within 16 forest patches in the Lacandona rainforest, Mexico. We assessed these effects at 13 spatial scales (from 300 to 1500 m radius, at 100 m intervals) for three metrics of effective number of species considering α - and β -diversity. We found that α -diversity was largely affected by landscape composition and β -diversity by landscape configuration. On the one hand, the amount of secondary forest influenced α -diversity. Additionally, species richness increased in landscapes with highly aggregated forest patches. On the other hand, β -diversity was affected positively by forest fragmentation and negatively by the edge contrast of forest patches with the surrounding matrix. Our findings indicate that landscape configuration is a strong driver of seedling diversity in highly deforested rainforests. Promoting forest patches and secondary forests through payment for ecosystem services' programs, favoring matrix quality within land-sharing schemes of smallholder agriculture and secondary forest management, and identifying restoration opportunities for assisted or unassisted natural regeneration are urgently needed for conservation of seedling diversity in human-modified tropical landscapes.

Introduction

Tropical rainforests harbor half of known animal and plant species of the planet [1], yet accelerated forest conversion to cattle pastures and large-scale plantations is restraining them to

since each response is influenced at different spatial and temporal scales [39,42]. For example, it is hypothesized that biological variables shaped by species colonization and extinction dynamics (i.e. species occurrence) should have larger scales of effect than those affected by local drivers (i.e. species abundance) [43]. Thus, the scale of effect is expected to be larger for lower-density populations (rare species) than for higher-density populations (typical or dominant species) [42]. This is particularly important in tropical rainforests, where a high number of species coexist at very low-density populations, making seedling communities highly α - and β -diverse in space [17,44].

Multi-scale approaches are a hierarchical approach useful to increase the possibility to detect the scale of effect of landscape structure on biological responses [45,46]. In these approaches, it is necessary the use of a wide range of scales, as assessments performed using very few scales within narrow ranges overlook the scale of effect [41,45]. To our knowledge, studies that have employed multi-scale analysis on plant communities in the tropics are scant [30,47], and none of them have assessed the effect of landscape structure on seedling communities in HMTLs (but see [30]). In this study, we employed a multi-scale approach to assess the contribution of landscape composition and configuration on tree seedling α - and β -diversity within rainforest patches. We particularly focused on the following question: Which components of landscape structure are influencing seedling α - and β -diversity and at which spatial scales?

Furthermore, we tested the following predictions: (1) α -diversity will increase in landscapes with high forest cover and β -diversity will increase in fragmented landscapes with low edge contrast; and (2) there should be larger scales of effect for richness-based metrics than for abundance-based metrics of seedling α - and β -diversity (i.e. dominant < typical < all species).

Material and methods

Study site

We conducted the study at the Marqués de Comillas region in the Lacandona rainforest in southeastern Mexico (Fig 1A). The monthly temperature oscillates between 24 and 26°C, and the annual precipitation ranges from 2500 to 3500 mm [48]. This region encompasses the largest rainforest of the Mesoamerican biodiversity hotspot and the Montes Azules and Lacantún Biosphere Reserves, which protect 393,000 ha of old-growth forest [49–51]. Land-cover change to cattle pastures has reduced its forest cover outside the reserve by more than 50% of its original extension [51,52]. Despite deforestation in the last 40 years, large mammals, birds, and primate species are still present in forest patches [6,53,54]. We selected 16 old-growth forest patches, ranging from 1 ha to 63 ha (Fig 1B). We only considered only patches located in floodplains to keep adult tree species composition, soil type and abiotic factors relatively constant. The land tenure of the Marqués de Comillas region is private, and land property is distributed among the inhabitants [51,52]. Therefore, we conducted our study only in those forest patches where permission was granted by the owners.

Tree seedling sampling

Patches were sampled by 1-ha blocks placed at the center of each forest patch from February to June of 2014 (Fig 1C). Each block contained ten 1-m² plots randomly arranged in groups of two or three plots along five equidistant transects (20 m apart; Fig 1D). For 1-ha forest patches, we never positioned plots at the edge in order to have a 20-m buffer zone of vegetation that protect them from edge effects as much as possible (Fig 1D) [13,15]. Within each 1-m² plot we counted and identified all tree seedlings (10–100 cm height) to the lowest possible taxonomic level with the help of a local parataxonomist and field guides [56,57]. When field identification

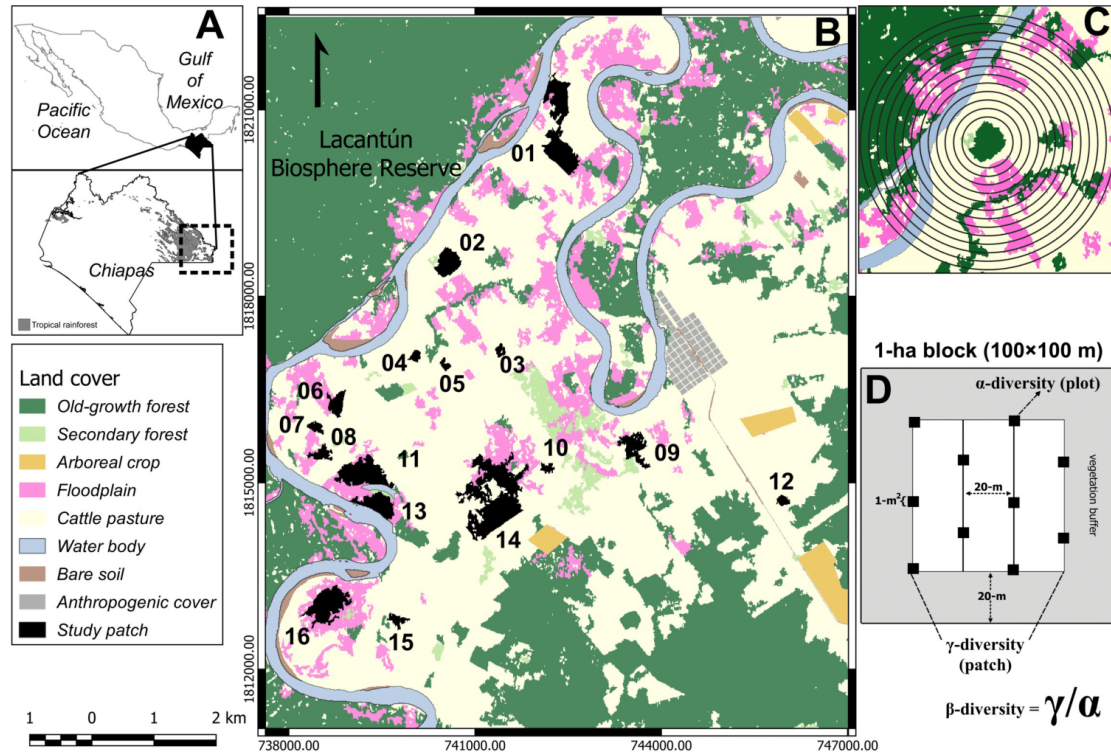


Fig 1. Study site and sampling design of tree seedlings. The figure shows the location of the study area (A) and the 16 study patches (black) in the Marqués de Comillas region (B). The 13 buffer sizes around the geographic center of the sampling block (C). Sampling method of tree seedlings (see [material and methods](#)), and the source of γ -diversity (species located in the 1-ha block), α -diversity (the average number of species per 1-m² plot) and β -diversity (the effective number of completely distinct assemblages within each 1-ha block) (D). Reprinted from [55] under a CC BY license, with permission from Sergio Nicasio-Arzeta, original copyright 2019.

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was not possible, we took samples for identification at several herbariums (MEXU, ECO-SC-H). We determined the dispersal syndrome of each species, based on their fruits and seed morphology [58,59]. This study did not involve the extraction or damage of endangered species. We based our analyses on animal-dispersed tree species only because they comprise up to 90% of the seed rain in rainforests [20,60]. In our case, 94% of the seedling species were animal-dispersed (see [Results](#)). Plant nomenclature followed the Missouri Botanical Garden database Tropicos [61].

Diversity estimation

We assessed the sampling completeness with the sample coverage estimator of Chao and Shen [62]. We combined the data of the 10 sampling plots within each patch. Then, we estimated the proportion of the total number of individuals that belong to the species represented in the sample [63]. The sample coverage among patches was high ($91.07 \pm 7.47\%$; mean \pm SD), indicating that our sampling effort was adequate for estimating seedling species diversity [64]. We then estimated the effective number of species for seedlings using the diversity decomposition

of effective numbers of species (the Hill numbers [65,66]). This method assesses the relative importance of species occurrence and abundance on α - and β -diversity [65,67]. Assessing these diversity metrics is important in the tropics because most tree species have small populations, and the scale of effect of landscape structure is expected to differ among species with varying abundances [60,68]. Consequently, diversity metrics that account for species abundance are expected to have smaller scales of effect than those measured by species occurrence only (presence-absence) [42,43]. Hill numbers (qD) are in units of species, which permit the characterization of species abundance distribution of a community and provide complete information about the community diversity [66,67]. We first calculated the patch gamma (γ) diversity of order q as follows:

$${}^qD_\gamma = \left(\sum_{i=1}^S \bar{p}_i^q \right)^{\frac{1}{1-q}}$$

where \bar{p}_i denotes the mean relative abundance of the i th species in the N 1-m² plots [65,69], and q is a parameter that determines the sensitivity of the measure to the relative abundances. Since this measure is undefined for $q = 1$, the γ -diversity of order one can be estimated with the following formula:

$${}^1D_\gamma = \exp\left(-\sum_{i=1}^S \bar{p}_i \log \bar{p}_i\right)$$

The 0D represents the species richness (hereafter called all species), which is not sensitive to individual abundances [65,69]. The 1D (equivalent to the exponential of Shannon’s entropy index) weights each species according to its abundance in the community, and could be interpreted as the number of equally common species which does not favor rare nor abundant species (hereafter called typical species) [65]. Finally, the 2D (equivalent to the inverse Simpson index) gives a high weight to abundant species and could be understood as the effective number of the most abundant species (hereafter called dominant species) [67,70]. We then considered the 1-m² plots to calculate the alpha (α) diversity with the formula:

$${}^qD_\alpha = \left(\frac{1}{N} \sum_{i=1}^S p_{i1}^q + \frac{1}{N} \sum_{i=1}^S p_{i2}^q + \dots \right)^{\frac{1}{1-q}}$$

Where p_i denotes the relative abundance of the i th species in each of the N 1-m² plots. We estimated α -diversity when $q = 1$ as:

$${}^1D_\alpha = \exp\left\{-\frac{1}{N} \left(\sum_{i=1}^S (p_{i1}^q \ln p_{i1}^q) + \sum_{i=1}^S (p_{i2}^q \ln p_{i2}^q) + \dots \right)\right\}$$

Afterward, we used the resulting α_{plot} - and γ_{patch} -diversity to calculate the “effective number of completely distinct assemblages” within each patch (β -diversity) as follows [65]:

$${}^q\beta_{\text{plot}} = \frac{{}^q\gamma_{\text{patch}}}{{}^q\alpha_{\text{plot}}}$$

This β -diversity ranges between one (when the assemblages of all 1-m² plots are identical) and N (when the assemblages of the N 1-m² plots are completely different from each other). We used the package *vegan* in R [71].

Landscape metrics and multi-scale assessment

We employed a multispectral SPOT-5 satellite image of 10 × 10 m pixel resolution recorded in March 2013 to carry out a supervised classification using the GRASS GIS software [72]. We

Table 1. Description, metric type and ecological relevance of the landscape metrics measured at class level employed in the study.

Metric	Description	Metric type	Ecological relevance	References
Forest cover (FC)	Percentage of landscape area covered by old-growth forest	Composition	Indicator of the landscape-scale habitat amount. It is positively associated with availability of propagules and seed dispersal	[31,60]
Secondary forest (SF)	Percentage of landscape area covered by secondary forest (regrowth of ≤ 15 years)	Composition	Measurement related to impacts on the dynamics and trajectories of floristic change within forest patches	[32,74]
Patch density (PD)	Number of forest patches per landscape area (n/ha)	Configuration	Fragmentation metric associated with increasing edge effects, the number of seed sources in the landscape and with landscape connectivity of seed dispersers in the tropics	[47,75]
Aggregation index (AI)	Percentage of like-adjacencies between forest patches. Maximum aggregation indicates a single, compact patch	Configuration	Aggregated patches facilitate the inter-patch movement, promoting dispersal at smaller spatial scales. These conditions allow the persistence of species in highly fragmented landscapes through complementation/supplementation dynamics	[8,76,77]
Patch isolation (PI)	Mean distance between forest patches within the landscape	Configuration	Metric employed to estimate connectivity on seed dispersal, altering seed abundance/richness and floristic differentiation among patches	[31,78]
Edge contrast index (EC)	Average degree of edge contrast between forest patches and their immediate neighborhoods	Configuration	High percentages of contrasting edges reduce the connectivity of vertebrate seed dispersers and terrestrial mammals in tropical forests	[6,79]

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used sampling points representing seven land-cover classes, including the following: 1) old-growth forest (undisturbed rainforest), 2) secondary forest (rainforest regrowth), 3) floodplains (swamp forests of bamboo and palm trees subjected to periodic flooding), 4) arboreal crops (oil-palm plantations), 5) cattle pasture (induced grassland for cattle raising), 6) anthropogenic cover (roads and urban settlements), and 7) water bodies (rivers and permanently flooded areas). The overall classification accuracy was 79%.

We then calculated six landscape metrics (Table 1) considered as drivers of seed and seedling community diversity [47,60]. The composition metrics included the percentages of old-growth forest (FC) and secondary forest (SF) covers, whereas the configuration metrics were the number of forest patches per landscape area (patch density; PD), the adjacency among forest patches (aggregation index; AI), the mean distance among forest patches (patch isolation; PI), and the edge contrast index (EC). We calculated the EC using quality values for matrix covers that describe both their capacity to reduce edge effects and the permeability for terrestrial mammals' movement. These values were based on the assumption that edge effects increase and mammals' presence declines along a gradient of habitat loss and relates the percentage of each land-cover type within the landscape matrix to its relative quality [2,6]. We ranked the relative quality of each land-cover type based on the suitability of vegetation structure for regulating edge microclimate and for mammal feeding, movement and/or habitat on a seven-point scale, including the following: 1 (water bodies, with the lowest suitability); 2 (anthropogenic cover); 3 (cattle pasture); 4 (arboreal crops); 5 (floodplains); 6 (secondary forest); and 7 (old-growth forest, representing the highest suitability). To obtain a more robust and realistic representation of landscape structure effects, we estimated the area-weighted mean of the EC index [73].

We neither considered patch size nor patch shape because both variables were highly correlated ($r = 0.81$), and none of them had significant effects on any of the α - and β -diversity metrics (Table A in S1 Text).

We followed the recommendation about the landscape radius that best predicts biological responses, which is 0.3–0.5 times the maximum dispersal distance of seed dispersers and herbivore mammals foraging behavior [41,45]. The maximum dispersal distance of arboreal seed dispersers and terrestrial mammals in the study region ranges between 500 and 4000 m [54,80–82]. Thus, we estimated six landscape metrics within 13 circular buffers (300 to 1500 m

radius, at 100 m intervals) from the center of each focal patch (Fig 1C). We only found correlations between AI and FC between the 900–1500 m radius (Table C in S1 Text). The distance among sampling sites was of 3602.56 ± 1932.21 m (mean \pm standard deviation), and were randomly aggregated (Clark-Evans $R = 1.02$; $p = 0.86$) [83]. The overlapping between buffers of nearby sampling points increased with buffer sizes (Table D in S1 Text). However, we did not find spatial autocorrelation between the distance of sampling sites and the diversity metrics (Table E in S1 Text), nor between the distance of sampling sites and the landscape metrics across the 13 buffer sizes (Table F in S1 Text).

Statistical analyses

Firstly, we estimated the scale of effect of each landscape metric using linear models. We fitted a diversity metric with a single landscape metric. We repeated this process on each buffer size and obtained 13 linear models of the same landscape metric. We verified variable normality with a Shapiro-Wilk test [84]. Then, we assessed the predictive power of each model using a leave-two-out cross-validation. This analysis splits the dataset in two sections; the first split contains the diversity data from 14 patches as calibration data and fits a linear model which is employed to predict the diversity values of the second split, which contains the two remaining values (validation data). We calculated the sums of squares between the estimated and observed diversity values employed as validation data. We repeated this procedure for each possible split of the data set and calculated the average sum of squares (\overline{SS}_{cv}). Next, we employed the \overline{SS}_{cv} to calculate the proportion of the variation that can be predicted by the model using the leave-two-out coefficient of determination (R^2_{CV}) as follows:

$$R^2_{CV} = 1 - \frac{\overline{SS}_{cv}}{\frac{1}{n} \sum_{i=1}^n (y_i - \bar{y})^2}$$

where \overline{SS}_{cv} is the average sum of squares obtained from the leave-two-out cross-validation, y_i is the diversity value for the i^{th} patch, \bar{y} is the mean value of a given diversity metric, and n is the number of patches. The R^2_{CV} ranges between $-\infty$ (indicating the model has a worse prediction power than the null model) and one (indicating the model predicts the validation data perfectly), and can be used to compare between response variables and scales of measurement [85–87]. Finally, we plotted the resulting 13 R^2_{CV} values and selected the one with the strongest response as the scale of effect for that landscape metric (Figs A and B in S1 Text [45]). We repeated this procedure for each diversity and landscape metric ($N = 468$ models).

Flawed study design, such as selecting a narrow range of spatial extents (smallest to largest) or by establishing wide distances among the analyzed extents, can lead to inaccurate estimations of the scale of effect [41,43]. Therefore, we estimated the uncertainty around the selected scale of effect through bootstrapping [88]. We randomly resampled the data from n patches, with resampling, from the set of 16 patches 1000 times. Then, we estimated the scale of effect for each resampled data set as described above and summed the number of times (out of 1000) that each buffer size was selected as the scale of effect. Although the uncertainty around the selected scale of effect varied among landscape- and diversity-metrics, the most frequently selected scale of effect based on the bootstrapped analysis matched with the scale of effect that we selected based on the highest R^2_{CV} value (Figs C and D in S1 Text). Additionally, we employed the data from the bootstrapped analysis to assess if the scale of effect differed among α - and β -diversity metrics and among landscape metrics. We only employed the data from the selected landscape metrics in the previous step.

Thereafter, we evaluated the effects of landscape metrics on diversity metrics through multiple linear models with a normal distribution. We constructed a global model for each of the

Table 2. Global model of each diversity metric employed for the model selection assessment.

α -diversity	β -diversity
$\alpha_{all} \sim AI_{600} + PI_{1200} + SF_{600}$	$\beta_{all} \sim EC_{800} + PD_{1400}$
$\alpha_{typical} \sim AI_{600} + PI_{1200} + SF_{600}$	$\beta_{typical} \sim PD_{300} + SF_{1300}$
$\alpha_{dominant} \sim AI_{600} + PI_{1200} + SF_{600}$	$\beta_{dominant} \sim AI_{500} + PD_{500}$

The landscape metrics are the aggregation index (AI), the edge contrast index (EC), patch isolation (PI), and the percentage of secondary forest (SF). The subscript numbers indicate the scale of effect (radius meters) of each landscape metric.

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α - and β -diversity metrics using only the landscape metrics at the scale of effect identified in the previous step (Table 2). All models were additive due to our limited sample size of 16 forest patches.

We did not find spatial autocorrelation between the distance of sampling sites and the model residuals of the selected landscape metrics (Table G in S1 Text). Therefore, we assured samples independency because spatial autocorrelation of model residuals—an indicator of pseudoreplication—is associated with the proximity between sampling sites [89,90]. We estimated the variance inflation factor (VIF) of landscape metrics beforehand with the *car* package [91]. Since we did not find significant collinearity ($VIF \geq 4$) between the explanatory variables [92], we employed the *dredge* function of the *MuMIn* package [93] to create all possible combinations of explanatory variables plus the null model (only the intercept). We ranked the models using the Akaike's information criterion corrected for small samples (AICc) and selected those models with a AICc difference lower than two ($\Delta AICc < 2$) as the best supported by the data [94].

Finally, we assessed the importance and the relative effect of each landscape metric measured at the scale of effect on each diversity value using an information theory approach and multimodel inference [94]. For this, we selected the subset of models that had 95% probability of containing the best model using the summed Akaike weights (w_i) of ranked models until $\sum w_i \leq 0.95$. We employed the w_i of the model's subset to calculate the relative importance and the model-averaged parameter estimates of each explanatory variable. We considered as influential variable those for which the unconditional standard error (USE) did not include zero in the averaged parameters. We carried out all statistical analysis in the R 3.5.2 statistical computing environment [95].

Results

We recorded a total of 1334 tree seedlings from 29 families, 51 genera and 72 species in 160 m² (Table H in S1 Text). Most seedlings were animal-dispersed species (1258 individuals; 94.3%), belonging to 24 families, 42 genera and 58 species. Mean species density of the animal-dispersed seedlings were 11.37 ± 2.96 species/10 m² (range 6–16 species; mean \pm standard error), mean seedling density was 78.62 ± 34.03 individuals/10 m² (range 10–129 individuals). The most abundant species was *Inga punctata*,—an early successional and disturbance tolerant tree species—, which represented 33% of all individuals sampled, followed by the late-successional tree species *Ampelocera hottlei* (13%) and *Brosimum alicastrum* (11%). Less than half of the species (ca. 31%) were restricted to one patch. We found that α - and β -diversity metrics were significantly higher than the other metrics when species abundances were not considered (Fig E in S1 Text).

Scale of effect and importance of landscape structure on seedling diversity

We observed that scale of effect of landscape composition and configuration was similar for all α -diversity metrics (Fig 2A–2C). The bootstrapped data showed the scale of effect of α -

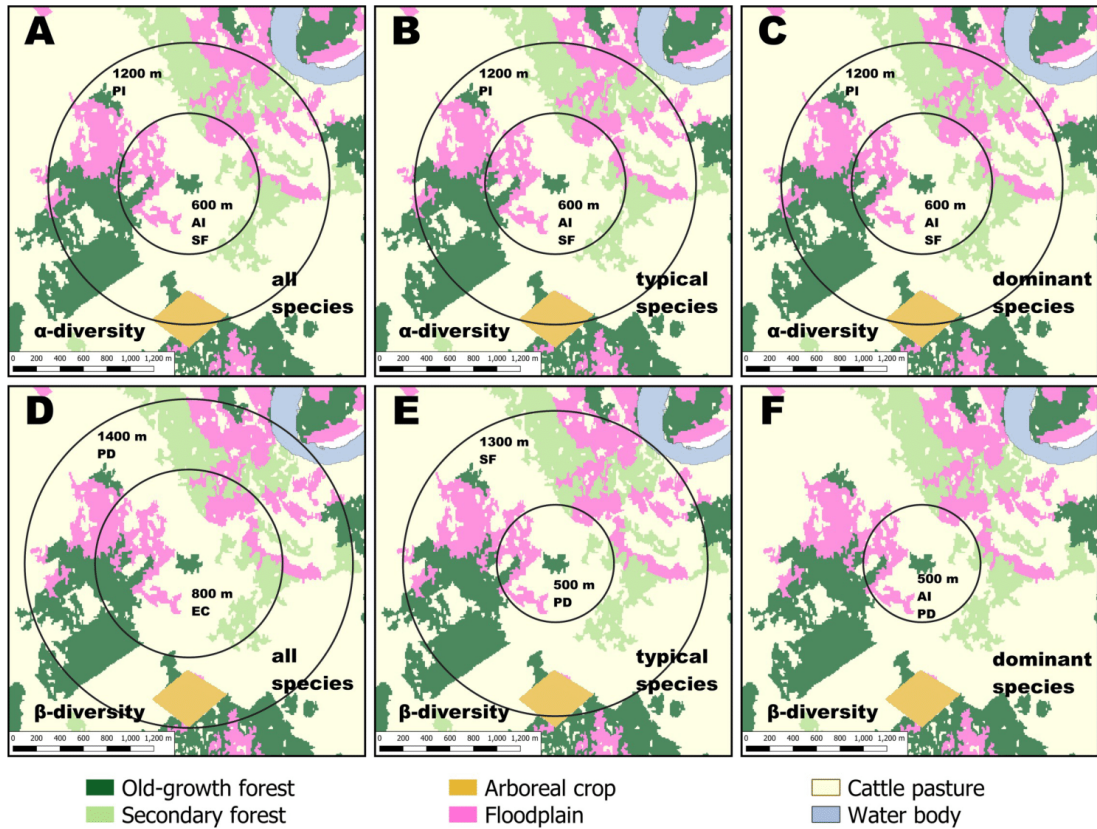


Fig 2. Scale of effect of landscape metrics on seedling diversity within forest patches in a fragmented tropical rainforest in southern Mexico. (A-C) Scale of effect on α -diversity. (D-F) Scale of effect on β -diversity. The landscape radius in meters and the associated landscape metrics are within each buffer. The landscape metrics are the aggregation index (AI), the edge contrast index (EC), the patch density (PD), patch isolation (PI), and the percentage of secondary forest (SF).

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diversity was the same for typical (807.52 ± 140.03 m; mean \pm standard error) and dominant species (801.73 ± 132.57 m), whereas the scale of effect for all species was slightly lower (764.27 ± 144.06 m). The best-fitting models showed a positive effect of patch aggregation, and a negative effect of patch isolation and secondary forest cover (Table 1 in S1 Text). When assessing the importance and relative effect of these landscape metrics, we found that secondary forest was strongly associated ($\sum w_i > 0.75$; Fig 3A) to α -diversity loss, regardless diversity metrics (Fig 3C). We also found an important contribution of patch aggregation in species richness (Fig 3C).

In contrast, we found that scale of effect varied among β -diversity metrics (Fig 2D–2F). The bootstrapped data indicated that the scale of effect decreased in the predicted order (i.e. dominant < typical < all), where the scale of effect was higher for β -diversity of all species (1021.07 ± 133.74 m) than for typical (826.54 ± 84.72 m) and dominant species (577.91 ± 101.79 m). The three β -diversity metrics were positively associated with patch

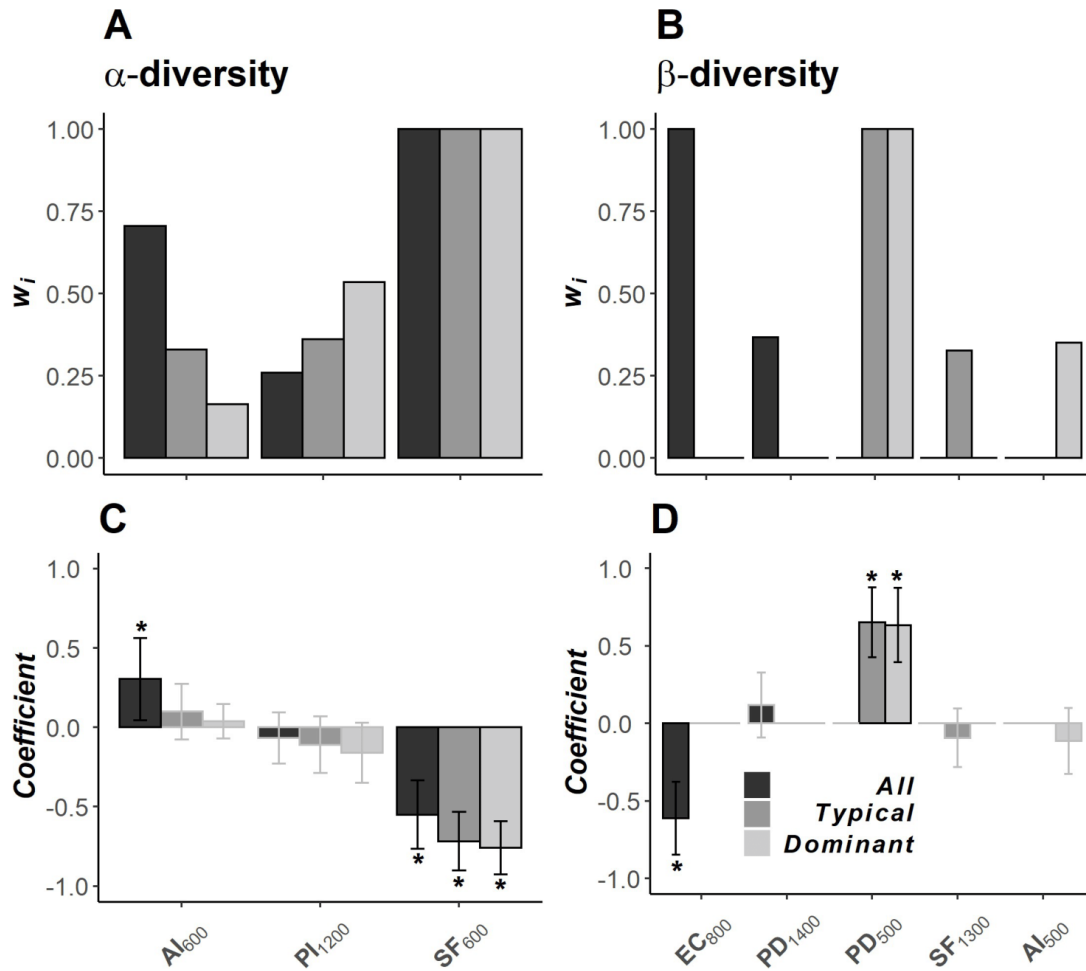


Fig 3. Importance and relative effects of the landscape metrics on seedling diversity within forest patches in a fragmented tropical rainforest in southern Mexico. We included the 95% confidence set of models for (A and C) α - and (B and D) β -diversity. The importance of each variable is represented by the sum of the Akaike weights ($\sum w_i$). The effects of each covariate were estimated through a model-averaged parameter estimate of information-theoretic-based model selection and multimodel inference. The whiskers represent the unconditional standard error (USE) and the asterisks indicate the influential variables (those for which the USE did not include zero). The landscape metrics are the aggregation index (AI), the edge contrast index (EC), the patch density (PD), patch isolation (PI), and the percentage of secondary forest (SF). The subscript numbers indicate the scale of effect of each variable.

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density, and negatively affected by secondary forest cover, patch aggregation, and edge contrast (Table I in S1 Text). The Akaike weighted values showed that landscape configuration was important for β -diversity (Fig 3B). Patch density was particularly relevant for β -diversity of typical and dominant species, while edge contrast was more important for β -diversity of all species (Fig 3D).

Discussion

Landscape effects on seedling diversity

This is the first study to assess the effects of landscape structure on the α - and β -diversity of the animal-dispersed tree seedling community at multiple spatial scales (but see [30]). As expected, we found that landscape composition affected α -diversity whereas β -diversity was influenced by landscape configuration. We also observed that the scale of effect varied among β -diversity metrics in the predicted order.

Effects on α -diversity. We found a decrease of α -diversity in forest patches surrounded by secondary forests, which are documented to exert a significant influence in the composition of tree communities within forest patches [32,33,96]. Altered microclimate and the opening of canopy gaps in small patches promote the proliferation of early-successional tree species while reducing the richness and abundance of old-growth species [74,97]. The arrival of early-successional species is driven by short-distance dispersal of generalist seed dispersers [98–100]. This may explain the high abundance of *I. punctata* seedlings, a species from seasonally flooded rainforests that is commonly found in secondary forests and as isolated trees in pastures [101–103]. Additionally, secondary forests restrict the arrival of some forest-specialist seed dispersers to forest patches at similar landscape sizes (100-ha or 564-m radius) [53,104] than the observed scale of effect of secondary forest on α -diversity (600-m radius). Thus, our findings support the contribution of the surrounding matrix on tree diversity within forest patches [32,33]. This results should be taken cautiously, because secondary forests are fundamental for forest regeneration and restoration in HMTLs [105,106], whether by increasing habitat amount, attracting pollinators and seed dispersers, buffering edge effects, and by providing seed sources that maximize the large-scale regeneration of disturbed abandoned fields [107,108]. Thus, conservation strategies in HMTLs should preferably promote secondary forests than agricultural fields in the surrounding matrix of forest patches [105,109].

Contrary to our predictions, we observed that α -diversity was also affected by landscape configuration, supporting the fragmentation threshold hypothesis [38]. The positive effect of patches aggregation on seedling richness suggests that seed sources and seed dispersal can be sustained where patches are highly aggregated. Previous studies predicted that dispersal distance of species may be limited in highly deforested landscapes, favoring species response to local characteristics, such as patches aggregation across the landscape [110,111]. In our study, the selected landscapes presented a low percentage of forest cover across the 13 buffer sizes ($19.01 \pm 5.1\%$; mean \pm standard deviation). The aggregation of forest patches facilitates the inter-patch movement of forest-dependent species in landscapes with low forest cover, where the distribution of species home range comprises several patches [77,112]. For example, frugivore birds can adjust their movement behavior according to landscape configuration, flying larger distances and visiting more forest patches in landscapes with decreasing forest cover [113]. These changes in the spatial configuration of forest patches also affect species foraging—which relies in resources from nearby patches (i.e. landscape supplementation dynamics [7])—and therefore, seed dispersal patterns. Some studies indicate that fruit removal increased and dispersal distance decreased where plants aggregation is high [114]. Thus, colonization of tree species in highly disturbed landscapes may be operating at smaller spatial scales [42].

Effects on β -diversity. Landscape configuration affected β -diversity, but seedling responses varied among diversity metrics. The β -diversity of all species was negatively associated with edge contrast of old-growth forests in the 800-m radius landscapes. The loss of β -diversity by edge contrast indicates an homogenization of seedling assemblages by edge effects, as well as by a limited heterogeneity of microclimate conditions, a processes observed in HMTLs with low forest cover [18,115]. Additionally, edge contrast can influence the presence

of herbivorous mammals within patches [6], which promote β -diversity of tree seedlings through secondary seed dispersal, and seedling herbivory and trampling [22,23]. Accordingly, the association between seedling β -diversity and edge contrast suggests that low exposure to edge effects, a more heterogeneous microclimate and the presence of density-independent factors (i.e., terrestrial herbivores) promotes a broader species coexistence [35].

The overall increment of β -diversity for the typical and dominant species by patch density is not surprising. According to the “biotic differentiation hypothesis”, the limited exchange of seeds and differences in disturbance regimes across forest patches promote floristic differentiation in highly deforested landscapes [68,96]. Spatial patterns that limit the species exchange influence β -diversity in the tropics [17,116]. These effects are intensified in fragmented rainforests, where differences in forest cover amount, disturbance source (i.e., slash-and-burn practices or tree removal by machinery), degree of fragmentation and connectivity, lead to contrasting regeneration dynamics within patches [96,117]. Consequently, the establishment of a wider array of species is facilitated by landscape configuration and a less evenly distributed number of individuals among dominant species [34,77]. These conditions can prevent competitive exclusion (i.e. excluding strong competitors that are weak dispersers), spread the risk of simultaneous extinctions and increase landscape complementation, favoring β -diversity [8,68]. This is consistent with the mechanisms involved in β -diversity increase through the colonization of different opportunistic species and the extinction of shared species among sites [118]. Thus, fragmentation is having positive effects on seedling diversity by facilitating the arrival and establishment of species otherwise poorly represented in old-growth forests [8].

Scale of landscape effect on diversity metrics

Our results did not support the predicted order of the scale of effect (dominant < typical < all) for α -diversity metrics, which also showed a substantial uncertainty in the selected scale of effect (Fig C in S1 Text). These variations reinforce the findings that scale of effect does not vary in a predictable order among response variables [43,88]. The inconsistencies observed on α -diversity metrics may be associated to the temporal scale and landscape composition and configuration variables assessed [88]. Temporal scale variations regarding tree phenology and movement/foraging patterns of seed dispersers in fragmented HMTLs [54,119] may be obscuring the expected differences in the scale of effect. A study in the same region found that scale of effect of landscape composition on seed rain followed the predicted order [31,60]. The sampling period however, comprised a whole year in order to control for the temporal variation in the seed rain [60]. Additionally, the scale of effect was possibly influenced by the low forest cover of the study area (<25%), which can reduce the expected scale of effect to local factors, such as patch size [42]. However, comparative studies in HMTLs have found non-significant differences in the scale of effect among regions with contrasting disturbance levels [54,60]. Further studies including comparative assessments among regions are needed to fully understand the importance of regional context in landscape metrics and the scale of effect.

On the contrary, the scale of effect observed among β -diversity metrics did follow the predicted order. The scale of effect of edge contrast is similar to those observed for matrix contrast on stem density of understory vegetation in los Tuxtlas, Mexico (798 m), and for matrix openness in seed abundance and species richness of animal-dispersed seeds and forest generalist saplings (798–977 m) [30,31,47]. These studies suggested that matrix contrast operates at smaller scales, where it drives local edge effects involved in the mortality of animal-dispersed seedling species, and favors the dispersal and establishment of wind-dispersed seeds [31,47]. This also holds true for the scale of effect of patch density in our study (500 m) and those observed for richness of old-growth forest specialist- saplings (500 m) in the study area [30].

Patch density is positively related to habitat heterogeneity and to the number of subpopulations in the landscape, increasing community dissimilarity [8,35]. These findings suggest that matrix contrast is limiting the dispersal and recruitment of lower-density species, whereas forest fragmentation is promoting the coexistence of higher-density species.

Study limitations

Our results are conservative as a larger number of landscapes are desirable to making further inferences. Also, the lack of information related to species composition of secondary forest trees and seed rain within forest patches limit our understanding about the contribution of secondary forests to seedlings α -diversity [60,120]. Finally, our study did not consider a long-term monitoring of tree seedlings to elucidate the contribution of patch microclimate, fruiting events, foraging behavior of seed dispersers, disturbance regimes, and density-dependent mortality factors, which strongly influence the composition and dynamics of seedling communities in the tropics [16,121,122].

For seedlings β -diversity, our findings should be taken cautiously, because we inferred the presence of mammals only through landscape metrics, regardless mammal species have different responses to landscape structure [6]. Furthermore, herbivorous mammal effects on seedling diversity vary according to the incidence, relative abundance and body mass [23,24], as well as by population fluctuations and food quality and availability [123]. In turn, these biological responses (i.e. abundance, richness, incidence and body size) are affected by landscape structure at varying spatial scales [39,42]. Finally, the mechanisms of β -diversity loss are highly sensitive to the ratio of winner-loser species and the spatial scale at which communities are assessed [118]. Nevertheless, the use of structural equation modeling in multi-scale assessments have permitted analyzing the direct and cascading effects of landscape structure on different biological responses in HMTLs [47]. This represents a promising approach to evaluate the response of interactions-mediated patterns and processes to landscape composition and configuration. Thus, further studies regarding cascading effects of landscape structure on different biological responses of herbivorous mammals and tree seedling communities are needed to fill these knowledge gaps.

Conclusions and conservation implications

Contrary to studies performed in the same area [30], we found that forest fragmentation is a strong driver of seedling diversity at smaller spatial scales in highly deforested rainforests. Severe fragmentation of rainforests will occur in the short term [4], urging economic incentives and innovations in policy and governance that favor forest regrowth, matrix quality and patches number.

Implementation of payment for ecosystem services' schemes and REDD+ programs can prevent further loss of old-growth and secondary forests while promoting agroforestry management [124,125]. Favoring matrix quality and configurational patterns that increase habitat heterogeneity are feasible in landscapes with land-sharing schemes of smallholder agriculture [126]. In these rural landscapes, diversification of small-scale agroecosystems, intensification of agriculture and reduction of cattle ranching promotes forest regeneration and food security [127,128]. Additionally, enactment of programs of enrichment planting of trees with local commercial/ecological importance, and forest management policies that permit local governance for harvesting timber and non-timber products are needed to enhance the management and diversity of secondary forests [129]. Finally, is necessary to map and classify the biophysical and socio-ecological land use dimensions of land covers and naturally regenerated forests to identify restoration opportunities for assisted or unassisted natural regeneration [129,130].

Novel institutional and policy approaches in Costa Rica have proven that decreasing meat exports while supporting payment for environmental services, agricultural intensification and native species plantations lead to a forest transition in less than 15 years [131,132]. Incorporation of these conservation, management and restoration guidelines are urgently needed to preserve regeneration of forest patches and future conservation value of HMTLs.

Supporting information

S1 Text. Supporting file.
(DOCX)

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Writing – review & editing: Sergio Nicasio-Arzeta, Isela E. Zermeño-Hernández, Susana Maza-Villalobos, Julieta Benítez-Malvido.

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Supplemental information

Table A. Coefficient of determination (R^2) and p value (in parenthesis) between α - and β -diversity metrics and patch size and shape.

Diversity	Diversity order	Patch size	Patch shape
α	All (${}^0\alpha$)	0.071 (0.16)	0.054 (0.195)
	Typical (${}^1\alpha$)	0.103 (0.122)	0.037 (0.229)
	Dominant (${}^2\alpha$)	0.098 (0.128)	0.001 (0.33)
β	All (${}^0\beta$)	-0.007 (0.358)	0.023 (0.264)
	Typical (${}^1\beta$)	0.043 (0.217)	0.047 (0.208)
	Dominant (${}^2\beta$)	-0.027 (0.45)	-0.01 (0.373)

Table B. Abbreviation and description of the landscape metrics employed in the analysis

Landscape metric	Abbreviation	Formula	Description
Aggregation index	AI	$AI = 100 \times \left[\frac{g_{ii}}{\max \rightarrow g_{ii}} \right]$	Where g_{ij} is the number of joins between pixels of forest patch i and $\max \rightarrow g_{ii}$ is the maximum number of joins between pixels of forest patch i based on the single-count method
Edge contrast index	EC	$EC = 100 \times \left[\frac{\sum_{k=1}^m p_{ijk} \cdot d_{ik}}{p_{ij}} \right]$	Where p_{ijk} is the edge length (m) of patch ij adjacent to cover type k , d_{ik} is the edge contrast weight between patch i and cover k , and p_{ij} is the length of perimeter of patch ij
Euclidean-nearest neighbor distance	ENN	$ENN = \frac{\sum_{i=1}^n h_{ij}}{n_h}$	Where h_{ij} is the edge-to-edge distance between two patches j of the land cover i , and n_h is the number of distances recorded
Forest cover	FC	$FC = 100 \times \left[\frac{\sum_{j=1}^n a_i}{A} \right]$	Where a is the area (m^2) of forest patch i and A the is the total landscape area
Secondary vegetation	SV	$SV = 100 \times \left[\frac{\sum_{j=1}^n a_i}{A} \right]$	Where a is the area (m^2) of cover patch i and A the is the total landscape area
Patch density	PD	$PD = 100 \times \left[\frac{n_i}{A} \times (10,000) \right]$	Where n_i is the total number of patches in the landscape of the land cover i and A is the total landscape area (m^2)

Table C. Correlation values among landscape metrics across the 13 spatial scales employed in the multi-scale analysis in a fragmented tropical forest in southeastern Mexico. Strong correlations ($-0.7 > r > 0.7$) are in bold.

Landscape radius		<i>AI</i>	<i>EC</i>	<i>PD</i>	<i>FC</i>
300m	<i>EC</i>	-0.475			
	<i>PD</i>	-0.582	0.283		
	<i>FC</i>	0.497	-0.621	-0.303	
	<i>SF</i>	0.333	-0.102	-0.464	0.202
400m	<i>EC</i>	-0.071			
	<i>PD</i>	-0.618	-0.373		
	<i>FC</i>	0.573	-0.477	-0.145	
	<i>SF</i>	-0.066	-0.278	0.186	0.127
500m	<i>EC</i>	0.171			
	<i>PD</i>	-0.57	-0.568		
	<i>FC</i>	0.303	-0.44	0.256	
	<i>SF</i>	-0.188	-0.156	-0.119	-0.273
600m	<i>EC</i>	0.084			
	<i>PD</i>	-0.135	-0.334		
	<i>FC</i>	0.155	-0.459	0.413	
	<i>SF</i>	-0.413	-0.123	0.076	-0.417
700m	<i>EC</i>	-0.051			
	<i>PD</i>	-0.088	-0.339		
	<i>FC</i>	0.079	-0.474	0.503	
	<i>SF</i>	-0.528	-0.073	-0.189	-0.438
800m	<i>EC</i>	-0.102			
	<i>PD</i>	-0.299	-0.382		
	<i>FC</i>	-0.062	-0.492	0.67	
	<i>SF</i>	-0.307	0.033	-0.498	-0.398
900m	<i>EC</i>	-0.331			
	<i>PD</i>	-0.031	-0.225		

	<i>FC</i>	0.875	-0.33	0.148	
	<i>SF</i>	-0.38	0.056	-0.237	-0.451
1000m	<i>EC</i>	-0.32			
	<i>PD</i>	-0.355	-0.226		
	<i>FC</i>	0.882	-0.354	-0.124	
	<i>SF</i>	-0.282	0.068	-0.145	-0.426
1100m	<i>EC</i>	-0.195			
	<i>PD</i>	-0.142	-0.373		
	<i>FC</i>	0.838	-0.264	0.293	
	<i>SF</i>	-0.265	0.128	-0.211	-0.371
1200m	<i>EC</i>	-0.16			
	<i>PD</i>	-0.039	-0.687		
	<i>FC</i>	0.817	-0.376	0.291	
	<i>SF</i>	-0.244	0.27	-0.48	-0.36
1300m	<i>EC</i>	-0.321			
	<i>PD</i>	-0.014	-0.551		
	<i>FC</i>	0.859	-0.498	0.207	
	<i>SF</i>	-0.291	0.444	-0.291	-0.427
1400m	<i>EC</i>	-0.361			
	<i>PD</i>	-0.02	-0.609		
	<i>FC</i>	0.906	-0.525	0.019	
	<i>SF</i>	-0.439	0.526	-0.233	-0.507
1500m	<i>EC</i>	-0.34			
	<i>PD</i>	0.193	-0.597		
	<i>FC</i>	0.932	-0.511	0.209	
	<i>SF</i>	-0.539	0.583	-0.218	-0.56

Table D. Percentage of landscape overlap among circular buffers of different radius (300 to 1500-m radius, at 100 m intervals) in a fragmented tropical forest in southeastern Mexico. We divided the total overlapped area by the sum of the 16 buffers' area for each landscape radius.

Landscape radius	Overlapping (%)
300m	2.139
400m	7.56
500m	13.159
600m	19.078
700m	25.026
800m	31.037
900m	37.676
1000m	44.865
1100m	52.491
1200m	60.964
1300m	70.349
1400m	80.433
1500m	91.091

Table E. Moran's I autocorrelation tests between the distance of sampling sites of tree seedling community and the reported α - and β -diversity (0D = all species, 1D = typical species and 2D = abundant species) in a fragmented tropical forest in southeastern Mexico.

Diversity	Order	Moran's I \pm standard deviation	P value
α	${}^0\alpha$	-0.043 \pm 0.065	0.721
	${}^1\alpha$	-0.055 \pm 0.065	0.853
	${}^2\alpha$	-0.063 \pm 0.065	0.957
β	${}^0\beta$	-0.077 \pm 0.059	0.861
	${}^1\beta$	-0.161 \pm 0.065	0.147
	${}^2\beta$	-0.175 \pm 0.063	0.088

Table F. Moran's I autocorrelation tests between the distance of sampling sites of tree seedling community and landscape metrics across the 13 spatial scales employed in the multi-scale analysis in a fragmented tropical forest in southeastern Mexico.

Landscape metric	Landscape radius	Moran \pm standard deviation	P value
AI	300m	-0.025 \pm 0.065	0.523
	400m	-0.021 \pm 0.065	0.485
	500m	-0.033 \pm 0.062	0.59

	600m	0.058 ± 0.064	0.054	
	700m	0.053 ± 0.066	0.068	
	800m	-0.092 ± 0.066	0.698	
	900m	-0.004 ± 0.062	0.31	
	1000m	-0.002 ± 0.062	0.293	
	1100m	-0.044 ± 0.061	0.71	
	1200m	-0.036 ± 0.06	0.607	
	1300m	-0.047 ± 0.063	0.75	
	1400m	-0.047 ± 0.064	0.762	
	1500m	-0.066 ± 0.063	0.99	
EC	300m	-0.035 ± 0.064	0.624	
	400m	-0.044 ± 0.064	0.722	
	500m	-0.062 ± 0.065	0.938	
	600m	-0.083 ± 0.065	0.806	
	700m	-0.111 ± 0.065	0.493	
	800m	-0.125 ± 0.065	0.373	
	900m	-0.111 ± 0.065	0.5	
	1000m	-0.071 ± 0.066	0.951	
	1100m	-0.01 ± 0.066	0.385	
	1200m	0.044 ± 0.066	0.093	
	1300m	0.051 ± 0.067	0.077	
	1400m	0.057 ± 0.067	0.063	
	1500m	0.034 ± 0.066	0.131	
	PD	300m	-0.024 ± 0.056	0.446
		400m	-0.035 ± 0.066	0.635
500m		-0.076 ± 0.064	0.887	
600m		-0.114 ± 0.065	0.469	
700m		-0.028 ± 0.063	0.541	
800m		-0.15 ± 0.061	0.176	
	900m	-0.017 ± 0.063	0.432	

	1000m	-0.023 ± 0.065	0.507
	1100m	0.04 ± 0.063	0.091
	1200m	0.038 ± 0.065	0.109
	1300m	-0.124 ± 0.066	0.386
	1400m	-0.038 ± 0.059	0.629
	1500m	-0.038 ± 0.061	0.638
FC	300m	0.067 ± 0.067	0.044*
	400m	0.04 ± 0.064	0.097
	500m	-0.018 ± 0.058	0.4
	600m	-0.043 ± 0.048	0.625
	700m	-0.051 ± 0.04	0.7
	800m	-0.052 ± 0.032	0.639
	900m	-0.024 ± 0.045	0.346
	1000m	0.005 ± 0.06	0.228
	1100m	0.009 ± 0.065	0.246
	1200m	-0.001 ± 0.065	0.319
	1300m	-0.012 ± 0.065	0.402
	1400m	-0.029 ± 0.064	0.557
	1500m	-0.05 ± 0.063	0.793
SF	300m	-0.018 ± 0.066	0.459
	400m	-0.044 ± 0.066	0.73
	500m	-0.038 ± 0.065	0.657
	600m	-0.038 ± 0.064	0.652
	700m	-0.069 ± 0.06	0.971
	800m	-0.075 ± 0.06	0.886
	900m	-0.071 ± 0.059	0.946
	1000m	-0.071 ± 0.058	0.935
	1100m	-0.073 ± 0.058	0.911
	1200m	-0.073 ± 0.059	0.911

1300m	-0.074 ± 0.06	0.903
1400m	-0.078 ± 0.061	0.855
1500m	-0.069 ± 0.062	0.972

* Significant autocorrelation values

Table G. Moran's I autocorrelation tests between the distance of sampling sites of the biotic-dispersed tree seedling community and the residuals of the linear models between α - and β -diversity metrics (0D = all species, 1D = typical species and 2D = dominant species) and landscape metrics in a fragmented tropical forest in southeastern Mexico. The numbers in parenthesis indicate de standard deviation. The subscript numbers indicate the scale of effect of each variable.

Diversity	Order	Landscape metric	Moran's I	P value
α	$^0\alpha$	AI ₆₀₀	-0.036 (0.064)	0.632
		SF ₆₀₀	-0.064 (0.066)	0.967
		PI ₁₂₀₀	-0.067 (0.065)	0.999
	$^1\alpha$	AI ₆₀₀	-0.045 (0.064)	0.74
		SF ₆₀₀	-0.1 (0.065)	0.61
		PI ₁₂₀₀	-0.032 (0.064)	0.588
	$^2\alpha$	AI ₆₀₀	-0.057 (0.064)	0.877
		SF ₆₀₀	-0.096 (0.063)	0.643
		PI ₁₂₀₀	-0.023 (0.065)	0.496
β	$^0\beta$	EC ₈₀₀	-0.134 (0.057)	0.24
		PD ₁₄₀₀	-0.055 (0.053)	0.827
	$^1\beta$	AI ₅₀₀	-0.073 (0.065)	0.922
		SF ₁₃₀₀	-0.164 (0.064)	0.129
	$^2\beta$	AI ₅₀₀	-0.123 (0.064)	0.383
		PD ₅₀₀	-0.173 (0.064)	0.098

Table H. Tree seedlings recorded during the sampling period in the Lacandona rainforest, at southeastern Mexico. Seedling dispersal syndromes were categorized whether by abiotic (A) or biotic (B) dispersion.

Family	Genus	Species	Dispersal	Number
ACANTHACEAE	<i>Bravaisia</i>	<i>B. integerrima</i>	A	29
ANACARDIACEAE	<i>Spondias</i>	<i>S. mombin</i>	B	6
		<i>S. radlkoferi</i>	B	4
	<i>Astronium</i>	<i>A. graveolens</i>	B	1
ANNONACEAE	<i>Annona</i>	<i>A. sp1</i>	B	4
	<i>Xylopia</i>	<i>X. frutescens</i>	B	1
APOCYNACEAE	<i>Stemmadenia</i>	<i>S. galeottiana</i>	B	2
BIGNONIACEAE	<i>Tabebuia</i>	<i>T. rosea</i>	A	1

BORAGINACEAE	<i>Cordia</i>	<i>C. odorata</i>	B	4	
		<i>C. bicolor</i>	B	2	
BURSERACEAE	<i>Bursera</i>	<i>B. simaruba</i>	B	1	
	<i>Protium</i>	<i>P. copal</i>	B	1	
CAPPARACEAE	<i>Capparis</i>	<i>C. quiriguensis</i>	B	1	
CHRYSOBALANACEAE	<i>Hirtella</i>	<i>H. americana</i>	B	15	
		<i>Licania</i>	<i>L. hypoleuca</i>	B	8
		<i>L. platypus</i>	B	1	
CLUSIACEAE	<i>Calophyllum</i>	<i>C. brasilense</i>	B	10	
	<i>Garcinia</i>	<i>G. intermedia</i>	B	4	
EUPHORBIACEAE	<i>Croton</i>	<i>C. schiedeanus</i>	B	11	
FABACEAE	<i>Acacia</i>	<i>A. mayana</i>	B	11	
		<i>Albizia</i>	<i>A. leucocalyx</i>	A	5
		<i>Andira</i>	<i>A. inermis</i>	B	4
		<i>Cojoba</i>	<i>C. arboreum</i>	B	4
		<i>Dialium</i>	<i>D. guianense</i>	B	10
		<i>Inga</i>	<i>I. edulis</i>	B	3
			<i>I. pavoniana</i>	B	1
			<i>I. punctata</i>	B	418
			<i>I. vera</i>	B	2
			<i>I. sp1</i>	B	4
			<i>I. sp2</i>	B	2
		<i>Lonchocarpus</i>	<i>L. cruentus</i>	A	2
			<i>L. pubescens</i>	A	8
		<i>Platymiscium</i>	<i>P. yucatanum</i>	A	10
		<i>Pterocarpus</i>	<i>P. hayesii</i>	A	1
<i>P. yucatanum</i>	A		1		
ICACINACEAE	<i>Calatola</i>	<i>C. costaricanum</i>	B	3	
LASCISTEMATACEAE	<i>Lacistema</i>	<i>L. aggregatum</i>	B	2	
LAURACEAE	<i>Licaria</i>	<i>L. capitata</i>	B	8	
	<i>Nectandra</i>	<i>N. ambiens</i>	B	17	
		<i>N. reticulata</i>	B	5	
MALVACEAE	<i>Hampea</i>	<i>H. stipitata</i>	B	4	
	<i>Luehea</i>	<i>L. seemannii</i>	A	4	
	<i>Quararibea</i>	<i>Q. funebris</i>	B	4	
MELIACEAE	<i>Cedrela</i>	<i>C. odorata</i>	A	3	
		<i>Guarea</i>	<i>G. excelsa</i>	B	8
		<i>G. glabra</i>	B	9	
		<i>G. grandifolia</i>	B	5	
	<i>Trichilia</i>	<i>T. havanensis</i>	B	3	
MORACEAE	<i>Brosimum</i>	<i>B. alicastrum</i>	B	147	

		<i>B. latescens</i>	B	20
	<i>Castilla</i>	<i>Castilla elastica</i>	B	32
	<i>Maclura</i>	<i>M. tinctoria</i>	B	3
	<i>Trophis</i>	<i>T. racemosa</i>	B	93
PRIMULACEAE	<i>Ardisia</i>	<i>A. paschalis</i>	B	27
MYRTACEAE	<i>Eugenia</i>	<i>E. edulis</i>	B	4
		<i>E. mexicana</i>	B	57
		<i>E. nigrita</i>	B	5
POLYGONACEAE	<i>Coccoloba</i>	<i>C. barbadensis</i>	B	1
RUBIACEAE	<i>Faramea</i>	<i>F. occidentalis</i>	B	20
	<i>Posoqueria</i>	<i>P. latifolia</i>	B	13
		<i>P. sp1</i>	B	5
SALICACEAE	<i>Pleuranthodendron</i>	<i>P. lindenii</i>	B	18
SAPINDACEAE	<i>Cupania</i>	<i>C. dentata</i>	B	8
		<i>C. glabra</i>	B	14
	<i>Sapindus</i>	<i>S. saponaria</i>	B	1
SAPOTACEAE	<i>Chrysophyllum</i>	<i>C. mexicanum</i>	B	10
	<i>Pouteria</i>	<i>P. durlandii</i>	B	4
ULMACEAE	<i>Ampelocera</i>	<i>A. hottlei</i>	B	173
VOCHYSIACEAE	<i>Vochysia</i>	<i>V. guatemalensis</i>	A	7
Unknown		Morpho 1		1
		Morpho 2		3
		Morpho 3		1

Table I. Model selection results of the linear models that explain each α - and β -diversity metric (0D = all species, 1D = typical species and 2D = dominant species) of the animal-dispersed tree seedling community in a fragmented tropical rainforest at southern Mexico. The models log-likelihood (LL), the corrected Akaike index criterion ($AICc$), the Akaike difference from the best model (Δ), the Akaike weight (w_i) and the coefficient of determination (R^2). The landscape metrics are the aggregation index (AI), the edge contrast index (EC), patch isolation (PI), and the percentage of secondary forest (SF). The subscript numbers indicate the scale of effect (radius meters) of each landscape metric. The superscript symbols indicate the positive (+) or negative (-) effects of landscape metrics. The values in italic correspond to the null model.

	Diversity metric	Model	R^2	LL	$AICc$	Δ	w_i
α -diversity	All species	${}^0\alpha \sim +AI_{600} + \cdot SF_{600}$	0.6	-8.38	28.39	0	0.52
		${}^0\alpha \sim \cdot SF_{600}$	0.45	-11.46	30.91	2.52	0.15
		${}^0\alpha \sim \cdot PI_{1200} + \cdot SF_{600}$	0.52	-9.87	31.38	2.98	0.12
		${}^0\alpha \sim +AI_{600} + \cdot PI_{1200} + \cdot SF_{600}$	0.6	-7.69	31.39	2.99	0.12
		${}^0\alpha \sim +AI_{600}$	0.39	-12.33	32.66	4.27	0.06
		${}^0\alpha \sim +AI_{600} + \cdot PI_{1200}$	0.43	-11.25	34.13	5.74	0.03

	${}^0\alpha \sim \text{PI}_{1200}$	0.24	-14.1	36.21	7.82	0.01
	${}^0\alpha \sim 1$	0	-16.84	38.61	10.22	0
Typical species						
	${}^1\alpha \sim \text{PI}_{1200} + \text{SF}_{600}$	0.67	-1.84	15.31	0	0.32
	${}^1\alpha \sim \text{Al}_{600} + \text{SF}_{600}$	0.67	-1.93	15.5	0.19	0.29
	${}^1\alpha \sim \text{SF}_{600}$	0.61	-3.81	15.63	0.32	0.27
	${}^1\alpha \sim \text{Al}_{600} + \text{PI}_{1200} + \text{SF}_{600}$	0.69	-0.78	17.57	2.26	0.1
	${}^1\alpha \sim \text{Al}_{600}$	0.29	-8.64	25.29	9.98	0
	${}^1\alpha \sim \text{PI}_{1200}$	0.26	-8.99	25.98	10.67	0
	${}^1\alpha \sim \text{Al}_{600} + \text{PI}_{1200}$	0.36	-7.25	26.13	10.82	0
	${}^1\alpha \sim 1$	0	-11.94	28.8	13.49	0
Dominant species						
	${}^2\alpha \sim \text{PI}_{1200} + \text{SF}_{600}$	0.74	2.42	6.8	0	0.49
	${}^2\alpha \sim \text{SF}_{600}$	0.67	0.03	7.95	1.14	0.28
	${}^2\alpha \sim \text{Al}_{600} + \text{SF}_{600}$	0.7	1.23	9.17	2.37	0.15
	${}^2\alpha \sim \text{Al}_{600} + \text{PI}_{1200} + \text{SF}_{600}$	0.73	2.88	10.23	3.43	0.09
	${}^2\alpha \sim \text{PI}_{1200}$	0.27	-6.39	20.79	13.99	0
	${}^2\alpha \sim \text{Al}_{600}$	0.23	-6.83	21.65	14.85	0
	${}^2\alpha \sim \text{Al}_{600} + \text{PI}_{1200}$	0.33	-5.19	22.02	15.22	0
	${}^2\alpha \sim 1$	0	-9.49	23.91	17.11	0
β -diversity All species						
	${}^0\beta \sim \text{EC}_{800}$	0.4	-11.04	30.08	0	0.56
	${}^0\beta \sim \text{EC}_{800} + \text{PD}_{1400}$	0.45	-9.77	31.17	1.08	0.32
	${}^0\beta \sim \text{PD}_{1400}$	0.25	-12.79	33.59	3.5	0.1
	${}^0\beta \sim 1$	0	-15.68	36.27	6.19	0.03
Typical species						
	${}^1\beta \sim \text{PD}_{500}$	0.44	-13.71	35.41	0	0.63
	${}^1\beta \sim \text{PD}_{500} + \text{SF}_{1300}$	0.48	-12.61	36.86	1.44	0.3
	${}^1\beta \sim \text{SF}_{1300}$	0.24	-16.19	40.37	4.96	0.05
	${}^1\beta \sim 1$	0	-18.94	42.8	7.39	0.02
Dominant species						

${}^2\beta \sim +PD_{500} + -SF_{1300}$	0.45	-9.81	27.62	0	0.56
${}^2\beta \sim -AI_{500} + +PD_{500} + -SF_{1300}$	0.49	-8.6	28.84	1.23	0.3
${}^2\beta \sim -AI_{500} + -SF_{1300}$	0.34	-11.32	30.64	3.02	0.12
${}^2\beta \sim 1$	0	-15.17	35.26	7.64	0.01

Figure A. Association between buffer size and the coefficient of prediction (R^2_{CV}) between each landscape metric and each α -diversity metric (all, typical and dominant species) of the animal-dispersed tree seedling community in a fragmented rainforest in southeastern Mexico. Each point represents the R^2_{CV} value of a simple linear regression between a diversity metric and single landscape metric measured at each buffer size. The points above the horizontal grey line indicate the proportion of variation that can be predicted by the model, whereas those points below the horizontal line indicate the prediction power of the model is worse than for a null model. The scale of effect is indicated within each panel with vertical lines.

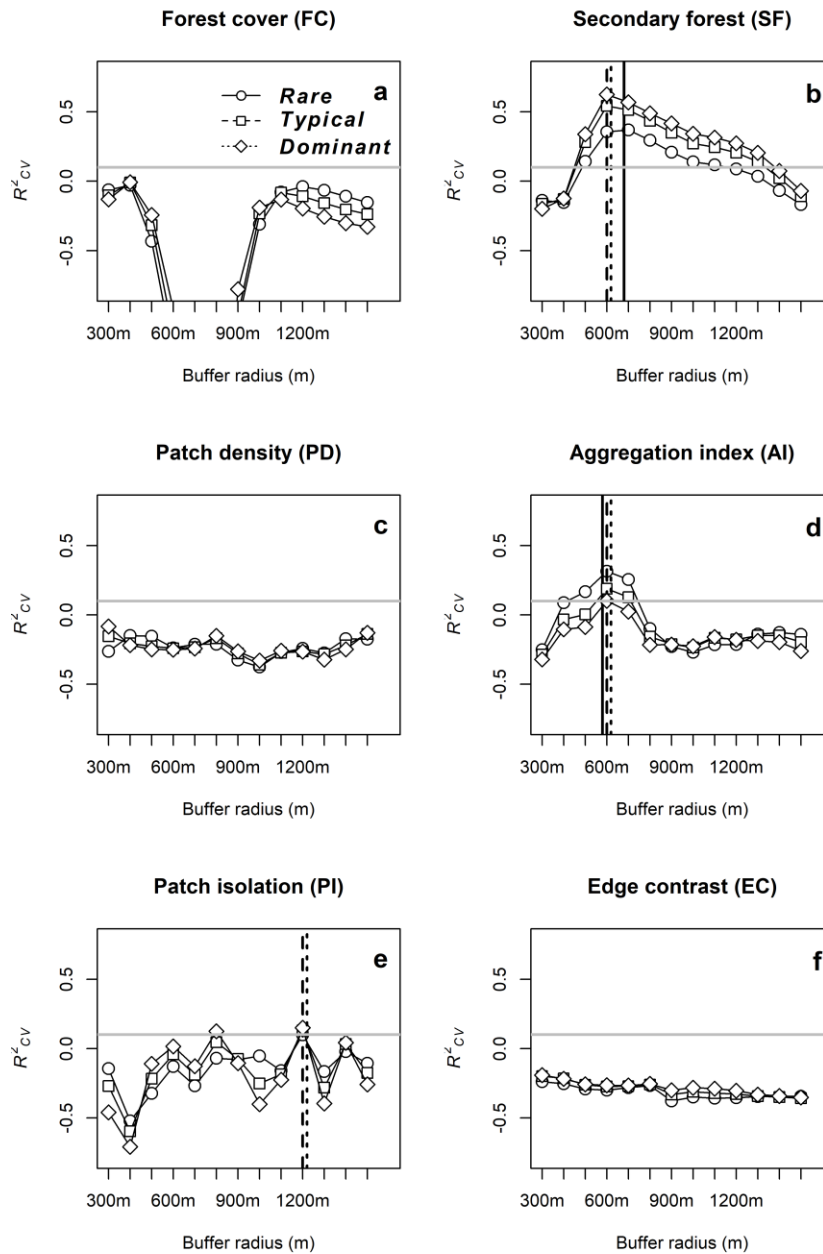


Figure B. Association between buffer size and the coefficient of prediction (R^2_{CV}) between each landscape metric and each β -diversity metric (all, typical and dominant species) of the animal-dispersed tree seedling community in a fragmented rainforest in southeastern Mexico. Each point represents the R^2_{CV} value of a simple linear regression between a diversity metric and single landscape metric measured at each buffer size. The points above the horizontal grey line indicate the proportion of variation that can be predicted by the model, whereas those points below the horizontal line indicate the prediction power of the model is worse than for a null model. The scale of effect is indicated within each panel with vertical lines.

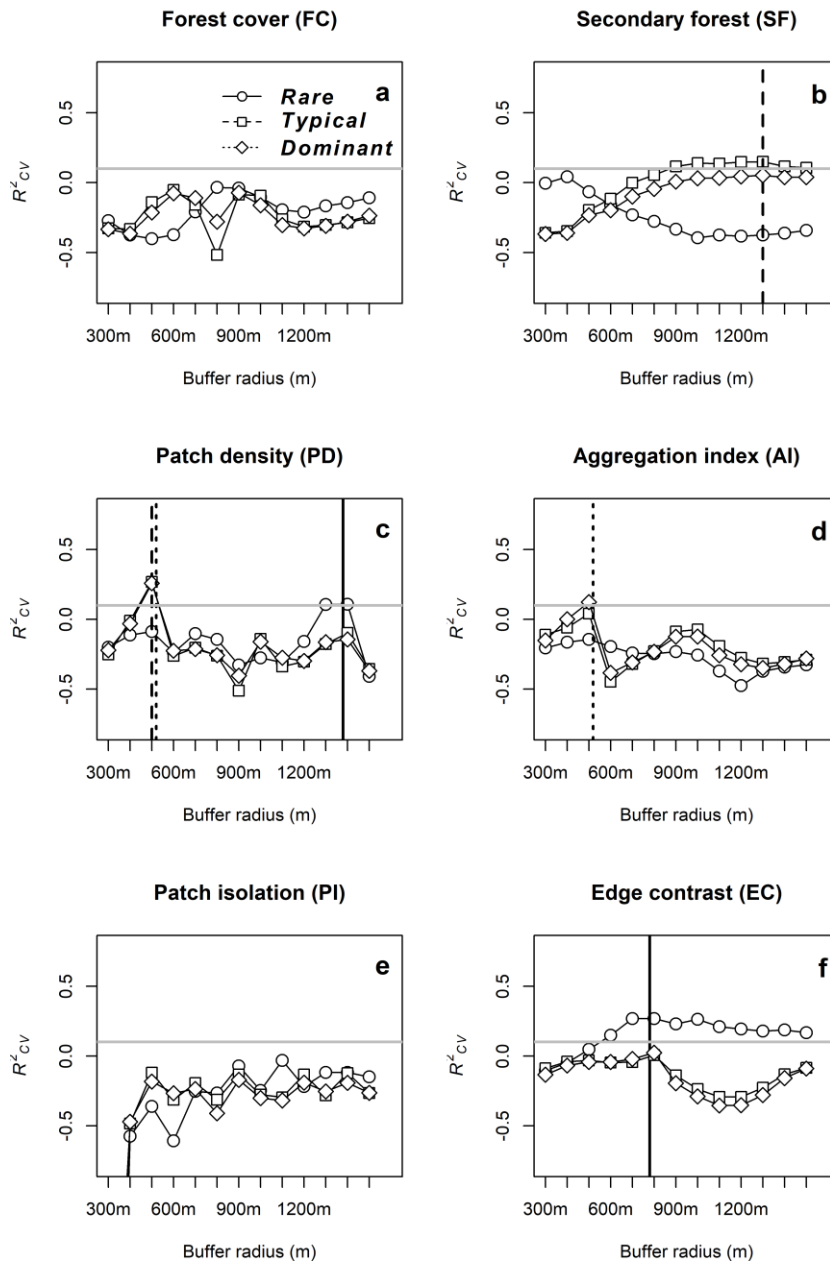


Figure C. Uncertainty around the estimated scale of effect between each landscape metric and each α -diversity metric (all, typical and dominant species) of the animal-dispersed tree seedling community, estimated by bootstrapping. For each landscape-diversity combination we randomly re-sampled the data from n patches, with replacement, from the set of the 16 forest patches, 1000 times. We then estimated the scale of effect for each resampled data. The total number of times (out of 1000) that each buffer size was selected as the scale of effect is plotted. The estimated scale of effect (red arrow) is the buffer size with the highest R^2_{CV} for the complete data set (see Fig A).

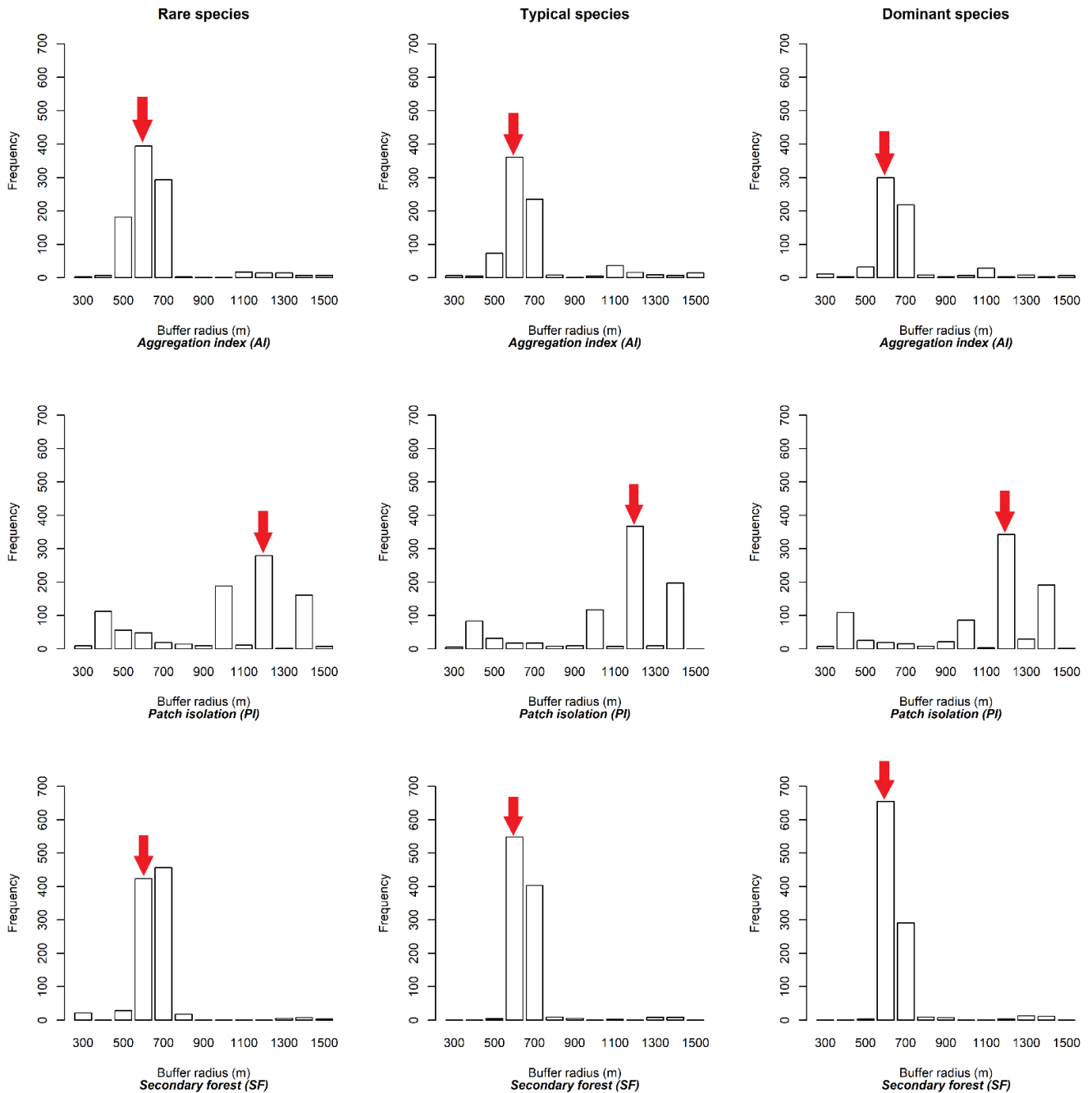


Figure D. Uncertainty around the estimated scale of effect between each landscape metric and each β -diversity metric (all, typical and dominant species) of the animal-dispersed tree seedling community, estimated by bootstrapping. For each landscape-diversity combination we randomly re-sampled the data from n patches, with replacement, from the set of the 16 forest patches, 1000 times. We then estimated the scale of effect for each resampled data. The total number of times (out of 1000) that each buffer size was selected as the scale of effect is plotted. The estimated scale of effect (red arrow) is the buffer size with the highest R^2_{CV} for the complete data set (see Fig B).

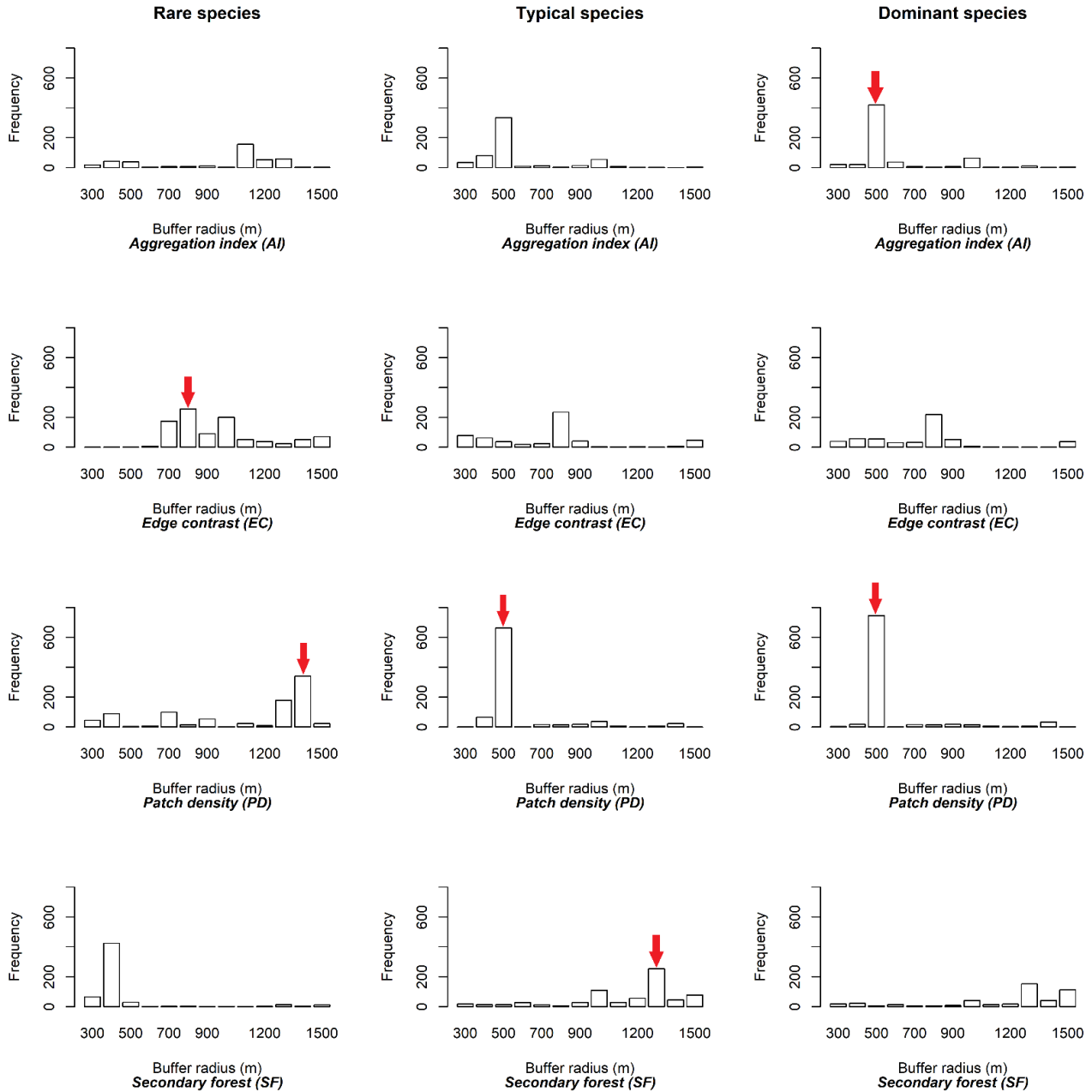


Figure E. Values of a) α - and b) β -diversity orders for tree seedlings within the patches' sampling plots in a fragmented tropical forest in southeastern Mexico. We observed ${}^0\alpha$ -diversity was significantly higher (2.703 ± 0.716) than ${}^1\alpha$ (2.295 ± 0.527) and ${}^2\alpha$ (2.114 ± 0.452) within sampling plots ($\chi^2 = 6.99$; $d.f. = 2$; $P = 0.0302$). This also held true for β -diversity ($\chi^2 = 32.511$; $d.f. = 2$; $P < 0.001$; Figure 2b), which had ${}^0\beta$ values higher (4.303 ± 0.665) than ${}^1\beta$ (2.258 ± 0.816) and ${}^2\beta$ (1.617 ± 0.644).

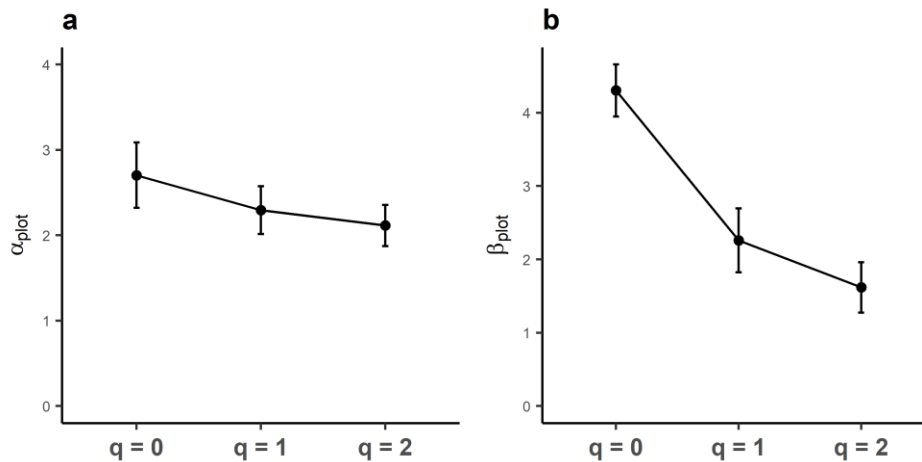
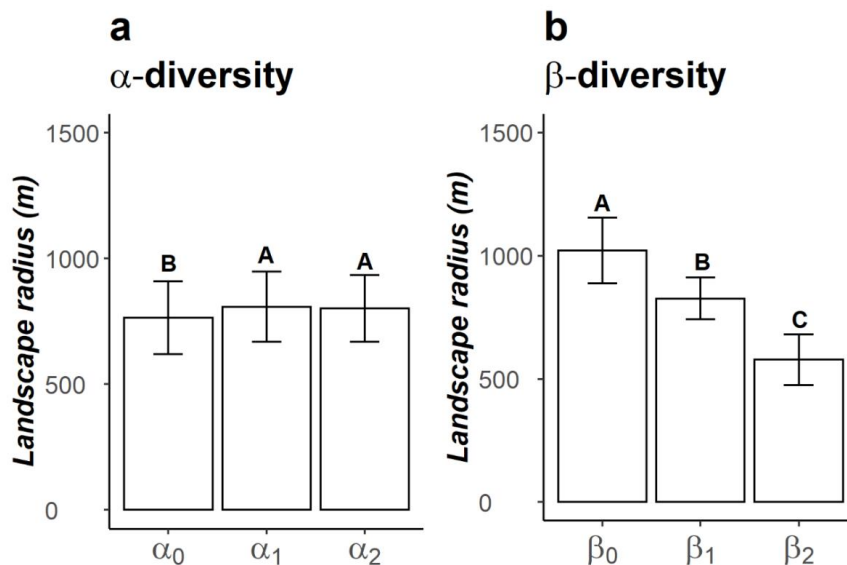


Figure F. Differences in the scale of effect between each landscape metric and each α - and β -diversity metric of the animal-dispersed tree seedling community. The values employed were estimated by bootstrapping. For each landscape-diversity combination we randomly re-sampled the data from n patches, with replacement, from the set of the 16 forest patches, 1000 times. We then estimated the scale of effect for each resampled data. The bars represent the mean and the whiskers the standard error.



**CAPÍTULO IV. THE ROLE OF LANDSCAPE MATRIX EFFECTS ON
MAMMAL OCCUPANCY WITHIN RAINFOREST PATCHES**

The role of landscape matrix effects on mammals' occupancy within rainforest patches

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Abstract

Landscape matrix plays a key role to sustain patch occupancy of terrestrial mammals in fragmented habitats, whether by contributing to habitat amount or by restraining the inter-patch movement. Nevertheless, the magnitude and spatial extent at which these matrix effects operate remains unexplored. We assessed the relative importance of matrix effects on habitat amount and resistance to movement on patch occupancy of four terrestrial mammals in a fragmented rainforest in southeastern Mexico. We monitored the incidence of each species within 16 forest patches. We then employed landscape composition metrics to measure matrix effects on habitat amount and textural variables from satellite imagery to estimate matrix resistance. We assessed both matrix effects on patch occupancy at four spatial scales (200, 300, 400 and 500 ha). We found a negative effect of both matrix effects on patch occupancy. However, matrix effects on habitat amount were more important for mammals with small home-range sizes than matrix effects on resistance to movement. Consequently, we observed that matrix effects on habitat amount operated at smaller spatial scales than matrix resistance. Our results suggest that the

magnitude and contribution of matrix effects differ among species and spatial scales, urging to assess these effects separately in fragmentation studies. We also highlight the key role of remote sensing techniques to estimate matrix resistance at multiple spatial scales in the tropics.

Introduction

The conversion of natural habitats to different land uses accounts for ca. 70% of forest loss, threatening biodiversity worldwide (Leadley et al., 2014; Song et al., 2018). Tropical rainforests are particularly vulnerable, because 10% of their remaining area persists as small and isolated patches within human-modified landscapes (Gibbs et al., 2010; Hansen et al., 2013; Lewis et al., 2015; Taubert et al., 2018). Biodiversity conservation in these habitats urges to understand how landscape structure promote the persistence of the remaining species within forest patches (Chazdon et al., 2009; Gardner et al., 2009).

Landscape matrix plays a key role in biodiversity conservation within fragmented habitats (Kupfer et al., 2006; Watling et al., 2011). The matrix influences species persistence mainly through effects on resources' availability (Driscoll et al., 2013) and functional connectivity (Eycott et al., 2010; Prevedello & Vieira, 2010). These effects are driven by matrix traits and species-specific requirements. On the one hand, matrix effects on resources availability (i.e., species breeding/foraging requirements) is associated to the composition of its land covers (Tscharntke et al., 2012; Watling et al., 2011). On the other hand, matrix can facilitate/impede the inter-patch movement throughout the structural resemblance of matrix vegetation with forest patches, as well as species movement requirements (Daniel et al., 2010; Driscoll et al., 2013; Prevedello & Vieira, 2010). Accordingly, understanding the role of matrix effects on the remaining species is crucial for reinforcing conservation strategies in fragmented landscapes (Driscoll et al., 2013; Lindenmayer et al., 2008; Watling et al., 2011).

Studies that appropriately differentiate among matrix effects, however, are lacking, leading to ineffective management practices (but see Watson et al. 2014). Untangling matrix effects can be achieved using composition metrics from landscape studies (Garmendia et al., 2013; Umetsu et al., 2008) and resistance measurements based on circuit theory (McRae et al., 2008; McRae & Beier, 2007). The latter has been successfully employed to measure the inter-patch movement resistance in a variety of scenarios derived from vegetation data (S. J. Watson et al., 2014). Nonetheless, resistance measurements are usually derived from expert-based estimations of categorical data (Ayram et al., 2015; Zeller et al., 2012), leading to inaccurate representations of connectivity (Koenig & Bender, 2018; Muratet et al., 2013). These shortcomings can be avoided by using satellite sensor data (Cavender-Bares et al., 2020; Muratet et al., 2013). The spectral values of satellite imagery are associated with different vegetation components (Lausch et al., 2020; Madritch et al., 2020; Pettorelli et al., 2014; Vihervaara et al., 2017) and are widely employed to map forest structure, disturbance, and suitability for species persistence in a variety of ecosystems (Baccini et al., 2012; M. Pfeifer et al., 2016; Strassburg et al., 2010; Turner et al., 2003). Textural variables are of particular importance in the tropics, since they have proven to overcome the limitations associated with the saturation of spectral bands at high biomass levels (Castillo-Santiago et al., 2010; Gallardo-Cruz et al., 2012; Gallardo-Cruz et al., 2018; Solórzano et al., 2017).

Furthermore, there is also a lack of knowledge regarding the spatial extent at which matrix effects are operating. The scale effect (SE) represents the spatial extent where landscape structure exerts its strongest effects on particular species (Fahrig, 2013; H. B. Jackson & Fahrig, 2012). The SE can be influenced by species traits (i.e. ranging behavior, habitat specialization, dispersal movements) and landscape predictors (habitat amount/fragmentation measures), suggesting that SE of landscape structure may vary among species and landscape patterns (Jackson & Fahrig, 2012, 2015; Miguet et al., 2016). This is a critical issue since the species'

responses can be overlooked if assessed at an incorrect scale, misleading conservation plans (Lenore Fahrig, 2013; H. B. Jackson & Fahrig, 2015). Nonetheless, the SE of landscape structure remains unexplored for most species (Lenore Fahrig, 2013; H. B. Jackson & Fahrig, 2012; McGarigal et al., 2016), and to our knowledge, no one has ever tested between matrix effects (Driscoll et al., 2013; Miguet et al., 2016).

Terrestrial mammals play a key role in the long-term functioning of rainforests (Galetti & Dirzo, 2013; Stoner et al., 2007; Terborgh, 2013). Nevertheless, they are highly vulnerable to habitat loss and fragmentation (Ahumada et al., 2011; Lyra-Jorge et al., 2010; M. Pfeifer et al., 2017; Wearn et al., 2017). Several studies have found that the proportion of tree cover in the surrounding matrix is associated with the abundance, occupancy and richness of mammals within forest patches at varying spatial scales (García-Marmolejo et al., 2015; Garmendia et al., 2013; Nagy-Reis et al., 2017; Umetsu et al., 2008). These studies, however, have overlooked the identity and the scale of matrix effects involved in the persistence of terrestrial mammals. Therefore, the aim of this study is to assess the role of matrix effects on (1) habitat amount and (2) resistance to movement on patch occupancy by four large terrestrial mammals using a multi-scale spatial approach. We untangled matrix effects by using composition metrics to represent matrix effects on habitat amount and matrix resistance estimations derived from satellite imagery to represent resistance to movement. Thereafter, we assessed the contribution of each parameter through occupancy models' selection and their variables' importance at different spatial extents. Following the hypothesis described by Miguet et al. (2016), we expected that the scale of effect of matrix effects on habitat amount should be smaller than those on resistance movement.

Material & Methods

Study area

We conducted the fieldwork at the Lacandona rainforest in southeastern Mexico (Fig 1a). The land-cover change to cattle pastures has shrunk its extension by two thirds in the last 40 years (De Jong et al., 2000; Kolb & Galicia, 2018; Mendoza & Dirzo, 1999). This region, however, encompasses the largest tropical rainforest of Mesoamerica (González-Espinosa et al., 2005). The vegetation is dominated by lowland tropical forest, reaching up to 40 m in height in alluvial terraces (Pennington & Sarukhán, 2005). The average monthly temperature is 24 to 26 °C, and the annual precipitation ranges from 2500 to 3500 mm (Mendoza & Dirzo, 1999). We selected 16 fragments of floodplain forest, ranging between 1 ha and 63 ha (Fig 1b).

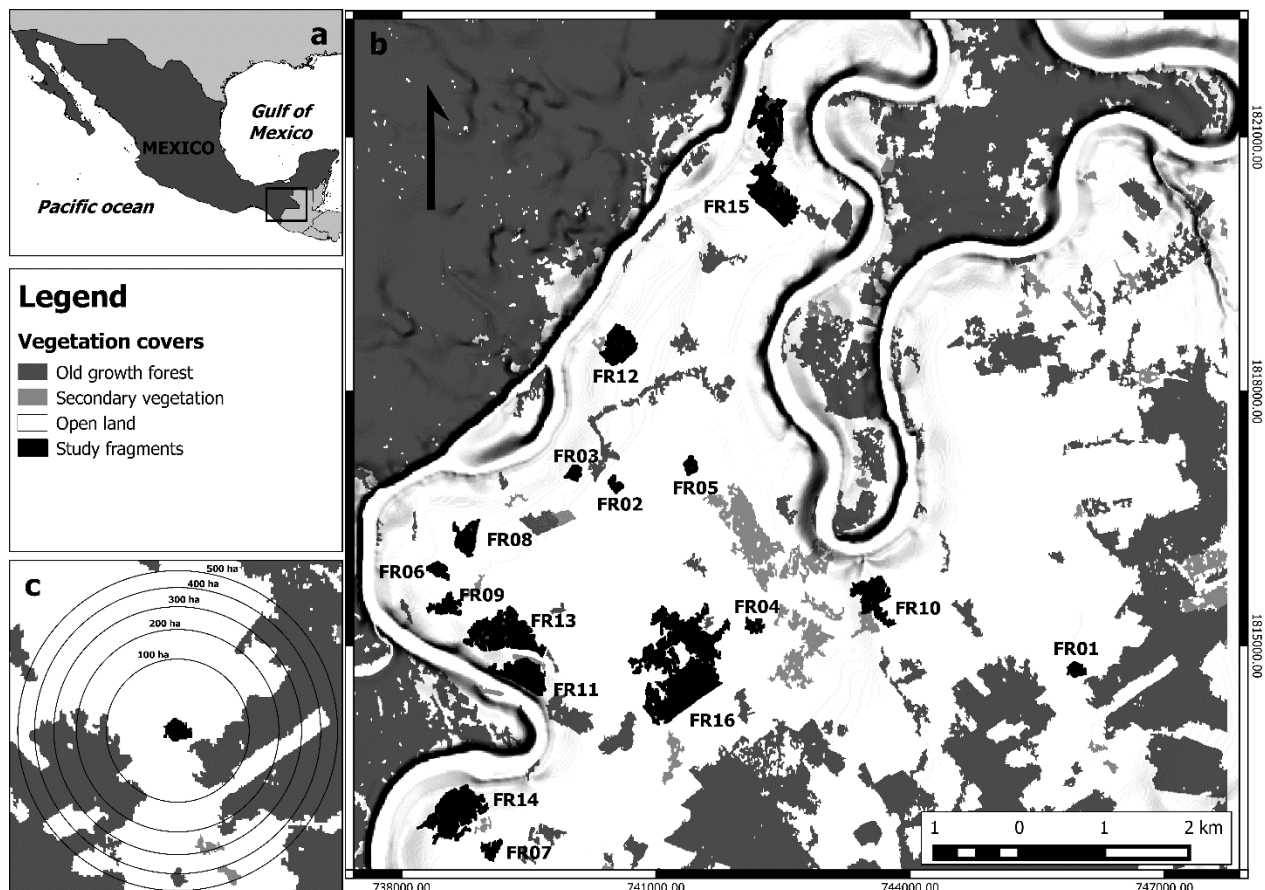


Fig 1. Location of the study area in the Lacandona rain forest in Chiapas, southeastern Mexico (a). We show the location of the 16 study forest fragments (black) in the Marqués de

Comillas Region (b). The five buffer sizes around the geographic center of the focal patch are also indicated (c).

Experimental design

We surveyed the terrestrial mammal community from February to June of 2012, during the peak of mammal activity (Haugaasen & Peres, 2007). We combined two sampling methods to obtain a more accurate estimation of species number within each fragment in a short period of time. Firstly, we used camera traps (Cuddeback IR) for 30 consecutive days. At the end of the sampling period, we identified each mammal species following field guides (Reid, 2009).

Secondly, we established two 500 × 3 m trails within each fragment and recorded the presence of mammals through sightings and indirect evidence (i.e., footprints, feces, feeding sites) three times a week for four weeks. We only considered terrestrial mammals with a body mass ≥ 1 kg (Medellin, 1994). We detected 15 species (Table S1), but we focused our analysis on 4 large-sized species: the white-tailed deer (*Odocoileus virginianus*), the ocelot (*Leopardus pardalis*), the white-collared peccary (*Pecari tajacu*) and the jaguar (*Panthera onca*).

Image classification

We employed a multispectral SPOT-5 satellite image recorded in March 2012. We carried out a supervised classification by using ground control points evenly distributed across different cover classes categorized as old-growth forest, secondary forest, arboreal crops, cattle pasture, anthropogenic features, bare soil, semiaquatic vegetation and water bodies. We used the GRASS GIS software (GRASS Development Team, 2017) for the complete procedure. The overall classification accuracy was 79%.

Landscape metrics

We measured three landscape metrics relevant to terrestrial mammal persistence in fragmented tropical forests (Garmendia et al., 2013; Goulart et al., 2009; Thornton et al., 2011; Umetsu et al., 2008) and one resistance distance within four spatial extents (buffers with sizes of 200, 300, 400 and 500 ha; Fig 1c) around the centroid of each patch (Table 1). Although the overlapping between buffers of nearby sampling points was high in the larger spatial extents (400 and 500 ha), the degree of spatial autocorrelation in the model residuals is not necessarily associated with the extent of landscape overlap but rather with the proximity between sampling sites (Cleary et al., 2016; Zuckerberg et al., 2012). According to Moran's I autocorrelation tests, we did not find spatial autocorrelation between the distance of sampling sites and the number of detection records of each species (Table S2).

Table 1. Landscape metrics and resistance estimators employed for the analysis

Landscape metric	Formula	Description
FC	$FC = 100 \times \left[\frac{\sum_{j=1}^n a_i}{A} \right]$	Where a is the area (m^2) of forest patch i and A the total landscape area
PD	$PD = \frac{n_i}{A} (10,000)(100)$	Where n_i is the total number of patches in the landscape of the land cover i and A is the total landscape area (m^2)
MH	$MH = 100 \times \left[\sum_{i=1}^6 \left(\frac{a_i \cdot Q_i}{A} \right) \right]$	Where a_i is the total area (m^2) of the land cover i , Q_i is a relative quality index obtained by dividing the suitability values of each land cover (as seen in Methods) by seven (the maximum suitability

		value, corresponding to the old-growth forest cover) and A is the total landscape area (m^2)
MR	$VAR = \sum_{i,j=0}^{N-1} P_{i,j} (i - MEAN)^2$	Where $P_{i,j}$ is the (i,j) element of the GLCM, and represents the probability of finding the reference pixel value i in combination with a neighbor pixel value j

We selected two metrics related to landscape composition and one related to landscape configuration. The composition metrics were the percentage of old-growth forest cover (FC) and the percentage of habitat accounting the amount high-quality matrix (MH). The FC is associated to habitat available, whereas MH is an indicator of additional habitat when accounting for high-quality matrices (Umetsu et al., 2008). The configuration metric was the density of patches (PD), which is an accurate descriptor of habitat fragmentation (Lenore Fahrig, 2013, 2017; Rayfield et al., 2011).

Matrix effect on habitat amount

We constructed the MH metric by assigning quality values to each land cover of the study area. The quality values were based on the assumption that overall species presence declines along a land-use gradient (Gardner et al., 2009). We ranked the quality of each land-cover type based on the suitability of vegetation structure for terrestrial mammals' feeding, movement and/or habitat on a six-point scale (Garmendia et al., 2013): 1 (water bodies, with the lowest suitability); 2 (anthropogenic cover); 3 (cattle pasture); 4 (arboreal crops); 5 (semiaquatic vegetation); and 6 (secondary forest, with the highest suitability). Then, we calculated the relative quality of land covers by dividing each value by seven (the quality value of old-growth

forest). Finally, we related the relative quality values to the area of each land cover to recalculate the percentage of total matrix (Table 1).

Matrix effect on resistance movement

We constructed the matrix resistance metric (MR) through the “variance” texture variable of the red (*R*) and near-infrared (*NIR*) spectral bands from the SPOT-5 satellite image (Table 1). The spectral values of the *R* and *NIR* layers are strongly associated to the canopy cover and aboveground biomass of woody vegetation of tropical rainforests, respectively (Castillo-Santiago et al., 2010; José Alberto Gallardo-Cruz et al., 2018; Sarker & Nichol, 2011). The texture variables were estimated by constructing a gray-level co-occurrence matrix (GLCM). Firstly, we transformed the pixel values to 64 levels of gray, and then we calculated the texture metrics within a window size of 9×9 pixels using an offset of one pixel in four angle directions (0° , 45° , 90° and 135°). We averaged the metrics calculated in the four directions to obtain a directionless metric. These are the optimal features for assessing the vegetation structure in the study site (Castillo-Santiago et al., 2010). We applied a mask to each band to eliminate water bodies and bare ground from the analysis. We employed the *glcm* package for the entire procedure (Zvloff, 2016).

Next, we employed a natural breaks classification to divide the distribution of the resulting cell values into 20 resistance classes (from 10 to 200). We added a buffer around each resistance layer to minimize the overestimation of resistance distances caused by artificial boundaries (Koen et al., 2010). Finally, we estimated patch isolation by matrix resistance using the Circuitscape software (McRae et al., 2008). We modeled the resistance distance across all pairs of forest patches within each buffer and obtained the mean resistance distance for each buffer size.

Statistical analysis

We used occupancy modeling with a likelihood-based approach (D. I. MacKenzie et al., 2006) to estimate patch occupancy (ψ) of each species while accounting for detection probability (p). We regarded occupancy as a measure of mammal habitat use instead of true occupancy because the home range size of each species exceeds that of the sampling units (Hines et al., 2010; D.I. MacKenzie & Nichols, 2004; Nagy-Reis et al., 2017). This supposition did not violate the assumption that the sampling sites are closed to changes in occupancy during repeated surveys (D.I. MacKenzie, 2005). We employed a single-season approach, wherein data from camera trapping and trail sampling were included in different sampling occasions.

We determined a general model for p by developing models where ψ remained constant ($\psi(\cdot)$) while allowing p to vary as a function of either a single or a combination of the following observation covariates: survey method of each sampling occasion (camera trapping/trail sampling), patch use by cattle (presence/absence), and patch flooding during the rainy season (flooded/not flooded). We assessed the goodness of fit of these models using the Mackenzie and Bailey goodness of fit test for single-season occupancy (Darryl I. MacKenzie & Bailey, 2004). First, we calculated the Pearson χ^2 from the observed and expected values based on the detection histories and single-season occupancy model, respectively (D. I. MacKenzie et al., 2006; Darryl I. MacKenzie & Bailey, 2004). Then, we employed a parametric bootstrap approach (100 simulation per model) to obtain the p-value of the χ^2 statistic and the overdispersion parameter (\hat{c}) of each model using the *AICcmodavg* package (Mazerolle, 2020). We selected the best-fit models (i.e., those with non-significant χ^2 values and $\hat{c} < 2$; Table S3) and ranked them using the corrected version of the Akaike information criterion for small sample sizes (AICc). We accounted for those models with an AICc difference lower than two ($\Delta\text{AICc} < 2$) from the top-

ranking model as best-supported by the data (Burnham & Anderson, 2002). The model of survey method ($p(s)$) provided enough evidence of changes in detection probability for three species, whereas models that where p remained constant (null model) and that included patch flooding ($p(f)$) were the best supported by data for *P. tajacu* (Table S4).

We determined the SE of site covariates (Table 1) for each species by developing a general model for p while allowing ψ to be constant or to vary as a function of the same covariate measured at the different buffer sizes (D I MacKenzie, 2006). We previously assessed the resistance distances that best represented the MR metric, and we selected the R_v values for all species (Table S5). We estimated the Nagelkerke's R^2 index (Trust, 1991) and selected the model with the highest R^2 to determine the scale of effect of the covariate when different spatial extents were equally plausible (Table S6). We employed the *unmarked* package for the entire procedure (Fiske & Chandler, 2011). Once selected the SE of each covariate, we assessed the contribution and relative importance of landscape metrics on patch occupancy. We constructed a global model and developed the set of sub-models using the dredge function of the *MuMIn* package (Barton, 2018). We excluded those models that did not converge or had abnormal convergences (Welsh et al., 2013). Furthermore, we ranked the models as mentioned above and selected the subset of models for which the sum of the Akaike weights ($\sum w_i$) were ≤ 0.95 (Burnham & Anderson, 2002). We employed the selected models to estimate the relative contribution of each covariate through a model-average approach, and assessed their relative importance by summing the Akaike weights (w_i) of the models where the covariate was present (Burnham & Anderson, 2002).

Finally, we tested for differences in the SE between landscape metrics and among species with a nonparametric Kruskal-Wallis test using the R 3.5.2 statistical computing environment (R Development Core Team, 2018).

Results

Relative importance and effects of landscape metrics

We observed a high importance of patch density and forest cover on patch occupancy of all species. On one hand, density of patches was present in the best-supported models (Table 2) and represented the most important variable ($\sum w_i > 0.5$) for *O. virginianus*, *L. pardalis* and *P. onca* occupancy (Figs 2a, 2e and 2g). According to the averaged estimators, patch occupancy increased as the number of patches per unit area increased (Figs 2b, 2f and 2h). This was not true for *P. tajacu*, whose occupancy was not affected by the density of patches (Figs 2c, and 2d). The percentage of forest cover represented the second most important variable for *O. virginianus*, *P. tajacu* and *P. onca* (Figs 2a, 2c and 2g), and promoted patch occupancy between the 300-500ha spatial scales (Figs 2b, 2d and 2h).

Table 2. Set of occupancy models of four terrestrial mammals employed for the model selection assessment and model-average approach in the Lacandona rainforest in Mexico.

Specie	Model	LL	AICc	Δ_i	w_i	R^2	χ^2	P	\hat{c}
<i>O. virginianus</i>	$p(s)\psi(FC_{300}+PD_{400})$	-21.53	59.07	0	0.59	0.77	84.4	0.39	1.24
	$p(s)\psi(MH_{300}+PD_{400})$	-22.96	61.92	2.85	0.14	0.73	84.37	0.36	1.43
	$p(s)\psi(PD_{400})$	-25.37	62.38	3.31	0.11	0.62	84.21	0.37	1.12
	$p(s)\psi(.)$	-27.47	62.94	3.87	0.08	0.51	84.18	0.37	1.25
	$p(s)\psi(MR_{200}+PD_{400})$	-24.3	64.61	5.54	0.04	0.67	84.17	0.48	0.9
	$p(s)\psi(MR_{200})$	-27.29	66.21	7.14	0.02	0.52	84.18	0.34	1.17
	$p(s)\psi(MH_{300}+MR_{200}+PD_{400})$	-22.68	66.69	7.62	0.01	0.74	84.42	0.48	1.13
	$p(.)\psi(MH_{300}+PD_{400})$	-28.56	68.75	9.68	0	0.43	131.76	0.75	0.72
	$p(.)\psi(PD_{400})$	-30.92	69.84	10.77	0	0.23	131.27	0.7	0.71
	$p(.)\psi(.)$	-33	70.92	11.86	0	0	131.07	0.63	0.82
$p(.)\psi(MR_{200}+PD_{400})$	-29.84	71.31	12.24	0	0.33	131.07	0.64	0.72	

	$p(.)\psi(MR_{200})$	-32.81	73.63	14.56	0	0.02	131.05	0.74	0.65
<i>P. tajacu</i>	$p(.)\psi(MR_{500})$	-51.39	110.78	0	0.24	0.21	346.05	0.02	1.48
	$p(.)\psi(.)$	-53.23	111.38	0.6	0.18	0	345.92	0.02	1.38
	$p(.)\psi(FC_{500})$	-51.79	111.59	0.81	0.16	0.16	345.84	0.03	1.42
	$p(.)\psi(FC_{500}+MR_{500})$	-50.84	113.31	2.53	0.07	0.26	345.92	0.02	1.43
	$p(.)\psi(FC_{500}+MH_{400})$	-50.92	113.47	2.69	0.06	0.25	345.98	0.05	1.4
	$p(.)\psi(FC_{500}+PD_{400})$	-50.97	113.58	2.8	0.06	0.25	355.69	0.05	1.48
	$p(.)\psi(MR_{500}+PD_{400})$	-51.15	113.94	3.16	0.05	0.23	346.05	0.02	1.41
	$p(.)\psi(PD_{400})$	-53.02	114.04	3.26	0.05	0.03	345.98	0.04	1.46
	$p(.)\psi(MH_{400})$	-53.08	114.16	3.38	0.04	0.02	345.91	0.05	1.41
	$p(.)\psi(MH_{400}+MR_{500})$	-51.36	114.35	3.57	0.04	0.21	346.04	0.03	1.47
	$p(.)\psi(FC_{500}+MH_{400}+MR_{500})$	-50.17	116.34	5.56	0.01	0.32	346.12	0.03	1.41
	$p(.)\psi(FC_{500}+MR_{500}+PD_{400})$	-50.72	117.44	6.66	0.01	0.27	346.03	0.07	1.42
	$p(.)\psi(MH_{400}+PD_{400})$	-52.93	117.49	6.71	0.01	0.04	345.79	0.04	1.44
	$p(.)\psi(FC_{500}+MH_{400}+PD_{400})$	-50.91	117.82	7.04	0.01	0.25	355.9	0.06	1.49
	$p(.)\psi(MH_{400}+MR_{500}+PD_{400})$	-51.14	118.29	7.51	0.01	0.23	345.98	0.02	1.43
$p(.)\psi(FC_{500}+MH_{400}+MR_{500}+PD_{400})$	-50.02	121.38	10.6	0	0.33	345.99	0.03	1.44	
<i>L. pardalis</i>	$p(s)\psi(MH_{300}+PD_{400})$	-30.02	76.04	0	0.58	0.72	102.12	0.57	0.64
	$p(s)\psi(PD_{400})$	-33.48	78.59	2.55	0.16	0.56	101.18	0.53	0.74
	$p(.)\psi(MH_{300}+PD_{400})$	-34.12	79.88	3.83	0.08	0.53	182.9	0.6	0.91
	$p(s)\psi(.)$	-36.02	80.03	3.99	0.08	0.4	101.15	0.54	0.73
	$p(s)\psi(MR_{500}+PD_{400})$	-32.65	81.3	5.26	0.04	0.61	101.13	0.58	0.56
	$p(s)\psi(MR_{500})$	-35.46	82.55	6.51	0.02	0.44	101.25	0.56	0.75
	$p(.)\psi(PD_{400})$	-37.52	83.04	6.99	0.02	0.27	184.26	0.67	0.85
	$p(.)\psi(MR_{500}+PD_{400})$	-36.69	85.01	8.97	0.01	0.35	184.52	0.64	0.84
	$p(.)\psi(.)$	-40.05	85.03	8.99	0.01	0	184.39	0.65	0.84
	$p(.)\psi(MR_{500})$	-39.5	86.99	10.95	0	0.07	184.28	0.63	0.89
<i>P. onca</i>	$p(s)\psi(FC_{400}+PD_{400})$	-15.3	46.59	0	0.43	0.81	10.93	0.63	0.82
	$p(s)\psi(.)$	-19.8	47.59	1	0.26	0.65	10.91	0.67	0.87
	$p(s)\psi(PD_{400})$	-18.48	48.6	2.01	0.16	0.7	10.95	0.59	0.9

$p(s)\psi(MR_{500})$	-19.13	49.9	3.31	0.08	0.68	10.92	0.64	0.85
$p(s)\psi(FC_{400}+MR_{500})$	-17.14	50.29	3.7	0.07	0.76	10.64	0.68	0.84
$p(.)\psi(.)$	-27.67	60.26	13.67	0	0	116.61	0.74	0.7
$p(.)\psi(PD_{400})$	-26.37	60.74	14.15	0	0.15	117.15	0.72	0.7
$p(.)\psi(MR_{500})$	-27.01	62.02	15.43	0	0.08	116.82	0.76	0.66
$p(.)\psi(MR_{500}+PD_{400})$	-25.4	62.44	15.85	0	0.25	117.46	0.78	0.64
$p(.)\psi(MH_{400}+MR_{500}+PD_{400})$	-24.01	64.03	17.44	0	0.38	116.85	0.73	0.72

The covariates are the percentage of forest cover (FC), patch density (PD), the percentage of habitat from matrix (MH) and matrix resistance (MR). The subscript numbers indicate the buffer size (in hectares) of each covariate. The null model does not include a covariate (.). The models' log-likelihood (LL), Akaike Information Criterion (AIC), Akaike differences (Δ_i), Akaike weights (w_i), the Nagelkerke's R^2 (R^2), the models' Pearson chi-square (χ^2), significance (P) and overdispersion parameter (\hat{c}) of each model are shown.

The relative contribution of matrix effects was lower than for forest cover and patch density. Interestingly, the importance between matrix effects varied among species, and were characterized for presenting contrasting values. The percentage of matrix habitat was more important than matrix resistance for *O. virginianus* and *L. pardalis* (Figs 2a and 2e), and affected negatively patch occupancy (Figs 2b and 2f). Contrarily, matrix resistance was more important than matrix habitat for *P. tajuacu* and *P. onca* (Figs 2c and 2g), whose occupancy decreased as patch isolation by resistance increased (Figs 2d and 2h). We must emphasize that matrix resistance had significant effects for all species, with the exception of *O. virginianus*.

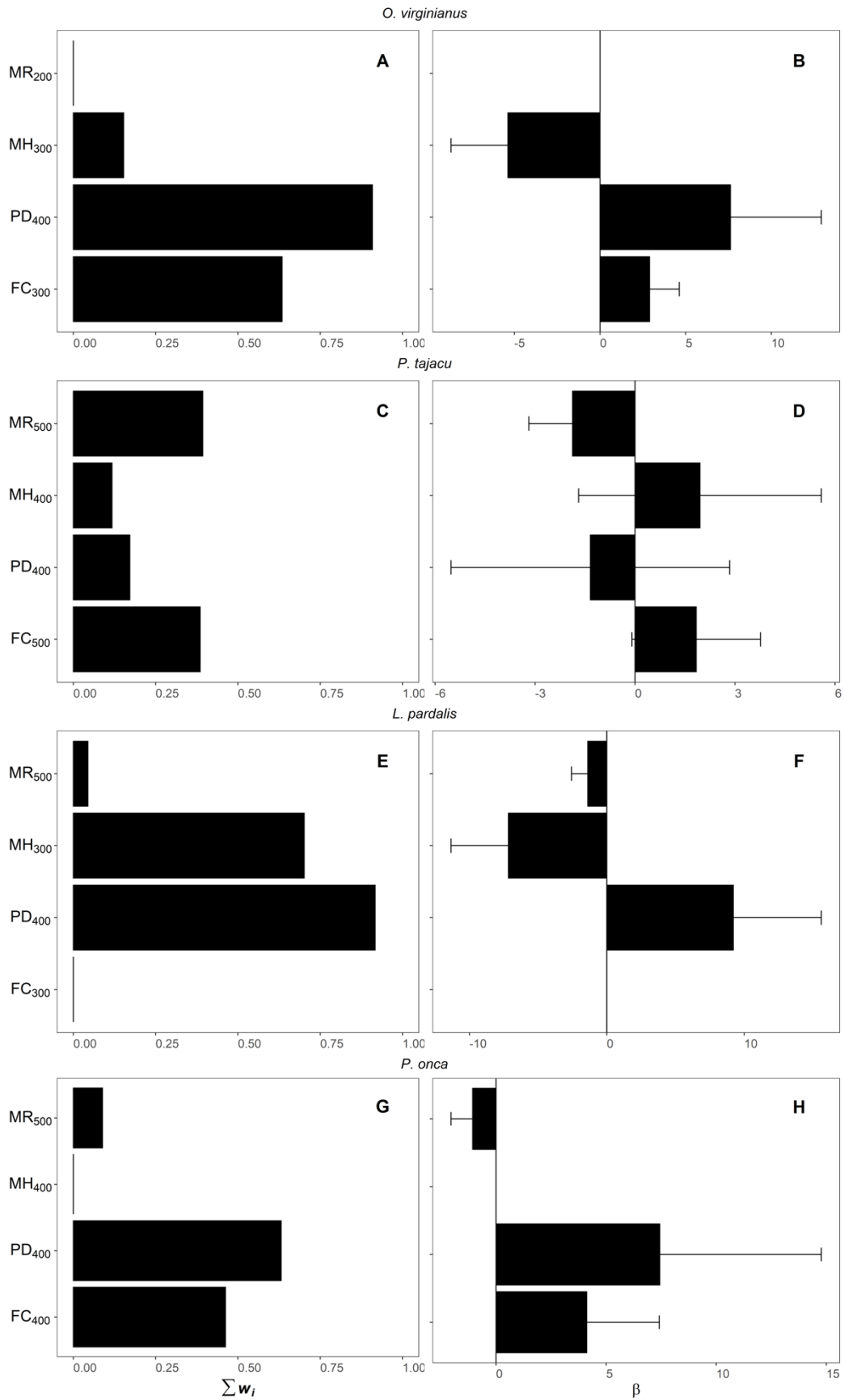


Fig 2. Relative importance and effects of covariates on patch occupancy of four terrestrial mammals in the Lacandona rainforest in Mexico. The importance of each covariate is represented by the sum of the Akaike weights ($\sum w_i$). The effects of each covariate were estimated through a model-averaged parameter estimate (β) of information-theoretic-based model selection and multimodel inference for occupancy. The whiskers represent the unconditional standard error (USE) and those for which the USE did not include zero are the influential covariates. The covariates are the percentage of forest cover (FC), patch density (PD), the percentage of habitat when accounting for suitable matrix (MH) and matrix resistance (MR). The subscript numbers indicate the scale of effect of each covariate.

Differences in the scale of effect among landscape metrics

Although we did not find differences in the scale of effect among species ($\chi^2 = 8.4$, d.f. = 3, $P = 0.26$), the scale of effect varied among landscape metrics ($\chi^2 = 8.25$, d.f. = 3, $P = 0.04$) when accounting for those with significant effects (Fig 2). The scale of effect of landscape metrics increased as follows: $MH < FC < PD < MR$. We also observed a consistent scale of effect for percentage of matrix habitat (300ha), patch density (400ha) and matrix resistance (500ha), regardless species.

Discussion

We aimed to investigate the mechanisms that sustain patch occupancy by terrestrial mammals at multiple spatial scales by untangling two matrix effects in the landscape. Our results showed matrix effects had negative effects on patch occupancy at varying spatial scales. Matrix effects on habitat amount affected patch occupancy at smaller spatial scales than matrix resistance. We also observed that the importance of these effects shifted among species. Such differences may be associated with the scale of interaction between landscape structure and species' home range, habitat requirements and dispersal distance (Miguet et al., 2016).

Matrix effects on habitat amount

Previous studies have associated matrix covers that provide food resources or improve habitat quality within species home range with the provision of complementary/supplementary resources near habitat patches (Asensio et al., 2009; Garmendia et al., 2013). In our study, the amount of habitat provided by the matrix increased along the proportion of semiaquatic vegetation and secondary forests, suggesting these covers are critical for the persistence of the herbivore mammals assessed. It is well documented that floodplains provide fruits and suitable conditions for collared peccaries (A. Keuroghlian & Eaton, 2008). Secondary forests are preferred by the white-tailed deer because of the high availability of leaves and stems, which represent most of their diet (Arceo et al., 2005; García-Marmolejo et al., 2015; Tejeda-Cruz et al., 2009). Nevertheless, they require old-growth forest landscapes for their long-term persistence (García-Marmolejo et al., 2015). The latter is consistent with the high importance of forest cover and patch density observed. We must emphasize that our study site is highly deforested (< 25% of forest cover) and the dominant land covers in the matrix are cattle pastures, which explains the negative relationship between forest cover and matrix habitat ($r = -0.82$). Accordingly, the negative effect of the percentage of matrix habitat on the occupancy of *O. virginianus* may be related with a low habitat quality and depauperate food resources from the dominant matrix covers within their home-range, suggesting the role of landscape matrix to sustain complementation/supplementation processes at smaller scales (Dunning et al., 1992).

The negative effect of matrix habitat on *L. pardalis* occupancy and its null effect for *P. onca* supports their condition as forest-dependent species. For *L. pardalis*, the use of matrix as complimentary habitat is possibly restricted to secondary forests, as ocelots tend to avoid flooded areas (Trolle & Kéry, 2005) and prefer large patches of densely-forested vegetation (Garmendia et al., 2013; V. L. Jackson et al., 2005; Pérez-Irineo & Santos-Moreno, 2016). This

also holds true for the jaguar, a species with large home range requirements and a preference for primary forest over human-modified habitats (Cullen-Junior et al., 2013; de la Torre et al., 2017; Morato et al., 2016). We must emphasize that jaguar records were restricted in landscapes near to large patches of undisturbed old-growth forest (> 100 ha), a pattern observed in similar cattle ranching areas (de Souza et al., 2018; Zimbres et al., 2017). These conditions, as well as the null association of matrix habitat to patch occupancy, suggest that jaguar presence may result from spillover effects.

Matrix effects on resistance to movement

Matrix resistance was important only at larger spatial scales (500ha) and decreased patch occupancy. These findings, along the positive effects of patch density at similar spatial scales (400ha) suggest that functional connectivity is driven by vegetation quality of matrix and by fragmentation *per se* (Bowman, Cappuccino, et al., 2002; Holzschuh et al., 2010; Radford et al., 2005; Saura & Rubio, 2010; Tischendorf & Fahrig, 2000).

An increase of patches in the landscape promotes the availability of habitat across the landscape, as well as the amount of forest edges, which had positive effects in mammals' richness in the study area at the same spatial scales (Garmendia et al., 2013). The same study also found the positive effects of forest patches and edges increased when accounting the positive effect of matrix quality at smaller spatial scales. The latter suggests that matrix quality promotes functional connectivity at large spatial scales, as observed elsewhere (Powney et al. 2011, 2012; Muratet et al., 2013). It is interesting that matrix effects on resistance movement were more important for mammals with large home-range sizes and dispersal distances, such as ocelots, collared-peccaries and jaguars (de la Torre et al., 2017; Di Bitetti et al., 2006; Alexine Keuroghlian et al., 2004). This is supported by Miguet et al. (2016), who predicted that scale of

effect increases with the mean dispersal distance, a trait strongly related with home range size (Bowman, Jaeger, et al., 2002).

In this sense, our findings indicate that connectivity loss has a negative effect on terrestrial mammals at large scales in fragmented rainforests and that matrix resistance plays a key role in driving the inter-patch movement of mammals with large dispersal distances (Thornton et al. 2011; Garmendia et al. 2013). Nonetheless, since we observed these effects in the largest spatial scale, more studies are needed to accurately find the scale of effect of matrix resistance, which might be operating at larger spatial scales (H. B. Jackson & Fahrig, 2015).

The role of remote sensing techniques to estimate matrix resistance

Our results indicate that resistance estimations from satellite can provide of realistic connectivity measurements. Although quality indices has been employed as indicator of matrix permeability in fragmented rainforests (Arroyo-Rodríguez, González-Perez, et al., 2013; Galán-Acedo et al., 2018, 2019; Garmendia et al., 2013), it does not represent isolation distances and/or movement rates (Rayfield et al., 2011; Tischendorf & Fahrig, 2000). Additionally, the use of textural variables on spectral data allow to obtain a more heterogeneous range of values associated to the structure of matrix vegetation than vegetation indices or spectral bands alone. This is critical in the tropics, where textural variables outperform vegetation indices when estimating forest structure (J. V. Solórzano et al., 2018, 2017).

Mammal species were only affected by R_v matrix resistance. In the tropics, the red band captures the reflectance of the crown area of canopy vegetation (Bourgoin et al., 2018; Steininger, 2000), and the textural variables are associated with the spatial variability of canopy structure (J. Alberto Gallardo-Cruz et al., 2012; M. Pfeifer et al., 2016; J. V. Solórzano et al., 2018). Tree canopy plays a key role in sustaining biodiversity in human-modified landscapes by

regulating both a forest's understory and edge microclimate (Ewers & Banks-Leite 2013; Hardwick et al. 2015). The association between forest canopy with terrestrial mammal's composition suggests that canopy cover also promote animal movement along human-modified landscapes (Pardo et al., 2018). Accordingly, our results indicate that matrix resistance for our species is driven by canopy structure through microclimate regulation.

Estimating more heterogeneous resistance values improves connectivity assessments. Connectivity networks that derived from NDVI values improved model fit more than their homogeneous counterparts when assessing community dissimilarity among habitat patches in an urban landscape (Muratet et al., 2013). We must emphasize, however, that the use of satellite imagery in connectivity assessments is considered in its infancy, and more studies are needed to evaluate its suitability to measure, map and understand functional connectivity for a variety of species in human-modified landscapes (Muratet et al., 2013; Pettorelli et al., 2014). Estimations of vegetation structure and matrix heterogeneity can be affected by the scale of the site, the image resolution and the temporal range of the image generation (Gamon et al., 2020; Hernando et al., 2017).

Forthcoming approaches aimed at improving our understanding regarding the role of landscape matrix on biodiversity conservation should consider untangling matrix effects at varying spatial scales. We must emphasize that further studies are needed to fully understand how mammals respond to canopy structure, as well as to perform analysis at larger spatial scales to determine the scale of effect of landscape matrix on connectivity. We also encourage more detailed connectivity assessments using satellite imagery for long-term monitoring purposes.

Conclusions

We conclude that the interaction between landscape matrix with species home range and dispersal requirements drives the scale of effect and importance of the different matrix effects. Mammals with small home-range sizes rely on the nearby matrix covers to fulfill their food and habitat requirements within their home range. Species with larger home ranges and long dispersion distances rely on the canopy structure of matrix vegetation to facilitate their inter-patch movement at larger spatial scales. Our findings also represent an important step in conservation planning because mammal's conservation in human-modified landscapes should favor the preservation of matrix covers that increase habitat amount and quality. Meanwhile, management practices that favor canopy cover at larger spatial extents –such as alley cropping, extensive silvopasture or riparian buffers– would reduce matrix resistance for vagile species (Martínez-Ramos et al., 2016). These recommendations support a land-sharing strategy to improve the conservation value of the remaining forest patches (Lenore Fahrig, 2017), particularly where forest is already hyper-fragmented (Hernández-Ruedas et al., 2018).

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Supplemental information

Table S1. Terrestrial mammals recorded during the field surveys in the Lacandona rainforest, Mexico

Common name	Scientific name
Striped hog-nosed skunk	<i>Conepatus semistriatus</i>
Lowland paca	<i>Cuniculus paca</i>
Common opossum	<i>Didelphis virginiana</i>
Ocelot	<i>Leopardus pardalis</i>
Margay	<i>Leopardus wiedii</i>
Red brocket	<i>Mazama temama</i>
White-nosed coati	<i>Nasua narica</i>
White-tailed deer	<i>Odocoileus virginianus</i>
Jaguar	<i>Panthera onca</i>
White-collared peccary	<i>Pecari tajacu</i>
Northern raccoon	<i>Procyon lotor</i>
Jaguarundi	<i>Puma yagouaroundi</i>
Tapeti	<i>Sylvilagus brasiliensis</i>
Baird's tapir	<i>Tapirus bairdii</i>
White-lipped peccary	<i>Tayassu pecari</i>

Table S2. Moran's I autocorrelation tests between the sampling sites distance and the number of detection records of four terrestrial mammal species in the Lacandona rainforest, Mexico.

Specie	Moran's I	P value
<i>O. virginianus</i>	- 0.0284 (0.0635)	0.547
<i>P. tajacu</i>	- 0.0684 (0.0669)	0.978
<i>L. pardalis</i>	0.0075 (0.0640)	0.246
<i>P. onca</i>	- 0.0879 (0.0626)	0.734

Table S3. Goodness of fit of detectability models of four terrestrial mammals in the Lacandona rainforest, Mexico.

Specie	Model	χ^2	P	\hat{c}
<i>O. virginianus</i>	$\psi(.)p(.)$	131.07	0.63	0.75
	$\psi(.)p(c)$	127.24	0.82	0.65
	$\psi(.)p(f)$	153.14	0.36	1.12
	$\psi(.)p(s)$	84.18	0.47	0.81
	$\psi(.)p(f+s)$	101.57	0.2	1.5
	$\psi(.)p(c+s)$	80.99	0.49	0.85
	$\psi(.)p(c+f)$	157.03	0.46	0.99
	$\psi(.)p(c+f+s)$	81.11	0.51	0.9
<i>P. tajacu</i>	$\psi(.)p(.)$	345.92	0.03	1.45
	$\psi(.)p(c)$	347.42	0	1.46
	$\psi(.)p(f)$	419.47	0.02	1.79
	$\psi(.)p(s)$	793.3	0.01	6.53
	$\psi(.)p(f+s)$	842.78	0.01	9.17

	$\psi(.)p(c+s)$	795.26	0	7.35
	$\psi(.)p(c+f)$	415.15	0	1.76
	$\psi(.)p(c+f+s)$	907.25	0.02	6.62
<i>L. pardalis</i>	$\psi(.)p(.)$	184.39	0.75	0.86
	$\psi(.)p(c)$	184.33	0.73	0.9
	$\psi(.)p(f)$	214.47	0.51	1.01
	$\psi(.)p(s)$	101.15	0.52	0.62
	$\psi(.)p(f+s)$	116.65	0.42	0.75
	$\psi(.)p(c+s)$	98.74	0.58	0.75
	$\psi(.)p(c+f)$	226.39	0.38	1.05
	$\psi(.)p(c+f+s)$	121.44	0.45	0.7
<i>P. onca</i>	$\psi(.)p(.)$	116.61	0.72	0.71
	$\psi(.)p(c)$	134.99	0.52	0.9
	$\psi(.)p(f)$	126.81	0.56	0.94
	$\psi(.)p(s)$	10.91	0.59	0.88
	$\psi(.)p(f+s)$	11.62	0.63	0.82
	$\psi(.)p(c+s)$	12.34	0.46	0.95
	$\psi(.)p(c+f)$	144.77	0.66	0.85
	$\psi(.)p(c+f+s)$	15.2	0.41	1

The covariates are the survey method used in each sampling occasion (s), the use of the patch by cattle (c), and patch flooding during the rainy season (f). The null model does not include a covariate (.). The models' Pearson chi-square (χ^2), significance (*P*) and overdispersion parameter (\hat{c}) are shown.

Table S4. Model selection analysis for detection probability (p) covariates for four terrestrial mammals in the Lacandona rainforest, Mexico.

Specie	Model	df	LL	$AICc$	Δ_i	w_i
<i>O. virginianus</i>	$\psi(.)p(s)$	3	-27.47	62.94	0	0.68
	$\psi(.)p(f+s)$	4	-27.21	66.06	3.11	0.14
	$\psi(.)p(c+s)$	4	-27.26	66.16	3.22	0.14
	$\psi(.)p(c+f+s)$	5	-27.26	70.51	7.57	0.02
	$\psi(.)p(.)$	2	-33	70.92	7.98	0.01
	$\psi(.)p(f)$	3	-32.67	73.34	10.4	0
	$\psi(.)p(c)$	3	-32.93	73.85	10.91	0
	$\psi(.)p(c+f)$	4	-32.62	76.88	13.94	0
<i>P. tajacu</i>	$\psi(.)p(.)$	2	-53.23	111.38	0	0.59
	$\psi(.)p(f)$	3	-52.67	113.35	1.97	0.22
	$\psi(.)p(c)$	3	-53.23	114.45	3.07	0.13
	$\psi(.)p(c+f)$	4	-52.24	116.11	4.72	0.06
<i>L. pardalis</i>	$\psi(.)p(s)$	3	-36.02	80.03	0	0.58
	$\psi(.)p(f+s)$	4	-35.21	82.05	2.02	0.21
	$\psi(.)p(c+s)$	4	-36	83.63	3.6	0.09
	$\psi(.)p(.)$	2	-40.05	85.03	5	0.05
	$\psi(.)p(c+f+s)$	5	-34.87	85.73	5.7	0.03
	$\psi(.)p(f)$	3	-39.19	86.38	6.35	0.02

	$\psi(\cdot)p(c)$	3	-40.05	88.11	8.08	0.01
	$\psi(\cdot)p(c+f)$	4	-38.83	89.3	9.27	0.01
<i>P. onca</i>	$\psi(\cdot)p(s)$	3	-19.8	47.59	0	0.7
	$\psi(\cdot)p(c+s)$	4	-19.61	50.86	3.27	0.14
	$\psi(\cdot)p(f+s)$	4	-19.66	50.97	3.37	0.13
	$\psi(\cdot)p(c+f+s)$	5	-18.91	53.83	6.24	0.03
	$\psi(\cdot)p(\cdot)$	2	-27.67	60.26	12.67	0
	$\psi(\cdot)p(c)$	3	-27.42	62.85	15.26	0
	$\psi(\cdot)p(f)$	3	-27.51	63.02	15.43	0
	$\psi(\cdot)p(c+f)$	4	-26.92	65.48	17.89	0

The covariates are the survey method used in each sampling occasion (s), the use of the patch by cattle (c), and patch flooding during the rainy season (f). The null model does not include a covariate (\cdot). The models' degrees of freedom (*df*), log-likelihood (*LL*), Akaike Information Criterion for small samples (*AICc*), Akaike differences (Δ_i), and Akaike weights (w_i) are shown. The selected models for further analysis are in bold.

Table S5. Model selection analysis for occupancy (ψ) covariates measured at different spatial extents (buffer sizes) for four terrestrial mammals in the Lacandona rainforest, Mexico.

Specie	Scale	Model	K	AICc	Δ AICc	w_i	LL	R ²
<i>L. pardalis</i>	200ha	$\psi(\cdot)p(s)$	3	79.23	0	0.76	-35.53	0
		$\psi(R_v)p(s)$	4	82.76	3.53	0.13	-35.38	0.02
		$\psi(NIR_v)p(s)$	4	83.05	3.81	0.11	-35.52	0
	300ha	$\psi(\cdot)p(s)$	3	80.03	0	0.86	-36.02	0
		$\psi(R_v)p(s)$	4	83.65	3.61	0.14	-36	0
		$\psi(NIR_v)p(s)$	4	100.28	20.25	0	-44.32	-1.84
	400ha	$\psi(\cdot)p(s)$	3	80.03	0	0.65	-36.02	0
		$\psi(NIR_v)p(s)$	4	82.57	2.54	0.18	-35.47	0.07
		$\psi(R_v)p(s)$	4	82.78	2.75	0.16	-35.57	0.05
	500ha	$\psi(\cdot)p(s)$	3	80.03	0	0.65	-36.02	0
		$\psi(R_v)p(s)$	4	82.56	2.53	0.18	-35.46	0.07
		$\psi(NIR_v)p(s)$	4	82.77	2.74	0.17	-35.57	0.06

<i>O. virginianus</i>	200ha	$\psi(.) p(s)$	3	62.35	0	0.77	-27.08	0
		$\psi(\text{NIR}_v) p(s)$	4	66.1	3.75	0.12	-27.05	0
		$\psi(\text{R}_v) p(s)$	4	66.12	3.77	0.12	-27.06	0
	300ha	$\psi(.) p(s)$	3	62.94	0	0.85	-27.47	0
		$\psi(\text{R}_v) p(s)$	4	66.36	3.42	0.15	-27.36	0.01
	400ha	$\psi(.) p(s)$	3	62.94	0	0.65	-27.47	0
		$\psi(\text{NIR}_v) p(s)$	4	65.05	2.11	0.23	-26.71	0.09
		$\psi(\text{R}_v) p(s)$	4	66.35	3.41	0.12	-27.36	0.01
	500ha	$\psi(.) p(s)$	3	62.94	0	0.65	-27.47	0
		$\psi(\text{NIR}_v) p(s)$	4	65.02	2.08	0.23	-26.69	0.1
		$\psi(\text{R}_v) p(s)$	4	66.24	3.3	0.12	-27.3	0.02
	<i>P. onca</i>	200ha	$\psi(.) p(s)$	3	47.18	0	0.76	-19.5
$\psi(\text{R}_v) p(s)$			4	50.65	3.47	0.13	-19.32	0.02
$\psi(\text{NIR}_v) p(s)$			4	51	3.82	0.11	-19.5	0
300ha		$\psi(.) p(s)$	3	47.59	0	0.86	-19.8	0
		$\psi(\text{R}_v) p(s)$	4	51.22	3.63	0.14	-19.79	0
400ha		$\psi(.) p(s)$	3	47.59	0	0.69	-19.8	0
		$\psi(\text{NIR}_v) p(s)$	4	50.25	2.66	0.18	-19.31	0.06
		$\psi(\text{R}_v) p(s)$	4	51.03	3.44	0.12	-19.7	0.01
500ha		$\psi(.) p(s)$	3	47.59	0	0.68	-19.8	0
		$\psi(\text{R}_v) p(s)$	4	49.92	2.33	0.21	-19.14	0.09
		$\psi(\text{NIR}_v) p(s)$	4	51.24	3.64	0.11	-19.8	0
<i>P. tajacu</i>		200ha	$\psi(.) p(s)$	3	83.5	0	0.55	-37.66
	$\psi(\text{R}_v) p(s)$		4	84.33	0.83	0.36	-36.16	0.18
	$\psi(\text{NIR}_v) p(s)$		4	87.07	3.57	0.09	-37.54	0.02
	300ha	$\psi(.) p(s)$	3	84.77	0	0.75	-38.39	0
		$\psi(\text{R}_v) p(s)$	4	86.95	2.18	0.25	-37.66	0.09
	400ha	$\psi(\text{R}_v) p(s)$	4	82.75	0	0.7	-35.56	0.3
		$\psi(.) p(s)$	3	84.77	2.02	0.26	-38.39	0
		$\psi(\text{NIR}_v) p(s)$	4	88.4	5.65	0.04	-38.38	0
	500ha	$\psi(\text{R}_v) p(s)$	4	84.73	0	0.46	-36.55	0.21
		$\psi(.) p(s)$	3	84.77	0.04	0.45	-38.39	0
		$\psi(\text{NIR}_v) p(s)$	4	87.93	3.2	0.09	-38.15	0.03

The covariates are resistance distances derived from the textural variance of the red (R_v), and near-infrared (NIR_v) spectral bands. The null model does not include a covariate (.). The models' number of parameters (K), Akaike Information Criterion for small samples (AICc), Akaike differences (ΔAICc), Akaike weights (w_i), log-likelihood (LL), and Nagelkerke's R^2 index of (R^2) are shown.

Table S6. Model selection analysis for occupancy (ψ) covariates for four terrestrial mammals in the Lacandona rainforest, Mexico.

Specie	Metric	Model	LL	$AICc$	Δ_i	w_i	R^2	χ^2	P	\hat{c}
<i>O. virginianus</i>	FC	$\psi(FC_{300})p(s)$	-22.59	56.82	0	0.33	0.47	92.92	0.3	0.89
		$\psi(FC_{200})p(s)$	-22.74	57.11	0.29	0.29	0.46	84.27	0.4	1.05
		$\psi(FC_{400})p(s)$	-22.8	57.24	0.42	0.27	0.46	90.73	0.28	0.67
		$\psi(FC_{500})p(s)$	-23.85	59.33	2.52	0.09	0.38	99.02	0.17	1.39
		$\Psi(.)p(.)$	-27.47	62.94	6.13	0.02	0	84.18	0.37	1.19
	PD	$\psi(PD_{400})p(s)$	-25.37	62.38	0	0.34	0.24	84.21	0.42	1.07
		$\psi(PD_{500})p(s)$	-25.51	62.65	0.27	0.3	0.23	84.21	0.37	1.13
		$\Psi(.)p(.)$	-27.47	62.94	0.57	0.26	0	84.18	0.34	1.14
		$\psi(PD_{200})p(s)$	-27.23	66.1	3.72	0.05	0.03	84.18	0.36	1.14
		$\psi(PD_{300})p(s)$	-27.37	66.37	3.99	0.05	0.01	84.19	0.4	1.28
	MH	$\psi(MH_{300})p(s)$	-22.85	57.33	0	0.63	0.45	91.14	0.27	1.23
		$\psi(MH_{400})p(s)$	-24.13	59.9	2.57	0.17	0.35	81.13	0.4	1.15
		$\psi(MH_{200})p(s)$	-24.33	60.29	2.96	0.14	0.34	84.69	0.36	1.15
		$\Psi(.)p(.)$	-27.47	62.94	5.62	0.04	0	84.18	0.34	1.17
		$\psi(MH_{500})p(s)$	-26.72	65.08	7.75	0.01	0.09	84.24	0.37	1.05
MR	$\Psi(.)p(.)$	-27.47	62.94	0	0.57	0	84.18	0.4	1	
	$\psi(MR_{200})p(s)$	-27.29	66.21	3.27	0.11	0.02	84.18	0.43	1.16	
	$\psi(MR_{500})p(s)$	-27.29	66.22	3.28	0.11	0.02	84.18	0.33	1.34	

		$\psi(MR_{300})p(s)$	-27.32	66.27	3.32	0.11	0.02	84.18	0.39	1.01
		$\psi(MR_{400})p(s)$	-27.39	66.41	3.47	0.1	0.01	84.18	0.41	1.05
<i>P. tajacu</i>	FC	$\psi(FC_{500})p(\cdot)$	-23.26	54.53	0	0.26	0.44	121.66	0.45	0.92
		$\psi(FC_{400})p(\cdot)$	-23.33	54.67	0.14	0.25	0.43	116.38	0.5	0.83
		$\psi(FC_{300})p(\cdot)$	-23.7	55.4	0.87	0.17	0.4	115.47	0.61	0.7
		$\psi(FC_{200})p(\cdot)$	-24.03	56.05	1.52	0.12	0.38	117.44	0.79	0.65
		$\psi(FC_{400})p(f)$	-22.97	57.58	3.05	0.06	0.46	117.34	0.55	0.78
		$\psi(FC_{300})p(f)$	-23.11	57.86	3.33	0.05	0.45	115.46	0.81	0.64
		$\psi(FC_{500})p(f)$	-23.12	57.88	3.35	0.05	0.45	119.79	0.57	0.79
		$\psi(FC_{200})p(f)$	-23.87	59.38	4.85	0.02	0.39	120.79	0.7	0.66
		$\Psi(\cdot)p(\cdot)$	-27.67	60.26	5.73	0.01	0	116.61	0.76	0.68
		$\psi(\cdot)p(f)$	-27.51	63.02	8.5	0	0.02	126.81	0.69	0.77
	PD	$\Psi(\cdot)p(\cdot)$	-27.67	60.26	0	0.3	0	116.61	0.73	0.62
		$\psi(PD_{400})p(\cdot)$	-26.37	60.74	0.49	0.24	0.15	117.15	0.77	0.63
		$\psi(PD_{500})p(\cdot)$	-27.15	62.31	2.05	0.11	0.06	116.49	0.7	0.68
		$\psi(PD_{200})p(\cdot)$	-27.22	62.44	2.18	0.1	0.06	116.8	0.65	0.73
		$\psi(PD_{300})p(\cdot)$	-27.44	62.88	2.62	0.08	0.03	116.49	0.77	0.64
		$\psi(\cdot)p(f)$	-27.51	63.02	2.77	0.08	0.02	126.81	0.68	0.79
		$\psi(PD_{400})p(f)$	-26.23	64.09	3.83	0.04	0.17	130.85	0.61	0.87
		$\psi(PD_{200})p(f)$	-27.09	65.82	5.56	0.02	0.07	123.55	0.62	0.81

	$\psi(PD_{500})p(f)$	-27.09	65.82	5.56	0.02	0.07	120.15	0.65	0.87
	$\psi(PD_{300})p(f)$	-27.31	66.26	6	0.01	0.04	125.5	0.59	0.8
MH	$\psi(MH_{400})p(\cdot)$	-23.78	55.57	0	0.46	0.4	117.34	0.51	0.78
	$\psi(MH_{300})p(\cdot)$	-24.86	57.72	2.15	0.16	0.31	116.27	0.51	0.79
	$\psi(MH_{500})p(\cdot)$	-25.38	58.75	3.18	0.09	0.26	138.24	0.3	0.97
	$\psi(MH_{400})p(f)$	-23.77	59.17	3.6	0.08	0.4	111.91	0.67	0.66
	$\psi(MH_{200})p(\cdot)$	-26.01	60.02	4.45	0.05	0.19	115.93	0.73	0.69
	$\psi(MH_{500})p(f)$	-24.24	60.11	4.54	0.05	0.36	132.1	0.39	0.96
	$\Psi(\cdot)p(\cdot)$	-27.67	60.26	4.69	0.04	0	116.61	0.71	0.63
	$\psi(MH_{300})p(f)$	-24.32	60.27	4.7	0.04	0.35	117.05	0.72	0.72
	$\psi(\cdot)p(f)$	-27.51	63.02	7.46	0.01	0.02	126.81	0.64	0.82
	$\psi(MH_{200})p(f)$	-25.77	63.17	7.6	0.01	0.22	125.37	0.55	0.82
MR	$\Psi(\cdot)p(\cdot)$	-27.67	60.26	0	0.38	0	116.61	0.69	0.69
	$\psi(MR_{500})p(\cdot)$	-27.01	62.02	1.76	0.16	0.08	116.82	0.74	0.67
	$\psi(MR_{200})p(\cdot)$	-27.3	62.61	2.35	0.12	0.05	116.51	0.63	0.71
	$\psi(\cdot)p(f)$	-27.51	63.02	2.77	0.1	0.02	126.81	0.56	0.86
	$\psi(MR_{400})p(\cdot)$	-27.58	63.17	2.91	0.09	0.01	116.55	0.73	0.64
	$\psi(MR_{300})p(\cdot)$	-27.67	63.33	3.08	0.08	0	116.62	0.74	0.62
	$\psi(MR_{500})p(f)$	-26.94	65.51	5.25	0.03	0.09	120.07	0.66	0.77
	$\psi(MR_{200})p(f)$	-27.18	66	5.74	0.02	0.06	123.51	0.61	0.79

		$\psi(MR_{400})p(f)$	-27.47	66.58	6.32	0.02	0.03	123.71	0.74	0.7
		$\psi(MR_{300})p(f)$	-27.51	66.66	6.4	0.02	0.02	127.15	0.63	0.8
<i>L. pardalis</i>	FC	$\psi(FC_{300})p(s)$	-28.67	68.98	0	0.42	0.61	102.62	0.51	0.73
		$\psi(FC_{400})p(s)$	-28.85	69.33	0.36	0.35	0.6	103.65	0.65	0.6
		$\psi(FC_{500})p(s)$	-29.56	70.76	1.78	0.17	0.56	102.45	0.61	0.65
		$\psi(FC_{200})p(s)$	-30.59	72.82	3.84	0.06	0.5	103.71	0.53	0.73
		$\Psi(.)p(.)$	-36.02	80.03	11.05	0	0	101.15	0.63	0.71
	PD	$\psi(PD_{400})p(s)$	-33.48	78.59	0	0.44	0.27	101.18	0.56	0.71
		$\psi(PD_{500})p(s)$	-34.15	79.94	1.35	0.22	0.21	101.22	0.59	0.64
		$\Psi(.)p(.)$	-36.02	80.03	1.44	0.21	0	101.15	0.63	0.63
		$\psi(PD_{300})p(s)$	-35.33	82.3	3.71	0.07	0.08	101.04	0.57	0.57
		$\psi(PD_{200})p(s)$	-35.41	82.45	3.86	0.06	0.07	101.17	0.67	0.65
	MH	$\psi(MH_{300})p(s)$	-28.75	69.14	0	0.7	0.6	101.79	0.52	0.63
		$\psi(MH_{400})p(s)$	-29.72	71.08	1.93	0.27	0.55	106.2	0.48	0.63
		$\psi(MH_{200})p(s)$	-32.55	76.73	7.59	0.02	0.36	103.24	0.48	0.64
		$\psi(MH_{500})p(s)$	-33.22	78.07	8.92	0.01	0.3	103.97	0.46	0.66
		$\Psi(.)p(.)$	-36.02	80.03	10.89	0	0	101.15	0.56	0.6
	MR	$\Psi(.)p(.)$	-36.02	80.03	0	0.52	0	101.15	0.54	0.8
		$\psi(MR_{500})p(s)$	-35.46	82.55	2.52	0.15	0.07	101.25	0.66	0.59
		$\psi(MR_{200})p(s)$	-35.6	82.84	2.8	0.13	0.05	101.1	0.49	0.6

		$\psi(MR_{400})p(s)$	-35.65	82.93	2.9	0.12	0.05	101.07	0.54	0.57
		$\psi(MR_{300})p(s)$	-35.98	83.6	3.56	0.09	0	101.14	0.58	0.59
<i>P. onca</i>	FC	$\psi(FC_{400})p(s)$	-15.89	43.41	0	0.37	0.42	11.35	0.63	0.84
		$\psi(FC_{200})p(s)$	-16.13	43.89	0.48	0.29	0.4	10.98	0.65	0.87
		$\psi(FC_{300})p(s)$	-16.27	44.17	0.76	0.25	0.39	10.59	0.6	0.84
		$\Psi(.)p(.)$	-19.8	47.59	4.18	0.05	0	10.91	0.55	0.89
		$\psi(FC_{500})p(s)$	-18.12	47.88	4.46	0.04	0.21	10.69	0.74	0.82
	PD	$\Psi(.)p(.)$	-19.8	47.59	0	0.43	0	10.91	0.58	0.95
		$\psi(PD_{400})p(s)$	-18.48	48.6	1.01	0.26	0.17	10.95	0.67	0.9
		$\psi(PD_{500})p(s)$	-19.29	50.21	2.62	0.12	0.07	10.89	0.66	0.82
		$\psi(PD_{200})p(s)$	-19.34	50.32	2.72	0.11	0.06	10.9	0.64	0.88
		$\psi(PD_{300})p(s)$	-19.57	50.78	3.19	0.09	0.03	10.87	0.62	0.87
	MH	$\psi(MH_{400})p(s)$	-16	43.64	0	0.65	0.41	11.31	0.64	0.81
		$\psi(MH_{300})p(s)$	-17.48	46.6	2.95	0.15	0.27	10.69	0.64	0.79
		$\Psi(.)p(.)$	-19.8	47.59	3.95	0.09	0	10.91	0.71	0.84
		$\psi(MH_{200})p(s)$	-18.16	47.96	4.32	0.07	0.2	10.84	0.61	0.87
		$\psi(MH_{500})p(s)$	-18.85	49.33	5.69	0.04	0.12	10.76	0.71	0.86
	MR	$\Psi(.)p(.)$	-19.8	47.59	0	0.53	0	10.91	0.68	0.87
		$\psi(MR_{500})p(s)$	-19.13	49.9	2.31	0.17	0.09	10.92	0.64	0.89
		$\psi(MR_{200})p(s)$	-19.44	50.51	2.92	0.12	0.05	10.89	0.65	0.91

$\psi(MR_{400})p(s)$	-19.72	51.07	3.48	0.09	0.01	10.92	0.63	0.88
$\psi(MR_{300})p(s)$	-19.8	51.23	3.64	0.09	0	10.91	0.58	0.91

The covariates are the percentage of forest cover (FC), patch density (PD), the percentage of habitat from matrix (MH) and matrix resistance (MR). The subscript numbers indicate the buffer size (in hectares) of each covariate. The null model does not include a covariate (.). The models' log-likelihood (LL), Akaike Information Criterion (AIC), Akaike differences (Δ_i), Akaike weights (w_i), the Nagelkerke's R^2 (R^2), the models' Pearson chi-square (χ^2), significance (P) and overdispersion parameter (\hat{c}) of each model are shown. The selected models (and therefore, the scale of effect of each metric) for further analysis are in bold.

CAPÍTULO V. DISCUSIÓN Y CONCLUSIONES GENERALES

Conservar la diversidad de plántulas y mantener los factores que la promueven son cruciales para la regeneración de los bosques tropicales húmedos. Es por ello que en esta tesis identificamos y evaluamos qué factores de las estructuras del fragmento y el paisaje mantienen la diversidad de la comunidad de plántulas y la presencia de mamíferos terrestres en un bosque tropical húmedo hiper fragmentado en la Selva Lacandona.

A pesar de que esta región aún tiene un alto porcentaje de bosque remanente (Arasa-Gisbert et al., 2021; San-José et al., 2019), la deforestación se concentra en los bosques de terraza aluvial adyacentes al río Lacantún, donde la mayor parte de la cobertura arbórea ha sido eliminada para la expansión de pastizales ganaderos y cultivos temporales y agroindustriales (Berget et al., 2021; Carabias et al., 2015). El bosque primario de estas zonas está reducido a pocos fragmentos de tamaños menores a las 100 ha y la intensificación de las actividades agrícolas y ganaderas limitan su regeneración a partir de bosques secundarios (M. Lohbeck et al., 2022). Considerando que los bosques tropicales húmedos hiper fragmentados (aquellos con una cobertura de bosque primario < 30%, dominados por matrices agropecuarias fragmentos menores a las 50 ha; Oliveira et al., 2008; Tabarelli et al., 2012) serán dominantes en el mediano y largo plazo (Taubert et al., 2018), los paisajes analizados en nuestro sitio son representativos de las condiciones actuales y futuras de los PMH tropicales, así como idóneos para determinar los factores que actúan a múltiples escalas sobre la regeneración del bosque primario dentro de fragmentos.

Tomando en cuenta la baja cobertura de bosque primario remanente en el sitio de estudio (< 20%) y de acuerdo con la hipótesis del umbral de fragmentación (Andren, 1994), hipotetizamos que la diversidad de plántulas y la presencia de mamíferos terrestres se encuentran fuertemente influenciadas por factores a escala de fragmento (i.e., tamaño, estructura/diversidad del bosque y nivel de disturbio), la matriz circundante (i.e., porcentaje de bosques secundarios y sus efectos sobre la complementación/conectividad del paisaje en

mamíferos) y la configuración del paisaje (i.e., densidad, agregación y asilamiento de los fragmentos, así como su contraste de borde con las matrices adyacentes).

El área basal tiene efectos en cascada sobre la composición y la diversidad β de plántulas

A escala local, encontramos que la composición y la diversidad β de especies comunes y dominantes de plántulas fueron favorecidas por la reducción en la abundancia de plántulas demandantes de luz en fragmentos con una alta área basal. También observamos que la diversidad α de todas las especies de plántulas (riqueza) está determinada por la abundancia de plántulas tolerantes a la sombra, las cuales fueron afectadas negativamente en fragmentos con una alta abundancia de plántulas demandantes de luz y dentro de paisajes con una baja agregación de fragmentos.

El área basal influencia la abundancia y composición de plántulas a través de la producción de frutos (Chapman et al., 1992), la disponibilidad de luz, agua y nutrientes en el suelo (J S Denslow, 1987; Madelon Lohbeck et al., 2015; Martinez-Ramos & Soto-Castro, 1993; Turner, 2004) y la mortalidad conespecífica de plántulas (Wright, 2002). En nuestro estudio, el área basal solamente afectó la abundancia de plántulas demandantes de luz, sugiriendo que fragmentos perturbados (menor abundancia de árboles grandes) promueven el establecimiento y sobrevivencia de estas especies al incrementar la disponibilidad de luz (J S Denslow, 1987; Julie S. Denslow, 1996). Estas condiciones promueven el establecimiento de un subconjunto de especies que son ecológicamente redundantes, que carecen de relación filogenética y que dan lugar a ensamblajes de especies empobrecidos y dominados por pocas especies con altas abundancias (Oliveira et al., 2008; Tabarelli et al., 2012). Estos ensamblajes indican un proceso de homogenización biótica, caracterizado por una baja composición de la comunidad (i.e.,

menor disimilitud entre comunidades), como una reducción en la diversidad β (i.e., menor recambio de especies) a escala local y de paisaje (Olden & Rooney, 2006; Solar et al., 2015; Tabarelli et al., 2012).

Lo anterior es congruente con nuestras observaciones, donde una mayor abundancia de plántulas demandantes de luz redujo tanto la composición de plántulas, como la diversidad β de especies típicas y dominantes, variables sensibles a las fluctuaciones de abundancias (Jost, 2007; Roden et al., 2018). Además, la ausencia de efectos directos por parte de la estructura del paisaje, respaldan las hipótesis que establecen que las especies generalistas y las poblaciones con una alta densidad de individuos son principalmente afectadas por factores a escalas pequeñas (Miguet et al., 2016). Esto apoya los efectos en cascada de la estructura del paisaje sobre la abundancia de una palma competidora de luz, la cual representó el factor local tuvo el mayor impacto sobre respuestas biológicas afectadas a escalas pequeñas, tales como la densidad de especies e individuos (Hernández-Ruedas et al., 2018).

Por el contrario, la abundancia de plántulas tolerantes a la sombra fue mayormente explicada por la agregación de fragmentos en el paisaje, lo cual respalda las hipótesis de Miguet et al. (2016) arriba mencionadas. Las plántulas tolerantes a la sombra tienen una especialización de hábitat mayor que las especies demandantes de luz, así como una menor densidad de población (Julie S. Denslow, 1996; Turner, 2004). Además, sus abundancias son reguladas por la mortalidad conoespecífica de plántulas y el forrajeo de frutos, semillas y plántulas por mamíferos terrestres (Camargo-Sanabria et al., 2014; Comita et al., 2007, 2010; Dirzo & Miranda, 1990). Lo anterior sugiere que la estructura del paisaje afecta la abundancia de las plántulas tolerantes a la sombra al facilitar o limitar la llegada de dispersoras de semillas y de depredadores de semillas/plántulas.

Este supuesto está respaldado por estudios previos que encontraron una contribución significativa del porcentaje de bosque primario en el paisaje sobre la riqueza y abundancia de la

lluvia de semillas y los brinzales de especies de árboles especialistas de dispersión biótica (Arasa-Gisbert et al., 2021; San-José et al., 2019). Estos estudios concluyeron que la pérdida de especies tolerantes a la sombra es resultado de la falta de dispersión de semillas en paisajes cuya baja cobertura de bosque limita el movimiento de fauna, la cual dispersa el 90% de las especies de árboles de bosques tropicales (Jordano, 2000). No obstante, el tamaño del ámbito hogareño de la fauna puede variar en función de la agregación de fragmentos en paisajes con bajo porcentaje de hábitat (Gardiner et al., 2019; Hodgson et al., 2012; Radford et al., 2005), aminorando el efecto de la pérdida de hábitat sobre el movimiento de fauna.

En este trabajo, encontramos que la abundancia de plántulas tolerantes a la sombra aumentó en paisajes con una alta agregación de fragmentos, lo que sugiere que este patrón configuracional promueve la llegada de dispersores de semillas y de depredadores de semillas y plántulas. Las funciones de dispersión y depredación, aunadas a la mortalidad conespecífica de plántulas que promueven su diversidad (Comita et al., 2010; Harms et al., 2000; Wright, 2002) explican el efecto positivo que la abundancia de plántulas tolerantes a la sombra tiene sobre la diversidad α de todas las especies de plántulas. Resulta interesante que la abundancia de plántulas tolerantes a la sombra fuera afectada por la abundancia de plántulas demandantes de luz, lo cual sugiere que las segundas ejercen una presión competitiva por recursos. La exclusión competitiva puede jugar un papel importante en la reducción de la diversidad α dentro de bosques con una alta disponibilidad de luz y una menor presión por herbívoros (Wright, 2002). Esto respalda el efecto indirecto que observamos de la abundancia de plántulas demandantes de luz sobre la riqueza de especies.

La diversidad α y β responden a distintos componentes de la estructura del paisaje

A su vez, la diversidad α y β de las plántulas fueron afectadas por la composición y configuración del paisaje, respectivamente. Por un lado, la α disminuyó en paisajes dominados por bosques secundarios. Por otro lado, la diversidad β de especies comunes y dominantes fue promovida en paisajes con una alta densidad de fragmentos, mientras que la diversidad β que considera a todas las especies (incluyendo aquellas especies raras con bajas abundancias) disminuyó en paisajes con un alto contraste de borde.

En la sección anterior quedó establecido que el incremento en la riqueza de plántulas en paisajes con una mayor agregación de fragmentos está asociado con una mayor llegada de dispersores de semillas y de mamíferos terrestres que promueven su diversidad. Esto resalta el papel que tiene la configuración del paisaje para promover la dispersión de semillas y la persistencia de funciones ecológicas en paisajes muy fragmentados (Fahrig, 2017; Haan et al., 2020).

Por el contrario, el efecto negativo del porcentaje de bosques secundarios sobre los tres órdenes de diversidad α sugieren la llegada de un grupo de especies reducido en la lluvia de semillas. La composición de la vegetación secundaria puede ejercer una fuerte influencia en la diversidad de especies que se establecen en los bordes y claros de los fragmentos (Laurance et al., 2006, 2007; Nascimento et al., 2006). A su vez, esta composición es resultado de la historia de uso de suelo y la intensidad, duración y frecuencia de la perturbación previa al crecimiento de la vegetación secundaria (Laurance et al., 2007; Zermeño-Hernández et al., 2015).

En la región de estudio, las comunidades de plántulas que se establecen en coberturas con suelos fuertemente compactados, afectados por incendios severos y/o frecuentes presentan una menor densidad de individuos y especies, así como bajos valores de diversidad (Zermeño-Hernández et al., 2015). La cobertura asociada a este tipo de manejo son los pastizales ganaderos, los cuales son dominantes en los bosques de planicies aluviales (Berget et al., 2021; Carabias et al., 2015; Zermeño-Hernández et al., 2015). Por lo tanto, los bosques

secundarios que se desarrollan en estas condiciones presentan una estructura y diversidad menores que en bosques secundarios establecidos en usos de suelo con una menor severidad de disturbio como los sitios que fueron utilizados en este trabajo, reduciendo su potencial de regeneración (Chazdon, 2014).

Estos bosques a su vez, tienen una fuerte influencia en la trayectoria de regeneración de la vegetación dentro de fragmentos, las cuales divergen en función de la composición de especies de los bosques adyacentes (Laurance et al., 2007). A esto hay que agregar la influencia de las coberturas dominantes en el paisaje circundante, debido a que la diversidad arbórea está fuertemente afectada por la dominancia de coberturas de uso de suelo con un alto grado de perturbación (Tenius Ribeiro et al., 2019; Zermeño-Hernández et al., 2016). Como resultado, se espera que la dominancia de pastizales ganaderos y sus bosques secundarios derivados tengan un efecto negativo sobre la diversidad α de las plántulas.

El incremento de la diversidad β de las plántulas en paisajes con bajos contrastes de borde y una mayor densidad de fragmentos podría sugerir que la heterogeneidad espacial de las especies de plántulas es promovida ya sea por sea la llegada de especies desde fragmentos con distintos ensambles de árboles, la reducción de los efectos de borde o la llegada de dispersores de semillas y mamíferos terrestres que contribuyen a mantener su diversidad.

La alta heterogeneidad de hábitats y condiciones ambientales entre fragmentos promueven el establecimiento de comunidades de árboles con composiciones muy diferentes entre sí, lo que aumenta la diversidad β a escala regional (Arroyo-Rodríguez et al., 2013; Bennett & Gilbert, 2016; Hernández-Ruedas et al., 2014). Además, se espera que la dispersión de semillas se incremente en paisajes con un mayor número de fragmentos, los cuales tienen un efecto positivo sobre la lluvia de semillas (San-José et al., 2019). Por lo tanto, la densidad de parches puede incrementar la diversidad β de las plántulas al facilitar la llegada de semillas de

fragmentos con ensamblajes de árboles heterogéneos. Los resultados de los capítulos II y III nos muestran que la diversidad β de especies típicas y dominantes es afectada por el área basal del fragmento a través de la fluctuación de abundancias de plántulas demandantes de luz y por la configuración del paisaje que facilita la llegada de especies nuevas.

Finalmente, el efecto negativo del contraste de borde sobre la diversidad β de todas las especies (incluyendo aquellas con bajas abundancias), puede estar asociado a una mayor exposición a los efectos de borde o a una barrera que impide la llegada de mamíferos terrestres. La homogenización del paisaje está fuertemente vinculada con la magnitud de los efectos de borde, los cuales promueven la dominancia de pocas especies tolerantes a la perturbación (Oliveira et al., 2008; Tabarelli et al., 2010). Los efectos de borde incrementan la mortalidad de plántulas tolerantes a la sombra y desplazan a las especies raras, al mismo tiempo que promueven el recambio de especies demandantes de luz (Benítez-Malvido et al., 2018; Krishnadas et al., 2018). Por lo tanto, la diversidad β de plántulas raras puede mantenerse en fragmentos donde los efectos de borde son aminorados a través de un bajo contraste de borde.

Sin embargo, se esperaría que estos efectos ocurrieran a una escala espacial menor a la observada en nuestro trabajo (200 ha). La escala del efecto registrada para el contraste de borde fue similar a la observada para la proporción de matrices abiertas sobre la riqueza y diversidad de especies de brinzales especialistas en paisajes hiper fragmentados (Arasa-Gisbert et al., 2021). En este estudio, se plantea que la apertura de matrices incrementa el aislamiento de los fragmentos al limitar el movimiento de dispersores de semillas, dando como resultado una menor abundancia y riqueza (Galán-Acedo et al., 2019; San-José et al., 2019).

Además la riqueza y similitud de la comunidad de mamíferos terrestres entre fragmentos está fuertemente vinculada con la dominancia de matrices estructuralmente similares a la vegetación del fragmento (Arce-Peña et al., 2022; Garmendia et al., 2013), lo que sugiere que

las matrices con cobertura arbórea facilitan el movimiento de fauna entre fragmentos (Arroyo-Rodríguez et al., 2020). La actividad de estos mamíferos regula la abundancia y distribución espacial de especies de plántulas dominantes, lo que promueve la diversidad β y la persistencia de especies raras (Bleher & Böhning-Gaese, 2001; Fragoso et al., 2003; Paine & Beck, 2007; Villar et al., 2019). Corroborar este argumento requerirá de analizar los efectos en cascada del porcentaje de matrices abiertas/permeables o del porcentaje bordes contrastantes en el paisaje sobre la diversidad β a través de la ocupación, abundancia o índice de defaunación del fragmento.

El papel de la densidad de fragmentos y la resistencia de la matriz sobre la conectividad del paisaje en mamíferos terrestres

Finalmente, la presencia de mamíferos terrestres -cuyos patrones de actividad y forrajeo promueven la diversidad de plántulas- fue favorecida tanto por la densidad de fragmentos, como por la conectividad proporcionada por la matriz circundante. En este caso, encontramos que la probabilidad de ocupación de los fragmentos para tres de las cuatro especies de mamíferos evaluadas aumentó en los paisajes que tuvieron una menor resistencia al movimiento a través de la matriz.

Cabe destacar que los la escala de los efectos de la matriz sobre la disponibilidad de hábitat fue menor respecto a los efectos que tuvo sobre la resistencia al movimiento de fauna. Esto respalda la predicción de Miguet et al. (2016) planteando que la escala del efecto en variables que afectan el éxito reproductivo/forrajeo es menor que aquellas variables que influyen la dispersión. Este resultado muestra que la matriz afecta la presencia de mamíferos a través de procesos que operan a distintas escalas de manera simultánea.

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