CONSERVATION ECOLOGY – ORIGINAL RESEARCH

Agroforestry orchards support greater butterfy diversity than monoculture plantations in the tropics

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Abstract

Large-scale deforestation in the tropics, triggered by logging and subsequent agricultural monoculture has a signifcant adverse impact on biodiversity due to habitat degradation. Here, we measured the diversity of butterfly species in three agricultural landscapes, agroforestry orchards, oil palm, and rubber tree plantations. Butterfy species were counted at 127 sampling points over the course of a year using the point count method. We found that agroforestry orchards supported a greater number of butterfy species (74 species) compared to rubber tree (61 species) and oil palm plantations (54 species) which were dominated by generalist (73%) followed by forest specialists (27%). We found no significant difference of butterfy species composition between agroforestry orchards and rubber tree plantation, with both habitats associated with more butterfy species compared to oil palm plantations. This indicates butterfies were able to persist better in certain agricultural landscapes. GLMMs suggested that tree height, undergrowth coverage and height, and elevation determined butterfy diversity. Butterfy species richness was also infuenced by season and landscape-level variables such as proximity to forest, mean NDVI, and habitat. Understanding the factors that contributed to butterfy species richness in an agroecosystem, stakeholders should consider management practices to improve biodiversity conservation such as ground vegetation management and retaining adjacent forest areas to enhance butterfy species richness. Furthermore, our fndings suggest that agroforestry system should be considered to enhance biodiversity in agricultural landscapes.

Keywords Agricultural landscapes · Biodiversity · Conservation · Tropical forest · Oil palm · Rubber tree

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Introduction

The large-scale conversion of tropical forests into agricultural areas raised concerns among conservationists due to its major detrimental impacts on biodiversity (Barnes et al. [2014](#page-10-0); Ahrends et al. [2015;](#page-10-1) Azhar et al. [2015a;](#page-10-2) Warren-Thomas et al. [2020](#page-12-0); Alroy [2017;](#page-10-3) Giam [2017](#page-11-0); Pashkevich et al. [2020](#page-11-1); Potapov et al. [2020\)](#page-11-2). Most land-use change and intensifcation lead to landscape simplifcation and fragmentation, which result in signifcant biodiversity losses due to habitat destruction (Ewers et al. [2009\)](#page-11-3), as well as disruption of ecosystem services such as pollination, pest control, and nutrient cycling (Landis [2017](#page-11-4); Power [2010](#page-12-1)). With the loss of ecosystem services, human well-being will be in jeopardy (Diaz et al. [2006](#page-10-4)). Nevertheless, agroforestry system that includes productive and protected areas can reduce the impact of tropical deforestation by promoting a mosaic of diferent land-use types (Stanturf et al. [2019](#page-12-2); Schwarz et al. [2021](#page-12-3)).

In Malaysia, about 33.1% of the total land area has been converted for agricultural purposes (MPIC [2015](#page-11-5)); agriculture sectors account for approximately 23% of total exports and 7.2% of Malaysia's GDP (Rozhan [2015](#page-12-4)). The land conversion motivated by agricultural intensifcation in Malaysia began during the British colonial era with the introduction of monoculture rubber tree (*Hevea brasilensis*) and oil palm (*Elaeis guineensis*) plantations from Brazil and South Africa, respectively (Athukorala and Loke [2009;](#page-10-5) Nath and Chaudhuri [2010](#page-11-6); Sayer et al. [2012](#page-12-5)). To date, approximately 5,865,290 ha of land have been planted with oil palm trees and 1,106,861 ha with rubber trees (MPIC [2015](#page-11-5)). Furthermore, a mix of native fruit trees is planted in agroforestry orchards to provide farmers with additional income (Salma et al. [2006;](#page-12-6) Abdullah [2011\)](#page-10-6), covering approximately 116,369 ha of land (DOSM [2019](#page-10-7)). Thus, changes in habitat structure (both at the local and landscape scale) caused by diferent agricultural practices in each respective farmlands may harbor diferent composition of associated wildlife (Pogue and Schnell [2001;](#page-11-7) Azhar et al. [2015b](#page-10-8)).

Butterfies are well-documented taxa for agricultural biodiversity research and are frequently paired with other taxa such as birds (Posa and Sodhi [2006;](#page-11-8) Koh [2008](#page-11-9); Azhar et al. [2015b;](#page-10-8) Salek et al. [2018](#page-12-7)), spiders (Salek et al. [2018](#page-12-7)), bats (Azhar et al. [2015b\)](#page-10-8), and ants (Lucey and Hill [2012](#page-11-10)). Besides offering a wide range of ecosystem services, butterfies are good indicators for environmental changes as their diversity is related to other taxa such as birds and spiders (Ekroos et al. [2013](#page-11-11); Salek et al. [2018\)](#page-12-7). Butterfies play major roles in the ecosystem as food resources to other animals, pollinators by transferring pollen to fowers, and defoliators on numerous plants (Stokes et al. [1991;](#page-12-8) Cleary [2004](#page-10-9); Karen-Chia [2014\)](#page-11-12).

Previous research has shown that the richness of butterfy species in oil palm agroecosystems is infuenced by landscape and local scale-level habitat attributes (Azhar et al. [2015b](#page-10-8); Asmah et al. [2017](#page-10-10)). At the landscape level, mixedcrop agriculture supports more diverse arthropod communities compared to monoculture (Azhar et al. [2015b](#page-10-8); Ghazali et al. [2016](#page-11-13); Ashraf et al. [2018\)](#page-10-11). The polyculture system's heterogeneity provides more food resources and habitat for various butterfy species (Collinge et al. [2003](#page-10-12)). Besides, the proximity of agricultural areas to forests increases the diversity of butterfy species (Lucey and Hill [2012;](#page-11-10) Koh [2008](#page-11-9)). On a local scale, butterfy species richness has been linked to ground vegetation cover and height (Azhar et al. [2015b](#page-10-8); Asmah et al. [2017](#page-10-10)). This ground vegetation provides adults and immature butterfies with food and a breeding ground (Koh [2008](#page-11-9)).

Until now, butterfy research in Malaysian agroecosystems has been limited to oil palm plantations (Koh [2008](#page-11-9); Lucey and Hill [2012](#page-11-10); Azhar et al. [2015b;](#page-10-8) Asmah et al. [2017](#page-10-10)). No research has included butterfy species survey in rubber tree plantations or agroforestry orchards. As a result, farmland biodiversity in agroecosystems is poorly understood, and the role they play in agricultural landscapes in combination with either monoculture or polyculture farming systems, as well as encapsulating protected areas or forest reserves, is not well known.

In this study, we quantified butterfly presence and species richness in three different agricultural landscapes (agroforestry orchards, rubber tree plantations, and oil palm plantations) using point count methods. We also measured local- and landscape-level parameters and butterfy species distribution to answer three major questions;

- 1. Does the diversity of butterfy species difer between agroforestry orchards, rubber tree plantations, and oil palm plantations?
- 2. What local- and landscape-level factors are important in determining butterfy species richness in diverse agricultural landscapes?
- 3. Are there any changes in butterfy community structure across agricultural landscapes?

This study is critical for promoting associated biodiversity in agricultural ecosystems, with a focus on butterfies as they play important ecological roles in the ecosystem (Brown and Freitas [2003\)](#page-10-13) and are potentially afected by agricultural intensifcations (Koh [2008](#page-11-9); Lucey and Hill [2012](#page-11-10)). The study's fndings will inform farmers and conservationists about the necessity of enhancing habitat quality at local- and landscape-level to increase biodiversity within agricultural landscapes.

Materials and methods

Study sites

We conducted our research between Rembau District (2° 35′ 25.8900'' N, 102° 5′ 34.7532'' E), Negeri Sembilan, on the west coast of Peninsular Malaysia (Fig. [1](#page-2-0)). The study area (approximately 41,512 ha) was converted from lowland dipterocarp forests to agricultural areas at least 60 years ago (Denan et al. [2019\)](#page-10-14). We selected this area because it has a diverse land use that includes oil palm plantations, rubber tree plantations, and agroforestry orchards. Both oil palm plantations and rubber tree plantations are monoculture system planted uniformly with a single species. For this study, oil palm and rubber tree plantations with an average age of 10 years old were selected for the sampling. Both monoculture plantations have rigorous management practices which involved systematic applications of chemical pesticides to control weeds, pests, and diseases (Azhar et al. [2015a](#page-10-2)). On the other side, agroforestry orchards were mainly made up

Fig. 1 Map of study location in Pedas, Negeri Sembilan, Peninsular Malaysia. The left map indicates the location of the study site at Negeri Sembilan and the right map indicates the location of sampling

point: yellow circle=agroforestry orchards, red square=oil palm plantation, and blue triangle=rubber tree plantations

of native fruit trees, with an average of fve species of fruit trees planted at each orchard consisted mainly of durian (*Durio zibethinus* L.), rambutan (*Nephelium lappaceum* L.), langsat (*Lansium domesticum* Corr.), mangosteen (*Garcinia mangostana* L.), jering (*Archidendron pauciforum* (Benth.) I. C. Nielsen), petai/ stink bean (*Parkia speciosa* Hassk.), and cempedak (*Artocarpus integer* Merr.). Typically, agroforestry orchards practice less intensive management approaches including mechanical removal of understory vegetation one or twice a year before fruiting seasons.

Sampling design

Systematic sampling with a random starting point was used in this experiment to ensure better spatial coverage and lower variance (Morrison et al. [2008;](#page-11-14) Thomas et al. [2010](#page-12-9)). The butterfy and vegetation structure data were collected from 49 sampling points in oil palm plantations, 41 in rubber tree plantations, and 37 in agroforestry orchards. The distance among sampling points was at least 300 m apart to reduce the chances of counting the same individual at the sites. Each sampling point was visited once.

Butterfies sampling

Butterfies were observed using a point count method (Reynold et al. [1980\)](#page-12-10) from February 2019 to February 2020. At each point, we recorded butterfies for 10 min in a 30 m plot radius from 9.00 a.m. to 3.00 p.m. using binoculars and sweep nets. Binocular (PENTAX Papilio 8.5×21) was used to detect the butterfy wing patterns. We identifed the species of each butterfly using guidebooks by Kirton ([2014\)](#page-11-15) and Khoon [\(2015\)](#page-11-16). Butterfy species that could not be identifed were captured using a net and taken to the laboratory for species-level identifcation using a book by Corbet and Pandlebury [\(1992,](#page-10-15) [2020\)](#page-10-16). Equal number of sites from each type of treatment were sampled each month. We counted the species richness of butterfies and categorized them

according to their taxonomy (family, genus, and species) and distribution range (1- coastal mangrove, 2- secondary plant growth below 760 m, 3- primary forest below 760 m, and 4- all area above 760 m (Corbet and Pandlebury [2020](#page-10-16))). We grouped butterfy species based on their distribution range into generalist or specialist category. Butterfies encountered in area 1 and 2 as well as area 2, 3 and 4 were classifed as generalists, while those specifcally found in area 3 and 4 were classifed as specialists.

Measurement of local‑level variables

In-situ habitat quality variables such as vegetation structure were recorded for each sampling within the area of 1000 m². Variables measured at the local-level were: (i) tree height; (ii) number of tree species; (iii) diameter at breast height (dbh); (iv) canopy cover; (v) undergrowth coverage; (vi) undergrowth height; and (vii) elevation. Variables (iv), (v), and (vi) were measured in 1 m radius in four directions (east, north, west and south). We measured tree height and dbh using a laser range fnder (Nikon LASER 350) and dbh meter, respectively. Tree species with more than 10 cm dbh were identifed and recorded. We used HabitApp version 1.1 mobile application to measure canopy cover and Canopeo version 1.1.7 for undergrowth coverage, respectively. The undergrowth height was measured using measuring tape. We used GARMIN Global Positioning System (GPS) to provide the georeferenced location and determine the elevation at each sampling point.

Measurement of landscape‑level variables

We measured the normalized diference vegetation index (NDVI) of the crops since butterfly species richness is related to the vegetation structure and productivity (Bailey et al. [2004\)](#page-10-17). The NDVI was determined using ServEO version 1.3 mobile application. We also measured the distance between the sampling point and the nearest forest because butterfy species were afected by forest proximity especially for forest specialist (Lucey and Hill [2012\)](#page-11-10). The nearest forests to our sites were Angsi Forest Reserve (12,435 ha), Sungai Menyala Forest Reserve (1,280 ha), and Gunung Tampin Forest Reserve (5,541 ha). The distance to the nearest forest was measured using Google Earth Pro measuring tools.

Measurement of seasonal variation

We measured the effect of seasonality (wet and dry) on butterfy species richness. We assigned sampling months into the wet and dry seasons based on data provided by MET Malaysia; wet seasons (southwest monsoon and inter-monsoon) between May and November, and dry seasons between December and April (MET [2019](#page-11-17)).

Statistical analysis

To compare butterfy species richness in agricultural landscapes, we used iNEXT online which a non-asymptotic approach based on interpolation and extrapolation (Colwell et al. [2012;](#page-10-18) Chao et al. [2016](#page-10-19)). Using the diversity order $q=0$, we created sample-size-based rarefaction and extrapolation sampling curves (species richness). The default settings were used, with the number of bootstraps set to 50 and the level of confdence interval set to 95%.

The similarity percentage (SIMPER) analysis was used to determine the contribution of each species of butterfy assemblage in three diferent agricultural landscapes (i.e., agroforestry orchard, oil palm plantation and rubber tree plantation). The result showed the most persistence species in the pattern of similarity percentage. We used Bray–Curtis distance method to calculate the similarity metric between each landscape. The cutoff was set at 90% of the species accumulations.

We compared butterfy species composition between oil palm plantations, rubber tree plantations and fruit orchards using the analysis of similarity (ANOSIM). A resemblance matrix between samples was calculated using Bray–Curtis similarity. Percentages of similarity were used to determine the contribution of each butterfy species assemblage in three separate land uses. All multivariate analyses were performed in PRIMER Version 6 (PRIMER-E Ltd, Plymouth) (Clarke and Gorley [2006\)](#page-10-20).

We used generalized linear mixed models (GLMMs) (Schall [1991](#page-12-11)) to examine the relationships between the richness of butterfy species and habitat quality characteristics (local and landscape-level variables). Spearman's rank correlation coefficient tests were used to assess the multicollinearity of the explanatory variables. To prevent bias in model estimation, strongly correlated variables $(|r| > 0.7)$ were checked (Dormann et al. [2013\)](#page-10-21). However, none of the variables was strongly correlated. In the modelling process, Poisson distribution and log-link function were used with the number of butterfy species at each sampling point as the response variable and sampling month as a random model. Models with the lowest Mallow's Cp were identifed as the most parsimonious models. The modeling was conducted using Genstat (VSNI Hemel, UK).

Results

General patterns of butterfy biodiversity

We recorded 1567 butterfly individuals comprising five families (Hespiriidae, Lycaenidae, Nymphalidae, Papilionidae, and Pieridae), 68 genera, and 104 species (Supplementary Information). Out of 104 butterfy species, 28 (27%) were

specialists, while the remaining 76 (73%) were generalists. We also recorded two protected species in Malaysia under the Wildlife Conservation Act 2010 (Act 716), namely the banded peacock (*Papilio palinurus*) and common birdwing (*Troides helena).*

In agroforestry orchards, we recorded 472 butterfly individuals from 74 species (mean observation per point = 7.838) with 58 (78%) generalists and 16 (22%) forest specialists. For oil palm plantations, 615 butterfies from 54 species (7.102) were recorded with 44 (81%) generalists and 10 (19%) forest specialists while, for the rubber tree plantations we recorded 480 butterfies comprising 61 species (6.610) with 44 (72%) generalists and 17 (28%) forest specialists (Fig. [2\)](#page-4-0). Of the total butterfly species observed, 31 were found across all agricultural landscapes, 25 exclusively in agroforestry orchards, 17 in rubber tree plantations, and only nine in oil palm plantations (Fig. [3\)](#page-4-1). The rarefaction curves showed no overlap between three agricultural landscapes, suggesting that the observed diferences in species diversity and number of individuals were not caused by varying sampling coverages (Fig. [4\)](#page-4-2). Based on rarefaction and extrapolation results, our sampling effort managed to achieve 95% coverage of the butterfy communities across three agricultural landscapes (Table [1\)](#page-5-0).

SIMPER analysis (Supplementary Information) indicated 13 species of butterflies contributed to 90% of species composition for agroforestry orchards (average similarity=22.46%) leading by *Yptima baldus* (26.71%), *Appias libythea* (10.09%), *Eurema hecabae* (9.10%), *Junonia almana* (7.94%) and *Eurema sari* (6.34%). For the oil palm plantations (average similarity $= 26.29\%$), 11 species of butterfies contributed to 90% of species accumulation including *Elymnias hypermnestra* (16.23%), *Leptosia nina* (14.57%), *Appias libythea* (13.48%), *Ypthima baldus* (13.33%) and *Eurema hecabe* (10.60%). Eleven species of butterfies contributed to 90% of species accumulation in rubber tree plantations (average similarity = 21.34%) with *Ypthima baldus* (31.42%), *Leptosia nina* (10.78%), *Appias*

Fig. 2 The percentages of butterfy species richness of generalist and specialist observed at three agricultural landscapes

libythea (8.72%), *Eurema hecabe* (8.54%) and *Eurema sari* (7.13%) .

Overall, there was a significant difference of butterfy species composition with a high overlapping pattern (ANOSIM: number of permutations = 999; Global $R = 0.064$; $p = 0.001$). We found no significant difference in terms of butterfy species composition in agroforestry orchards and rubber tree plantations (ANOSIM $R = 0.003$; $p=0.387$). Butterfly species composition in oil palm plantations was signifcantly diferent compared to agroforestry orchards (ANOSIM $R = 0.075$, $p = 0.001$) and rubber tree plantations (ANOSIM $R = 0.099$, $p = 0.003$).

Fig. 3 Butterfy species richness and species overlap observed at three agricultural landscapes. The number in parenthesis represents the total number of species, and the square box represents species overlap in each habitat

Fig. 4 Rarefaction curves of butterfy species at three diferent agricultural landscapes

Table 1 Estimation of species diversity and sample coverage at specifed sampling eforts developed through samplesize-based rarefaction and extrapolation

m sample size, *qD* estimated diversity, *qD.LCL* bootstrap lower confdence limit, *qD.UCL* bootstrap upper confdence limit, *SC* estimated sample coverage

Determinants of butterfy species richness

We found that eight predictor variables of tree height, undergrowth coverage, undergrowth height, elevation, season, NDVI, habitat, and distance to forest infuenced butterfy species richness in three focal tropical agricultural landscapes. For the local level, a model with fve predictor variables (tree height, undergrowth coverage, undergrowth height, elevation, and season) was the most parsimonious combination with the lowest Mallow's Cp of 5.55 (Table [2;](#page-5-1) Fig. [5\)](#page-6-0). The model explained 30.23% of the variation in butterfy species richness. Our data revealed that butterfy species richness increased with increasing undergrowth coverage (slope=0.007681) and undergrowth height (slope=0.4057) (Table [3](#page-7-0)). On the other hand, we found out that butterfy species richness decreased with increasing tree height (slope $= -0.03052$) and elevation $(slope = -0.002891)$ (Table [3](#page-7-0)). Butterfly species richness was lower during the wet season compared to the dry season (slope = -0.2038 -0.2038 -0.2038) (Table 3).

At the landscape level, a model with four predictor variables (distance to forest, NDVI, habitat, and season) was the most parsimonious combination with the lowest Mallow's Cp of 22.52 (Table [2](#page-5-1); Fig. [6\)](#page-7-1). The model explained 20.41% of the variation in butterfy species richness of butterfy. We found that butterfy species richness increased as the distance to the forest decreased (slope $= -0.042$) (Table [3\)](#page-7-0). The species richness had a negative relationship with the NDVI value (slope $= -0.5194$). Oil palm plantations (slope $= -0.3109$) and rubber tree plantations $(slope = -0.1988)$ supported lower species richness in comparison to agroforestry orchards (Table [3](#page-7-0)). During the wet season (slope $= -0.2341$), butterfly species richness was lower than in the dry season (Table [3\)](#page-7-0).

The best model is shown in bold. *TH* tree height, *UC* undergrowth coverage, *S* season, *E* elevation, *UH* undergrowth height, *TS* tree species, *DBH* diameter at breast height, *DF* distance to forest, *H* habitat, *NDVI* mean NDVI

Fig. 5 Diferent responses to local-level parameters by butterfy species richness. Scatter plots have 95% confdence intervals (blue line) on the regression (red line)

Discussion

General patterns of butterfy diversity

There are about 1051 species of butterflies recorded in Peninsular Malaysia (Corbet and Pendelbury [2020\)](#page-10-16) and in this study we recorded a total of 104 species of butterfies, which representing approximately 10% of the total butterfy species reported. This fnding is consistent with previous research indicating that the richness of butterfy species is impoverished in tropical agricultural landscapes (Koh [2008](#page-11-9); Azhar et al. [2015b;](#page-10-8) Asmah et al. [2017\)](#page-10-10) compared to forest areas (Willott et al. [2000;](#page-12-12) Benedick et al. [2006](#page-10-22)). This may indicate that only a small proportion of butterfly species is able to persist well in agricultural landscapes, especially generalist species, which represent about 73%, followed by forest specialists at 27%.

We found that the richness of butterfy species was higher in agroforestry orchards in comparison to rubber tree and oil palm plantations. This fnding showed that agroforestry

Table 3 GLMMs models of butterfy species richness within diferent agricultural landscapes at local and landscape levels

Predictor variable	Slope	Standard error
Tree height	-0.0305	0.103
Undergrowth coverage	0.0077	0.002
Undergrowth height	0.4057	0.165
Elevation	-0.0029	0.003
Season		0.181
(1) Dry	0.000	
(2) Wet	-0.2038	
Distance to forest	-0.0420	
Mean NDVI	-0.5194	
Habitat		0.092
(1) Agroforestry orchard	0.000	
(2) Oil palm plantation	-0.1988	
(3) Rubber tree plantation	-0.3109	
Season		0.166
(1) Dry	0.000	
(2) Wet	-0.2341	

orchards with polyculture practices would offer better habitats for butterfies compared to rubber tree and oil palm plantations that are monocultures. In our case, agroforestry orchards were integrated with more than 5 species of fruit tree species consisting mainly of large-sized trees (e.g., durian, jering, petai), as well as other medium-sized trees (e.g., rambutan, mangosteen) forming a more complex habitat compared to monocultures. As such, polyculture systems improve habitat heterogeneity and vegetation structure (Horak et al. [2014;](#page-11-18) Elings et al. [2017\)](#page-11-19) and are expected to provide more resources for an insect (Jones and Gillet [2005](#page-11-20); Elba et al. [2014\)](#page-10-23). Agroforestry systems maintain biodiverse habitat and landscape heterogeneity (Bhagwat et al. [2008](#page-10-24); Yahya et al. [2022](#page-12-13)), in contrast to large-scale monoculture plantations, which have a lower potential for biodiversity conservation due to lower heterogeneity (Azhar et al. [2015a\)](#page-10-2). Native plants provide unlimited food resources for butterfies (Burghardt et al. [2009](#page-10-25)) as such, integrating multiple species of fruit trees would facilitate more butterfy species within an agriculture landscape (Asmah et al. [2017](#page-10-10)). We also observed the Painted Jezebel butterfy (*Delias hyparete*) feeding on rambutan fruit in the agroforestry orchards.

However, the species composition of butterfies between agroforestry orchard and rubber plantations are similar but diferent to oil palm plantations. The similarity of species richness is likely related to management practices, as most of the crops belonged to small-scale growers, except for a few rubber tree sites owned by the plantation company. On the other hand, oil palm plantations are mainly owned by plantation companies. Small-scale farmers typically practice less intensive management and are less dependent on modern machinery (Azhar et al. [2015a](#page-10-2)). This means that practices such as chemical weeding and agrochemical applications are rarely consistent compared to large-scale plantations. As a result, it will enhance the growth of understory vegetation and provide suitable habitats and food resources for the butterfies. Butterfies utilized (i.e., feed and breed) on naturally grown understory

Fig. 6 Diferent responses to landscape level parameters by butterfy species richness. Scatter plots have 95% confdence intervals (blue line) on the regression (red line)

vegetation across the agricultural landscape, regardless of land use. Broadleaved shrubs and tree (whiteweed—*Ageratum conyzoides*, Chinese violet—*Asystasia gangetica*, rhododendron—*Melastoma malabathricum*, Soapbush— *Clidemia hirta*) and grasses (wiregrass- *Eleusine indica*, congongrass—*Imperata cylindrica*) are the common weed species in the Malaysian agricultural landscape in (Barnes and Chan [1990;](#page-10-26) Maziatul-Suriza and Idris [2012;](#page-11-21) Fee et al. [2017](#page-11-22)). Broadleaved shrubs and tree provide a source of nectar for low-fying butterfies such as Psyche, Rings (*Ypthima*), Grass yellows and others. Grass is an important breeding host for Rings (*Ypthima*) since their caterpillars feed on grass (Poaceace) (Corbet and Pandelbury [2020\)](#page-10-16). Thus, this contributed to the higher abundance of the *Y. baldus* in agricultural landscapes. On the other hand, the abundance of *E. hypermnestra,* a common species in oil palm plantations (Kirton [2014](#page-11-15)) which is associated with various palm species (Corbet and Pandelbury [2020\)](#page-10-16). Oil palm plantations support fewer animal species than forest and other tree crops (Fitzherbert et al. [2008](#page-11-23)) due to intensive management practices such as the use of agrochemicals to increase yields (Azhar et al. [2015a\)](#page-10-2).

Oil palm and rubber tree plantations support lower butterfy species richness compared to agroforestry orchards. This fnding implies that agroforestry orchards would provide the butterfy with better habitat and food resources than oil palm and rubber tree plantations. Rubber tree and oil palm plantations are primarily renowned for a monoculture system with uniform stand age, which is unfavorable to farmland biodiversity (Ghazali et al. [2016](#page-11-13); Yahya et al. [2022](#page-12-13)). The structure and composition of vegetation infuence the diversity of butterfies (Kremen et al. [1993](#page-11-24); New et al. [1995\)](#page-11-25). The uniform canopy strata in monoculture plantations are unfavorable to butterfies due to their natural behavior of inhibiting diferent canopy strata of trees (Asmah et al. [2017](#page-10-10)). Butterfies of the Morphinae and Satyrinae subfamilies only fy in the understory on the lower vegetation layer (Schulze et al. [2001\)](#page-12-14) while Apaturinae, Charaxinae and Nymphalinae subfamilies can be found fying on the upper canopy layer (DeVries and Walla [2001](#page-10-27)). However, Apaturinae and Charaxinae were not detected by our sampling method (i.e., point count), as both are forest specialists and attracted to rotten fruit (Corbet and Pandlebury [2020\)](#page-10-16). Aside from that, ground cover vegetation in fruit orchards may provide a food source and breeding ground for butterfies. The majority of butterfies in Southeast Asia rely on monocotyledonous host plants for breeding and feeding (Schulze et al. [2001](#page-12-14); Corbet and Pandlebury [2020\)](#page-10-16). Aside from that, agroforestry orchards exhibit diferent phenology during the fowering season providing a constant supply of nectar for butterfies and other nectivorous insects (Elings et al. [2017\)](#page-11-19).

Butterfy species richness and infuencing attributes

Our data suggested that the tree height negatively afects the richness of the butterfy species. A similar result was also recorded in previous studies, which reported a negative association between tree height and butterfy diversity (e.g., Kumar et al. [2009;](#page-11-26) Wettstein and Schmid [1999;](#page-12-15) Thomas and Mallorie [1985](#page-12-16)). The biology and physiology of butterfies may have contributed to this negative relationship since butterfies are poikilothermic species. The solar radiation needed by butterfies for thermoregulation may be restricted by the shading of high trees (Kumar et al. [2009\)](#page-11-26).

Our fndings indicate that undergrowth coverage and undergrowth height had a positive impact on the richness of the butterfy species. This positive association is expected since undergrowth vegetation provide resources for the butterfies. Similar fndings were reported by Azhar et al. ([2015b](#page-10-8)), who discovered that decreases in ground vegetation structure and coverage reduced butterfy species richness. Both adult and larvae of butterfies require a certain host plant that serves as food source and their richness is closely linked to host plant distribution and nectar availability (Collinge et al. [2003\)](#page-10-12). The diversity of butterfies is also afected by the structure and composition of ground vegetation (Kremen et al. [1993;](#page-11-24) New et al. [1995\)](#page-11-25). Most of the butterfies, particularly those in the Morphinae and Satyrinae subfamilies, are flying at low to moderate height (Schulze et al. [2001;](#page-12-14) Kumar et al. [2009\)](#page-11-26) to fnd their resources (e.g., host plants, mates and foods) (Vane-Wright [2015](#page-12-17)).

A negative relationship was observed between butterfy species richness and elevation. This negative association was expected since butterfy diversity, species richness, and species abundance are generally higher in low-elevation habitats than in high-elevation habitats (Lien and Yuan [2003\)](#page-11-27). Several factors contribute to this negative relationship, including a decreasing trend in plant species richness and habitat heterogeneity, which decreases with elevation and eventually leads to lower insect diversity (Lieberman et al. [1996](#page-11-28); Grytnes and Beaman [2006](#page-11-29); Achrya and Vijayan [2015](#page-10-28)). Moreover, butterfy distributions in Malaysia are constrained by altitude and plant associations, with approximately half of all butterfy species occurring below 750 m above sea level (asl) (Corbet and Pandelbury [2020\)](#page-10-16). In addition, at elevations above 1000 m (asl), there are fewer butterfy species, but butterfy species that exist at higher elevations do not live at lower levels (Kirton [2014](#page-11-15)).

During the wet season, butterfy diversity is lower as rainfall afects humidity, temperature, and local microcli-mate (Speight et al. [2008](#page-12-18)). Most of the butterflies tend to be inactive during this period since low temperatures and precipitation may limit their activities. Butterfies are coolblooded organisms. Even small changes in ambient temperature afect their abundance, foraging activity, mating,

and physiology (Kearns [2012](#page-11-30); Klok [2013](#page-11-31)). Aside from that, heavy rain can impact the survival of butterfly larvae and pupae (Hill et al. [2003](#page-11-32)). Thus, lower butterfy species richness during the wet season is caused by environmental factors that are not measured in this study.

The mean NDVI and butterfly species richness were found to have a negative relationship. In this study, the mean NDVI in oil palm plantations was the highest followed by the rubber plantations, but the lowest when it came to butterfy species richness. Such results are expected, chemical fertilizers are systematically used in conventional plantations, especially in oil palm plantations to boost yield productions where more than 85% of the operation cost went to fertilizers alone (Goh [2005\)](#page-11-33). Fertilizer increases the chlorophyll content of plants, infuencing the NDVI reading (Gómez et al. [2019](#page-11-34)). Aside from that, the phenological diference in rubber tree plantations may infuence the NDVI reading especially during foliation stages resulting in higher NDVI reading (Yeang [2007;](#page-12-19) Janatul et al. [2018;](#page-11-35) Hazir et al. [2020](#page-11-36)). Rubber trees produce new leaves during this period, and the canopy recovers from defoliation, increasing the NDVI value (Dong et al. [2013](#page-10-29); Janatul et al. [2018\)](#page-11-35).

Our study also showed that species richness of the butterfy increased with decreasing distance to the nearest forest as the majority of the butterfy species observed in this study are associated with forest habitat. This fnding aligned with Lucey and Hill [\(2012\)](#page-11-10), and Panjaitan ([2020\)](#page-11-37), which reported that the spillover of butterfies of forest specialists from the nearest forest will improve diversity in the agriculture areas. Natural forests and secondary forest remnants adjacent to plantations serve as an important source of complementary breeding and feeding for butterfy species, especially forest dwellers, due to the availability of host plants (Koh [2008](#page-11-9)), and contribute to conservation on the landscape level (Veddeler et al. [2005\)](#page-12-20).

Our fndings suggested that ground vegetation cover and proximity to a forest were important in supporting butterfy species richness in agricultural landscapes. To improve butterfy diversity, the common practices in mainstream agriculture such as using synthetic herbicides for controlling overgrown shrubs and weeds should be reduced. Alternatively, livestock grazing which is more environmentally friendly can be integrated with the existing agricultural landscapes (Tohiran et al. [2017,](#page-12-21) [2019](#page-12-22); Azhar et al. [2017,](#page-10-30) [2021](#page-10-31)). Aside from that, preserving forest areas within agricultural landscapes is critical for facilitating butterfy dispersal and mitigating species loss due to land conversion (Lucey and Hill [2012;](#page-11-10) Lucey et al. [2014;](#page-11-38) Panjaitan et al. [2020](#page-11-37)). Small-scale farmers should practice alley-cropping system to improve biodiversity on their farmlands. This strategy could enhance the foristic composition and stand structural complexity for the arthropods, which eventually reduce the microclimate effects in the farmlands (Ashraf et al. [2018,](#page-10-11)

[2019\)](#page-10-32). Based on our data, agroforestry orchards showed the signifcance of habitat quality to boost biodiversity in agricultural landscapes and served as an example for key stakeholders in Southeast Asia's oil palm and rubber tree sector to improve biodiversity and ecosystem services in conventional plantations.

Our observational data are restricted to point counts of butterfy species. Other sampling methods, such as bait trapping, should be considered to obtain a comprehensive picture of the butterfy species community in future studies (Checa et al. [2019](#page-10-33); Hebel et al. [2022](#page-11-39)). However, due to disturbance from long-tailed macaques (*Macaca fascicularis*), bait trapping is not appropriate for use in our study area. Aside from that, future research on butterfy species communities in agricultural landscapes should include other factors that may infuence butterfies, such as the number of blossoms, water availability, host plants, and rotten fruits.

Conclusions

Based on the evidence presented in this study, tropical agricultural landscapes, particularly agroforestry orchards are instrumental for biodiversity conservation outside protected areas. Conservationists must consider the conservation value of agroforestry orchards for biodiversity to make conservation programs more efective. Because agroforestry orchards afect a substantial portion of the terrestrial region in Southeast Asia amid monoculture plantations, their contribution to biodiversity is critical for future conservation efforts.

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Data availability Data generated from this study will be provided upon reasonable request by the corresponding author.

Declarations

Conflict of interest The authors have no relevant competing fnancial interests or personal relationships that could have appeared to infuence the work reported in this paper.

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