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Tropical insectivorous birds' predation patterns that promote forest-farmland trophic connectivity for integrated top-down pest biocontrol

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Although conversion of natural forest to agriculture can negatively impact biodiversity in many ways, some affected species may respond by dispersing across the forest-farmland eco-zone, thereby facilitating functional connections through food-web interactions beneficial to crop production and forestry. This study examined patterns of Lepidoptera (butterflies and moths), Hemiptera (bugs), and Coleoptera (beetles) herbivory, and insectivorous bird predation within forestadjacent farms in western Kenya, and how these processes trophically connect the two ecosystems to promote pest biocontrol. Through δ^{13} C and δ^{15} N stable isotope analyses, proportions of maize, farmland legumes and forest trees in pest diets, and pest-prey in bird's diets were estimated. Birds' habitat associations and diet specializations' influence on pest consumption and basal plant carbon levels in birds' tissues were determined to evaluate birds' pest-biocontrol potential. Maize was the mostly consumed plant especially by Lepidoptera, but forest trees were peimarily consumed by Coleoptera and Hemiptera. In turn, Lepidoptera were mainly consumed by forest-associated birds, whereas Hemiptera and Coleoptera were mostly consumed by farmland-associated birds. Thus, birds showed crosshabitat pest consumption tendencies, though diet-specialization was unimportant in predicting those tendencies. Muscicapidae (flycatchers and allies); Hirundinidae (swifts and swallows); Motacillidae (pipits and wagtails); and Ploceidae (weavers) birds showed the highest contributory potential for pest biocontrol of Lepidoptera pests, but Estrildidae (manikins and waxbills), Muscicapidae, and Malaconotidae (boubous and gonoleks) birds showed the best potential against Hemiptera and Coleoptera. Furthermore, more maize basal carbon was assimilated by forest-associated compared to farmlandassociated birds, whereas most basal carbon from farmland legumes and forest trees were assimilated by farmland birds, suggesting that unlike pestprey choice, basal plant carbon pathways to avian insectivorous consumers did not strongly mirror birds' habitat associations. Lepidoptera and Hemiptera were potentially the most significant interhabitat trophic connector arthropods, and for birds, Muscicapidae, Ploceidae, and Estrildidae. These findings show that such functional connectivity may be enhanced through increasing structural cover elements that promote insectivorous birds' dispersal between farmland and adjacent forests to boost their pest-regulation ecosystem service contribution. The results serve to inform effective management practices by agronomists, foresters, and land-use planners toward promoting landscape-scale-integrated pest management for sustainable agriculture and biodiversity conservation.

KEYWORDS

ecotone, trophic linkage, biocontrol, ecological networks, connectivity, insectivorous birds, stable isotopes



1 Introduction

The rapid loss of forest cover to agricultural expansion is a major driver of ecological pressure on forest biodiversity due to habitat deterioration and diminishing foraging opportunities in the remnant forest patches (Bélanger and Grenier, 2002; Newmark and Stanley, 2011; Peter et al., 2015; Gonzalez et al., 2017). Furthermore, centuries of intensification of commercial agriculture, along with most farmers' heavy dependency on chemical inputs for crop production, have done little to guarantee global food security or eliminate impacts of arthropod pests on crop yields. Instead, such agronomic practices have continued to adversely affect wildlife and human health, in addition to fostering resistance by some target arthropod crop herbivores (Gavrilescu, 2005; Tiryaki and Temur, 2010; Deguine et al., 2021). In addition, forest loss and fragmentation have rendered many forest-associated species more dependent on adjacent farmlands for survival (Berg and Pärt, 1994; Newmark and Stanley, 2011; Garbach et al., 2012). Likewise, many farm-dwelling organisms are adversely impacted not only by pollution from agrochemical inputs but also through anthropogenic disturbance and unpredictable fluxes in resource availability occasioned by agronomic and structural habitat management practices (Tscharntke et al., 2008; Rincon-Parra et al., 2022).

As a consequence of these factors, only species adapting to such perturbations can persist in forest-farm ecotones and have to disperse across habitat to obtain resources necessary for their survival (Bos et al., 2007; Gonzalez et al.;., 2017; Alvarez-Alvarez et al., 2022). For instance, bees and wasps normally establish colonies in forests but often forage for nectar on neighboring farmlands (Hill and Webster, 1995) and many butterfly species oviposit and breed on croplands but roost and hibernate in woodlands (Sosa-Aranda et al., 2018). Similarly, several avian insectivores like flycatchers and granivores like grosbeaks nest in wooded areas but forage on croplands (Douglas et al., 2014), whereas warblers, sparrows, stocks, cranes and many raptors, which feed predominantly within the farmland space, frequently utilize adjacent forests for roosting and nesting (Berg and Part, 1994). Under circumstances of profound anthropogenic disturbance such as loss or deterioration in habitat quality, such cross-boundary dispersal and functional tendencies become significantly enhanced.

There is a wide range of negative consequences of such enhanced dispersal by species. These include arthropod pest outbreaks on croplands following clear-cutting of the adjacent forest stands, crop damage by forest-associated pest birds (Peter et al., 2015), or livestock depredation by carnivorous forest mammals and birds of prey (Goswami et al., 2015). Dispersing individuals may also transport pollutants and other harmful chemical material into forests (Fujita and Koike, 2007). On the other hand, positive

consequences and benefits of species' dispersal within the forest-farm matrix may occur through enhanced ecological connections between the two adjacent ecosystems. This can support broader scale functional networks of fragmented habitat patches within the matrix (Pejchar et al., 2018; Grass et al., 2019) and may take the form of expanded food-webs arising from increased trophic interactions (Dias et al., 2019) or genetic linkages through breeding opportunities (Klinga et al., 2019). In turn, such processes can enhance biodiversity's provisioning of important ecological services for both habitat zones, including natural pest biocontrol, crop pollination, seed dispersal, nutrient transport, and overall interhabitat energy subsidies or exchange (Decocq et al., 2016; Nyffeler et al., 2018; Grass et al., 2019). Based on such ecological processes and opportunities, biodiversity conservation practitioners now recognize that landscape management decisions for promoting sustainability of such benefits are likely to yield more long-term rewards for agricultural and silvicultural enterprises because they promote landscape resilience while also conserving biodiversity. Therefore, integrated pest management systems and field practices to conserve farm-forest landscape matrix connectivity can significantly boost cross-boundary food-web networks, and this can increase the potential for sustaining net benefits for both agriculture and forest management (Landis, 2017; Pejchar et al., 2018; Nelson et al., 2022).

Although many frugivorous and granivorous bird species mediate significant economic losses to farmers through crop damage (Firake et al., 2016; Lindell et al., 2016; Ernst et al., 2019; Kross et al., 2019), many insectivorous and omnivorous counterparts provide vital services by removing arthropod pests from many varieties of row crops (Girard et al., 2012; Garcia et al., 2018; Garfinkel et al., 2020). Depending on their foraging and ranging patterns, flocking behavior, diet specializations, or daily time budgets, some of these avian insectivores may provide pest removal services across many crop types and spatial scales (Shakelford and Corner, 1997). Nyffeler et al. (2018) estimated the annual global value of insects consumed by insectivorous birds on croplands to be in the region of 30 million metric tons. Insectivorous bird species inhabiting transitional habitat regions can be especially important in contributing to pest regulation from multiple plant hosts, whether agricultural or wild-growing, or across habitat boundaries such farmland-prairie, as farmland-wetland, or farmland-forest ecotones (Exnerová, et al., 2003; Massemin et al., 2006). Through this, such bird guilds can make significant contributions to reducing losses associated with crop damage, forest-tree defoliation, or pasture-land degradation (Exnerová, et al., 2003).

Majority of past studies on impacts of arthropod pests have focused on single crops types, single arthropod herbivore species, taxa, functional guild, or trophic relationships to a single specific suite of natural enemies or avian predator (Bendell et al., 1981; Tremblay et al., 2001; Amudavi et al., 2008; Hartman et al., 2016; Koch et al., 2016; Milligan et al., 2016; Ferger et al., 2012; Garcia et al., 2018). Few studies have extensively examined arthropod herbivory or insectivorous birds' predation patterns within nonagricultural habitats or herbivorous arthropods' impacts on specific non-crop host plants. Yet, in many countries in both temperate and tropical regions, substantial losses are incurred from arthropod pest damage in forestry and timber production systems, which have economic significance comparable to that of agriculture (Wylie and Speight, 2012; FAO, 2018; Li et al., 2019; Forest research, 2021). Furthermore, past studies on roles of birds in contributing to arthropod crop pest regulation in East Africa are still much fewer (Milligan et al., 2016; Otieno et al., 2019b) than those conducted in other regions (Tremblay et al., 2001; Karp et al., 2013; Garfinkel et al., 2020). Similarly, very few studies have applied stable isotopic techniques to assess the role of birds as contributors to arthropod pest regulation (Girard et al., 2012; Pagani-Núñez et al., 2017), even though a number of researchers have used them to assess diets of passerine birds in general (Hobson and Clark, 1992; Robbins et al., 2005; MacLellan, 2012; Ruhl et al., 2020). The advantage of stable isotope analyses (SIAs) over exclusion-cage, stomach-content analysis of, or other molecular techniques such as DNA barcoding, is that SIA facilitates explicit deciphering of consumers' diet options as well as the estimation of proportional contribution of each food option to consumers' diets, when combined with Bayesian mixing model procedures (Phillips et al., 2005; Parnell et al., 2013; Stock et al., 2018).

This study's overall aim was to apply $\delta^{13}C$ and $\delta^{15}N$ stable isotope analyses in assessing patterns and extent to which trophic interactions of common passerine insect-eating birds with three herbivorous arthropod taxa mediate the interflow of diet carbon subsidies between forest and adjacent farmland habitats, with implication for natural pest biocontrol in both habitats. The specific objectives were as follows:

- 1) To estimate dietary proportions of maize, farmland legume plants, and forest-trees assimilated in tissues of Lepidoptera, Hemiptera, and Coleoptera pests
- 2) To assess dietary proportions of the three pest-food sources in the common passerine birds of the farmland-forest ecotone
- To determine whether birds' habitat associations and diet specializations relate to choice and dietary proportions of arthropod pests they consume
- 4) To evaluate birds for their relative significance in contributing to top-down pest regulation based on their dietary pest-food proportions and levels of basal plant carbon (diet carbon derived from the bottom of the food chain)

To achieve those objectives, we interrogated the following two working hypotheses: 1) farmland- and forest-associated birds predominantly consume pest-prey primarily associated with the respective habitat zones and 2) proportions of plant basal carbon assimilated into tissues of bird families directly reflect the bird's dietary proportion of the pest-prey predominantly consuming that plant food resource.

In the first hypothesis, we assumed that forest and farmland habitats are functionally distinct with little or no trophic exchange, while the second one assumed that birds' arthropod pest-prey were essentially monophagous with significant trophic fidelity to specific plant food sources or types.

The present study is therefore significant in presenting an integrated examination of the arthropod pest management question from a perspective of multiple consumer taxa, trophic levels and host plants within two different land use types. It is also unique in being the first in the region to use SIA to address the objectives outlined.

2 Materials and methods

2.1 Study area

The study was conducted in western Kenya's County of Kakamega, on the farmland to the eastern side of Kakamega Forest (00° 11' 09"N-00° 26' 08"N and 34° 44' 30"E-34° 51' 26"E), an area of mid-level elevation c. 1,600 m asl with cropfield matrices, human settlements, small towns, and water courses, with the main crop grown being non-Bt maize (Farwig et al., 2008; Laube et al., 2008). The average temperature in this area is 21.4°C, and rainfall is approximately 1,800 mm per annum, with longerperiod seasonal rains peaking between March and May and shorterperiod rains peaking around November (The Kenya Meteorological Department, 2021). Kakamega Forest is the only true tropical rainforest in Kenya, which has undergone substantial cover loss form anthropogenic encroachment and conversion to agriculture and settlement over the past decades, leaving only 85 km² of closedcanopy primary forest from 300 km² nearly a century ago (Farwig et al.). As a relic of a previously contiguous continental Guineo-Congolean rainforest belt, it retains considerably high endemism of original flora and fauna, particularly birds, including two globally endangered and 15 regionally threatened species (BirdLife International, 2012; IUCN, 2023). Partly as a result of such significance, the forest is listed among Kenya's network of Important Bird Areas and as a UNESCO World Heritage Centre (Bird Life International, 2012). Except from the last two and a half decades when official forest protection enforcement began to improve, the trends in forest's destruction, degradation, and fragmentation had traditionally been tied to the fact that it is surrounded by farmland and a dense human population practicing mainly subsistence agriculture and small-scale livestock husbandry but with considerable dependence on the forest for wood, fiber, and other products, though this is now regulated by forest management authorities (Farwig et al., 2008). The main crop on the eastern side of the forest includes mainly non-Bt maize, which is commonly intercropped with legume crops (Otieno et al., 2019a). Farmers also typically maintain some stand of woodlots and fruit orchards, and also plant trees along property boundaries, and occasionally establish fast-growing agroforestry trees alongside row-crops (Diwani et al., 2013).

Field sampling was carried out during two maize cropping seasons (long-rain and short-rain) in 2007 at three cropping stages each between October and the following July. The cropping stages were early crop (from germination to first weeding), midcrop (from second weeding through flowering to ear formation), and mature crop (from cob hardening to harvesting). This was aimed at minimizing potential sampling bias on the temporal scale (Lancaster et al., 2006; Meadows et al., 2012).

2.2 Study farms selection and characterization

Farm selection was based primarily on the cropping system and proximity to forest such that a total of 15 farms were selected, which had fields of maize and at least some legume plants either intercropped with maize, or planted in separate field plots, or growing wild within fallow field patches. The forest's proximity to the crop fields had to be at most 1 km to ensure there was reasonable chance of trophic interactivity between birds and arthropods across the two habitats through dispersal (Ding et al., 2011). A 2-km minimum interfarm distance was maintained to ensure sampling independence with regards to the more mobile bird and arthropod groups (Lancaster et al., 2006). Furthermore, only farms in which maize or maize with other legume crops had been grown for at least two consecutive seasons were selected, in order to reduce significant short-term probabilistic fluctuations in arthropod and bird assemblage patterns.

2.3 Arthropod sampling

Plant and arthropod samples were taken during the three crop-growing stages outlined previously. Arthropods were collected using both standard sweep nets (using the standardized 100 sweeps along transects in each farm) and pitfall traps, which comprised the standard 70-mm diameter and 120-mm high plastic cups inserted upright and flush with the ground surface (Sekamatte et al., 2003). The pitfall traps were filled to one-third volume with 25% sodium chloride solution for preservation and to maintain samples' isotopic integrity, with and conical plastic rain shields propped above each trap. Three replicates were randomly placed along a diagonal line running across each maize field, at distance intervals which depended on the farm size. The traps were collected after 3 days. The samples were discarded, and traps were reset if flooding occurred from rainfall. Arthropods were additionally sampled on selected individual plants using a search-and-pick procedure on the leaves, stems, and flowers using forceps. At trap collection, the samples were transferred to airtight bags and frozen. Using illustrated field guides and identification keys tools, the arthropod samples were later identified to family, with only primary consumers (herbivores) retained. Subsequently, in order to achieve a sufficiently clear analytical resolution of isotopic signatures, the arthropod samples were pooled into three taxonomic groups (Orders), namely, Coleopteran (leaf beetles, flower beetles, sap beetles, and weevils), Hemiptera (leaf bugs, true bugs, and stink bugs), and Lepidoptera (moths and butterflies) (Hyodo, 2015).

2.4 Bird sampling

Bird samples were collected by the use mist nets. Bird mistnetting was conducted once on each farm, at the onset of the long rains in early April, and in each case, two 12-m mist nets were set up running perpendicular to the forest edge, one starting at the forest edge itself and a second one 25 m into the forest interior, ensuring that the two net lines were not directly aligned to each other (Newmark and Stanley, 2011; Ralph et al., 2014). The nets were set up and left out for 2 hours from 0700–0900 at each study farm, with inspections conducted every 15 min to extract captured individuals and to avoid any stressful impacts or injuries to the birds. At net inspection, the birds were identified to species, and subsequently a small portion was clipped off the tip of a primary wing feather (Hobson and Clark, 1992). Birds awaiting processing were kept in dark seclusion inside well-aerated wide-fitting cotton-fabric bags. To further prevent birds from being stressed out, it was ensured that the sampled birds were subsequently released back at the exact place of capture, within a maximum of 10 min after capture (Ralph et al., 2014). Clipped feather samples were immediately sealed in paper envelopes to minimize contamination, labelled, and sent to the laboratory.

2.5 Plant sampling

From each study farm, 12 plant-leaf samples were collected randomly (four each from maize, legume plants, and forest trees). On croplands, systematic randomization was achieved by sampling from the fifth plant of every 10th row, whereas in the forest, samples were collected from the central plant within each of 2-m radius quadrats established from forest edge toward forest interior at 10-m intervals, hence ~100 m into forest (Ilić et al., 2018). The plants were divided into three categories for purposes of SIA, so as to establish three isotopically distinct basal food-source origins for consumers (Ostrom et al., 1997; Layman et al., 2012). The three groups were: 1) maize, a crop plant which follows a C₄ photosynthetic pathway, 2) farmland legumes, field crops or farmland plants which follow a C₃ photosynthetic pathway, and 3) forest trees, which follow the C3 photosynthetic pathway. Immediately after collection, the samples were sealed in paper envelopes to reduce moisture loss or air contamination, labelled, and sent to the laboratory.

2.6 Pre-test processing of samples

In the laboratory, bird feathers were rinsed in 4% solution of formaldehyde to remove excessive oil and impurities before further handling (Paritte and Kelly, 2009; Haché et al., 2012). All bird-feather, arthropod, and plant-leaf samples were then oven-dried at 60°C to constant mass before being ground to fine powder. Arthropods from each of the taxonomic group were ground whole. For each farm, two 5-mg subsample replicate portions were then prepared per sample of each arthropod and plant group. The subsamples were subsequently transferred into tinfoil capsules for subsequent isotopic analyses.

2.7 SIA for consumer diet compositions

To identify the array of consumers' food sources, the processed samples were analyzed at the Environmental Isotope Laboratory of the iThemba Laboratory for the Accelerator Based Sciences (LABS) in Johannesburg, South Africa. These tests were conducted to determine signatures of δ^{13} C and δ^{15} N isotopes in the consumer and food source tissues. This was achieved using Flash HT Plus elemental analyzer coupled to a Delta V Advantage isotope ratio mass spectrometer by a *ConFloIV* interface combusting at 1 020°C (*ThermoFisher, Bremen, Germany*). These models are founded on the isotopic mass balance equation: $\delta X = [(R_{sample}/R_{standard}-1)] *$

1 000, where *X* refers to 13 C or 15 N and *R* is the corresponding ratio between ${}^{13}C/{}^{12}$ C and ${}^{15}N/{}^{14}$ N, respectively, and the standards based on the International Atomic Energy Agency standard sample stock in Vienna (Phillips et al., 2005).

2.8 Data analyses

Analyses of data for arthropod consumers (pests) were conducted at three taxonomic group (order) levels (Coleoptera, Hemiptera, and Lepidoptera). Each of the bird species was assigned to one of the two diet specialization guilds and a taxonomic family. The two guilds of diet specialization were obligate insectivore (predominantly consumes arthropods) and omnivore (consumes both arthropods and plant food material). The birds were further classified into two habitat association categories, that is, those primarily associated with the forest habitat and those primarily associated with the farmland. This categorization combines the classification by Bennun et al. (2009) for forest birds and that of the field guide to the birds of Kenya and Northern Tanzania (Zimmerman et al. (1999). Bird families represented by individual species that were encountered only once were excluded from the analyses. Thus, of the total 79 species encountered representing 24 families, 70 species from 16 families were subjected to analyses (Supplementary Table S1, listing according BirdLife International and NatureServe, 2011).

2.8.1 Food source contributions to consumers' diets

From the stable isotopic signatures incorporated in their tissues, estimates were made of relative contributions of the plant food sources to arthropod pest diets and arthropod pestfood sources to birds' diets. This was modeled using the mixing model procedure with the MixSIAR tool in R v.4.0 (R Core Team, 2021). In addition to identifying the range of various food source options to consumer's diets, MixSIAR incorporates appropriate fractionation discrimination values (or trophic enrichment factor (TEF)) to calculate the relative contribution of each food source to the consumer diets (Caut et al., 2009; Layman et al., 2012; Parnell et al., 2013; Stock and Semmens, 2016). TEFs serve to correct for any significant errors attributable to a consumerspecific metabolic process (Hobson and Clark, 1992; Layman et al., 2012; Parnell et al., 2013). The TEF values applied here were averages relevant to herbivorous arthropod consumers from a review by Spence and Rosenheim (2005), while for avian consumers, the values applied were those specific to avian feathers as applied by Ferger et al. (2012). In addition, evaluation was made of proportions of the plant basal carbon assimilated into birds' tissues via the arthropod pests to track the broader pathway for the farmland-forest trophic carbon subsidy and exchange (Decocq et al., 2016). For this purpose, the TEFs of the two trophic steps (plants to pests and pests to birds) were each combined by summing up into one final value. The MixSIAR model platform further facilitates testing of the role of environmental or other habitat explanatory factors that underpin the proportional food source contributions to

consumer diets (Stock and Semmens, 2016). Accordingly, in this study, the role of bird habitat association in driving dietary proportions of the various pest-food sources into birds' diets was analyzed.

The models were run separately for pest-to-plant (herbivory) and bird-to-pest (predation) trophic linkages. In running each model, mixture data (consumer file) were input, followed by source files (food sources data) subsequently incorporating respective TEFs (Stock and Semmens, 2016). Further analyses involved selecting and running uninformative priors, designing Jags models with specific error structures (residual and process, in the present case) (Stock and Semmens, 2016) followed by running a series of Jags models starting with a test Markov Chain Monte Carlo (MCMC) chain (1,000 length, 500 burn-in iterations) and building up chain lengths in subsequent runs through "very short," "normal," or "long" (chain length and burn-in iterations of 1,000, 5,000; 50,000, 25,000; and 100,000, 50,000, respectively) until convergences on true posterior distributions of estimated food source proportion contribution to consumer diets were achieved from the Jags outputs. This was confirmed using the Gelman-Rubin diagnostic procedure (Stock et al., 2018). The final outputs of summary statistics and posterior density plots of consumers' dietary proportions of food sources were then evaluated for median and mean quantile distributions and 95% credible intervals of such estimates.

Assessment was also made of the relationship between observed dietary proportions of birds' pest-food sources, to bird family identity, arthropod taxa identity, birds' habitat association, and birds' diet specialization. This was achieved using permutational analysis of variance, PERRMANOVA in Primer v.7 with PERMANOVA + add-on (Anderson, 2017; PRIMER-E, 2019). Here, the models were run for main tests with 9,999 permutations. The results of the analysis were plotted using principal coordinate ordination (PCO). Finally, canonical analyses of principal ordinates, CAP, was used to evaluate and plot relative importance of bird's habitat association and pest taxa identity, in relation to pests' contribution to birds' diets. All tests in PRIMER were performed on log (X+1) transformed Euclidean distance resemblance matrices derived from respective dissimilarity matrices of consumers' dietary proportions data.

3 Results

A total of 248 individual birds were captured and of the 70 bird species identified, 11 families were obligate insectivores (six farmland-associated and five forest-associated) and five families were omnivores (three farmland-associated and two forestassociated) as outlined in Supplementary Table S1.

3.1 Plant food sources in arthropod consumer diets

Each of the three arthropod pest taxa derived diet carbon from all the three plant food sources (Table 1). Nonetheless, in general, the bulk of pest consumer's plant food carbon originated from the farmland rather than from the forest, though maize was more targeted by the pests than were farmland legume plants, with each of the three pest groups deriving more than half of its plant carbon from maize (Table 1). Maize was a particularly important food source for Lepidoptera and Coleoptera pests (Figure 1A), while forest trees and farmland legumes were consumed mainly by Hemiptera pests (Table 1; Figure 1B). Although Lepidoptera pests incorporated the highest proportions of plant food in general, with a major trophic focal bias toward maize, Coleoptera and Hemiptera pests showed a more widespread trophic linkage to all three plant food options (Figures 1A–C).

3.2 Arthropod pest food in birds' diets

3.2.1 Food source proportions overall

Lepidoptera was disproportionately the most important pestprey for birds overall (median = 0.777, 95% CI = 0.656–0.913), followed by Hemiptera pests (median = 0.136, 95% CI = 0.011-0.289), while Coleoptera was the least preferred food source (median = 0.062, 95% CI = 0.003-0.248), making Lepidoptera almost six times more preferred by birds as compared to the other two options combined (Figure 2A).

3.2.2 Arthropod food proportions at bird families and habitat association scales

At the family level, although every bird family consumed Lepidoptera pests proportionately more than they did the other pest items (Table 2; Figure 2A), the four most significant consumers of Lepidoptera were muscicapids (flycatchers), hirundinids (swallows, martins, and saw-wings), ploceids (weavers), and motacillids (pipits and wagtails), each of these consuming a medial value of at least 85% of this pest-prey item. The four most important consumers of Coleoptera pests were estrildids (manikins, bluebills, and waxbills), muscicapids, ploceids, and malaconotids (boubous, gonoleks, and tchagras) (Table 2).

From the PERMANOVA results, these patterns were also revealed through the ordination analysis, with a 49% Eigenvalue in the first axis of PCO alone (Figure 3A). Nonetheless, bird's family identities in general (F = 0.101, p = 0.676, and df = 15), their habitat associations (F = 0.432, p = 0.711, and df = 1), and diet specializations (F = 0.178, p = 0.928, and df = 1) were not as significant in explaining variations in dietary pest food proportions as were identities of pest-prey items (F = 147.7, p < 0.001, and df = 2) (Figure 3B). Furthermore, ordination plots from CAP analyses showed that birds' pest-food choice had greater significance than habitat association in accounting for variations in birds' dietary pest food proportions (Figure 3B), with birds targeting prey items largely regardless of their own habitat associations.

Birds associated with farmland consumed higher proportions of Coleoptera and Hemiptera pests, which they derived mostly from forests (Table 3; Figure 2B). Forest-associated birds, on the other hand, incorporated higher proportions of Lepidopteran pest-prey items, mainly associated with the farmland habitat (Figure 2C).

3.3 Plant diet carbon in bird tissues

Of the three basal plant food options, maize showed the highest proportion in tissues of birds in general (median = 0.889 and CI = 0.744-0.979) followed by forest trees (median = 0.086 and

Otieno and Mukasi

CI = 0.006–0.240) and farmland legumes (median = 0.015 and CI = 0.001–0.079).

3.3.1 Proportions of basal plant diet carbon in birds by habitat association

Counterintuitively, higher proportions of maize diet carbon were detected in tissues of forest-associated birds compared to those detected in farmland birds (Table 4; Figures 4B, C). Conversely, higher proportions of forest-tree carbon were evident in farmland-associated birds, even though the farmland-associated birds also assimilated higher proportions of legume plants (Table 4; Figure 4B).

3.3.2 Proportions of basal plant diet carbon in birds at family scale

Whereas proportions of maize diet carbon was the highest in muscicapid, cisticollid (warblers and allies), ploceids, and stenosterid (blue flycatchers) tissues, diet carbon from forest trees occurred in highest proportions in tissues of muscicapids, estrildids, malaconotids, and acrocephalids (yellow warblers), while the leading assimilators of farmland legume carbon were estrildids, muscicapids, ploceids, and malaconotids (Table 5). Just as was seen for the overall case, of the three basal plant food sources, the highest proportion of dietary carbon across all bird species was contributed by maize.

These patterns were also depicted by PCO ordination results (Figure 5B), with PERMANOVA test outcomes further showing that variations in proportions of basal plant diet carbon in bird's tissues may be significantly explained by birds' diet specialization (F = 7.520, p = 0.002, and df = 1) and plant food choices (F = 171.4, p < 0.01, and df = 2), but not by birds' family identity (F = 0.881, p = 0.628, and df = 15) or habitat associations (F = 0.812, p = 0.415, and df = 1). The PCO ordination here showed a total of 47% of these similarity patterns explained by the Eigenvalues of the first two axes (Figure 5B).

4 Discussion

Even though the location of agricultural land in forest neighborhoods may not always compensate for the loss of native forest cover (Douglas et al., 2014; Kross et al., 2020), this study found nearly equitable representation of bird families between the two adjacent ecosystems. This is significant as it indicates that this farm-forest ecotone bears correspondingly equal importance to the respective avian communities (Deikumah et al., 2017). Within the prevailing land use systems in the study area, much of the original old-growth forest has been significantly reduced and fragmented over the decades to give way for agriculture and human settlement (Farwig et al., 2008). This study, therefore, has revealed that the surrounding farmland is important as a non-permanent but dynamic buffer providing dispersal and foraging opportunities to many affected non-specialist but forest-dependent insectivorous birds (Laube et al., 2008; Newmark and Stanley, 2011). Sekercioğlu (2002) demonstrated that, in fragmented forests, opportunity for insectivorous birds to utilize adjacent non-forest habitat or to disperse to the nearest forest patch is critical for longterm persistence. This is due to the necessity for such birds to compensate for declines in abundance and variety of prey resources occasioned by fragmentation-driven declines in sizes of foraging habitat and ranges. Conversely, the forest habitat may be of strategic significance to farmland insectivores for roosting, nesting, and as refuge from predation risk (Berg and Pärt, 1994). These complementary values of the two adjacent habitat types may thus promote the conservation of birds and other taxa, thereby contributing to overall landscape-scale functional diversity through expanded ecological networks underpinned by trophic interactions (Grass et al., 2019; Kross et al., 2020). This can, in turn, potentially promote provisioning of the ecological service of crop pest regulation through cross-boundary spillover effects of vertebrate and invertebrate predators associated with either habitat zone (Blitzer et al., 2012; Lucey and Hill., 2012; Gonzalez et al., 2017). However, the observed dominance of obligate insectivores within the assemblage of birds assessed in this ecotone primarily confirms that many species of this guild actually disperse widely within the forest-farmland mosaic to track the variety of arthropod prey or other proximate resources across habitat boundaries (Jones et al., 2005a). The predominance of obligate insectivores may further be attributed to the general dominance of avian insectivory in most ecosystems when compared to other avian diet guilds (Nyffeler et al., 2018). The significance of this finding is that insectivorous birds of both habitat zones bear a greater potential for facilitating the functional connectivity of the two habitat types despite the inherent anthropogenic perturbations in both zones.

4.1 Plant food sources in arthropod pest diets

Although a number of individual arthropod pest species are known to be monophagous and thus exhibit trophic fidelity to specific crop or non-crop host plants (Gates, 1980), it was evident from this study that at the taxa level, each of the three pest groups obtained food from both crops and non-crop plants. From the perspective of spatial and temporal fluxes in resource abundance such as the cycles of crop planting seasons followed by after-harvest transitions, such taxon-level food resource diversification and flexibility are important for the herbivores' survival and persistence (Driscoll et al., 2013). However, the herbivores 'apparent overall preference of maize over the other two plant food sources likely owed to the position of maize as the predominant and most abundant subsistence row-crop, as well as being the most common across these particular farming landscapes (Laube et al., 2008; Otieno et al., 2019a) For this reason. therefore, of the three plant food options assessed here, maize was the most readily available to most herbivorous species of the three arthropod taxa. An additional plausible explanation is that maize was more frequently encountered than forest trees or farmland legumes within the ecotone region, especially since legumes are typically planted primary as intercrops in between maize rows. In comparison to the other two plant food source options, maize is also more likely to be selected by arthropod herbivore consumers due to relatively higher interaction likelihood. This is because in addition to its value as a food source, maize's structural attributes also offer a wide variety of niches and microhabitats for various herbivorous arthropods that can utilize these for oviposition, aestivation, or refuge from predation risk (Broad et al., 2008; Suckling et al., 2017). For these reasons, the leading role of Lepidoptera pests as maize

| Plant food source | Arthropod herbivore | Median quantile | 95% credible interval (CI) |
|-------------------|---------------------|-----------------|----------------------------|
| Maize | Coleoptera | 0.652 | 0.556-0.742 |
| | Hemiptera | 0.505 | 0.367-0.630 |
| | Lepidoptera | 0.835 | 0.527-0.975 |
| Forest plants | Coleoptera | 0.265 | 0.041-0.392 |
| | Hemiptera | 0.422 | 0.049 0.595 |
| | Lepidoptera | 0.098 | 0.007-0.376 |
| Farmland legumes | Coleoptera | 0.050 | 0.004-0.336 |
| | Hemiptera | 0.069 | 0.001-0.477 |
| | Lepidoptera | 0.032 | 0.001-0.285 |

TABLE 1 Summary of estimated proportions of plant food sources in arthropod pest diets.



consumers shows that Lepidoptera constitutes the most significant arthropod pests of maize in this region. This pattern is well documented form studies in many maize growing regions of the world (Amudavi et al., 2008; Suckling et al., 2017; Khan et al., 2022).

On the other hand, the results showed Hemiptera bugs to be potentially the most significant pests of forest trees and farmland legumes. For farmland legume plants, such hemipteran herbivores would mainly include leaf bugs such as stink bugs (Penatomidae) and plant bugs (Miridae) (Hartman et al., 2016; Garfinkel et al., 2020) and sap-sucking bugs such as aphids (Aphididae) (Koch et al., 2016), while for forest trees, these include mainly the sap-sucking bugs of the scale insect variety (Coccomorpha). As insect pests, scale insects are among the most economically important in timber production systems worldwide (MacLean, 2016; Ülgentürk and Dokuyucu, 2019; Marini et al., 2022), although some Lepidopteran pest species in the genus Hypsipyla, Condylorrhiza,



and Doratifera are also recognized for causing serious damage in tropical forestry through tree defoliation (Wylie and Speight, 2012).

It was noteworthy that despite being a widespread taxa across many habitats especially croplands, and although they essentially derived their diet carbon from all the three plant food sources, Lepidoptera showed the strongest trophic connection bias toward maize with only marginal dependence on farmland legumes and forest plants (Broad et al., 2008). Contrastingly, despite consuming lower proportions of maize when compared to the case for Lepidoptera, Hemiptera, and Coleoptera pests exhibited nearly proportionate trophic linkages across the range of three plant food resources. This further underscores the relatively higher threat that Lepidoptera herbivores pose to maize as compared to most coleopteran and hemipteran pests (Khan et al., 2022). It also indicates the relatively lower herbivory risk that Lepidoptera pose to farmland legumes and forest plants in this region. However, Wylie and Speight (2012) showed the risk of certain Lepidoptera pest species to specific timber varieties across the world. On the other hand, the finding also implies that by not being trophically biased in their plant food selection, Coleoptera and Hemiptera pests are likely more resilient herbivores and more capable than Lepidoptera, of thriving in diversified landscape matrices bearing row-crops and patches of natural habitat. For instance, Ülgentürk and Dokuyucu showed that some Hemiptera species within the scale insect family (Pseudococcidae and Hemiptera) have a wide range of polyphagy covering up to 300 host plant species which they target for herbivory. The authors observed that the host plants spanned a wide spectral range of land uses and husbandry types from agriculture, forestry to greenhouse horticulture, and even wild-growing species.

4.2 Arthropod pest food sources in bird diets

The dominance of Lepidoptera in bird's overall diets highlights this pest taxa's abundance, landscape spread, and availability to insectivorous bird consumers. This pattern has been reported from many past studies in many types of croplands (Janzen, 1988; Sosa-Aranda et al., 2018). As seen previously, this pest taxon derived its diet carbon from all of the three plant food sources across both the forest and farmland habitats. This observation should be particularly relevant in association with passerine bird's breeding periods during which the fledgling young are fed exclusively on easily available invertebrate prey such as butterfly and moth larvae (McKinnon et al., 2012; Sánchez-Fernández et al., 2020). In addition, of all arthropod herbivores in farmland and other ecosystems, Lepidoptera have one of the highest taxonomic diversities with a wide range of host food plants most of which are also used for habitation, oviposition, and aestivation (Goldstein, 2017; Suckling et al., 2017). However, the considerably low level of birds' consumption of Hemiptera and Coleoptera pests is not easy to explain, given that most species of these taxa also breed during the same periods as do Lepidoptera and should therefore be abundant and widely available to insectivorous birds (Johansson et al., 2020), albeit at somewhat lower scales compared to Lepidoptera. In fact, a number of studies have cited Coleoptera pests as being nearly as preferred as Lepidoptera to many insectivorous birds (Nyffeler et al., 2018).

In terms of pest regulation roles, by being the most avid predators on Lepidoptera pests, muscicapids, hirundinids, ploceids, and motacillids seemed to be the most important contributors to the removal of these pest taxa from the cropland. Similarly, muscicapids, estrildids, malaconotids, and ploceids seemed to be the most significant contributors to removal of Hemiptera and Coleoptera pests' from farm legumes and forest plants. Indeed, by consuming approximately more than double the proportions of Hemiptera and Coleoptera compared to any other bird species, muscicapids, estrildids, and ploceids would be the most efficient avian contributors for the regulation of these pest taxa in the forest-farmland zone as a whole. These findings are supported by some related past studies on avian-arthropod feeding relations in farmland habitats. For instance, Ferger et al. (2012) showed that muscicapids flycatchers, ploceid weavers, and blue flycatchers (Stenosteridae) were the leading overall arthropod pest consumers on the farmlands of Kakamega County's sugarcanedominated farming landscape, which is located to the western side of Kakamega Forest opposite to the area of the present

| Bird family | Arthropod pest food source | | | | | | | | |
|----------------|------------------------------|--------|------------------------------|-----------|--------|----------------|-------------|-----------------|---------|
| | Coleoptera | | | Hemiptera | | | Lepidoptera | | |
| | Median 95% credible interval | | Median 95% credible interval | | | Median 95% cre | | edible interval | |
| | | 2.5% Q | 97.5% Q | | 2.5% Q | 97.5% Q | | 2.5% Q | 97.5% Q |
| Acrocephalidae | 5.8 | 0.4 | 23.5 | 6.4 | 0.6 | 19.8 | 86.0 | 82.4 | 98.5 |
| Cisticollidae | 5.3 | 0.1 | 19.7 | 5.2 | 0.2 | 29.8 | 85.2 | 76.4 | 99.6 |
| Estrildidae | 15.9 | 0.3 | 77.1 | 23.6 | 0.3 | 77.1 | 71.2 | 23.5 | 99.3 |
| Hirundinidae | 2.5 | 0.1 | 19.9 | 4.4 | 0.2 | 17.3 | 87.1 | 78.4 | 99.7 |
| Laniidae | 4.1 | 0.1 | 33.6 | 4.4 | 0.2 | 29 | 79.3 | 67.8 | 99.6 |
| Malaconotidae | 9.6 | 0.2 | 56 | 12.4 | 0.3 | 46.2 | 69.6 | 46.8 | 99.4 |
| Motacillidae | 4.4 | 0.1 | 36.7 | 4.7 | 0.2 | 28.9 | 86.3 | 65.7 | 99.5 |
| Muscicapidae | 14.7 | 0.3 | 61.4 | 24.9 | 0.4 | 49.3 | 87.3 | 42 | 95.7 |
| Nectariniidae | 2.4 | 0.1 | 17.2 | 2.5 | 0.1 | 14.1 | 83.4 | 85.9 | 99.7 |
| Pellomeidae | 5.1 | 0.1 | 39.2 | 5.9 | 0.2 | 36.1 | 84.5 | 60.9 | 99.5 |
| Picidae | 4.4 | 0.1 | 31.9 | 4.6 | 0.2 | 28.2 | 84.3 | 69.3 | 99.6 |
| Platysteiridae | 3.4 | 0.1 | 25.9 | 3.6 | 0.2 | 21.5 | 80.3 | 77.6 | 99.6 |
| Ploceidae | 13.5 | 0.2 | 47.2 | 10.6 | 0.4 | 41 | 86.9 | 57.1 | 99.3 |
| Pycnonotidae | 4.6 | 0.2 | 28.9 | 5.7 | 0.2 | 26.8 | 85.9 | 74 | 99.6 |
| Stenostiridae | 2.7 | 0.1 | 20.8 | 2.8 | 0.1 | 19.6 | 84.7 | 76.9 | 99.6 |
| Turdoidae | 3.5 | 0.1 | 26.8 | 3.8 | 0.2 | 23 | 79.9 | 75.7 | 99.6 |

TABLE 2 Estimated dietary proportions of arthropod pest food sources assimilated by various bird families.

Proportions are expressed in percentages.



FIGURE 3

(A) Principal coordinate ordination (PCO) plots of dietary proportion distributions showing associational clustering of the various bird families to the three arthropod food source options, and (B) canonical analysis of principal ordinates (CAP) plot of the relationship between birds' habitat association and pest food proportions in birds' diets. *FaB* = Farmland-associated birds and *FOB* = Forest-associated birds. Ordination is based on log (X+normalized resemblance matrix of the Euclidean distance of the dietary proportions data.

| Birds by habitat association | Pest food source | Median quantile | 2.5% quantile | 97.5% quantile |
|------------------------------|------------------|-----------------|---------------|----------------|
| Farmland-associated | Coleoptera | 0.038 | 0.000 | 0.167 |
| Forest-associated | | 0.015 | 0.001 | 0.119 |
| Farmland-associated | Hemiptera | 0.023 | 0.000 | 0.107 |
| Forest-associated | | 0.010 | 0.001 | 0.076 |
| Farmland-associated | Lepidoptera | 0.926 | 0.792 | 0.990 |
| Forest-associated | | 0.967 | 0.849 | 0.997 |

TABLE 3 New estimated proportions of arthropod pest-prey in diets of birds by habitat association.

TABLE 4 Estimated proportions of plant diet carbon in tissues of birds by habitat association.

| Basal plant carbon in bird tissue | Birds' habitat association | Median quantile | 95% credible interval | |
|-----------------------------------|----------------------------|-----------------|-----------------------|----------------|
| | | | 2.5% quantile | 97.5% quantile |
| Maize | Farmland | 0.902 | 0.744 | 0.984 |
| | Forest | 0.933 | 0.731 | 0.995 |
| Forest trees | Farmland | 0.069 | 0.002 | 0.234 |
| | Forest | 0.044 | 0.001 | 0.255 |
| Farmland legumes | Farmland | 0.019 | 0.000 | 0.091 |
| | Forest | 0.009 | 0.000 | 0.080 |



study. A study within an east European forest-farmland mosaic that encompassed row-crops, pasture, and hedgerows also reported the importance of Hemiptera herbivores in diets of several muscicapid insectivorous bird species (Exnerová et al., 2003). However, so far, few quantitative studies have been conducted on the roles of hirundinids or motacillid birds in the removal of insect pests from agricultural crops. Similarly, no published reports have highlighted the significance of estrildid waxbills and manikins, or weavers in regulating Hemiptera or Coleoptera pests of forest plants, the latter two being more commonly associated with damage to cereal crops.

With regards to removal of specific arthropod pest taxa, a review by Jones et al. (2005b) showed that most caterpillars (Lepidoptera larvae) constituted substantial proportions of the diet of most insectivorous farmland birds across many parts of the United Sates. Garfinkel et al. (2020), however, demonstrated that the maize arthropod pests targeted most by insectivorous farmland birds were corn rootworms (*Diabrotica barberi*, Chrysomelidae:

| Bird family | Arthropod pest food source | | | | | | | | |
|----------------|----------------------------|-------------------------|---------|------------------------------|--------|---------------------|------------------|---------------|---------|
| | Maize | | | Forest trees | | | Farmland legumes | | |
| | Median | n 95% credible interval | | Median 95% credible interval | | Median 95% credible | | ible interval | |
| | | 2.5% Q* | 97.5% Q | | 2.5% Q | 97.5% Q | | 2.5% Q | 97.5% Q |
| Acrocephalidae | 93.3 | 79.9 | 98.4 | 5.2 | 0.2 | 19.0 | 1.0 | 0 | 6.0 |
| Cisticollidae | 96.7 | 84.4 | 99.7 | 2.1 | 0.1 | 14.1 | 0.5 | 0 | 5.6 |
| Estrildidae | 82.9 | 34.7 | 99.4 | 9.8 | 0.1 | 62.6 | 3.1 | 0 | 29.7 |
| Hirundinidae | 96.1 | 77.4 | 99.7 | 2.6 | 0.1 | 20.6 | 0.6 | 0 | 7.1 |
| Laniidae | 94.7 | 65.8 | 99.7 | 3.4 | 0.1 | 31.7 | 0.6 | 0 | 10.9 |
| Malaconotidae | 94.7 | 49.5 | 99.5 | 7.3 | 0.1 | 49.0 | 1.1 | 0 | 19.5 |
| Motacillidae | 94.6 | 66.5 | 99.6 | 3.5 | 0.1 | 30.4 | 0.7 | 0 | 11.1 |
| Muscicapidae | 97.2 | 44.5 | 95.1 | 25.5 | 0.2 | 53.9 | 2.1 | 0 | 25.0 |
| Nectariniidae | 94.3 | 80.0 | 99.7 | 2.3 | 0.1 | 18.7 | 0.5 | 0 | 6.3 |
| Pellomeidae | 91.9 | 53.5 | 99.5 | 5 | 0.1 | 43.6 | 0.9 | 0 | 14.2 |
| Picidae | 94.1 | 65.3 | 99.6 | 4 | 0.1 | 32.5 | 0.7 | 0 | 10.5 |
| Platysteiridae | 94.7 | 70.3 | 99.5 | 3.6 | 0.1 | 27.8 | 0.7 | 0 | 8.6 |
| Ploceidae | 96.4 | 61.5 | 99.4 | 7.9 | 0.1 | 36.1 | 1.1 | 0 | 14.8 |
| Pycnonotidae | 92.9 | 64.6 | 99.6 | 4.8 | 0.1 | 34.0 | 0.8 | 0 | 10.2 |
| Stenostiridae | 95.5 | 74.2 | 99.6 | 2.9 | 0.1 | 23.8 | 0.6 | 0 | 8.0 |
| Turdoidae | 95.7 | 71.2 | 99.7 | 3.2 | 71.2 | 27.6 | 0.6 | 0 | 8.5 |

TABLE 5 Estimated median dietary proportions of basal plant diet carbon incorporated in tissues of various bird families.

 $\mathbf{Q}^{\star}=$ quantile. Dietary food source proportions are expressed as percentages.



FIGURE 5

(A) Estimated median dietary proportion of basal plant food sources to diets of birds in general. Error bars represent standard deviations at 95% confidence intervals and (B) canonical analysis of principal ordinates (CAP) plot showing associational clustering of various birds to the three basal plant food sources based on diet specialization. Ordination is based on log (X+normalized resemblance matrix of the Euclidean distance of the dietary proportions data.

Coleoptera) and tarnished plant bugs (*Lygus lineolarus*, Miridae: Hemiptera) rather than the Lepidoptera group. Tremblay et al. (2001) further indicated that the levels at which lepidopteran cutworm pest (*Agotis* sp) populations were regulated in some American cornfield by insectivorous birds were not very different from those levels observed for corn leaf aphids (*Rhopalosiphum maidis*, Aphididae: Hemiptera), weevils (*Sphenophorus* sp., Curculionidae: Coleoptera), or corn rootworms. The authors did not, however, provide the pests' trophic linkages to any of the bird species.

Such variations in study findings, though partly latitudinal, also suggest that although Lepidoptera arthropods may be the most widely targeted pest group, insectivorous birds' dietary choices and trophic fidelity to the range of arthropod food is largely flexible and may be subjected to a wider array of opportunistic local factors. These may include the landscape structural context, regional climate, and finer-scale anthropogenic impacts such as farmers' selective and discretionary application of chemical inputs in crop production or weed control (Kross et al., 2019; Kross et al., 2020). Such diet flexibility is particularly important for the survival of forest-associated generalist insectivores, in the face of loss, fragmentation, or changes in quality of their forest habitat resulting from anthropogenic impacts (Şekercioğlu, 2002). Furthermore, certain cases are considered in which some insectivorous birds opportunistically predate on beneficial arthropod pest' natural enemies (Grass et al., 2017; Pejchar et al., 2018) resulting in trophic cascades that may undermine overall pest suppression, especially where the dominant arthropod pests occur at outbreak abundance scales. In such cases, for instance, farmland avian predators may consume more Hemiptera than Lepidoptera prey during a leaf aphid outbreak period (Gonzalez et al., 2017; Riedl et al., 2018). Similarly, they may predate on more Coleoptera prey on croplands next to a recently clear-cut forest stands due to local upsurges in such prey items dispersing from the disturbed habitat (Yard et al., 2002; Exnerová et al., 2003; Bakx et al., 2020).

Despite these individual trends and patterns in insectivorous bird' predation, family-level bird identity, habitat association, or their diet-guild specialization did not seem to be important predictors of bird communities' overall dietary pest-food proportions. In other words, whether Muscicapidae or Laniidae, Cisticollidae or Ploceidae, farmland- or forest-associated, did not matter in generally explaining how much pest food from the various options was likely to be consumed by birds overall. This means, in particular, that there was no remarkable dominance of any one bird family in the rate of consumption of arthropod pests which is, however, is contrary to findings by Maas et al. (2015) in Indonesian cacao agroforestry systems. Likewise, it did not seem to matter whether birds were obligate insectivores or omnivorous; hence each family had similar contributory potential to the total proportions of food incorporated from across the range of options into diets of birds overall. However, identities of pest-prey items were important in predicting overall pest-food proportions in birds' diets. Therefore, just as was as seen earlier on, muscicapid, hirundinid or ploceid birds would more likely consume Lepidoptera pests whereas,

estrildids, malaconotid and ploceid birds consume Coleoptera and Hemiptera pest food resources without regard to whether these were closely linked to farmland or forest. On the one hand, this implies that birds of either zone are likely to disperse across the boundaries in order to secure preferred pest-prey types in the right proportions. Furthermore, birds of particular habitat zones are unlikely to discriminate against prey associated with the opposite habitat, when they opportunistically encounter these, as long as such prey are of a suitable taxa. The implication here is that pest-prey items were generally available at abundance scales that justified choice discretion among most bird families. This could be presumably based on prey suitability or quality, such that Lepidoptera prey were the most preferred overall. This presumption is founded on the tendency of predatory passerine birds to exhibit diet selectivity when prey resources are non-limiting (Jones et al., 2005b). Furthermore, in their study, Razeng and Watson (2014) showed that such selectivity can be significantly driven by birds' perception of the relative nutritional value of prey items. This fact might partly explain the global decline in populations of insectivorous birds in small forest fragments whose sizes can no longer support many such consumers due to reducing biomass and variety of high quality insect prey (Şekercioğlu, 2002; Peter et al., 2015).

4.3 Plant basal carbon in bird tissues

Contrary to expectation, forest-associated birds exhibited higher proportions of tissue-assimilated maize-derived basal diet carbon when compared to the case for farmland-associated birds. This means that substantial quantities of arthropods consumed by forest-associated birds, in turn, derived their basal plant food from maize. This is confirmed by the aforementioned results that Lepidoptera had the strongest trophic connection to maize, which is a farmland crop plant. It is also supported by the fact that Lepidoptera were the most highly consumed pest by birds in general, including by forest birds. In contrast, the bulk of the forest-tree diet carbon was detected in farmland-associated birds, showing that these birds derive considerable basal plant carbon through arthropod herbivores of forest trees, especially Coleoptera and Hemiptera, most of which are also known to be abundant in forests and wooded habitats (Bos et al., 2007; Bellamy et al., 2018). These patterns suggest that the forest-associated birds observed in this study do play more significant roles than do their farmland counterparts in contributing to the removal of arthropod pests within the open farmland. This may be accounted for in two possible ways. First, Lepidoptera pestprey, which was mainly consumed by forest birds, may be so significant in the diets of most forest birds that such birds are prepared to disperse into the farmland to track them down. The second possibility is that some Lepidoptera herbivores within the farmland also disperse into the forest for various purposes, including breeding, refuge, or roosting, during which they are consumed by many forest birds.

Similarly, forest-tree basal carbon occurring in the highest proportions in farmland-associated birds points to the possibility that many farmland birds also cross habitat boundaries to forage on

forest-tree arthropod herbivores. This may particularly owe to the importance of forest and other wooded habitats adjacent to farms as roosting, nesting, breeding, or refuge habitats for such birds (Berg and Part, 1994). Indeed just as in the case for forest birds, while in the forest, such farmland birds may also predate on arthropod pestprey that primarily consume forest trees and thus acquire forestbased plant carbon, and in the process, contribute to the removal of forest-tree pests. Such dynamic dispersal opportunities and trophic interaction options may be pivotal in facilitating avian and arthropod functional roles in promoting ecological connectivity and energy exchange between the agricultural and forest ecosystems (Alvarez-Alvarez et al., 2022). Furthermore, maize's highest proportion in birds' tissues at both the overall and the family levels underscores its importance over legume plants in the overall sustenance of all insectivorous birds across the farm-forest edge zone in this broader agrarian landscape.

The high proportions of maize carbon assimilated by muscicapid flycatchers, cisticollid warblers, ploceid weavers, and hirundinid birds primarily confirms that these four families mainly consumed maize arthropod pests, and this has a number of explanations. For instance, majority of these bird families frequently occur in the farmland or open habitat and therefore have a higher likelihood of trophic interaction with maize arthropod pests. Additionally, ploceid weavers are omnivorous feeders and thus may obtain a part of their maize diet carbon through direct granivory in addition to preying on maize arthropod pests such as Lepidoptera. By comparison, in addition to muscicapids and ploceids, the highest proportions of basal plant carbon from forest trees and farmland legumes appeared in estrildid and malaconotid bird families, which indicate that besides their dietary intake of Lepidoptera, these bird families significantly depend on pests of legumes and forest trees. Therefore, estrildid and malconitid birds could be uniquely suited as important contributory agents of arthropod pest biocontrol in legume crops and in forest trees. Within the context of birds' roles in interhabitat connectivity, these patterns clearly demonstrate the pathway for the diet carbon exchange between the farmland and the forest habitats by trophic interactions. Thus muscicapid flycatchers and malaconotids are the forest-associated bird families that are most likely responsible for the high levels of farmland diet carbon in the overall forest birds. On the other hand, estrildids, ploceids, and hirundinids may be responsible for the high forest diet carbon in farmland birds' tissues. Therefore, in functional terms, muscicapids, ploceids, estrildids, and hirundinids, either by being the leading consumers of the most important pest-prey or by virtue of their well-known wide mobility within the farm-forest ecotone, are potentially the best avian agents in contributing to arthropod pest biocontrol across the two habitats. They may also be efficient ecological connectors of fragmented forest patches within the forest-farmland habitat mosaic (Garbach et al., 2012; Alvarez-Alvarez, et al., 2022). Ploceids and estrildids should be particularly instrumental for this role given their typically gregarious, highly-itinerant, and sporadic-dispersal foraging patterns (Yard et al., 2002). Kross et al. (2019) suggested that such connectivity may be enhanced even further thorough establishment of such seminatural

features as hedgerows, fallow-field patches, and other forms of woody vegetation within the farmland, which Lourenço et al. (2021) also showed to be important in supporting birds' provisioning of arthropod pest biocontrol services within vineyard orchards. Such features can additionally help in boosting the recovery of populations of beneficial parasitoid wasps following stochastic population crashes (Schindler et al., 2022).

Considering the dietary proportions of pest-prey against proportions of basal plant carbon assimilated in bird's tissues, some variations were evident. For instance, cisticollid warblers were not among the leading consumers of Lepidoptera pests which were the principal herbivores of maize, or of Hemiptera, the main pest of forest trees and legume plants. Yet, these warblers were among the groups which incorporated maize diet carbon in the highest proportions. This implies that cisticollid warblers derived much of their maize diet carbon through arthropod pest-prey belonging to taxa beyond the scope of the three that were examined here, but which also constituted maize pests. As such, despite not being as trophically dependent on Lepidoptera pest-prey, these warblers do not necessary play a role subliminal to that of flycatchers, swallows, or weavers, in contributing to maize pest biocontrol. Similarly, although motacillids evidently exhibited considerable Lepidoptera carbon in their tissues, they showed relatively insignificant proportions of assimilated maize carbon, implying that these birds derived Lepidoptera pest-prey from plant sources extending beyond maize. However, these discrepancies may also arise from the intraguild predation processes (Sanders et al., 2011), which were, however, not focal to the present study. For instance, cisticollid warblers could also derive maize carbon through consuming arthropod natural enemies of maize pests such as ants and spiders (Sanders et al., 2011; Otieno et al., 2019a).

From the ordination analysis results, the more significant role of arthropod prey taxa and diet specialization over habitat association or family identity in accounting for birds' basal plant food signatures demonstrates that birds consumed pest food based on significant fidelity to diet guild orientation. It also shows therefore, that bird's associations to each habitat zone did not necessarily translate to them incorporating significantly more basal plant food from that zone. Further, the finding points to availability of prey resources being in such sufficient abundances as to allow birds of both guild classes to feed on a broad spectrum of prey options in accordance to their specific diet orientation without omnivores species having to switch to herbivory at any time, for instance. This is unsurprising, considering the theory of relatively higher variety of prey items within most tropical ecosystems as compared to those in temperate latitudes (Willig et al., 2003; Andrew and Hughes, 2005). As is widely known, in colder temperate regions, the arthropod prey resource supplies undergo significantly higher seasonal fluctuations, with significant declines in distribution and abundances during winters. This imposes dietary plasticity in avian omnivores in such regions to rely less on arthropods but more on plant-based food during this time (McKinnon et al., 2012; Garfinkel et al., 2020; Kross et al., 2020).

5 Conclusion

The overall high proportion of maize in diets of all three pest taxa reflects how important the crop is in the diets of these arthropod herbivores across the study area, but especially for Lepidoptera, while Hemiptera and Coleoptera pests were more liberal in their diet choices. Similarly, Lepidoptera pests were the favorite prey source for most insectivorous birds regardless of bird's diet specialization but especially forest-associated birds, while Hemiptera and Coleoptera pests were consumed nearly proportionately by both farmland- and forest-associated birds. Thus, with regards to predation, prey taxa identity and habitat association were highly important in explaining the overall distribution of pest food proportions in birds' diets. Furthermore, through the comparative tests of arthropod-bird and plant-bird trophic connections, the study showed that birds incorporated significantly more basal plant diet carbon from maize than from other plant sources, and that as the key drivers of these patterns, five avian families, namely, muscicapids, estrildids, ploceids, malaconotids, and hirundinids, are the most important agents of trophic connectivity of the two habitat zones. Thus, integrated pest management practices for maximizing birds' functional roles as contributors to top-down arthropod pest suppression should include measure to conserve such keystone interhabitat connector species. Accordingly, assessing the proportions of basal plant carbon assimilated in predators' tissues to complement results from direct-predation assessments offers a more explicit underpinning of the predators' role in driving the herbivory component of the prevailing food-web linkages. In particular, such a procedure is necessary for evaluating predators' relative importance in biocontrolling pest-prey that target the respective plants for herbivory. In small-scale farming ventures, such as the one predominant in much of East Africa, collaborating with forest-edge farmers to embrace agronomic practices that enhance landscape structural complexity such as through agroforestry and maintaining hedgerows or uncultivated margins would be especially important (Maas et al., 2015; Lindell and Dayer, 2022). For example, native trees on farms have been shown to be attractive to insedtivirous birds at high densities and variety since they serve as feeding habitats as well as stepping-stone elements facilitating birds' dispersal across the landscape (Douglas et al., 2014), thus boosting the trophic component of the broader functional ecological linkage across the farm-forest mosaic (Bos et al., 2007; Stewart et al., 2019; Kross et al., 2020; Alvarez-Alvarez et al., 2022; Otieno and Pryke, 2022).

6 Limitations of the study and further research

This study examined arthropod pests at the Order taxa levels to assess their trophic linkages to plant food sources and to their avian predators from a stable isotopic perspective. In our opinion, this approach offered the most optimal analytical resolution for a spatio-temporal mapping of the pest-prey trophic interaction in the landscape. However, through this strategy, the study did not capture roles that individual pest species or genera might play in contributing to the overall dietary proportions assimilated. Second, because of the single-event effort in mist-netting, a number of bird species might have been missed out within the study sites, and therefore for a more complete picture of the trophic linkages, future studies in the study area should consider increasing effort in the birdcapture exercise. Third, a more complete assessment of bird's relative importance in contributing to the removal of arthropod pest in each of the three plant groups examined here could be achieved by assessing a broader spectrum of herbivore taxa than the three that are treated here. Finally, more in-depth studies incorporating both stable isotope analyses and a sampling protocol for exclusion of either birds or pests, and plant damage rate assessment, would facilitate a precise dollar-value quantification of the pest control services provided by the birds.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

Ethics statement

The animal study was reviewed and approved by National Museums of Kenya and Kenya Wildlife Service.

Author contributions

NO: methodology, formal analysis, investigation, data curation, data analyses, writing—original draft, and project administration. JM: field investigation, bird systematics and identification, data collation, and draft manuscript approval. All authors contributed to the article and approved the submitted version.

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Conflicts of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

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