


Identification and evaluation of agroecosystem compartments as forage sources for free-ranging goats in smallholder farming systems of western Democratic Republic of Congo

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ABSTRACT

Free-ranging goats forage across multiple agroecosystem compartments, thereby contributing to nutrient redistribution. Despite their importance in nutrient flows, little is known about how these compartments support goat diets or about their spatial structure. This study addresses this gap by identifying, mapping, and characterising the agroecosystems exploited by free-ranging goats as forage sources and by assessing the contribution of each compartment to their daily diet. Twelve adult female goats from local herds were monitored during both the dry and rainy seasons for three consecutive days at four sites, using GPS collars combined with drone surveys and direct observations. Two observers were assigned to each goat: one, equipped with a stopwatch and tracking sheet, recorded feeding stations, forages consumed, and biting times; the other, using a Sony HDR-CX405 camera, documented biting behaviour.

Drone-based mapping revealed five distinct compartments of the foraging area: croplands (6.1 %–14.8 %), fallow lands (2.7 %–12.5 %), rangelands (36.4 %–69.7 %), residue supply areas (9.8 %–46.7 %), and hedgerows (0.1 %–5.4 %). The agroecosystem landscapes were highly fragmented (24–54 patches/ha) and exhibited SHDI values ranging from 0.94 to 1.19. Goats consumed 57 forage species, dominated by grasses (17 %) and legumes (9 %), with the remaining 74 % spanning 31 botanical families. The integration of GPS tracks with land-use maps and direct observations revealed that rangelands and agricultural lands were the primary contributors to free-ranging goats' daily dry matter intake and forage species diversity. The marked heterogeneity and floristic richness of these agroecosystem compartments underpin their importance as key forage sources supporting smallholder free-range goat farming in western Democratic Republic of Congo.

1. Introduction

Terrestrial ecosystems have been under increasing human pressure since the Green Revolution, largely owing to rising global food demand (Foley et al., 2011; Barnosky et al., 2012; Bourban, 2019). Subtropical savannahs are among the most intensively exploited ecosystems because of their multifunctional role in supporting food production (Boval et al., 2017; IPBES, 2018; Kuyah et al., 2021). However, they are also highly vulnerable to degradation resulting from unsustainable practices such as slash-and-burn agriculture with short fallow periods (Boval et al., 2017; Mishra et al., 2021), high-intensity fires used for rangeland management, uncontrolled wildfires (McNew et al., 2023), and inappropriate afforestation strategies (Veldman et al., 2015; Fernandes et al., 2016).

These practices compromise key ecosystem services—such as nutrient cycling, biodiversity, and food web stability—threatening long-term crop and livestock productivity and the livelihoods of local communities in sub-Saharan Africa (SSA), especially under the pressures of climate change.

Barnosky et al. (2012) emphasise the urgent need to address the root causes of anthropogenic global change and to strengthen biodiversity and ecosystem service management, noting that ecosystems often fail to recover once critical disturbances occur.

In SSA, smallholder farmers rely heavily on extensive agriculture, and livestock production is embedded within natural savannah ecosystems (Callo-Concha et al., 2013; Losch, 2016; Orina et al., 2024). There is a pressing need for research focused on the multifunctional roles of

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these ecosystems (Lemaire et al., 2014; Boval et al., 2017) because significant nutrient export through crop and livestock production leads to soil degradation and loss of fertility. However, free-ranging livestock return a fraction of nutrients through dung and urine while foraging, thereby playing a significant role in maintaining soil fertility (Augustine et al., 2003; Rufino et al., 2006; Grillot et al., 2018; Huruba et al., 2018), although the extent of this contribution remains poorly documented.

Agroecological farming practices that enhance interactions among soil, plants, and animals through efficient nutrient recycling are considered viable strategies for preserving ecosystem health and ensuring sustainable food production (Muramoto et al., 2000; Bonaudo et al., 2014; Altieri et Nicholls, 2020; Altieri et al., 2024; Audouin et al., 2024). Integrated crop–livestock systems (ICLS) exemplify such agroecological practices, supporting sustainable intensification within agroecosystems by improving nutrient recycling through the management of manure and crop residues. These practices reduce dependence on external inputs and enhance resilience to the impacts of climate change (Rufino et al., 2009; Bonaudo et al., 2014; Stark et al., 2016; Grillot et al., 2018, 2018; Steinmetz et al., 2021).

However, livestock management strategies vary across agroecological zones, shaped by differences in land-use allocation and pressure, feed resource availability, and territorial planning (Rufino et al., 2006). This results in diverse ICLS configurations and a range of potential strategies and designs. In the free-ranging systems common in SSA, livestock govern nutrient flows and act as key contributors to soil fertility in both savannah and cultivated lands (Rufino et al., 2006; Audouin et al., 2024). Animal diet composition directly influences the quality and quantity of nutrients returned to the soil via excreta (Rufino et al., 2006). Therefore, assessing forage potential in terms of species diversity, nutritional value, and biomass productivity is critical for developing livestock management strategies that promote the sustainable use of tropical savannah ecosystems (Huruba et al., 2018). Such assessments are particularly important for improving understanding of nutrient transfer from grazing areas to cropland (Audouin et al., 2024), especially in the context of ICLS.

Among livestock, goats are a common species in rural areas of SSA because of their adaptability to diverse and harsh environments, as well as their compatibility with extensive farming systems that require low feed and management inputs (Dossa et al., 2008, 2015; Gasigwa Sabimana et al., 2018; Sejian et al., 2021; Andre Mataveia et al., 2023).

In the Democratic Republic of Congo (DRC), goat farming relies almost entirely on natural forages within agroecosystem landscapes, with herders occasionally providing limited supplementation using crop residues (Gasigwa Sabimana et al., 2018; Wasso et al., 2019; Ndong et al., 2024). Despite this dependence, little is known about the structure of these agroecosystem landscapes and their contribution to the daily diets of free-ranging goats. Addressing this knowledge gap is essential for evaluating the role of free-ranging goats as potential nutrient drivers within an agroecosystem, in line with ICLS principles. We therefore hypothesise that under free-range farming conditions, goats exploit multiple agroecosystem compartments but these compartments contribute unequally to their daily diets.

Accordingly, this study aimed (1) to map and characterise the agroecosystems exploited by free-ranging goats, with particular attention to identifying the main compartments of the foraging area (CoFA); (2) to assess and compare the individual contribution of these CoFA to the goats' daily dry matter intake (DMI); (3) to record and identify the forage species consumed; and (3) to evaluate the quantity and quality of the daily goat diet in terms of DMI, forage species diversity, acid detergent fibre (ADF), neutral detergent fibre (NDF), crude protein (CP) content, and digestibility.

On one hand, these parameters indicate diet quality and act as proxies for nutrient recycling, providing insights into the quantity and quality of nutrients returned to the soil through goat excreta. Such knowledge is crucial for developing sustainable management strategies that enhance nutrient flows within agroecosystem compartments. On

the other hand, free-ranging goats often damage crops and provoke land-use conflicts between herders and farmers, as observed in rural areas of the DRC and across SSA. Mapping CoFA and identifying the forages consumed could therefore help inform alternative management strategies that regulate access to these CoFA, secure animal nutrition, and mitigate potential farmer–herder conflicts.

2. Materials and methods

2.1. Study area

Four sites were selected for this study: the Masamuna, Mosango, Ngeba, and Kikola villages, located in the western provinces of the DRC (Fig. 1). The Ngeba village (5°11' S, 15°12' E) and the Kikola village (5°06' S, 15°06' E) are situated in Kongo Central province, while the Masamuna (4°48' S, 17°37' E) and Mosango (4°50' S, 18°06' E) villages lie within Kwilu province. These study areas share a similar ecosystem pattern characterised by partly cropped savannahs interwoven with gallery forests that thrive along riverine and wetland habitats (Ndong et al., 2024).

According to the International Vegetation Classification, the biome of these study areas belongs to the West-Central African Mesic Woodland and Savannah division (Faber-Langendoen, 2020). These savannahs constitute an important source of fodder and serve as a vital resource for extensive goat rearing. Following Köppen's classification (Lohmann et al., 1993), the climate is categorised as Aw4. The regions experience a tropical–humid Sudanian climate with two distinct seasons. The dry season lasts for approximately 4 months, from mid-May to mid-September, in Kwilu province (Mosango and Masamuna villages). In Kongo Central province (Ngeba and Kikola villages), it can extend up to 5 months, until mid-October. The rainy season spans roughly 8 months. Mean annual temperatures range between 25 °C and 28 °C, with an average annual rainfall of approximately 1400 mm (Gasigwa Sabimana et al., 2018; Vangu et al., 2023).

2.2. Experimental design

In each village, three female adult goats (27 ± 1.9 kg) were selected from local herds. All had given birth at least once and were raised under typical local free-range farming conditions. Each goat was fitted with a GPS collar (Fastrax UP 501; Fastrax Ltd., Finland) to record its location while foraging across the agroecosystem from 8:00 AM to 5:30 PM each grazing day, over 3 consecutive days, during both the dry (July–August) and rainy seasons (March–April). The GPS recorded the animal's position every 5 min. The University of Liège ethics committee (No. 19–2116, Liège, Belgium) approved the use of tracking devices as appropriate from an animal experimentation perspective (Dumortier et al., 2021).

Additionally, two observers were assigned to monitor and supervise each goat during the three free-grazing days, for approximately 9 h per day (Zampaligré et al., 2013), from 8:00 AM to 5:30 PM, following a 3-day familiarisation period as recommended in previous studies (Agréil & Meuret, 2004; Feldt & Schlecht, 2016; Vandermeulen et al., 2018; Chebli et al., 2020). One observer was equipped with a stopwatch and a daily monitoring sheet to record the duration of biting behaviour at feeding stations, the name and part of the forage plants consumed, and the CoFA where the foraging occurred (e.g. cropland, fallow land, grassland, etc.) (Barroso et al., 2000; Zampaligré et al., 2013). The observer also noted the duration of resting periods.

The second observer simultaneously recorded direct visual observations of goat biting behaviour using a Sony HDR-CX405 2.29 Mpix camera with 60 × optical zoom. Consumed forages were documented under their vernacular or local names as ethnobotanical data (Ayantunde et al., 2008), and specimens were collected for identification in the herbarium of the Plant Ecology and Systematics Laboratory at the University of Kinshasa, Lemba, DRC (Kembelo et al., 2021;

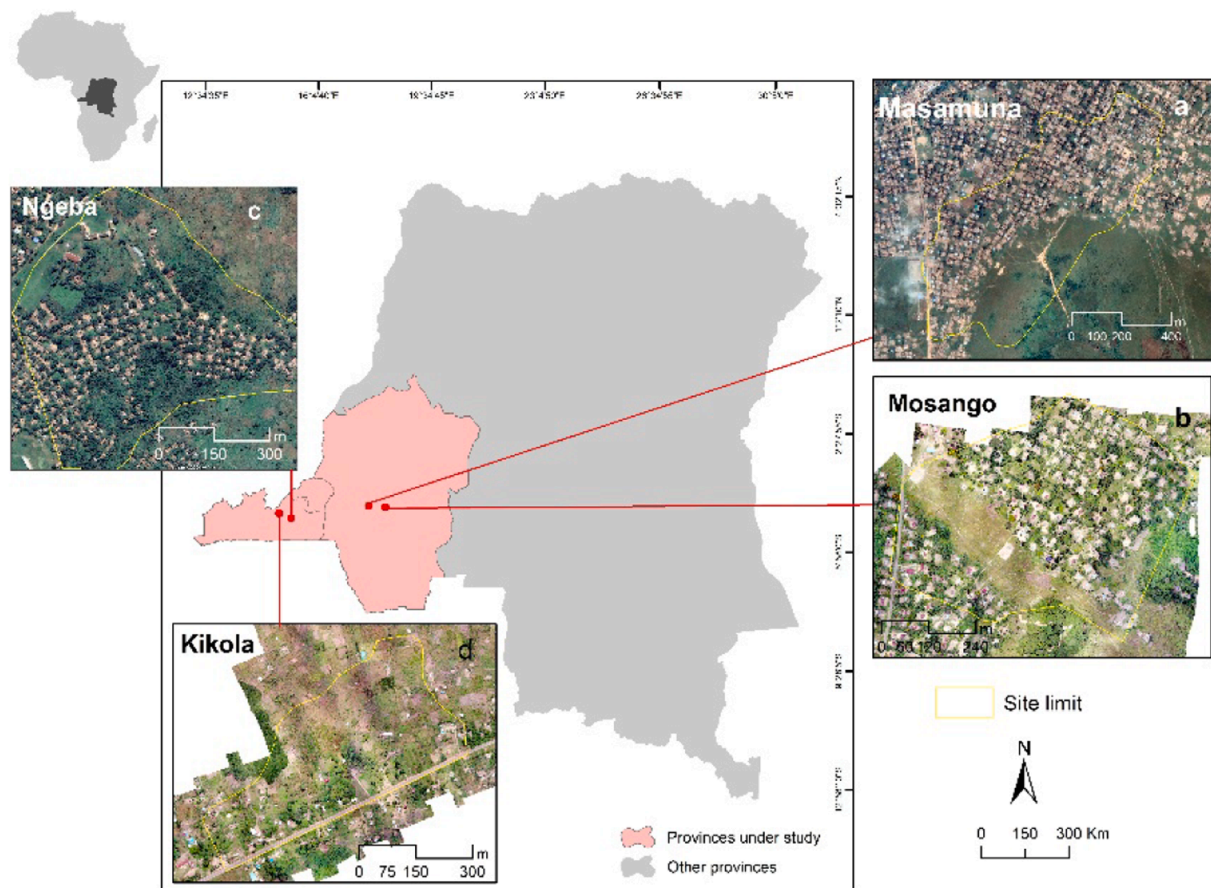


Fig. 1. Study area including the (a) Masamuna, (b) Mosango, (c) Ngeba, and (d) Kikola sites.

Ouachinou et al., 2017).

Finally, on the third day of data collection, an aerial photography survey was carried out at a flight altitude of 80 m using a DJI Mavic Pro 2 unmanned aerial system (UAS) equipped with a 1-inch 20-megapixel CMOS sensor and an adjustable focal aperture ranging from $f/2.8$ to $f/11$, along with a 77-degree field of view lens. The purpose of this aerial survey was to complement direct observations by producing high-resolution orthophotos that captured detailed imagery of the foraging areas across the agroecosystems exploited by the goats.

2.3. Data collection and processing

Aerial photographs were processed using Agisoft Metashape photogrammetry software to generate orthomosaics of the surveyed agroecosystems. The main steps included image alignment, generation of a dense point cloud, and extraction of the orthomosaic (Sherwood et al., 2018; Hatcher et al., 2020; Over et al., 2021). The resulting orthomosaics had a spatial resolution of 0.05 m based on the WGS 1984 UTM Zone 33S projected coordinate system. These were then used to map and characterise the agroecosystem compartments.

A land use and land cover (LULC) map of each agroecosystem was produced by digitising the UAS-derived orthomosaics. Automatic classification was not applied because the presence of non-essential objects (e.g. roofs with multiple colours) created radiometric classes that were difficult to distinguish, and some agroecosystem compartments showed strong spectral similarity (e.g. rangelands and fallow lands). Instead, objects were visually and manually labelled during polygon digitisation. A vector layer of polygons was created to represent objects or reference classes irrespective of pixel values (Dupuy et al., 2022; Jolivot & Dupuy, 2023). Houses, trees, and palm stands were digitised as distinct LULC

classes but were not considered as CoFA.

The CoFA classes were then characterised using the Patch–Matrix Model (Lausch et al., 2015), which describes the spatial structure of a landscape. Following Plexida et al. (2014), a limited set of six landscape metrics was applied to capture heterogeneity within the agroecosystems exploited by goats: patch density (the number of CoFA patches per hectare), CoFA index (the percentage of the agroecosystem area occupied by a CoFA class), mean patch size \pm standard deviation (the area of a CoFA class divided by its number of patches), minimum and maximum patch sizes per CoFA class, CoFA richness (the number of distinct CoFA classes), and Shannon Diversity Index (SHDI).

Goat GPS tracks were then superimposed onto the LULC maps. For each goat, the time spent on each CoFA was calculated from the georeferenced points automatically recorded at 5-minute intervals. Because of land-use dynamics, with frequent shifts between fallow and cropped field classes, both were grouped under the category “agricultural land”.

Daily DMI was estimated indirectly using the hand-plucking method, which mimics the bites taken by grazing goats (Bonnet et al., 2015; Chebli et al., 2020). This method considered three main parameters: feeding time (FT; duration of forage collection), bite mass (BM; average dry mass of forage per bite), and bite rate (BR; number of bites per unit time). DMI was then calculated using Eq. (1):

$$DMI(gd^{-1}) = BM \times BR \times FT \quad (1)$$

Each variable in the equation was determined experimentally. BM for each forage species was estimated using the hand-plucking simulation method, which involved manually collecting portions of forage to replicate goat grazing behaviour. For each recorded forage species, at least 100 hand-plucking repetitions were performed. The collected

samples were weighed fresh and then oven-dried at 60 °C until a constant dry weight was achieved. The average BM for each forage species was calculated by dividing the total dry mass by the number of hand-plucking replicates:

$$BM = \frac{\text{Mass of dried sample}}{\text{Number of hand - plucking simulations}} \quad (2)$$

The BR was determined from direct observation of the videos captured by the second observer. These videos were processed using VSDC professional video editing software (Lee & Kim, 2023). A minimum grazing sequence of 20 s on the forage plant consumed was randomly selected for each goat that had fed on it, resulting in at least 240 s of mixed footage per forage species. By replaying the footage in slow motion, the number of bites taken by the goats on the forage plant in question was accurately counted over the 240-second observation period.

The FT for collecting feed resources was defined as the period during which bites were taken on forage plants. This time excluded all other activities not related to forage collection, such as walking between locations within different CoFA, resting, or ruminating.

The CoFA within the agroecosystem were identified and mapped using UAS-generated orthomosaics. Geolocation data from the goats revealed the time spent on each CoFA, while direct observations confirmed the forage species consumed and recorded the time goats allocated to each species while grazing. Taking the CoFA into account, the data collected allowed the daily intake equation to be adapted as follows:

$$DMIT (gd^{-1}) = \sum_{i=1}^n DMICoFA_i \quad (3)$$

where

- DMIT (g d⁻¹) = total daily DMI (in grams per day)
- DMI CoFA_i = daily DMI within a single CoFA (in grams per day)

Within each CoFA, the daily intake was determined using Eq. (4):

$$DMICoFA (gd^{-1}) = \sum_{j=1}^m DMISp_j \quad (4)$$

where

- DMISp_j = daily intake of each forage species (grams per day) consumed within a single CoFA.

At this step, we referred to Eq. (1) to estimate DMI_{Sp_j}, expressed using Eq. (5) as follows:

$$DMISp_j = BTSp_j \times BFSp_j \times BMSp_j \quad (5)$$

where

- BTSp_j = total time spent by the animal collecting forage species *j* within the same CoFA during a grazing day (Eq. (6))
- BFSp_j = average bite frequency on forage plant *j* (Eq. (8))
- BMSp_j = mass of a bite on forage plant *j* (Eq. (9)).

The total time spent by the animal collecting forage species *j* within the same CoFA during a grazing day (BTSp_j) was calculated using Eq. (6):

$$BTSp_j = \left(\frac{BTvis Sp_j}{\sum_{j=1}^m BTvis Sp_j} \right) \times (T_{gps} CoFA_i - T_{rest} CoFA_i - T_{walk} CoFA_i) \quad (6)$$

where

- BTvis Sp_j = time recorded by the observer (visual observation) that the goat spent in CoFA_i collecting forage species Sp_j
- $\sum_{j=1}^m BTvis Sp_j$ = total time recorded by the observer (visual observation) spent in CoFA_i collecting all forage species consumed
- T_{gps} CoFA_i = time recorded by the goat's GPS collar within a single CoFA_i while foraging
- T_{rest} CoFA_i = non-grazing and/or resting time recorded by the observer within CoFA_i
- T_{walk} CoFA_i = time allocated to walking within CoFA_i (Eq. (7))

The walking time within a CoFA (T_{walk}CoFA_i) was calculated using Eq. (7):

$$T_{walk} CoFA_i = \frac{D_{gps}}{S} \quad (7)$$

where

- D_{gps} = distance walked within CoFA_i, recorded automatically by the goat's GPS collar
- S = walking speed of the goat when not collecting forage, estimated at 50 m min⁻¹ (Meuret et al., 1985)

The average bite frequency on forage plant *j* (BFSp_j) was then determined using Eq. (8):

$$BFSp_j = \frac{NBSp_j}{t} \quad (8)$$

where

- NBSp_j = number of bites counted on forage plant *j* by replaying the grazing footage in slow motion using VSDC Video Editor
- t = time spent making these bites

Finally, the mass of a bite on forage plant *j* (BMSp_j) was obtained using Eq. (9):

$$BMSp_j = \frac{\sum_{k=1}^l BMSimul}{l} \quad (9)$$

where

- l = number of hand-plucking repetitions performed on the consumed part of forage plant *j*
- $\sum_{k=1}^l BMSimul$ = total dry weight (g) of the *l* hand-plucking repetitions for forage plant *j*

2.4. Forage sample analyses

Hand-plucked forage samples, previously collected to estimate BM and oven-dried at 60 °C, were subsequently ground to pass through a 1-mm mesh sieve (Cyclotec 1093 Sample Mill; FOSS Electric, A/S, Hillerød, Denmark). The samples were then sent to the Laboratory of Precision Livestock and Nutrition, Gembloux Agro-BioTech (Liège University), for chemical analysis. NDF and ADF values were determined following the methodology outlined by Van Soest et al. (1991). The in vitro digestibility of organic matter was assessed using an enzymatic procedure involving pepsin and cellulase, according to the method described by Aufrere (1982). Total nitrogen content was determined using the Kjeldahl method (AOAC, 1990), and CP was calculated as N × 6.25.

2.5. Statistical analysis

Descriptive statistics were used to characterise the agroecosystems exploited by free-ranging goats. To assess the effects of site, season, and

their interaction on daily DMI, diet quality (CP, ADF, NDF, and digestibility), and the daily number of forage species consumed, a two-way analysis of variance (ANOVA) was performed, provided that the assumptions of normality and homogeneity of variances were met (using the Kolmogorov–Smirnov and Levene tests), with the *car* package implemented in R Core Team (2024) RStudio version 4.4.2. If these assumptions were violated, a two-way aligned rank transform (ART) ANOVA was conducted using the *ARTool* package as an alternative to the F-test (Kay & Wobbrock, 2016). Following either the ANOVA or ART ANOVA, Tukey’s post-hoc test was applied to identify significant

differences ($p < 0.05$) within site and season groups.

To evaluate and compare the contribution of different CoFA to daily DMI, grazing time, and consumed forage diversity, a three-way ART ANOVA was performed, considering season, site, and CoFA as fixed factors, including the interaction term season \times CoFA. Tukey’s post-hoc test was again used to detect significant differences ($p < 0.05$) between CoFA classes.

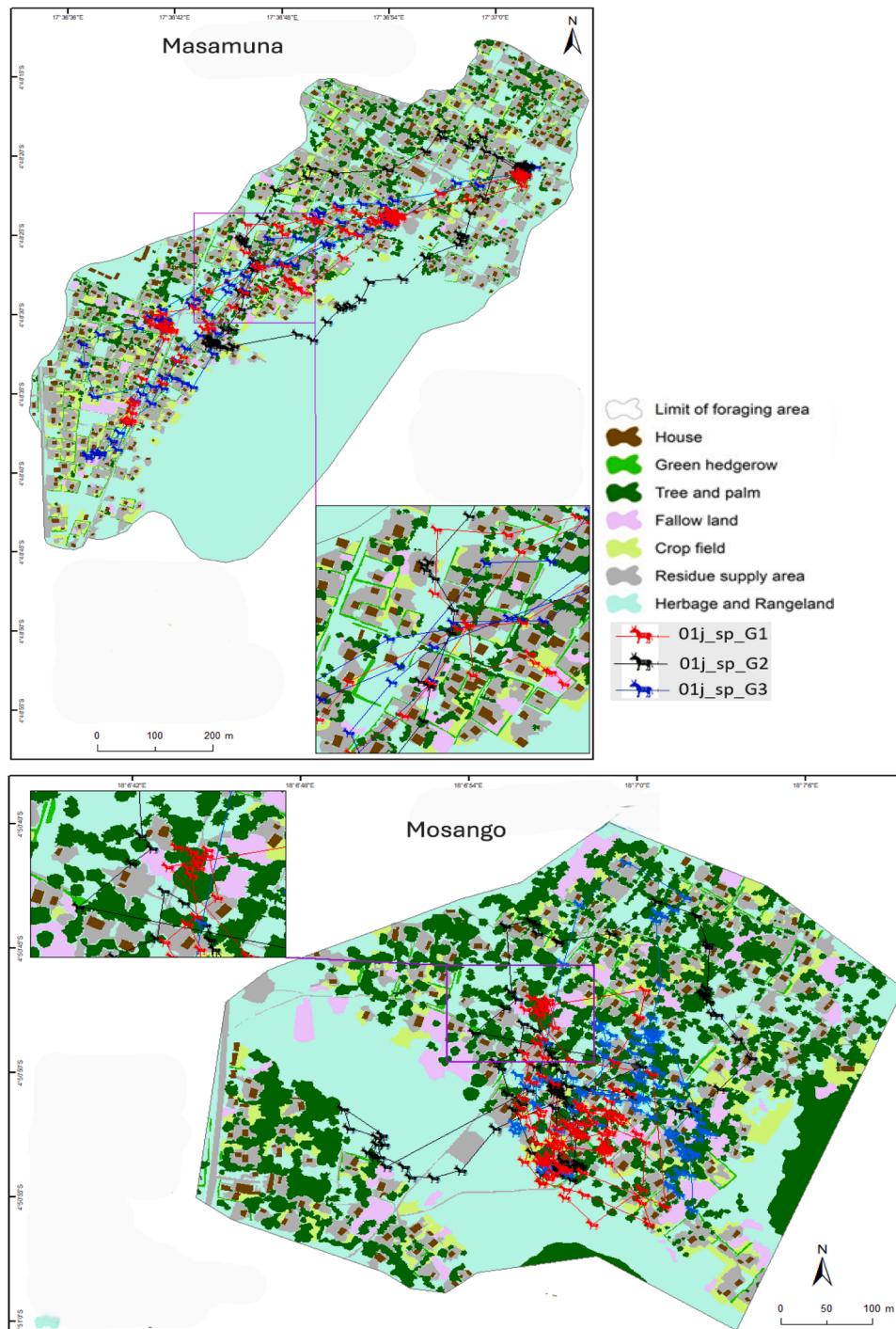


Fig. 2a. Land use and land cover maps illustrating the spatial configuration of compartment of foraging area (CoFA) within the agroecosystems exploited by free-ranging goats. The red, blue, and black lines exemplify georeferenced movement tracks recorded by GPS-collared goats (G1, G2, and G3) on day 1 of monitoring in rainy season at the Masamuna and Mosango sites.

3. Results

The spatial structure of the agroecosystems across the four study sites (Fig. 2a, Fig. 2b) revealed five distinct land-use classes exploited by free-ranging goats as CoFA. These compartments, described in Table 1, were croplands (comprising annual crops and vegetable gardens under active cultivation); fallow lands (post-cultivation areas), both of which constituted agricultural land; rangelands and natural vegetation (dominated by herbaceous species); green hedgerows (featuring woody or herbaceous perennial species used as plot borders); and residue supply areas (areas lacking vegetation cover, such as house halls, front or back yards, where goats consumed crop residues or household organic waste). Despite a moderate diversity of compartments, as indicated by SHDI values close to 1 (Table 1), the agroecosystems exploited by the goats were heterogeneous across all sites. Masamuna showed the highest fragmentation index (54 CoFA patches per hectare), whereas the other sites ranged between 24 and 31 patches per hectare (Fig. 2a).

Goats in free-range systems consumed a total of 57 forage species, including grasses (17 %, $n = 10$) and legumes (9 %, $n = 5$), which were the most frequently consumed botanical groups. The remaining 74 % ($n = 42$) of species were distributed across 31 distinct botanical families. Detailed information on life forms, plant parts consumed, seasonal consumption patterns, and nutritional profiles of these species is presented in Tables 2a and 2b for the Kongo Central and Kwilu provinces, respectively.

Goat diet composition varied across sites and seasons (Table 3). DMI, forage species diversity, and diet digestibility all showed significant effects of site and season ($p < 0.05$). By contrast, NDF content did not differ significantly between sites or seasons ($p > 0.05$), while ADF exhibited spatial but not seasonal variation ($p > 0.05$).

Goat diets were more species-diverse and contained higher DMI and CP content during the rainy season than during the dry season across all sites ($p < 0.05$). Across sites, Masamuna and Mosango showed the greatest species diversity in the goats' daily diets

($p < 0.05$), although CP content did not differ significantly among sites ($p > 0.05$).

The contribution of each CoFA to daily DMI, forage species richness,

and time spent foraging (Fig. 3) differed significantly among CoFA ($p < 0.01$). Within a given CoFA, species richness varied significantly between seasons ($p < 0.01$), whereas DMI and grazing time were more stable. Tukey's post-hoc test indicated that agricultural land contributed the highest proportion of daily DMI during the rainy season ($38 \% \pm 9 \%$), which declined in the dry season ($28 \% \pm 10 \%$), while the contribution from rangelands increased from $34 \% \pm 10 \%$ in the rainy season to $36 \% \pm 13 \%$ in the dry season. Residue supply areas became more important in the dry season ($31 \% \pm 12 \%$) than in the rainy season ($23 \% \pm 9 \%$), while green hedgerows contributed minimally in both seasons ($5 \% \pm 3 \%$ in the rainy season and $6 \% \pm 3 \%$ in the dry season).

The number of forage species consumed per CoFA was significantly higher in the rainy season, except in the residue supply areas and green hedgerows. The most species-rich compartments were agricultural land (8 ± 3 forage species in the rainy season and 5 ± 2 in the dry season) and rangelands (7 ± 2 and 5 ± 2 , respectively), compared with the residue supply areas (5 ± 2 in both seasons) and green hedgerows (2 ± 1 in both seasons). In terms of grazing time allocation, goats spent the most time in the rangelands ($38 \% \pm 14 \%$ of total time in the rainy season and $41 \% \pm 12 \%$ in the dry season) and agricultural lands ($38 \% \pm 10 \%$ in the rainy season and $29 \% \pm 7 \%$ in the dry season), with considerably less time spent in the other compartments.

4. Discussion

In free-range goat farming, little is known about how agroecosystem landscapes are structured and how they contribute to goats' daily diets. This study was conducted under typical traditional free-range goat farming conditions, characterised by small herd sizes (<15 goats) composed mainly of females and unweaned kids (Airs et al., 2023; Alexandre et al., 2012; Namonje-Kapembwa et al., 2022; Seleka et al., 2024; Tchouamo et al., 2005; Wasso et al., 2019). Adult males and growing young goats were not included because they are generally sold or consumed early. The limited sample size (12 female goats monitored for 3 days across 4 sites) constrains the generalisation of the findings to the wider goat population, given natural variability in grazing behaviour and nutritional requirements.

Table 1
Characterisation of CoFA and agroecosystem landscape heterogeneity metrics.

Site	CoFA Class	Frequency of patch	Patch area range (m ²) (Min-Max)	Average area (m ²) \pm Std	CoFA area (m ²)	Percentage (%)	CoFA Richness	Patch density (n/ha)	Shannon diversity index
Kikola	Cropland	63	3 - 4271	461.4 \pm 738.2	29,069	14.8	5	28	1.0705
	Fallow land	34	3 - 3222	720.6 \pm 922.1	24,501	12.5			
	Range land	372	5 - 86,048	330.6 \pm 4480	122,985	62.8			
	Residue supply area	78	4 - 9687	245.7 \pm 1095.9	19,164	9.8			
	Green hedgerow	9	4 - 68	25 \pm 19.2	225	0.1			
Masamuna	Cropland	347	4 - 519	79.5 \pm 79.7	27,581	8.9	5	54	1.1928
	Fallow land	58	14 - 1263	143.1 \pm 172.3	8301	2.7			
	Range land	372	4 - 17,486	106.7 \pm 660.2	113,185	36.4			
	Residue supply area	525	11 - 11,083	276.6 \pm 689.5	145,229	46.7			
	Green hedgerow	389	2 - 435	43.3 \pm 52.2	16,832	5.4			
Mosango	Cropland	152	4 - 1504	126.6 \pm 172	19,248	6.1	5	31	0.9427
	Fallow land	50	30 - 1758	363.6 \pm 374	18,178	5.7			
	Range land	470	4 - 98,988	469.8 \pm 4634.1	220,804	69.7			
	Residue supply area	212	10 - 3265	257.9 \pm 328.2	54,694	17.3			
	Green hedgerow	100	5 - 172	38.6 \pm 34.9	3857	1.2			
Ngeba	Cropland	185	5 - 9027	257.4 \pm 868.6	47,613	12.7	5	24	1.0027
	Fallow land	23	61 - 6034	861.3 \pm 1357.8	19,809	5.3			
	Range land	298	4 - 160,668	853.6 \pm 9328.4	254,381	67.8			
	Residue supply area	291	2 - 1631	166.2 \pm 181.6	48,375	12.9			
	Green hedgerow	117	4 - 234	43 \pm 47.3	5030	1.3			

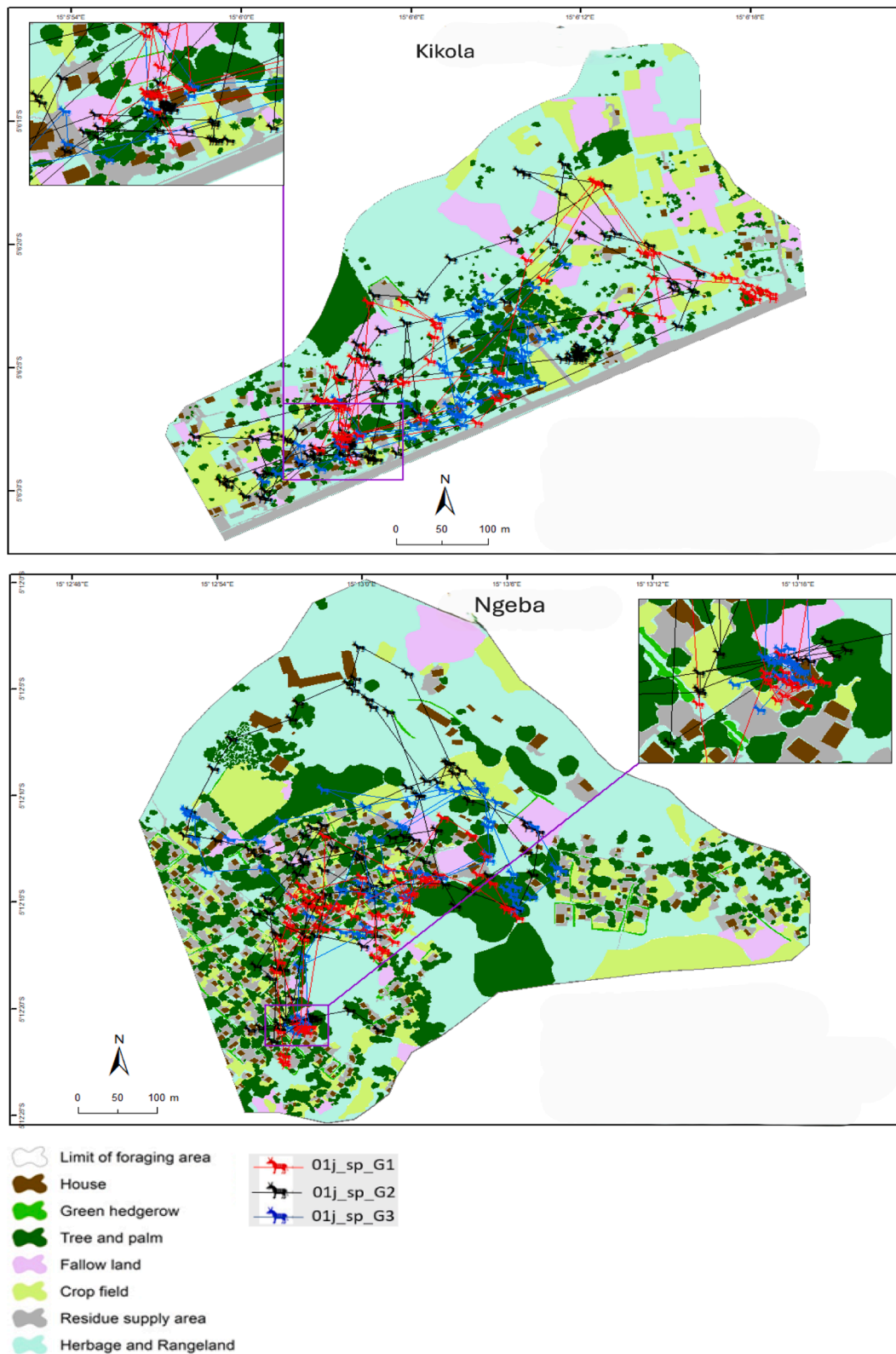


Fig. 2b. Land use and land cover maps illustrating the spatial configuration of compartment of foraging area (CoFA) within the agroecosystems exploited by free-ranging goats. The red, blue, and black lines exemplify georeferenced movement tracks recorded by GPS-collared goats (G1, G2, and G3) on day 1 of monitoring in rainy season at the Kikola and Ngeba sites.

Table 2a
Forage consumed by goats at Ngeba and Kikola sites, Kongo Central region, DRC.

Category ¹	Species ²	Part consumed ³	Botanical family ⁴	Live form ⁵	Season of consumption ⁶	CP ⁷ (% of DM)		NDF ⁸ (% of DM)		ADF ⁹ (% of DM)		Digestibility ¹⁰ (% of DM)	
						RS	DS	RS	DS	RS	DS	RS	DS
Grass and grass-like plant	<i>Cynodon dactylon</i> (L.) Pers.	leaves	Poaceae	Herbaceous	RS and DS	16.2	13.8	68.1	72.9	29.7	36.4	44.5	40.2
	<i>Cyperus esculentus</i> L.	leaves	Cyperaceae	Herbaceous	RS	25.3	–	47.4	–	26.6	–	75.6	–
	<i>Digitaria horizontalis</i> Willd.	leaves	Poaceae	Herbaceous	RS	9.9	–	45.6	–	30.5	–	47.5	–
	<i>Eleusine indica</i> (L.) Gaertn.	leaves	Poaceae	Herbaceous	RS and DS	19.8	14.4	61.8	68.8	28.3	32.1	57.5	49.9
	<i>Panicum maximum</i> Jacq.	leaves	Poaceae	Herbaceous	RS and DS	20.8	15.6	58.5	67.1	32.4	34.1	62.4	46.1
	<i>Paspalum notatum</i> Alain ex Fliiggé	leaves	Poaceae	Herbaceous	RS and DS	17.4	15.1	62.6	68.5	32.1	36.4	53.3	40.5
	<i>Centrosema virginianum</i> (L.) Benth.	leaves	Fabaceae	Herbaceous	RS	32.1	–	39.7	–	22.6	–	70.5	–
Legume	<i>Acacia auriculiformis</i> A. Cunn. ex Benth.	seed	Fabaceae	Tree	DS	–	27.2	–	34.3	–	25.3	–	68.2
	<i>Acacia auriculiformis</i> A. Cunn. ex Benth.	leaves	Fabaceae	Tree	RS and DS	16.9	16.3	56.5	57.9	48.7	50.1	39.4	36.5
	<i>Psophocarpus scandens</i> (Endl.) Verdc.	leaves	Fabaceae	Herbaceous	RS	39.9	–	38.7	–	17.8	–	81.2	–
	<i>Calopogonium mucunoides</i> Desv.	leaves	Fabaceae	Herbaceous	RS	30.0	–	43.2	–	26.7	–	58.2	–
	<i>Amaranthus blitum</i> L.	Leaves and stem	Amaranthaceae	Herbaceous	RS	32.2	–	25.8	–	13.9	–	69.7	–
	<i>Boerhavia diffusa</i> L.	leaves	Nyctaginaceae	Herbaceous	RS	30.9	–	32.9	–	18.1	–	70.5	–
	<i>Oncoba welwitschii</i> Oliv. Syn. <i>Caloncoba welwitschii</i> (Oliv.) Gilg	leaves	Salicaceae	Shrub	RS and DS	19.9	17.1	49.4	51.2	27.9	32.3	54.1	50.2
Others	<i>Carica papaya</i> L.	leaves	Caricaceae	Herbaceous	RS and DS	15.8	15.1	20.3	21.1	19.6	20.9	80.1	79.6
	<i>Chromolaena odorata</i> (L.) R.M. King & H. Rob.	leaves	Asteraceae	Herbaceous	RS and DS	32.0	21.4	33.6	39.9	17.1	26.9	82.3	63.1
	<i>Combretum racemosum</i> P. Beauv.	leaves	Combretaceae	Shrub	RS and DS	13.9	13.1	40.2	38.5	27.4	26.8	56.7	57.1
	<i>Commelina diffusa</i> Burm. f.	Leaves and stem	Commelinaceae	Herbaceous	RS and DS	27.5	25.1	37.5	39.1	22.9	23.4	80.2	79.4
	<i>Erigeron sumatrensis</i> Retz. Syn. <i>Coryza sumatrensis</i> (Retz.) E. Walker	leaves	Asteraceae	Herbaceous	RS and DS	20.3	16.6	32.7	35.1	28.2	29.7	53.4	51.4
	<i>Costus phyllocephalus</i> K. Schum.	leaves	Costaceae	Herbaceous	RS and DS	21.1	18.9	51.9	55.5	27.2	29.3	43.8	40.2
	<i>Croton hirtus</i> L'Hér.	Leaves and stem	Euphorbiaceae	Herbaceous	RS and DS	26.9	21.9	32.9	36.8	22.8	23.5	75.6	73.0
	<i>Cyathula prostrata</i> (L.) Blume	Leaves and stem	Amaranthaceae	Herbaceous	RS and DS	24.4	24.7	37.7	36.5	15.7	18.9	62.7	54.8
	<i>Dacryodes edulis</i> (G. Don) H.J. Lam	leaves	Bursaceae	Tree	RS and DS	11.3	9.8	43.7	46.5	35.8	37.1	40.5	39.6
	<i>Elaeis guineensis</i> Jacq.	leaves	Arecaceae	Herbaceous	RS and DS	19.8	17.9	59.4	61.9	33.3	35.7	52.5	49.3
	<i>Euphorbia hirta</i> L.	Leaves and stem	Euphorbiaceae	Herbaceous	RS	14.2	–	19.9	–	17.0	–	33	–
	<i>Ficus bubu</i> Warb.	leaves	Moraceae	Shrub	RS and DS	17.8	14.5	37.4	39.1	24.5	27.4	64.2	59.1
	<i>Gymnanthemum coloratum</i> (Willd.) H. Rob. & B. Kahn	leaves	Asteraceae	Shrub	DS	–	30.1	–	26.5	–	21.2	–	70.7
	<i>Hymenocardia ulmoides</i> Oliv.	leaves and twig	Phyllanthaceae	Shrub	RS and DS	29.9	25.7	48.6	50.5	31.6	35.3	68.7	65.2
	<i>Phragmanthera usuiensis</i> (Oliv.) M.G. Gilbert subsp. <i>Usuiensis</i> Syn. <i>Loranthus albizziae</i> De Wild.	leaves	Loranthaceae	Shrub	RS and DS	11.6	11.9	38.2	37.8	31.6	34.9	52.5	51.3
	<i>Mangifera indica</i> L.	leaves	Anacardiaceae	Tree	RS and DS	9.3	10	43.7	45.0	33.5	32.7	54.5	54.1
	<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	tuber	Euphorbiaceae	Shrub	RS and DS	1.1	0.9	1.1	1.1	0.7	0.6	99.1	98.9
	<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	leaves	Euphorbiaceae	Shrub	RS and DS	30.5	28.9	33.5	34.1	17.3	18.1	76.7	77.1
	<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	tuber peels	Euphorbiaceae	Shrub	RS and DS	5.8	6.1	20.8	21.6	17.2	17.5	73.6	74.2
	<i>Megaphrynium macrostachyum</i> (Benth.) Milne-Redh.	leaves	Marantaceae	Herbaceous	RS and DS	14.7	16.6	66.4	63.8	37.9	36.6	32.3	33.8

(continued on next page)

Table 2a (continued)

Category ¹	Species ²	Part consumed ³	Botanical family ⁴	Live form ⁵	Season of consumption ⁶	CP ⁷ (% of DM)		NDF ⁸ (% of DM)		ADF ⁹ (% of DM)		Digestibility ¹⁰ (% of DM)	
						RS	DS	RS	DS	RS	DS	RS	DS
	<i>Morinda morindoides</i> (Baker) Milne-Redh.	leaves	Rubiaceae	Shrub	RS and DS	24.4	24.9	49.2	50.5	40.6	39.9	56.1	55.4
	<i>Musa acuminata</i> Colla	leaves	Musaceae	Herbaceous	RS and DS	13.6	14.1	47.5	50.1	27.1	26.4	45.1	45.5
	<i>Passiflora edulis</i> Sims	Leaves and stem	Passifloraceae	Herbaceous	RS and DS	35.3	28.9	22.5	27.4	17.9	18.7	91.3	85.9
	<i>Persea americana</i> Mill.	leaves	Lauraceae	Tree	RS and DS	12.1	11.8	44.3	45.6	35.3	33.9	36.5	38.3
	<i>Psidium guajava</i> L.	leaves	Myrtaceae	Tree	RS and DS	13.4	12.9	44.8	46.3	34.3	35.1	37.2	35.6
	<i>Rhabdophyllum arnoldianum</i> (De Wild. & T. Durand) Tiegh.	leaves	Ochnaceae	Shrub	RS and DS	9.1	9.8	44.3	45.1	28.3	29.6	40.1	39.7
	<i>Sida acuta</i> Burm. f.	Leaves and stem	Malvaceae	Herbaceous	RS and DS	29.2	22.3	36.1	41.7	19.2	23.3	87.2	79.1
	<i>Smilax anceps</i> Willd.	Leaves	Smilacaceae	Herbaceous	RS and DS	12.3	13.9	52.5	50.1	37.3	36.5	38.7	41.5
	<i>Urena lobata</i> L.	Leaves and stem	Malvaceae	Herbaceous	RS and DS	19.4	18.8	44.9	45.7	29.9	31.1	50.5	49.7
	<i>Tithonia diversifolia</i> (Hemsl.) A. Gray	Leaves	Asteraceae	Shrub	DS	–	22.9	–	33.4	–	27.2	–	58.6

¹ Category denotes the primary taxonomic classification of the consumed forage species within plant community hierarchies.

² Species specifies the binomial nomenclature (genus and species) of the taxonomically identified forage consumed.

³ Plant part consumed indicates the specific anatomical structure of the forage ingested.

⁴ Botanical family classifies the consumed forage species according to its taxonomic family.

⁵ Live form describes the physiognomy of the consumed forage.

⁶ Season of consumption is the period of forage utilisation, defined as dry season (DS), rainy season (RS), or both (RS and DS).

⁷ CP is the crude protein content expressed as grams per 100 g of dry matter (g/100 g DM).

⁸ NDF (neutral detergent fibre) measures hemicellulose, cellulose, and lignin content, reported as grams per 100 g of dry matter (g/100 g DM).

⁹ ADF (acid detergent fibre): measures cellulose and lignin content reported as grams per 100 g of dry matter (g/100 g DM).

¹⁰ Digestibility represents the proportion of dry matter digested (DMD) (g/100 g DM ingested).

Despite these limitations, the combined use of GPS tracking, drone-based mapping, and direct observation provided valuable insights into smallholder free-range goat farming. This integrated approach linked goat movement patterns to the structural heterogeneity of agroecosystems, providing more accurate insights into compartment use and forage availability than traditional monitoring methods.

Agricultural lands (cropped fields and fallows) and rangelands emerged as the main compartments of the agroecosystem exploited by goats (Fig. 3). Their higher forage diversity and richness likely attracted goats, reflecting the animals' selective foraging behaviour and strategies to meet daily nutritional needs. Residue supply areas and hedgerows contributed less overall but still served as supplementary forage sources. The UAS-based maps and GPS tracks align with earlier findings from the DRC (Wasso et al., 2019; Ndong et al., 2024), which highlighted the importance of rangelands and crop residues as feed resources for extensive goat farming systems.

The drone-based maps revealed strong structural heterogeneity within the agroecosystems, with 24–54 CoFA patches per hectare, indicating high landscape fragmentation (Table 1) driven by anthropogenic activities (Molinario et al., 2017). Land use consisted of small crop plots, scattered farm structures, and livestock shelters interspersed with natural savannah. This fragmentation explains the landscape diversity indices (0.94–1.19) and species richness observed, with goats consuming 57 different forage species (Tables 2a, 2b). The plant diversity included both cultivated species (in croplands and hedgerows) and a predominance of spontaneous grasses and legumes (in rangelands and fallows), together contributing to a diverse forage base. The Aw4 Köppen climate subtype, characterised by an 8- to 9-month rainy season and high temperatures (Lohmann et al., 1993), favours and explains this floristic richness. Similar associations between rainfall, temperature, plant diversity, and biomass productivity have been reported in Venezuelan tropical ecosystems (Marín et al., 2001).

Savannah-type rangelands dominated the agroecosystem landscape, covering 36%–70% of the total area (Table 1). Their extensive coverage reflects the low population density in rural areas of the DRC, driven largely by rural–urban migration (Loola, 2025), rather than by

deliberate land-use management strategies. Such rangeland availability underpins extensive livestock farming systems, as shown in earlier studies (Augustine et al., 2003; Zampaligré et al., 2013; Boval et al., 2017; Ouédraogo et al., 2021).

GPS tracking data also confirmed the functional importance of rangelands and croplands, where goats spent most of their time and obtained the highest proportion of their daily DMI. These CoFA likely play a significant role in nutrient recycling, with nutrients returned directly via goat excreta during foraging or indirectly through manure collected from night corrals and later applied to croplands (Audouin et al., 2024). Although nutrient fluxes within these compartments were not directly quantified, this assumption of recycling is realistic. Schlecht et al. (2006) demonstrated that excreta deposition by free-ranging herbivores is proportional to the time spent in specific areas. Thus, prolonged goat presence in agricultural land, rangelands, and night corrals likely contributes to nutrient transfer through faeces and urine, supporting ICLS by enhancing soil fertility and reducing dependence on external inputs.

Green hedgerows, though marginal contributors to DMI, showed potential as valuable supplementary forage sources. Their contribution could be enhanced by enrichment with high-fodder-value perennials such as *Leucaena*, *Albizia*, *Gliricidia*, *Acacia* trees, and vetiver or elephant grasses. These species, known for their high productivity and year-round palatability, could increase both forage diversity and the DMI contribution from hedgerows. This finding aligns with Zampaligré et al. (2013) and Ouédraogo et al. (2021), who emphasised the agroecological importance of woody perennials in alleviating dry-season forage shortages. A greater availability of woody forages may also help improve the relatively low daily DMI observed in this study (20–25 g DM kg⁻¹ live weight [LW] across seasons) (Table 3). These values are close to the minimum requirement (19 g kg⁻¹ LW) but remain below the average (30 g kg⁻¹ LW) and maximum (41 g kg⁻¹ LW) reported under similar subtropical conditions in Brazil (Almeida et al., 2019). Increasing DMI through hedgerow forage supply could also allow for longer goat confinement, facilitating manure collection and enhancing nutrient recycling into croplands. Moreover, hedgerows restrict goat access to croplands, thereby helping to reduce crop damage and mitigate

Table 2b
Forage consumed by goats at Masamuna and Mosango sites, Kwilu region, DRC.

Category ¹	Species ²	Part consumed ³	Botanical family ⁴	Live form ⁵	Season of consumption ⁶	CP ⁷ (% of DM)		NDF ⁸ (% of DM)		ADF ⁹ (% of DM)		Digestibility ¹⁰ (% of DM)	
						RS	DS	RS	DS	RS	DS	RS	DS
Grass and grass-like plant	<i>Anthephora cristata</i> (Döll) Hack. ex De Wild. & T. Durand	leaves	Poaceae	Herbaceous	RS	19.9	–	45.5	–	26.6	–	65.2	–
	<i>Cynodon dactylon</i> (L.) Pers.	leaves	Poaceae	Herbaceous	RS and DS	16.7	8.1	63.9	77.2	30.8	35.5	49.1	29.3
	<i>Cyperus esculentus</i> L.	leaves	Cyperaceae	Herbaceous	RS and DS	16.3	16.1	62.1	63.6	31.8	33.7	46.2	43.8
	<i>Digitaria horizontalis</i> Willd.	leaves	Poaceae	Herbaceous	RS	9.9	–	44.5	–	31.5	–	47.1	–
	<i>Eleusine indica</i> (L.) Gaertn.	leaves	Poaceae	Herbaceous	RS and DS	18.2	13.6	55.6	61.7	30.2	32.5	62.6	43.8
	<i>Hyparrhenia diplandra</i>	leaves	Poaceae	Herbaceous	RS and DS	8.1	15.0	66.7	57.1	38.4	25.9	33.5	46.7
	<i>Panicum maximum</i> Jacq.	leaves	Poaceae	Herbaceous	RS and DS	16.5	15.7	64.7	67.5	36.4	39.1	50.3	44.1
	<i>Paspalum notatum</i> Alain ex Flügge	leaves	Poaceae	Herbaceous	RS and DS	15.8	11.1	55.7	60.1	31.7	33.3	55.1	49.3
	<i>Pennisetum setaceum</i>	leaves	Poaceae	Herbaceous	RS	13.9	–	57.7	–	38.2	–	65.2	–
	<i>Phyllostachys viridiglaucescens</i>	leaves	Poaceae	Herbaceous	RS and DS	27.9	22.3	58.1	65.9	28.3	32.7	55.9	47.5
	<i>Calopogonium mucunoides</i> Desv.	leaves	Fabaceae	Herbaceous	RS	30.1	–	30.7	–	24.3	–	58.8	–
	<i>Psophocarpus scandens</i> (Endl.) Verdc.	leaves	Fabaceae	Herbaceous	RS	31.1	–	49.7	–	30.9	–	55.1	–
	<i>Zornia latifolia</i>	Leaves and stem	Fabaceae	Herbaceous	RS	25.3	–	32.1	–	26.9	–	72.1	–
Others	<i>Aframomum albobolaceum</i> (Ridl.) K. Schum.	leaves	Zingiberaceae	Herbaceous	RS and DS	19.8	17.1	58.7	62.1	32.3	32.9	30.4	28.2
	<i>Amaranthus blitum</i> L.	Leaves and stem	Amaranthaceae	Herbaceous	RS	32.2	–	25.8	–	13.9	–	69.6	–
	<i>Asystasia gangetica</i>	Leaves and stem	Acanthaceae	Herbaceous	RS and DS	19.9	14.9	15.4	36.0	12.1	30.1	51.6	27.8
	<i>Boerhavia diffusa</i> L.	leaves	Nyctaginaceae	Herbaceous	RS	23.4	–	27.1	–	17.2	–	70.0	–
	<i>Breynia disticha</i> J.R. Forst. & G. Forst.	leaves and twig	Phyllanthaceae	Shrub	RS and DS	24.7	23.2	31.7	33.3	18.6	20.4	77.9	70.1
	<i>Carica papaya</i> L.	leaves	Caricaceae	Herbaceous	RS and DS	21.0	20	24.3	25.7	19.9	20.6	74.2	69.7
	<i>Chromolaena odorata</i> (L.) R.M. King & H. Rob.	leaves	Asteraceae	Herbaceous	RS and DS	27.6	24.6	35.8	39.8	33.9	35.8	53.5	52.8
	<i>Commelina diffusa</i> Burm. f.	Leaves and stem	Commelinaceae	Herbaceous	RS and DS	22.3	19.1	31.6	35.8	23.7	24.5	62.3	55.1
	<i>Croton hirtus</i> L' Hér.	Leaves and stem	Euphorbiaceae	Herbaceous	RS and DS	19.8	13.4	30.8	37.9	12.6	30.1	54.1	27.3
	<i>Cyathula prostrata</i> (L.) Blume	Leaves and stem	Amaranthaceae	Herbaceous	RS and DS	19.7	18.8	32.6	30.6	16.2	17.1	47.4	46.7
	<i>Dacryodes edulis</i> (G. Don) H.J. Lam	leaves	Burseraceae	Tree	RS and DS	10.9	9.8	41.2	44.5	36.3	37.6	41.3	40.9
	<i>Dracaena fragrans</i> (L.) Ker Gawl.	leaves	Asparagaceae	Herbaceous	RS and DS	19.8	20.1	44.1	43.4	32.3	33.7	55.7	51.2
	<i>Elaeis guineensis</i> Jacq.	leaves	Arecaceae	Herbaceous	RS and DS	16.5	13.2	56.4	64.7	34.8	37.2	45.3	39.1
	<i>Euphorbia hirta</i> L.	Leaves and stem	Euphorbiaceae	Herbaceous	RS	13.6	–	17.1	–	14.4	–	32.9	–
	<i>Ficus bubu</i> Warb.	leaves	Moraceae	Shrub	RS and DS	18.2	15.1	49.9	56.4	40.1	43.5	54.1	46.3
	<i>Gymnanthemum coloratum</i> (Willd.) H. Rob. & B. Kahn	leaves	Asteraceae	Shrub	DS and DS	29.5	20.2	24.5	31.9	20.7	24.9	71.6	59.4
	<i>Hymenocardia acida</i> Tul.	leaves and twig	Phyllanthaceae	Shrub	RS and DS	21.8	22.9	70.1	64.5	64.8	55.3	21.1	26.3
	<i>Ipomoea batatas</i> (L.) Lam.	leaves and twig	Convolvulaceae	Herbaceous	RS and DS	25.2	20.1	24.2	30.1	21.5	23.6	74.0	68
	<i>Phragmanthera usuiensis</i> (Oliv.) M.G. Gilbert subsp. <i>Usuiensis</i> Syn.	leaves	Loranthaceae	Shrub	RS and DS	10.9	10.5	55.1	53.9	49.6	51.1	34.3	32.1
	<i>Loranthus albizziae</i> De Wild.												
	<i>Mangifera indica</i> L.	leaves	Anacardiaceae	Tree	RS and DS	11.3	10.1	42.1	42.9	30.9	40.2	57.1	54.6
	<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	tuber	Euphorbiaceae	Shrub	RS and DS	0.8	1	1.7	1.9	1.5	1.5	97.9	98.7
	<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	leaves	Euphorbiaceae	Shrub	RS and DS	32.2	29.4	34.3	37.7	28.1	31.2	68.9	68.4
<i>Manihot esculenta</i> Crantz Syn. <i>Manihot utilissima</i> Pohl	tuber peels	Euphorbiaceae	Shrub	RS and DS	9.5	8.7	21.7	20.9	18.2	19.1	51.4	47.1	

(continued on next page)

Table 2b (continued)

Category ¹	Species ²	Part consumed ³	Botanical family ⁴	Live form ⁵	Season of consumption ⁶	CP ⁷ (% of DM)		NDF ⁸ (% of DM)		ADF ⁹ (% of DM)		Digestibility ¹⁰ (% of DM)	
						RS	DS	RS	DS	RS	DS	RS	DS
	<i>Megaphrynium macrostachyum</i> (Benth.) Milne-Redh.	leaves	Marantaceae	Herbaceous	RS and DS	15.3	13.1	69.3	65.1	44.2	41.8	22.1	24.5
	<i>Musa acuminata</i> Colla	leaves	Musaceae	Herbaceous	RS and DS	16.6	17.3	61.3	60.7	34.4	33.8	31.5	30.2
	<i>Parinari capensis</i> Harv.	leaves	Chrysobalanaceae	Shrub	RS and DS	9.5	11.1	71.9	69.4	64.7	60.8	15.4	18.8
	<i>Passiflora edulis</i> Sims	Leaves and stem	Passifloraceae	Herbaceous	RS and DS	41.3	40.5	20.8	21.5	18.5	19.9	86.6	81.3
	<i>Persea americana</i> Mill.	leaves	Lauraceae	Tree	RS and DS	12.2	11.9	42.1	40.4	32.6	31.9	41.2	42.5
	<i>Rumex usambarensis</i>	Leaves and stem	Polygonaceae	Herbaceous	RS and DS	27.1	29.4	31.5	30.9	20.2	20.7	75.4	72.8
	<i>Sida acuta</i> Burm. f.	Leaves and stem	Malvaceae	Herbaceous	RS and DS	26.5	23.7	22.4	34.0	18.1	17.1	74.3	76.2
	<i>Spermacoce latifolia</i> Aubl.	Leaves	Rubiaceae	Herbaceous	RS	20.9	–	23.6	–	18.9	–	73.1	–
	<i>Urena lobata</i> L.	Leaves and stem	Malvaceae	Herbaceous	RS and DS	22.2	13.9	33.4	40.1	28.8	31.5	65.4	40.5
	<i>Talinum triangulare</i> (Jacq.) Willd.	Leaves and stem	Talinaceae	Herbaceous	RS and DS	16.4	14.1	21.9	35.4	13.3	17.7	52.4	45.1
	<i>Tithonia diversifolia</i> (Hemsl.) A. Gray	Leaves	Asteraceae	Shrub	DS	–	25.4	–	34.5	–	33.2	–	58.6

¹ Category denotes the primary taxonomic classification of the consumed forage species within plant community hierarchies.

² Species specifies the binomial nomenclature (genus and species) of the taxonomically identified forage consumed.

³ Plant part consumed indicates the specific anatomical structure of the forage ingested.

⁴ Botanical family classifies the consumed forage species according to its taxonomic family.

⁵ Live form describes the physiognomy of the consumed forage.

⁶ Season of consumption is the period of forage utilisation, defined as dry season (DS), rainy season (RS), or both (RS and DS).

⁷ CP is the crude protein content expressed as grams per 100 g of dry matter (g/100 g DM).

⁸ NDF (neutral detergent fibre) measures hemicellulose, cellulose, and lignin content, reported as grams per 100 g of dry matter (g/100 g DM).

⁹ ADF (acid detergent fibre): measures cellulose and lignin content reported as grams per 100 g of dry matter (g/100 g DM).

¹⁰ Digestibility represents the proportion of dry matter digested (DMD) (g/100 g DM ingested).

Table 3

Goat's daily diet quantity and quality across seasons and sites.

		Dry mater intake (g DMI.kg ⁻¹ LW ⁻¹)	Number species consumed	CP (% of DM)	ADF (% of DM)	NDF (% of DM)	Digestibility (% of DM)
Kikola	Rainy season daily diet	22.1 ± 2.6	12±3	20.1 ± 1.4	24.2 ± 1.9	42.1 ± 2.9	66.1 ± 2.5
	Dry season daily diet	20.4 ± 1.6	9 ± 1	17.8 ± 2.6	24.6 ± 1.5	43.3 ± 2.1	63.9 ± 3.1
Ngeba	Rainy season daily diet	23.9 ± 1.7	13±3	20.2 ± 1.6	22.8 ± 1.8	40.4 ± 3	66.1 ± 3.3
	Dry season daily diet	20.6 ± 2.3	13±1	18.0 ± 1.4	24.3 ± 1.9	42.4 ± 5.3	64.7 ± 3.2
Masamuna	Rainy season daily diet	25.2 ± 1.3	22±2	19.3 ± 1.4	27.1 ± 1.3	39.6 ± 2.6	56.8 ± 2.4
	Dry season daily diet	21.7 ± 1.7	15±2	16.9 ± 1.2	32.9 ± 2.4	46.5 ± 5.3	52.9 ± 3.8
Mosango	Rainy season daily diet	25.4 ± 1.5	22±3	18.9 ± 0.9	28.7 ± 1.4	42.9 ± 1.5	53.9 ± 2.4
	Dry season daily diet	21.1 ± 1.5	20±2	18.4 ± 1.3	28.9 ± 1.2	43.6 ± 2.5	55.3 ± 3.2
P-value	Site	0.0018	< 0.001	0.205	< 0.001	0.347	< 0.001
	Season	< 0.001	< 0.001	< 0.001	0.062	0.067	0.0016
	Site x Season	0.1899	< 0.001	0.245	< 0.001	0.005	0.4339

farmer–herder conflicts—an ongoing challenge in African agro-pastoral systems (Barrière & Barrière, 2002; Azalou et al., 2023).

Because forage nutritional value is a key determinant of diet quality and, consequently, of the nutrient composition of excreta (Rufino et al., 2006; Piñero-Vázquez et al., 2017; Wang et al., 2018), the goats' daily diet also varied according to the nutritional value of the forage species (Tables 2a, 2b), the phenological stage of the consumed plant parts, the season, and the site. For instance, *Manihot esculenta* showed pronounced contrasts in CP content between tubers (approximately 1 % CP), peelings (6.0 %–9.5 % CP), and leaves (approximately 30 % CP). Legumes consistently exhibited the highest CP content (25 %–40 % CP), attributable to symbiotic nitrogen fixation (Prell & Poole, 2006; Barbieri et al., 2023). As ruminants use nitrogen inefficiently, excess intake of protein-rich legumes leads to nitrogen-rich excreta. Both unassimilated dietary protein and nitrogen released by ruminal microbiota are largely excreted as urinary urea (Rotz et al., 2006; Nasiru et al., 2014).

Surprisingly, several non-leguminous species (*Passiflora* spp., *Amaranthus blitum*, *Chromolaena odorata*, *Hymenocardia ulmoides*,

Gymnanthemum coloratum, and *Boerhavia diffusa*) also exhibited high CP contents (≥30 %), likely due to the presence of non-protein nitrogen in immature leaves with low C/N ratios (Lu et al., 2005; Zeng & Chen, 2018; Baath et al., 2024). The goats' selective feeding behaviour (Lee and Kim, 2023; Dias-e-Silva & Abdalla Filho, 2020)) favoured tender, nutrient-rich foliage, though they also consumed fibrous, mature plants (NDF/ADF > 30 %, digestibility < 70 %). This mixed foraging strategy helps balance rumen microbiome function and nutrient intake, consistent with findings from other studies on ruminant feeding behaviour (Fedele et al., 1993; Lu et al., 2005; Zampaligré et al., 2013; Claps et al., 2020).

The high CP values observed in grasses (Tables 1a, 1b) were comparable to those reported for fertilised pastures at early growth stages (Hirata et al., 2016; Muntiferung et al., 2000; Matta et al., 2023; F. D. Fernandes et al., 2014; Paciullo et al., 2017; Pedreira et al., 2024; Sokupa et al., 2024), possibly reflecting soil fertility and post-fire regrowth effects (McNew et al., 2023), which promote young, protein-rich shoots often preferred by grazers (Augustine et al., 2003).

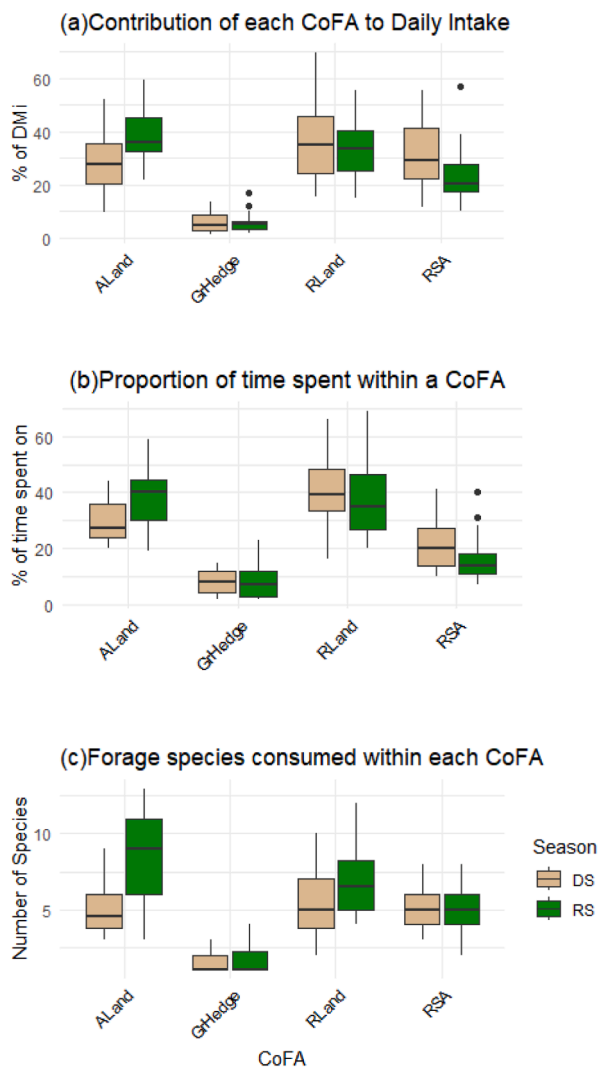


Fig. 3. The importance of each compartment of foraging area (ALand: agricultural land including crop and fallow land; GrHedge: living green hedgerow used as plot boundary or crop field protection; RLand: rangeland comprising savanna and/or natural herbage) in terms of their contribution to daily dry matter intake (a), proportion of time spent on each one (b) and number of different forage species consumed (c), during the rainy (RS) and dry (DS) season.

Conversely, lower CP values reported in other studies (Alsunaydi et al., 2024; Foster et al., 2011; Hare et al., 2015) may result from differences in climate, soil, or sampling protocols, as those studies often collected mature forages (>30 days post-emergence) with higher lignin content (Zeng & Chen, 2018; Baath et al., 2024).

Even under similar geo-climatic conditions, seasonal patterns in forage quality were evident. CP content and digestibility peaked during the rainy season, while NDF and ADF increased in the dry season due to rapid lignification under water stress and rising C/N ratios, as noted by Ephrem et al. (2015). These seasonal shifts negatively affected forage quality and, consequently, the goats' daily diets during the dry season.

Although CP contents were generally high (Tables 2a, 2b), they do not necessarily reflect protein bioavailability or digestibility because of the presence of antinutritional compounds such as alkaloids and tannins (Alavarse et al., 2022), which bind dietary proteins and inhibit enzymatic hydrolysis and subsequent nutrient assimilation (Ebrahimi et al., 2009; Sá et al., 2020). Moreover, non-protein nitrogen compounds—commonly found in immature or rapidly growing vegetative tissues—differ from those in other plant parts and may artificially inflate

CP estimates without increasing the availability of usable amino acids.

The proportion of plant parts consumed reflects individual differences in goats' selective grazing behaviour. Such selectivity directly shapes daily diet composition and, consequently, the nutrient profile of excreta (Rufino et al., 2006; Piñeiro-Vázquez et al., 2017; Wang et al., 2018). However, this study did not quantify the proportion of plant parts within total DMI. Combined with the small sample size (three goats per site) and short monitoring period (3-days across four sites), this limits the extent to which the findings can be generalised. In addition, the nutrient recycling potential within agroecosystems was inferred rather than directly measured through excreta collection and analysis, which introduces uncertainty due to individual variability in grazing behaviour and potential nutrient losses.

Despite these limitations, the integration of GPS tracking, direct observation, and drone mapping provided valuable insights into the spatial dynamics of free-ranging goats and their foraging patterns across agroecosystem compartments. Even with the availability of forage in rangelands, free-ranging goats obtained a substantial portion of their daily DMI from croplands. This dependence increases competition between livestock and humans for food resources and exacerbates farmer–herder conflicts.

Future research should expand the sample size to include other goat groups (e.g., adult males and young goats), extend the monitoring duration, and quantify both the proportions of plant parts consumed and the nutrient composition of excreta. Such work would offer a more comprehensive understanding of free-ranging goat diets and their potential contribution to nutrient recycling within ICLS.

5. Conclusions

This study examined the structure of agroecosystem landscapes and their contribution to the daily diets of free-ranging goats. The results show that, in smallholder farming systems of western DRC, free-ranging goats exploit multiple agroecosystem compartments and consume a diverse range of 57 forage species of varying nutritional quality, without relying on external feed inputs. Among the exploited compartments, agricultural lands and rangelands contributed most to daily DMI and forage diversity.

Although green hedgerows contributed little to the goats' daily diets, their role could be strengthened through enrichment with high-value forage species, such as woody and perennial grass fodders. Doing so could provide a dual benefit: reducing goat intrusion into croplands while supplying substantial forage biomass year-round.

The spatial heterogeneity of agroecosystem compartments, combined with their floristic diversity and variable nutritional quality, supports their role as key forage sources for free-ranging goats in western DRC.

These findings, however, should be interpreted with caution. The small sample size, limited representativeness of goat populations, and short observation period may have introduced uncertainties. Nonetheless, direct observations combined with GPS tracking and drone mapping have yielded valuable insights into smallholder free-range goat farming systems.

Future research should therefore involve larger and more diverse goat populations, extended monitoring periods, and a broader range of agroecosystem contexts. Such investigations would provide stronger evidence on the compartments exploited, forage diversity, and the nutritional quality of goats' daily diets. These parameters are critical for understanding free-ranging goat dynamics and their potential role in nutrient recycling, which is essential for designing appropriate ICLS. Advancing knowledge of nutrient fluxes within agroecosystem compartments will be key to developing sustainable ICLS tailored to the diverse farming systems of SSA.

Ethical statement

All animals were cared for in accordance with acceptable practices and experimental protocols approved by the University of Liège ethical committee (No. 19–2116, Liège, Belgium).

CRedit authorship contribution statement

Alain Ndong: Writing – original draft, Visualization, Validation, Software, Methodology, Investigation, Formal analysis, Data curation, Conceptualization. **Anthony Kikufi Batoba:** Writing – review & editing, Validation, Methodology, Investigation, Formal analysis. **Eric Lutete:** Writing – review & editing, Validation, Methodology, Formal analysis. **Bienvenu Kambashi Mutiaka:** Writing – review & editing, Validation, Investigation. **Charles-Henri Moulin:** Writing – review & editing, Visualization, Validation, Methodology, Data curation, Conceptualization. **Yves Beckers:** Writing – review & editing, Visualization, Validation. **Jérôme Bindelle:** Writing – review & editing, Visualization, Validation, Supervision, Resources, Project administration, Methodology, Data curation, Conceptualization.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Data availability

Data will be made available on request

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