

Predictive distribution modeling with enhanced remote sensing and multiple validation techniques to support mountain bongo antelope recovery

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Introduction

Restorative conservation measures are becoming increasingly needed as numerous species' populations decline (Griffith *et al.*, 1989; Ebenhard, 1995; Ceballos & Ehrlich, 2002). Successfully restoring a species requires understanding its fundamental ecological requirements (Sarrazin & Barbault, 1996), particularly the spatial distribution of habitat use, so that conservation measures can be targeted to appropriate areas. Spatial habitat use can be effectively estimated using predictive distribution models (PDM) provided datasets are adequately sized, represent important environmental gradients at relevant spatio-temporal scales, and are temporally and spatially transferable (Vaughan & Ormerod, 2003, 2005; Guisan & Thuiller, 2005). These requirements are often difficult to meet when studying rare and endangered species, whose low densities usually translate to small sample sizes (Guisan & Zimmermann, 2000; Rushton, Ormerod & Kerby, 2004). Rarity is often compounded by other factors (e.g. challenging terrain), which

Abstract

Understanding endangered species' spatial ecologies is fundamental to designing effective recovery strategies. Transferable predictive distribution models (PDMs), based on predictors describing the ranges and scales of relevant environmental gradients, can provide this understanding. Using such models for rare species such as the mountain bongo *Tragelaphus eurycerus isaaci*, an endangered antelope being restored within its endemic range in Kenyan montane forests, is difficult because the species' rarity and challenging terrain complicate data collection. To help overcome data limitations, we used advanced remote sensing (RS) and multiple validation techniques to improve bongo PDMs, which were developed using logistic regression and the information-theoretic approach. We derived predictors using RS, including a new technique for measuring micro-scale vegetation structure, and assessed predictive performance using bootstrapping and independent observations. Terrain ruggedness was the strongest habitat-use predictor, followed by soil moisture availability, distance from law enforcement outposts, vegetation structural complexity, and vegetation edge density. Prediction accuracy generally ranged between 73 and 89%, but terrain ruggedness limited model transferability. The more direct RS-based vegetation predictor improved model transferability. Bongo restoration efforts should focus on high probability areas delineated via a composite of all tested models. The techniques used – particularly RS – enhanced inference quality and the transferability of distribution models, and can be applied to other critical species and ecosystems.

may force a trade-off between maximizing observations of the focal species (e.g. Estes *et al.*, 2008) and the need to follow a comprehensive, fully stratified and multi-scaled sampling design (Vaughan & Ormerod, 2003). Furthermore, factors influencing a species' rarefaction can lead to atypical habitat use (e.g. Rubin *et al.*, 2002), complicating a generalized understanding of its ecology.

These research complications apply to the mountain bongo *Tragelaphus eurycerus isaaci* (hereafter 'bongo'), a critically endangered antelope sub-species (East, 1999) endemic to Kenyan montane forests (Fig. 1). An international conservation effort aims to restore the bongo within its former range, particularly in the Aberdare mountains and on Mount Kenya (MK). This initiative's success depends upon implementing appropriate recovery measures, which require an adequate understanding of the bongo's ecological requirements. These requirements are poorly understood (Estes *et al.*, 2008). Our objectives are to determine: (1) the primary factors influencing bongo habitat use; (2) where suitable bongo habitat is distributed; (3) where management

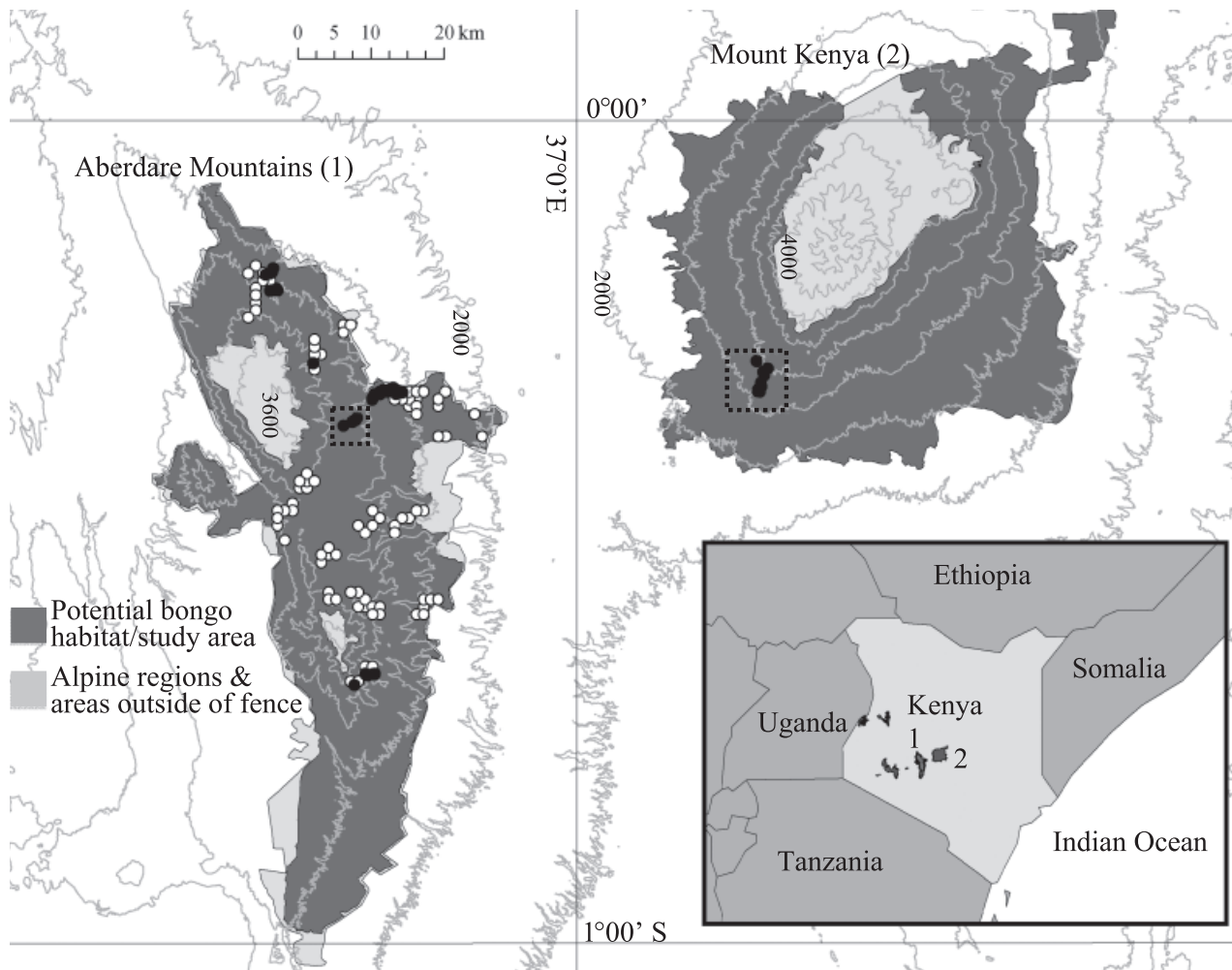


Figure 1 The Aberdares mountains (1), Mount Kenya (MK) (2), and the mountain bongo's *Tragelaphus eurycerus isaaci* historical distribution (dark gray patches in inset). Circles indicate bongo presence (black) and non-presence (white) plots. The gray lines denote elevation contours at 400 m intervals. The dashed boxes show the locations of bongo observations collected in follow-up surveys (MK: 7/2006; 2/2008; Aberdares: 8/2006).

actions could be most effective. We use logistic regression based on indirect bongo observations (e.g. tracks, dung, feeding sign) to develop predictive distributions models for the Aberdares and MK. We use remotely mapped habitat predictors representing spatial scales relevant to bongo (Estes *et al.*, 2008), including quantitative vegetation structure maps based on a new technique that is more accurate and more geographically transferable than approaches typically used in ecological models (Estes *et al.*, 2010). To minimize the high misidentification rates associated with interpreting the sign of bongo and other forest ungulates, we confirm bongo observations using mitochondrial DNA analysis of associated dung samples (Faria *et al.*, 2011; Bowkett *et al.*, 2009). We use independent observations from MK to assess the transferability of our predictive models, an important consideration because bongo populations in most of their habitat blocks (Fig. 1) are too small (if not absent) to provide adequate model calibration data.

The combination of improved remote habitat mapping and multiple validation techniques provides a template for enhancing modeling studies of other rare species.

Methods

Background

Until the 1960s, Kenya's bongo population likely exceeded 1000 animals, despite its narrow, restricted range, but since then the wild population has decreased to ~100 individuals confined primarily to the Aberdare mountains (Fig. 1, Stanley-Price, 1969; Kingdon, 1982; Reillo, 2002; Pettejohn, 2004). The precise cause of this decline is uncertain, but interactions between rapid human population growth, increased hunting pressure, habitat loss and epizootic events seem likely (Gathaara, 1999; Imbernon, 1999; Kock *et al.*,

1999; Lambrechts *et al.*, 2003; Prettejohn, 2008). The recovery project aims to (1) re-establish the MK (Fig. 1) population using captive bongo repatriated from North America; (2) rebuild the Aberdares' and other remaining wild herds through targeted management actions, with the broader goal of improving conservation of Kenya's important montane forests by leveraging the bongo's status as a flagship species (Caro & O'Doherty, 1999; Reillo, 2002, 2004; Estes *et al.*, 2008).

The Aberdares (3998 m) and MK (5199 m) are equatorial mountains of volcanic origin (Fig. 1). Climate is defined by the bi-modal precipitation pattern [annual rainfall range: ~950 mm (leeward north-western slopes) to ~2250 mm (windward south-eastern slopes)] caused by the shifting Inter-Tropical Convergence Zone, and temperature patterns that are shaped more by time of day and elevation than season (Schmitt, 1991; Bussmann, 1994).

Both mountains are protected by national parks and forest reserves administered by the Kenya Wildlife and Forest Services, and are encircled by dense rural settlements. A recently completed game fence encloses the Aberdares (Butynski, 1999; Rhino Ark, 2010). Our study focused on the forests lying within the Aberdares fenceline (1764 km²) or above lower forest reserve boundaries on MK (1963 km²). These forests contain a mix of bamboo and broadleaf forests, shrublands and open glades that are principally stratified by elevation (Schmitt, 1991; Bussmann, 1994). The bongo, which migrates seasonally along this habitat-elevation gradient, depends on a mosaic of primary forest, secondary forest and open glades (Kingdon, 1982), a configuration providing shelter as well as edge conditions where the bongo's preferred browse – tall forbs and lianes – can flourish (Estes *et al.*, 2008). More detailed descriptions of the study area and bongo ecology can be found in Estes *et al.* (2008, 2010).

Survey data

We conducted surveys in the Aberdares during three climatically representative periods in 2005–2006 (see Estes *et al.*, 2008, 2010 and supporting information Appendix S1 for details on survey data collection methods). We collected vegetation structural and compositional data in thirty-seven 0.04 ha circular plots centered on bongo habitat-use signs (e.g. browsing, resting) identified by expert trackers. An additional 90 'non-presence' plots representing the range in available habitats were collected from a point overlay established on a 1 km² grid (Fig. 1). Further surveys in 2006 and 2008 yielded 20 additional bongo-presence observations (five in the Aberdares, 15 on MK) without associated habitat information. Bongo observations were validated by analyzing mitochondrial DNA extracted from dung samples collected within or near sample plots (Faria *et al.*, 2011; supporting information Appendix S1). Using guidelines established by Dunn (1978), bongo observations were deemed independent based on the age differences between habitat-use signs recorded in adjacent presence plots (supporting information Appendix S1).

Supplemental surveys showed bongo signs were detected with a sufficiently high probability (0.88) such that imperfect detectability bias was negligible (MacKenzie, 2006; supporting information Appendix S1).

Predictors

Vegetation structural properties drive bongo habitat selection (Estes *et al.*, 2008). Another likely factor is avoiding predation, particularly from humans (other than park staff and tourists), whose use of the study area for hunting and other consumptive activities is pervasive (Kingdon, 1982; Gathaara, 1999; Lambrechts *et al.*, 2003; Prettejohn, 2008). We hypothesized that bongo habitat selection may be driven by (1) both vegetation properties and predation avoidance; (2) vegetation properties only; (3) predation avoidance only. We modeled these possibilities by selecting five variables related to these two habitat factors. To overcome data-collection constraints imposed by terrain and bongo rarity, we used spatial predictors derived primarily from remote sensing (RS).

Vegetation-related predictors

Bongo habitat selection is strongly influenced by micro-scale (0.04 ha) vegetation structure (Estes *et al.*, 2008). The difficulty of RS at this fine scale led us to develop a new technique (Estes *et al.*, 2010; supporting information Appendix S2) to improve microhabitat-mapping accuracy, which includes rigorous image pre-processing to minimize atmospheric and terrain-shadowing effects and positional errors (<17 m). We used a multiple regression model to predict and map values of a micro-scale (0.09 ha, corresponding to image pixel size) vegetation structural complexity index [*Vegetation structure* – range 0 (grassland) to ~80 (mature forest)], using five predictors derived from Advanced Spaceborne Thermal Emission and Reflection Radiometer images with spectral mixture and texture analysis (Adams, Smith & Gillespie, 1989; Tuceryan & Jain, 1993). In the resulting maps 99% (Aberdares) and 94% [MK, with clouds obscuring 13 km² (0.7%)] of predicted *Vegetation structure* scores fell within the observed range (0–80).

We used *Vegetation structure* maps to calculate edge density (*Edge*) at a 20.25 ha scale. We selected *Edge* [total boundary length (m) between vegetation types falling within a moving window divided by the window's area (McGarigal *et al.*, 2002)] because forest structure at this scale influences bongo habitat selection (Estes *et al.*, 2008), and bongo prefer an 'edgy' mixture of mature forest, secondary growth and open glades (Kingdon, 1982; Klaus-Hugi *et al.*, 1999). *Edge* calculations were based on three *Vegetation structure* classes: grassland/sparse shrubland (0–35); shrublands/bamboo forest (35–65); mature woody forest (>65) (the last two were reduced by 10 on MK to account for *Vegetation structure* underestimates due to color-balancing problems within the composited image).

RS analyses were unable to accurately predict understory structural properties, despite their importance to

bongo (Estes *et al.*, 2008; 2010). To provide information on forage quality, we employed a measure of soil moisture availability, the improved topographical relative moisture index (hereafter '*Moisture*', Parker, 1982; Manis, Lowry & Ramsey, 2001; supporting information Appendix S2), which varies between 1 (xeric) and 27 (mesic). We chose this measure because topography exerts strong controls on soil moisture, which is positively linked to forage nutrient content and persistence (Lambers, Chapin & Pons, 1998; Western *et al.*, 1999; Vázquez-de-Aldana *et al.*, 2000). Some African ungulates select habitat where terrain enhances moisture persistence and forage quality during dry seasons (e.g. Stelfox & Hudson, 1986). We calculated *Moisture* using a 90 m Shuttle Radar Terrain Mapping Mission (SRTM) digital-elevation model (DEM) interpolated to 30 m resolution.

Security variables

We derived two variables that can affect bongo's ability to escape predation: (1) distance to the nearest KWS ranger station (*Distance*), an indicator of anti-poaching effort, which we expect should be inversely related to bongo presence; (2) an index of terrain ruggedness (*Ruggedness*), which should correlate positively with bongo presence by facilitating predator escape. We calculated *Ruggedness* using the 90 m SRTM DEM, defined here as the mean squared elevation difference between each pixel and its eight neighbors (Riley, DeGloria & Elliot, 1999). We averaged resulting values using a 20.25 ha moving window, on the assumption that ruggedness influences bongo at the patch scale.

Habitat models

We tested 31 logistic regression models using all 132 Aberdares observations (42 presence, 90 non-presence). Model selection was based on the Akaike's Information Criterion for small sample sizes (AIC_c) and the associated Akaike difference (Δ) and weight (w) (Burnham & Anderson, 2002; supporting information Appendix S3). As we gave considerable *a priori* attention to variable selection, we tested all possible models (excluding interactions) to fully explore each predictor's influence. We used Hosmer & Lemeshow's (1989) test to assess the global model's goodness-of-fit. Correlation coefficients for each pairwise predictor combination were $<|0.5|$, and variance inflation factors were below 2.5 (Allison, 1999), indicating that multi-collinearity was not a concern. Tests to determine the effects of three potentially autocorrelated points revealed negligible model bias (supporting information Appendix S3).

We then created 30 m resolution predictive distribution maps representing the three combinations of the two overarching habitat determinants (predation avoidance; vegetation). Maps were created with model coefficients averaged (Burnham & Anderson, 2002, supporting information Appendix S3) across each set of models representing the three combinations (e.g. across the six models incorporating only vegetation predictors). 'Multi-model inference' provides more conservative error estimates by incorporating

uncertainty regarding 'best' model structure into predictors' confidence intervals (Burnham & Anderson, 2002). We used frequency distributions of mapped predicted presence probabilities (P) values to estimate potential habitat area under different P thresholds.

To assess model predictive performance, we calculated the area under the curve (AUC) of the receiver-operating characteristic [ROC, a threshold-independent measure of classification accuracy (Fielding & Bell, 1997)] for both individual and composite (averaged) models.

To examine the degree of model over-fitting, we used the '.632' bootstrap, which corrects accuracy measures (e.g. AUC) based on their variation when fit to re-sampled datasets (Efron, 1983; Harrell, 2005; supporting information Appendix S3). We further assessed model predictive capacity using the independent MK bongo observations. As these observations were exclusive to bamboo forests, we compared MK P values with those calculated for 10 Aberdares bongo plots recorded in comparable habitat, using the Mann-Whitney-Wilcoxon Rank Sum to assess differences in group mean P .

Results

Habitat models and predictor influence

We first ran the 31 individual models (Table 1), then created the three composite models (Table 2) by averaging: (1) all 31 models (FullMod); (2) six models based on vegetation variables only (VegMod); (3) the three models excluding vegetation variables (SecMod). The results show that *Ruggedness* was the strongest predictor of bongo habitat use, based on the strong AUCs of *Ruggedness*-based models (0.84–0.90; Tables 1 and 2) and AIC_c -based measures [e.g. summing w_i for individual models in which this predictor appeared indicates its relative influence (Burnham & Anderson, 2002), which in this case is 1 – the maximum possible value]. Soil moisture is nearly as important in influencing habitat use ($\sum w_i = 0.97$), followed by *Distance* ($\sum w_i = 0.76$), and *Vegetation structure* ($\sum w_i = 0.71$), but the patch-scaled measure of vegetation structural diversity (*Edge*) was least influential ($\sum w_i = 0.31$). However, all five predictors appeared in the top six individual models [the 96% confidence set, based on summing w_i (Burnham & Anderson, 2002)], whose low Δ_i and comparable w_i indicate substantial model selection uncertainty, which shows that both security and vegetation aspects (particularly micro-scale vegetation structure and soil moisture) determine bongo habitat use. The adequate predictive performances (AUCs >0.7 ; Pearce & Ferrier, 2000) of most individual models incorporating *Vegetation structure* and *Moisture* but lacking security variables (Table 1) and VegMod (Table 2) further support the importance of fine-scaled vegetation features in shaping bongo habitat use.

Model performance and transferability

Applying the .632 bootstrap (5000 replicates) showed that the individual models' predictive abilities were robust (Table 1), with a mean reduction in AUC of only 0.009.

Table 1 Results of the 31-tested logistic regression models, including the relative rank, model structure, number of parameters (K), AIC_c value, AIC_c differences (Δ_i), akaike weights (w_i), area under the curve (AUC) and the bootstrap-corrected AUC_{corr}

Rank	Model	K	AIC_c^a	Δ_i	w_i	AUC	AUC_{corr}
1	<i>Vegetation structure + Moisture + Distance + Ruggedness</i>	5	103.77	0.00	0.34	0.899	0.888
2	<i>Moisture + Distance + Ruggedness</i>	4	105.03	1.26	0.18	0.892	0.884
3	<i>Vegetation structure + Moisture + Ruggedness</i>	4	105.35	1.58	0.15	0.889	0.882
4	<i>Vegetation structure + Edge + Moisture + Distance + Ruggedness^b</i>	6	105.82	2.05	0.12	0.896	0.883
5	<i>Edge + Moisture + Distance + Ruggedness</i>	5	106.39	2.62	0.09	0.890	0.880
6	<i>Vegetation structure + Edge + Moisture + Ruggedness</i>	5	106.61	2.84	0.08	0.887	0.878
7	<i>Vegetation structure + Distance + Ruggedness</i>	4	110.22	6.45	0.01	0.881	0.872
8	<i>Distance + Ruggedness</i>	3	111.15	7.38	0.01	0.873	0.868
9	<i>Edge + Moisture + Ruggedness</i>	4	112.14	8.37	0.01	0.869	0.862
10	<i>Vegetation structure + Edge + Distance + Ruggedness</i>	5	112.20	8.42	0.00	0.878	0.866
11	<i>Edge + Distance + Ruggedness</i>	4	112.41	8.64	0.00	0.870	0.862
12	<i>Vegetation structure + Ruggedness</i>	3	114.28	10.51	0.00	0.866	0.861
13	<i>Vegetation structure + Edge + Ruggedness</i>	4	115.01	11.24	0.00	0.861	0.854
14	<i>Moisture + Ruggedness</i>	3	115.42	11.65	0.00	0.860	0.856
15	<i>Edge + Ruggedness</i>	3	119.67	15.90	0.00	0.847	0.841
16	<i>Ruggedness</i>	2	123.99	20.22	0.00	0.842	0.841
17	<i>Vegetation structure + Moisture + Distance</i>	4	142.53	38.76	0.00	0.779	0.764
18	<i>Vegetation structure + Moisture</i>	3	144.34	40.57	0.00	0.765	0.755
19	<i>Vegetation structure + Distance</i>	3	144.43	40.66	0.00	0.761	0.752
20	<i>Vegetation structure + Edge + Moisture + Distance</i>	5	144.53	40.76	0.00	0.784	0.759
21	<i>Vegetation structure + Edge + Moisture</i>	4	145.68	41.91	0.00	0.764	0.747
22	<i>Vegetation structure + Edge + Distance</i>	4	146.38	42.61	0.00	0.759	0.744
23	<i>Vegetation structure</i>	2	148.05	44.28	0.00	0.728	0.728
24	<i>Vegetation structure + Edge</i>	3	149.17	45.40	0.00	0.726	0.718
25	<i>Moisture + Distance</i>	3	151.08	47.31	0.00	0.731	0.722
26	<i>Edge + Moisture + Distance</i>	4	151.92	48.15	0.00	0.742	0.722
27	<i>Distance</i>	2	152.19	48.42	0.00	0.733	0.733
28	<i>Edge + Distance</i>	3	152.78	49.01	0.00	0.738	0.729
29	<i>Edge + Moisture</i>	3	160.04	56.27	0.00	0.681	0.664
30	<i>Edge</i>	2	162.36	58.59	0.00	0.645	0.644
31	<i>Moisture</i>	2	163.97	60.20	0.00	0.622	0.619

^a AIC_c of null model ($K=1$; intercept only) = 167.16.

^bHosmer–Lemeshow test results on full model indicate adequate fit: $\chi^2=5.42$, d.f. = 8, $P < 0.71$.

Both the independent MK bongo observations and the Aberdares bamboo zone bongo (ABZ) plots were predicted to have low occurrence probabilities (P) by all three composite models (particularly SecMod), such that 70–100% would be falsely classified as non-presences using a standard P threshold of 0.5 (Fig. 2), and a majority would be misclassified even with the threshold $P = 0.3$. However, FullMod and SecMod predicted much larger discrepancies between ABZ and MK P than VegMod, which found that bongo are as likely to use bamboo forests on MK as they are in the Aberdares (Fig. 2 and Table 2). These results indicate that VegMod was more transferable, and this pattern of between-mountain comparison was reflected in each set of models underlying the composite models (supporting information Table S2).

Habitat availability

We created habitat suitability maps (Fig. 3) using the spatial predictors and their corresponding model-averaged coefficients. FullMod predicts that 21% of the Aberdares and 8%

of MK are suitable at the 0.5 threshold, or 34 and 15% when $P = 0.3$ (Fig. 4). Under this model, deep valleys and escarpments with moderate to high vegetation structural complexity and soil moisture availability were predicted to have the highest probabilities of bongo occurrence. SecMod (not illustrated) delineated the most rugged areas as bongo habitat, making just 15% of the Aberdares and 6% of MK suitable at the 0.3 threshold. VegMod identified the greatest extents of suitable bongo habitat [32% in the Aberdares and 29% of MK ($P = 0.5$), or 53 and 55% ($P = 0.3$)], comprised of areas with high vegetation structural complexity and soil moisture availability, which are conditions that predominate in lower-elevation woody forests.

Discussion

Ecological findings

The performance of *Ruggedness*-based models indicates that mountain bongo strongly prefer difficult terrain (Tables 1 and 2). *Distance* was not as strong a predictor, but its

Table 2 The model averaged parameter estimates ($\bar{\beta}$), unconditional standard errors (SE) and upper and lower 95 per cent confidence limits for the three composite models: AllMod (average of 31 models), VegMod (average of six models based on non-security predictors) and SecMod (average of three models based on the two security variables)

Model	Intercept	Vegetation					AUC	MK	ABZ	MWW
		structure	Edge	Moisture	Distance	Ruggedness				
FullMod										
$\bar{\beta}$	-9.113	0.060	0.004	0.168	-0.175	0.065	0.892	μ 0.21	0.40	1.86 ^a
SE	2.876	0.031	0.005	0.061	0.087	0.013		σ 0.24	0.33	
Upper	-3.476	0.121	0.015	0.287	-0.004	0.090		\downarrow 0.03	0.04	
Lower	-14.750	-0.001	-0.007	0.048	-0.345	0.039		\uparrow 0.78	0.92	
VegMod										
$\bar{\beta}$	-6.681	0.085	0.004	0.104			0.766	μ 0.28	0.35	0.97
SE	1.457	0.021	0.004	0.046				σ 0.19	0.20	
Upper	-3.826	0.127	0.012	0.194				\downarrow 0.04	0.16	
Lower	-9.536	0.044	-0.004	0.015				\uparrow 0.64	0.83	
SecMod										
$\bar{\beta}$	-3.651				-0.237	0.060	0.873	μ 0.10	0.21	1.58
SE	0.820				0.069	0.011		σ 0.05	0.15	
Upper	-2.045				-0.101	0.082		\downarrow 0.04	0.04	
Lower	-5.258				-0.374	0.037		\uparrow 0.17	0.46	

^a($P < 0.1$).

The AUCs based on predicted probabilities (P) of the averaged models are presented, as well as the mean (μ), standard deviation (σ), minimum (\downarrow) and maximum (\uparrow) of P at Mount Kenya (MK) and Aberdares Bamboo Zone (ABZ) bongo plots. MWW is the Mann–Whitney–Wilcoxon Z score approximation for tests between MK and ABZ means, and its associated significance code.

relative importance ($\sum w_i = 0.76$) and negative association with bongo presence (Tables 2 and S1) suggests that bongo select areas closer to – thus more likely to be patrolled by – park rangers. Although these measures do not directly quantify predation risk, the most plausible explanation for their importance is that they relate to bongo's ability to evade predation. We believe this risk is primarily from humans, given the prevalence of poaching and other disruptive human activity (Kingdon, 1982; Butynski, 1999; East, 1999; Gathaara, 1999; Lambrechts *et al.*, 2003; Prettejohn, 2008), and that a *Distance* effect would be less likely in the absence of human hunting (ranger patrols do not deter other predators). Other large herbivores, such as mule deer and bighorn sheep, use rugged terrain to escape predators (e.g. Lingle, 2002; Mooring *et al.*, 2004; Pierce, Bowyer & Bleich, 2004), while populations of several large African herbivores have shown positive correlations with anti-poaching patrol effort (Leader-Williams & Albon, 1988; Hilborn *et al.*, 2006), and increased vigilance and habitat avoidance associated with hunting and other disruptive human activities (Laurance *et al.*, 2006; Buij *et al.*, 2007; Fischer & Linsenmair, 2007). Nevertheless, direct measures of predation risk are needed to confirm its importance in determining bongo habitat.

The proxies for predation risk do not explain all bongo habitat use. Model results showing the importance of *Moisture* and *Vegetation structure* confirmed that fine-scaled vegetation features also influence bongo habitat use (Estes *et al.*, 2008). *Moisture* does not measure browse abundance, but other studies suggest that it influences forage quality and hence the habitat selection of other large African ungulates (Stelfox & Hudson, 1986; Lambers *et al.*, 1998; Western

et al., 1999; Vázquez-de-Aldana *et al.*, 2000; supporting information Appendix S2). Given these relationships, it makes sense that bongo might prefer mesic sites, as the available forage is likely to have higher and longer lasting nutritive value. Additional field evidence also suggests that the abundance of preferred bongo food plants tends to increase with structural complexity (unpublished data), thus areas with both high *Vegetation structure* and *Moisture* values may provide optimal foraging conditions, or 'core' habitat, where forage quality is highest and most persistent.

Historical records and our data indicate that bongo use bamboo forests (Kingdon, 1982), but our models' predict low usage probabilities for this habitat. This is due to *Vegetation structure* describing bamboo's structural simplicity relative to other forest types (Estes *et al.*, 2010), meaning that vegetation-related models predicted lower P (when all other variables were equal) for bamboo than more complex woody forests. Assessing *Vegetation structure's* influence on habitat use within particular – rather than across all – vegetation types could resolve this contradiction, which is supported by our results showing that bamboo zone bongo plots occurred in areas of higher occurrence probabilities relative to their immediate surroundings (see inset boxes in Fig. 3b., where P for most plots is 0.2–0.6, compared with 0.0–0.2 in the wider neighborhood). However, this step depends on accurately mapping vegetation types, which is difficult. Moreover, it is possible that bamboo forests comprise less suitable bongo habitat than woody forests, as bamboo suppresses understorey growth over much of its life cycle (Wimbush, 1945; Schmitt, 1991), thereby reducing the relative abundance of forage plants. The bamboo zone may

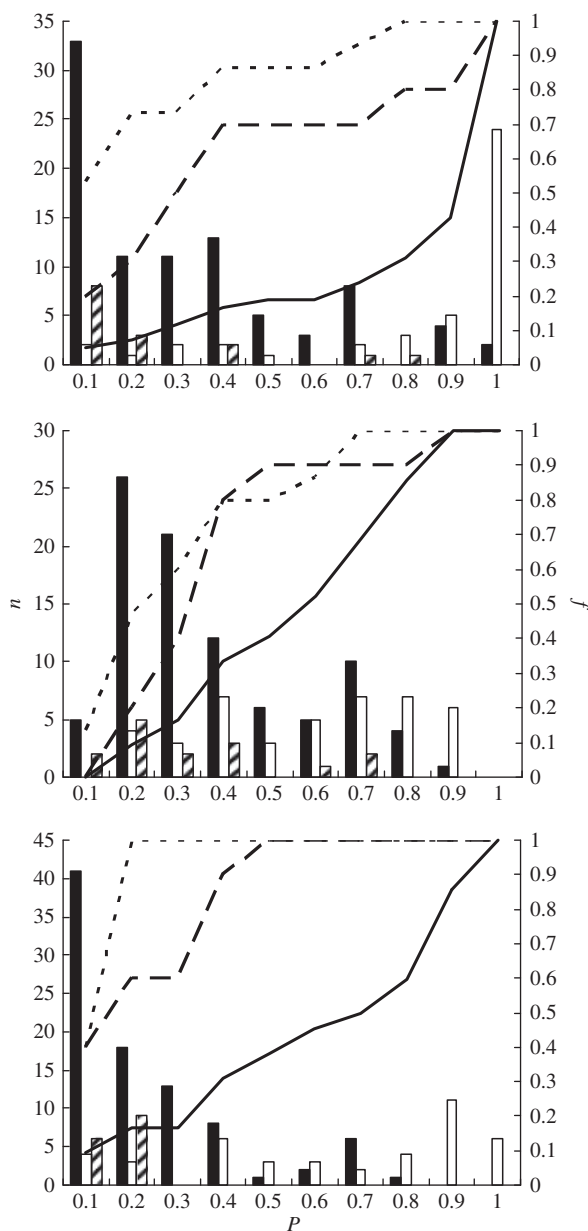


Figure 2 The distributions of occurrence probabilities (P) predicted by the FullMod (top), VegMod (middle) and SecMod (bottom) models for bongo presence [white columns=Aberdares; striped=Mount Kenya (MK)] and non-presence (black columns) observations. Lines indicate the false absence frequencies (n) at each P threshold for all Aberdares (solid), Aberdares bamboo zone (dashed) and MK bongo plots (dotted).

thus support lower bongo densities than woody forests, or serve primarily as a dry season refugium, when it benefits from high altitude, mist-derived moisture that is not received in the lower woody forests (Kingdon, 1982).

Edge density was less influential than expected. Previous work suggested that bongo prefer mosaic habitats (Kingdon, 1982), and that both patch- and micro-scale vegetation

structure influence bongo habitat use (Estes *et al.*, 2008). We thus expected *Edge* to be a stronger predictor. Although *Edge's* presence in four models of the 96% confidence set (Table 1) suggests a non-negligible influence, here we found that the finer-scaled vegetation variables were most important.

Based on these findings, the full composite model (Full-Mod) identified the realized niche (Austin, Nicholls & Margules, 1990) of a population striving to simultaneously avoid predation pressure and find suitable foraging conditions. These constraints mean that <20% of either mountain's forests are currently well suited for bongo. VegMod, which did not incorporate security-related variables, predicted occurrence in places currently avoided by bongo due to human activity (e.g. in lower forests far from KWS security bases). Although it effectively segregated bongo and non-presence observations, SecMod was unrealistic because it ignored vegetation-related requirements, leading it to delineate all rugged places as suitable bongo habitat, even non-forested areas that bongo would not use.

Model performance

Modeling rare species is difficult because sample sizes are usually small and subject to substantial errors, such as non-independence due to spatial clustering (Guisan & Zimmermann, 2000; Rushton *et al.*, 2004). Our bongo sample set was small and appears clustered, but was comparable in size to those used in other rare-species studies (e.g. Gibson *et al.*, 2004; Jeganathan *et al.*, 2004), and largely independent according to temporal criteria (Dunn, 1978; Estes *et al.*, 2008; supporting information Appendix S1), while model bias due to potentially autocorrelated points was negligible (supporting information Appendix S3). The dung sample mitochondrial DNA analysis allowed us to identify field identification errors and helped exclude false or potentially misidentified bongo observations that would have confounded our models (Estes *et al.*, 2008; Faria *et al.*, 2011; supporting information Appendix S1). A comparison of ordinary logistic regression and occupancy models run with simulated detection probabilities showed that our estimated DP of 0.88 would reduce P downwards by 0.02 at most, suggesting that missed detections did not unduly bias our results (supporting information Appendix S1).

These checks improved confidence in results, but the methods used to construct the checks could have error. For example, if the true detection probability was substantially lower (e.g. 0.61) than we estimated, our models could have under-predicted P by up to 0.1 (supporting information Figure S3).

Our predictors are also sources of potential model bias, as shown by our model transferability assessments. Effective PDMs must be transferable between habitats, but this property may be compromised by several factors, including over-parameterization and spatio-temporal variation in species–environment relationships (Guisan & Zimmermann, 2000; Vaughan & Ormerod, 2005). Bootstrapping revealed that models were not over-fit (Vaughan & Ormerod, 2005),

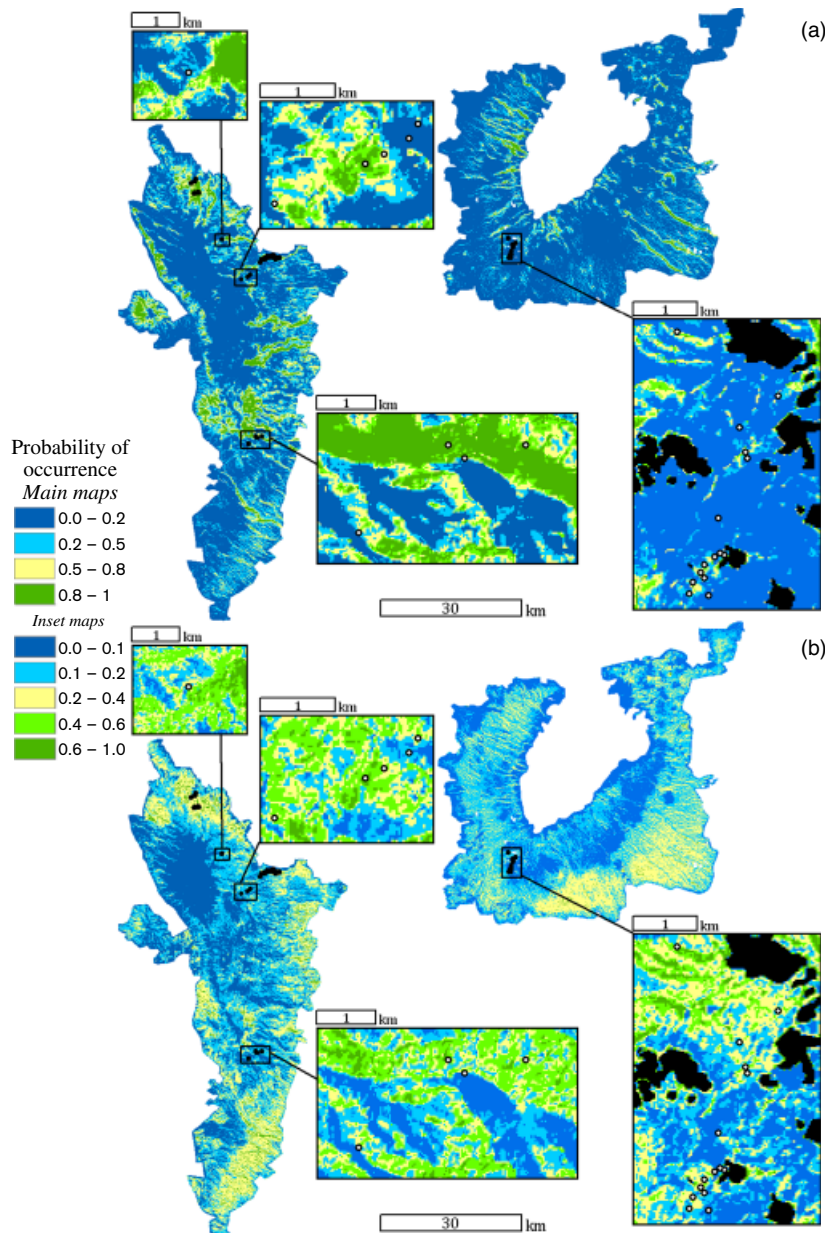


Figure 3 Bongo occurrence probabilities for the Aberdares (left) and Mount Kenya (right), as predicted by the FullMod (a) and VegMod (b) models. The inset boxes focus on Aberdares bamboo zone and Mount Kenya (black fill denotes no data) bongo plots (black circles with white fill).

but MK observations, although few and confined to one habitat type, showed that the modeled species-habitat relationship varied between mountains. Bongo habitat associations should not differ greatly between the Aberdares and MK, which are geographically, climatically and floristically close. The limited transferability of some models is instead related to between-mountain variations in certain predictors' environmental associations, particularly those of *Ruggedness*, the strongest but most locally biased predictor. This bias was due to the relative gentleness of MK's terrain relative to the Aberdares'. Model coefficients thus underestimated the ruggedness effect for MK, resulting in low *P* for bongo observations collected in terrain that in fact is relatively steep (the southern cluster in Fig. 3a inset). This

result highlights the inherent local bias of indirect topographic predictors (Guisan & Zimmermann, 2000), and is underscored by VegMod's better fit to MK plots, due to the dominance of *Vegetation structure* and *Moisture*, which are more directly related to habitat use, while their underlying derivation methods are also more transferrable (Austin, 2007; Estes *et al.*, 2010). Replacing *Ruggedness* (and *Distance*) with a variable quantifying predation risk could potentially increase model transferability.

Vegetation structure error also impacted between-mountain model predictions. Unlike the Aberdares, the imagery used to cover MK showed temporal variation (e.g. phenology and solar illumination) that increased brightness values in certain places (particularly around bongo samples),

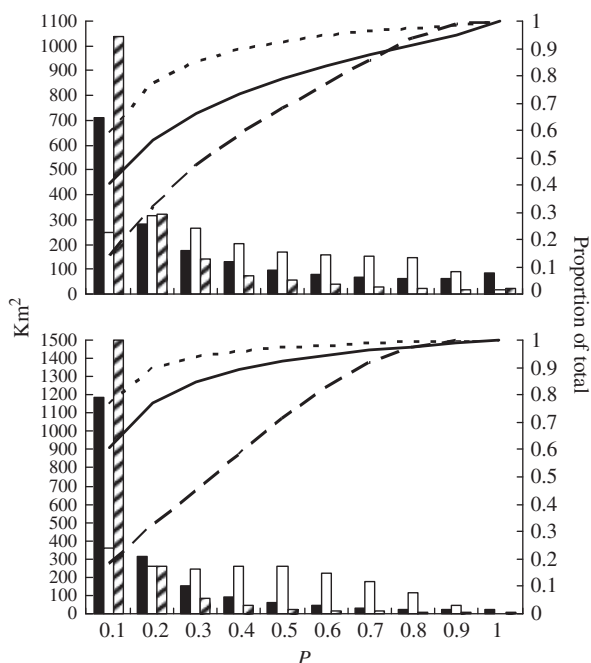


Figure 4 The area (km²) habitat on the Aberdares (top) and Mount Kenya (bottom) falling into suitability classes [based on occurrence probability (P) deciles] generated by the FullMod (black columns), VegMod (white columns), and SecMod (striped columns) models. Lines indicate the proportions of the two study areas falling below each P threshold (FullMod = solid; VegMod = dashed; SecMod = dotted).

causing *Vegetation structure*, and thus P , to be underestimated. The effect on model transferability was nevertheless minor compared with *Ruggedness*-based models, given the similarity between VegMod's ABZ and MK mean P values and estimates of habitat suitability on both mountains (Figs 3 and 4).

Moisture, a topographic index, may account for ~50% of the variance in measured soil moisture (Western *et al.*, 1999), while the correlation strength between soil moisture and forage quality is not known. Constructing models that predict the quality and abundance of preferred browse species is a logical next step, if additional data become available.

All but the last three individual models had adequate-to-good predictive ability, but showed high model selection uncertainty ($AUC \geq 0.73$; Table 1), meaning that the ranking of the 'best' model can change given new data (Burnham & Anderson, 2002). To avoid this model selection bias, we averaged models to predict bongo habitat distribution according to the three habitat use hypotheses. The relationship described by VegMod was more general to both habitats, but FullMod provided a 'truer' representation of bongo habitat use in an environment with substantial predation risk. Both models likely underestimated the extent of suitable bongo habitat, because they did not consider seasonal variations in use. These models instead identified what should be core bongo habitat suitable for the longest periods. Developing direct measures of predation risk and bongo food plant abundance, using a block effect to account

for structural complexity differences within vegetation types, and sourcing better RS imagery for MK could improve them.

Management implications

Despite FullMod's reduced performance on MK, we believe it is nevertheless useful – and most relevant – for informing bongo recovery plans. Management actions should be focused where FullMod shows the greatest extent of suitable habitat, based on a high P threshold to ensure that high quality habitat is identified, thereby improving the odds of success (Pearce & Ferrier, 2000). In the Aberdares, population recovery may be best achieved by concentrating anti-poaching efforts in the two or three largest blocks where $P > 0.5$ – 0.7 (e.g. in the north-western patch in Fig. 3a.). On MK, potential bongo reintroduction sites can be located using thresholds lowered to 0.3–0.4 to accommodate AllMod's poorer fit. Here results suggest a site on the western side of the mountain would be most suitable (Fig. 3a).

The suite of individual validated models could also be used to test different management goals. For instance, if cost constraints prevent the extension of protective measures to new areas, Model 17 (*Vegetation structure + Moisture + Distance*) could identify areas with optimal microhabitat conditions close to existing KWS bases.

Bongo surveys need to be continued at a broad scale throughout MK and the Aberdares, and should be targeted using FullMod. VegMod offers insight into overall habitat availability, and may help in estimating potential bongo population sizes as demographic information becomes available.

General implications

Besides providing important basic ecological information for this poorly understood animal, our study may help explain habitat use by other large forest herbivores – particularly the more widely spread lowland bongo (*Tragelaphus eurycerus eurycerus*) of Central and West Africa – as it presents a means to improve rare species modeling. Our RS techniques for vegetation-structure mapping improve on those typically used by ecologists (Estes *et al.*, 2008, 2010), allowing us to incorporate crucial microhabitat features in the multi-scale models, and augment sample size. The layered validation approach (mtDNA confirmation of observations; internal and external model performance assessments) builds on recent well-recognized rare-species modeling studies (e.g. Gibson *et al.*, 2004; Jeganathan *et al.*, 2004). Such generalizable, multi-level methodologies are increasingly needed to guide the management and recovery of rare species and ecosystems, where ecological data are difficult to obtain and swift conservation action must be based upon best-available inferences.

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Supporting Information

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

Table S1. Basic statistics.

Table S2. Full model results.

Appendix S1. Survey data details.

Appendix S2. Remote sensing methodology.

Appendix S3. Habitat modeling techniques.

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