CHAPTER 3

Vulnerability of South African animal taxa to climate change

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Abstract

The responsiveness of South African fauna to climate change events is poorly documented and not routinely incorporated into regional conservation planning. We model the likely range alterations of a representative suite of 179 animal species to climate change brought about by the doubling of CO2 concentrations. This scenario is expected to cause a mean temperature increase of 2°C. We applied a multivariate climate envelope approach and evaluated model performance using the most comprehensive bird data set. The results, based on distribution records from South Africa only, were encouraging although model performance was inconsistent in the eastern coastal area of the country. The levels of climate change-induced impacts on species ranges varied from little impact to local extinction. Some 17% of species expanded their ranges, 80% displayed range contraction (0 - 98%) and 3% showed no response. The majority of range shifts (41%) were in an easterly direction, reflecting the east-west aridity gradient across the country. Species losses were highest in the west. Substantially smaller westward shifts were present in some eastern species. This may reflect a response to the strong altitudinal gradient in this region, or may be a model artifact. Species range change (composite measure reflecting range contraction and displacement) identified selected species that could act as climate change indicator taxa. Red-data and vulnerable species showed similar responses but were more likely to display range change (58% vs. 43% for all species). Predictions suggest that the flagship Kruger National Park conservation area may lose up to 66% of the species included in this analysis. This highlights the extent of the predicted range shifts, and indicates why conflicts between conservation and other land uses are likely to escalate under conditions of climate change.

Introduction

Large-scale species extinctions and movements were precipitated by major climate events in the past (Gates 1993; Coope 1995; Roy et al. 1996). Consequently, prediction of the biodiversity consequences of current and predicted climate change is becoming increasingly important (McNeely et al. 1995), especially because such change is not likely to affect all species similarly. Some species are expected to benefit from and others to suffer under altered climatic conditions (Fajer et al. 1989; Freedman 1989; Cammell & Knight 1992; Davis et al. 1998a,b). Thus, not only are species likely to be affected by such change, but community composition is also likely to be altered because of individualistic species responses (see Coope 1995; Jablonski & Sepkoski 1996 for examples of such changes in the past). If biodiversity is to be conserved (Convention on Biodiversity, UNEP, Rio de Janeiro 1992), an understanding of how species and communities are likely to change under conditions of climate change is essential. Such understanding is particularly important because existing conservation networks, as well as theoretically selected areas, will not perform adequately if species temporal turnover is not taken into consideration (Rodrigues et al. 2000). This is likely to be true particularly if current and predicted climate change result in dramatic species movements, as seems likely to be the case (Parmesan et al. 1999, Pounds et al. 1999). Furthermore, interactions between habitat fragmentation, and changing species distribution patterns imply considerable future conservation conflicts (McNeely 1994, Fearnside 2000), making predictions of likely range shifts all the more important.

In South Africa, early qualitative (McDonald & Midgley 1996), and more recent quantitative (Rutherford et al. 2000) analyses of the effects of predicted climate change on the flora have suggested that the species rich biomes in the south-west of the country are likely to suffer substantial biodiversity losses, and that species loss from the existing protected area network will be significant. Despite these rather dramatic predictions, to date no attempts have been made to investigate the likely effects of predicted climate change on the South African fauna. Here we provide a first assessment of individual animal species responses to a climate change scenario in South Africa, by modeling the likely responses of the geographic ranges of a selection of 179 animal species from across the taxonomic spectrum.

The analytical approach we employ is generally referred to as the "climate envelope" approach (Sutherst & Maywald 1985; Sutherst et al. 1995; Markham & Malcolm 1996), and explores the extent to which species ranges might shift in response to changes in the

surrounding environment. Our use of this approach does not mean that we are unaware that climate is only one determinant of species distributions. Rather, we reasoned that while other factors, such as both horizontal and vertical linkages in ecosystems (Davis *et al.* 1998a, b; Buse *et al.* 1999; Hochberg & Ives 1999), and the phylogenetic history of taxa (see Myers & Giller 1988; Brown & Lomolino 1998), are likely to influence the geographic distributions of taxa, in the main, geographic distributions are likely to be determined to a large degree by climate (Jablonski *et al.* 1985; Root 1988; Rogers & Williams 1994; Coope 1995; Robinson *et al.* 1997a, b; Chown & Gaston 1999; Spicer & Gaston 1999). Nonetheless, we acknowledge that a full understanding of the consequences of climate-induced range changes for the functioning of local ecosystems is unlikely to be achieved without additional investigations. These include the development of an understanding of the influence of such changes on regional diversity, and in turn its influence on local communities (see Cornell & Lawton 1992; Ricklefs & Schluter 1993; Tokeshi 1999 for review of local and regional interactions), and the effect that novel species interactions, precipitated by range changes, will have at local scales.

However, given the speed with which climate change is taking place (IPCC 1992, 1995, 1997, 2000) and that, within the context of urgent local development needs (e.g. ANC 1994), conservation decisions have to be made (van Jaarsveld *et al.* 1998), the approach we use provides a rapid and initial best estimate of likely responses of species over broad areas (Guisan & Zimmerman 2000). This estimate can be used as a broad guide to the urgent conservation action that is required, while additional, more specific data on species interactions and their change under different climate scenarios are collected (e.g. Masters et al. 1998; Buse et al. 1999; Fielding et al 1997).

Materials and methods

Species screening and distribution data

Distribution data for birds, mammals, reptiles (snakes, lizards, and tortoises), butterflies, dung beetles (Scarabaeidae), antlions (Neuroptera), jewel beetles (Buprestidae) and termites were collated. The last four invertebrate taxa were grouped together under "other invertebrates" because they are generally poorly known in the subregion (Scholtz & Chown 1995; Koch et al. 2000). The species distribution data were obtained from: *birds* (Avian Demography Unit, University of Cape Town), *mammals* (Natural History Museum, Northern

Flagship Institution, Pretoria (NFI); Durban Natural Science Museum; Conservation Planning Unit, University of Pretoria (CUP)), reptiles (NFI; Port Elizabeth Museum), butterflies (NFI), antlions (National Insect collection, ARC Plant Protection Research Institute (NCI)), jewel beetles (NFI), dung beetles (NCI, NFI, CUP) and termites (CUP). Data were generalized to a 15'x 15' grid cell resolution (~25 X 25 km), the finest common resolution between taxa, to generate absence-presence maps. Because many southern African species remain poorly known, both taxonomically and geographically (Scholtz & Chown 1995; van Jaarsveld 2000), taxonomic experts were requested to identify a suite of representative species in their study taxon, to increase the likelihood of including accurate and representative distributions, based on the following criteria: (a) accurate but not necessarily comprehensive distribution patterns, (b) representative geographic range types (e.g. species restricted to winter or to summer rainfall regions), (c) taxonomic clarity, and (d) species known from a reasonable number of records. A total of 34 bird, 19 mammal, 50 reptile, 19 butterfly, and 57 "other invertebrate" species were finally selected (Appendix 1).

Climate data (historic and predicted)

Climate data for South Africa were provided by the Computing Centre for Water Research (CCWR, University of Natal, Pietermaritzburg). The historic data (30 year monthly and annual means: 1960 – 1989) were re-sampled to 15' x 15' grid cells from interpolated climate surfaces available at a minute by minute resolution to conform with the resolution of the species data. The variables employed in the present study are: minimum temperature, maximum temperature, and precipitation seasonality measured as the difference between February and August mean rainfall figures. These months were selected because a factor analysis of the 12 monthly rainfall means showed that February rainfall contributed most to Factor 1, which explained 56% of the variance, and August rainfall contributed most to Factor 2, which explained 37% of the variance in the data. Preliminary analyses with these data indicated that temporal variability is a dominant feature, with annual mean precipitation values tending to disguise seasonal aspects of climate change (Erasmus *et al.* 2000).

Equivalent precipitation and temperature variables were also obtained for a predicted climate change scenario (CCWR). These climate data were derived from climate surfaces produced from GCM (General Circulation Model) predictions. Climate changes values were derived by implementing the Hadley Centre Unified Model with no sulphates, which represents a worst-case scenario for South Africa (Hewitson 1998). This GCM predicts a

temperature increase of $2.5 - 3^{\circ}$ C for South Africa with atmospheric CO₂ doubling from preindustrial levels. Erring on the side of caution, this means that significant climate change can be expected by the year 2050 (Hewitson 1998), and quite possibly earlier.

Species distribution modeling and evaluation

The model we employed is conceptually similar to the one proposed by Jeffree & Jeffree (1994, 1996). The original model was modified to accept multivariate data as input, and to produce a probability surface as output, rather than a simple binary prediction (see Erasmus et al. 2000 for a full description). Essentially, on a scatterplot of climate variables (multivariate climate space), the values of climate variables at known records are used to construct a confidence region where there is a high probability that the records reflect the core range of the species. Points falling inside this confidence region are then mapped back to geographic space to represent an interpolated distribution (ID). Climate affected distributions are produced by altering the climate variables of individual cells according to the GCM predictions, i.e. shifting the climate space, and subsequently mapping the predicted climate affected species distributions (CAD) back into geographic space.

Changes in distribution patterns owing to predicted climate change were assessed by comparing interpolated distribution patterns (ID) with climate affected distribution patterns (CAD) using a number of variables:

- a. Species richness pattern number of species predicted per grid cell using interpolated distributions;
- b. Species range shift proportion of climate-affected distributions (CAD) inside the interpolated distribution (ID).
 - c. Species range contraction climate-affected distribution (CAD) as a proportion of the interpolated distribution (ID), and
 - d. Species range change the sum of interpolated distribution grid cells lost and additional grid cells added under climate change, expressed as a percentage of the original interpolated distribution (ID). This additional measure provides an integrated assessment of range alterations through the combined effects of range contraction and shift.

These analyses were conducted for each taxonomic group separately, collectively across all species and for Red-data (mammals, birds, butterflies) and vulnerable species (identified as such by taxon experts: reptiles, jewel beetles, dung beetles, termites and antlions). Areas of

the country that are currently more than 50% transformed by human activity (Fairbanks *et al.* 2000) were superimposed on the post climate change species richness map to assess the potential availability of pristine land for accommodating climate-driven species range shifts.

To evaluate our model (Oreskes et al. 1994, Guisan & Zimmerman 2000), we compared interpolated distributions with known records. Incomplete distribution data severely limited the extent to which absence data could be included in model evaluation procedures for most taxa. The bird data, resulting from extensive sampling, was the best approximation of true presence-absence data and was used for evaluation. Instead of dichotomising the predicted probability output to predicted presence-absence by applying a threshold probability, which results in loss of information (Deleo 1993, Fielding et al. 1997), the actual predicted probabilities were utilized using receiver operating characteristic analyses (ROC). ROC analysis has its origins in engineering in the context of measuring the ability of a detector to detect a particular signal. It has found wide application in clinical medical tests (Fielding et al. 1997) and here it is used to measure the performance of a predictive species' distribution model. Kraemer's (1988) critique of ROC application in biology concerns the subjectivity of observations and the inability of the observed characters to be controlled or known with absolute certainty. The bird data set has been subjected to extensive expert vouching procedures and as such presents a close approximation of objective observations (Allan et al. 1997, van Jaarsveld et al. 1998). Therefore, we regard ROC analyses as suitable for our purposes of evaluation.

ROC analysis primarily concerns the calculation of specificity and sensitivity values. Sensitivity is defined as the number of true occurrences of a species (true positive predictions, TP) divided by the total number of positive predictions, whether true (TP) or not (false negatives predictions, FN). Similarly, specificity is defined as the number of true negative predictions (TN) divided by the total number of negative predictions, whether TN or false positive (FP) predictions (Fielding *et al.* 1997). The area-under-curve (AUC) of a plot of 1-specificity against sensitivity at every given probability of occurrence value is used as a test statistic. The AccuROC® software (Vida 1993) uses the nonparametric method of De Long *et al.* (1988) to determine the statistical significance of the difference between an AUC of 0.5 (corresponding to a random test with no discriminatory ability) and the AUC of every bird species. The confidence interval for the AUC was calculated using the asymptotic method (Obuchowski *et al.* 1998). A random subsample (with no replacement) of 20% of the known records of each species were also taken and used as input to generate probabilities of occurrence. The AUC of the 20% subsample was compared with the AUC based on all

known records (De Long *et al.* 1998). Such cross validation should ideally involve two independent data sets, although subsampling is an acceptable alternative (Guisan & Zimmerman 2000). Owing to a lack of absence data, the ROC analyses were only applied to the bird dataset to evaluate the model's ability to match predicted and observed data. The predicted probabilities generated for the other taxonomic groups had to be dichotomized into presence-absence data after determining a probability of occurrence threshold level from the ROC analyses of the bird data. This threshold was determined by calculating the correct classification rate ([TP + TN]/[TP+FP+TN+FN]) for all possible thresholds and selecting an appropriate probability threshold at which to interpret distribution changes.

Results

Model evaluation

The model correctly classified TN and TP predictions at a mean correct classification rate of 0.79 (SD=0.09) using a 50% probability of occurrence threshold for all 34 bird species. Such a high correct classification rate suggests strongly that the model can make useful predictions. Further support for the usefulness of the model (Zweig & Campbell 1993) comes from the construction of a confusion matrix (Fielding & Bell 1997) for the bird data. The results are summarized in Table 1. Mean values of 60% for true positives and 88% for true negatives suggest a reasonable performance of the model. This result is further underscored by the narrow ranges recorded for the variables. Although the mean figures for false positives and false negatives may inspire confidence in the above conclusion, the extremely large ranges recorded for false positives (1 – 44%) suggest that the model is useful for predicting the true ranges of some species but less so for other bird species. For those species where the model performs poorly, the 50% probability of occurrence area may extend markedly beyond the recorded range of the species. The large mean figure obtained for false negative values (38%) can be explained by our application of the 50 % probability of occurrence rule, with many species being recorded at lower probabilities of occurrence than the benchmark 50% level.

Given these considerations, the 50% probability of occurrence figure was considered an acceptable probability of occurrence and was applied to all other taxa. Thus, species were considered present when the model predicted a probability of

Table 1 Percentage prediction reliability at the 50% probability of occurrence level, for the multivariate model (Erasmus *et al.* 2000) used in this study. These estimates are for the bird data only because true distributions of the other taxa are too poorly known.

que to 5 1 kg st	True positives	True negatives	False positives	False negatives
Mean ± S.D.	60.3 ± 5.2	87.8 ± 10.3	12.2 ± 10.3	39.7 ± 5.2
Range	49.1 – 69.3	56.0 – 98.6	1.4 - 44.0	30.7 - 50.9

occurrence of \geq 50%, and absent below 50%. The application of this general rule accounts for losses of species from many areas of the country. Thus, a 100% species loss in any one area in effect means that no species were present in that area at a \geq 50% probability of occurrence.

Although ROC analyses performed on the bird data could not be applied to other taxa due to a lack of absence data, the results form these analyses provided further support for the usefulness of the model's predictions. Model evaluation using the area under the ROC curve as a test statistic found that all 34 bird species in the ROC analyses had area-under-curve (AUC) values significantly different (p < 0.05, ranging from 0.6357 to 0.9665, mean 0.85) from an AUC of 0.5 (see Fig. 1 for an example). The number of known records needed to detect a significant difference (probability of type I error 0.1, probability of type II error, 0.5) between an AUC of 0.5 and 0.85 was found to be lower (44) than the number of known records for the species with the least number of records (49 for the Drakensberg Siskin). The remaining 33 bird species had larger ranges with the number of known records varying from 98 to 1573 with a mean of 697. This means that differences between the measured AUC values and an AUC of 0.5 are real and that the model performs significantly better than a random model with no discriminatory ability.

For 24 of the 34 bird species, the AUC values derived from analyses using a 20% subsample of known records were not significantly different (p < 0.05) from the AUC values derived from the complete data set of known records for each bird. The remaining 10 species (all non-passerine and of relatively large body size) had AUC values significantly different (p values ranging from 0.0553 to 0.9131) from the AUC values based on the complete data set of known records for each species. This result suggests that these 10 species might experience range limitation due to other factors than climate, such as resource quality or resource availability.

Species distribution changes

The species richness pattern for all taxa, produced using the interpolated distributions (ID), differs markedly from the richness pattern using climate affected data (CAD) (Fig. 2). The model predicts that the most species areas in the country will be concentrated on the eastern escarpment, with significant species losses occurring in the western, arid regions (Fig. 2b). A general decline in the size of species areas and a concentration of species around the eastern escarpment was a consistent outcome of the model across all taxonomic groups, and reflects

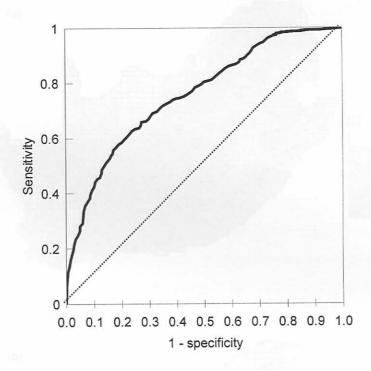


Fig. 1 An example of a ROC curve. The area under the dotted line represents an AUC of 0.5, which would translate into a test with discriminatory ability. The area under the solid black line represents the AUC for this particular species, the South African shelduck.

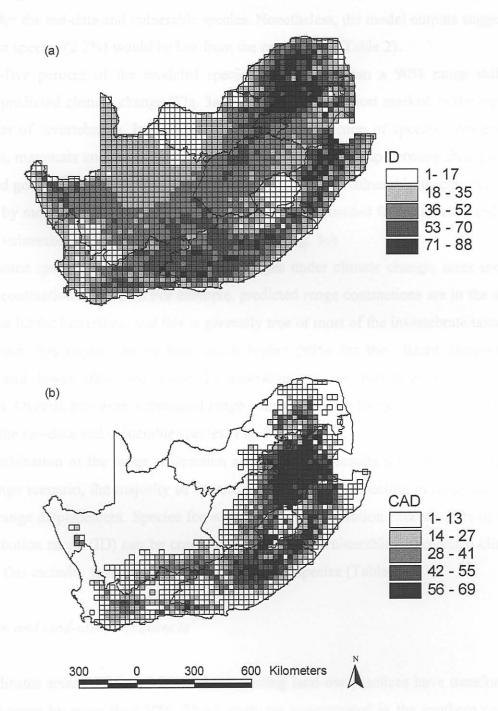


Fig. 2 Species richness patterns for (a) interpolated distributions (ID), (b) climate affected distributions (CAD) for all 179 species included in this study.

to some extent, the likely western movement of most species and also movement of species up altitudinal gradients towards the eastern and south-eastern highlands. Similar patterns were found for the red-data and vulnerable species. Nonetheless, the model outputs suggested that only four species (2.2%) would be lost from the country (see Table 2).

Twenty-five percent of the modeled species show more than a 90% range shift in response to predicted climate change (Fig. 3a). This pattern was most marked in the reptiles and a number of invertebrates. In contrast, a substantial proportion of species from groups such as birds, mammals and butterflies show more than a 50% overlap between their present and predicted geographic ranges. The proportion of Red-data and vulnerable species that shift their ranges by more than 90% approaches 40% vs. the 25% recorded for all species, and few Red-data or vulnerable species displayed no range shift (Fig. 3c).

While some species appear to expand their ranges under climate change, most species show range contractions (Fig. 3b). For example, predicted range contractions are in the order of 40% of the ID for butterflies, and this is generally true of most of the invertebrate taxa. On the other hand, this figure can be both much higher (98% for the lizard *Homopholis wahlbergi*) and lower (0%, see Table 2) depending on the particular taxon under investigation. Overall, however, substantial range contractions are likely to occur, and this is true also of the red-data and vulnerable species (Fig. 3c)

The combination of the range contraction and range shift results suggests that, under a climate change scenario, the majority of species will show both a decline in range size and substantial range displacement. Species for which this range alteration exceeds 50% of their extant distribution range (ID) can be considered particularly vulnerable to predicted climate change, and this includes several red-data and vulnerable species (Table 3).

Conservation and land-use consequences

Figure 4 indicates areas in South Africa where existing land-use practices have transformed natural land-cover by more than 50%. These areas are concentrated in the southern coastal belt (South-Western Cape), across the Northern Provinces (Free State, Gauteng, Mpumalanga, North-West and Northern Province) as well as parallel to the eastern shores (KwaZulu-Natal and Eastern Cape). The areas where most overlap between transformed areas and regions likely to form species richness 'hotspots' under conditions of climate change is likely to occur are in the northern Free State, Gauteng, Mpumalanga and Northern Province. This overlap between climate affected species richness 'hotspots' and areas

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Table 2. Species included in the present study that the model predicted would likely go locally extinct in South Africa following climate change (< 50% probability of occurrence throughout the country).

Species

Stolotermes – termite genus

Epirinus gratus - canthonine dung beetle

Chrysospalax trevelyani – golden mole

Cordylus macrophallus - armoured lizard

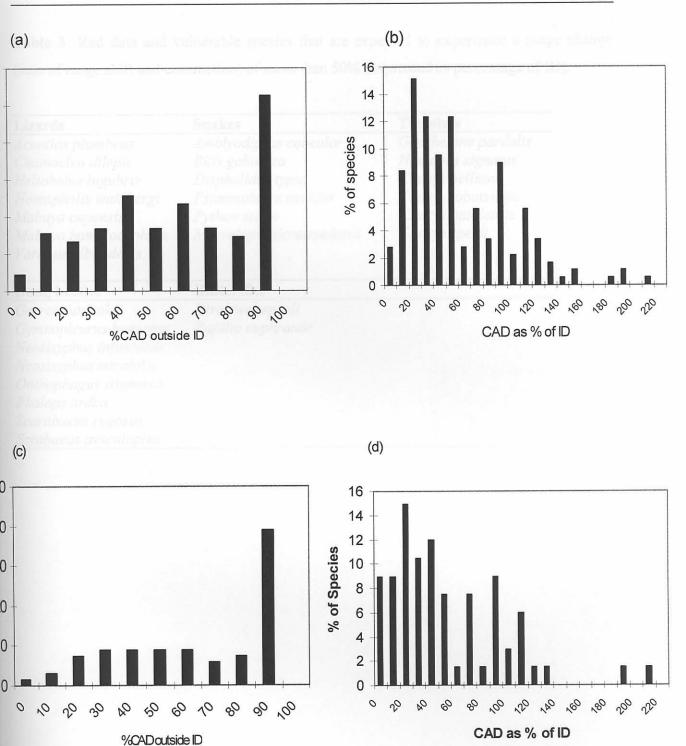


Fig. 3 Percentage of species that exhibit range shifts (expressed as percentage of climate affected distribution outside the interpolated distribution) for (a) all species and (c) red-data and vulnerable species. Percentage of species that exhibit range contraction (climate affected distribution expressed as a percentage of the interpolated distribution) predicted for (b) all species and (d) red data and vulnerable species.

Table 3 Red data and vulnerable species that are expected to experience a range change (sum of range shift and contraction) of more than 50% (expressed as percentage of ID).

Lizards	Snakes	Tortoises
Acontias plumbeus	Amblyodipsas concolor	Geochelone pardalis
Chamaeleo dilepis	Bitis gabonica	Homopus signatus
Heliobolus lugubris	Dispholidus typus	Kinixys belliana
Homopholis wahlbergi	Psammobates oculifer	Kinixys lobatsiana
Mabuya capensis	Python sebae	Kinixys natalensis
Mabuya homalocephala	Macrelaps microlepidotus	Kinixys spekii
Varanus albigularis	A	
Dung beetles	Butterflies	
Garreta unicolor	Dira swanepoeli	
Gymnopleurus humanus	Papilio euphranor	
Neosisyphus infuscatus		
Neosisyphus mirabilis		
Onthophagus stigmosis		
Phalops ardea		
Scarabaeus rugosus		
Scrabaeus aesculapius		

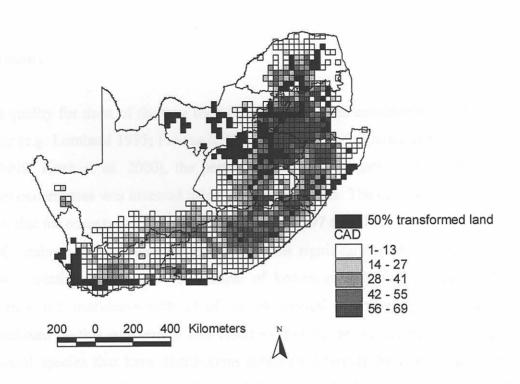


Fig. 4 Predicted climate affected (CAD) species richness patterns and extant land-cover patterns in South Africa. Areas coloured in black are at least 50% transformed and/or degraded.

presently subjected to cultivation and land transformation points to an increased potential for land-use conflict (Wessels *et al.* 2000) between agents of land transformation and conservation advocates.

Discussion

Model performance

Because data quality for most of the taxa used in this study varies considerably, and is often relatively poor (e.g. Lombard 1995; Freitag & Van Jaarsveld 1995; Freitag & Mansell 1997; Hull *et al.* 1998; Koch *et al.* 2000), the performance of the climate model in predicting known species occurrences was assessed using the bird atlas data. The quality of these data is far superior to that for other taxa (compare Harrison *et al.* 1997 and Hull *et al.* 1998).

The ROC analyses showed that the model performs significantly better than a random model. Cross validation with a 20% subsample of known records further supported the model's accuracy and usefulness with 24 of the 34 species' complete distributions being accurately predicted by this subsample. This result might thus be considered an estimate of the proportion of species that have distributions determined largely by climate, and hence provide insight into the broader applicability of the approach. If the model is valid for approximately 70% of the species in the taxa we have selected, then we consider the outcome of this exercise a useful first take at the likely impacts of climate change on animal distributions in South Africa. In some instances, the model did not perform well, especially in predicting presence/absence data for eastern parts of South Africa. This may well be due to the biogeographic complexity of the region and its strong altitudinal gradient that has a considerable influence on animal distribution patterns (see Low & Rebelo 1996; van Wyk 1996; Schulze et al. 1997; Oatley & Arnott 1998; Poynton 1961; White 1983; Poynton & Boycott 1996; Davis 1997; Eeley et al. 1999; Van Rensburg et al. 2000)

Notwithstanding these problems, we are of the opinion that the model used here represents a best estimate for a provisional study of the likely impacts of climate change on the South African higher invertebrate and vertebrate terrestrial fauna. We hold this view for a variety of reasons. First, despite what appear to be poor model performances on the east coast, the mean values for false presences and absences at a countrywide level were generally low, while those for true presence and absence data were not unreasonable (60, and 87%,

respectively). Second, given the resolution of the available data, and the quality of the data available for most taxa, this model is currently the only way in which the potential impacts of climate change can be investigated for a reasonable "shopping basket of taxa" (Hammond 1994).

The effects of climate change: model outputs and biological implications

Under the climate change scenario associated with a doubling in pre-industrial CO₂ levels, impacts on the fauna of South Africa range from minimal (six species showed no change in range size) to severe (four species were predicted to go extinct). Nonetheless, it appears that many species will show a range alteration characterized by a marked range shift from west to east, a somewhat smaller shift from east to west, and a substantial reduction in absolute range size.

Eastward range shifts

Range shifts from west to east are a common prediction of the model, and are characteristic of approximately 41% of all the taxa we examined. Given the pronounced aridity gradient in an east-west direction across the country (Rutherford & Westfall 1994; Schulze et al. 1997), the general decline in species richness in this direction (e.g. Drinkrow & Cherry 1995; Gelderblom et al. 1995; Freitag & Van Jaarsveld 1995, but also Gelderblom & Bronner 1995; Branch et al. 1995), and replacements over this gradient of species that differ markedly in their physiological tolerances (e.g. Le Lagadec et al. 1998), these changes are undoubtedly a realistic reflection of the likely impacts of climate change. In this context it is significant that range shifts as substantial as those predicted here have already been documented over the past century in western Europe (Parmesan et al. 1999), and similar range shifts are predicted elsewhere (e.g. Kerr & Packer 1998).

The extent to which such predicted eastward shifts in range will translate into realized alterations in range position will obviously also vary between taxa. For example, in species that are dependent on surface water for drinking, such as many southern African dove species, eastward shifts in range may not take place at all if agricultural practices continue to involve the surface provision of artesian water to livestock. Likewise, if species are capable of adapting to local conditions by behavioural or physiological means, realised range shifts may not be as pronounced as those predicted. However, information on the relationships

between species ranges and behavioural patterns and physiological tolerances, and the extent to which behavioural and physiological flexibility influence species ranges is limited to just a few South African taxa (e.g. Lovegrove & Wissel 1988; Richter *et al.* 1997; LeLagadec *et al.* 1998).

Landscape alterations in the eastern and central portions of South Africa will also have a marked impact on the extent to which the predicted changes will be realised. Extensive habitat alteration and fragmentation could prevent eastward range shifts from taking place because of the unavailability of suitable habitat patches (see also Hill et al. 1999; Parmesan et al. 1999), and because of large distances between suitable patches (see Brown & Lomolino 1998; Thomas et al. 1999; Tokeshi 1999; Channell & Lomolino 2000 for further discussion of the influence of these parameters on species occurrences). This interaction between species range alterations and habitat transformation is amongst the most significant consequences of climate change in a landscape that has been transformed by humans. If a species is unable to move into an area because of a lack of suitable habitat, or because that area is too distant from the closest source population of that species, then that area is effectively unavailable to the species and local extinction (or extirpation) is the most likely outcome. In effect this means that range contractions predicted by the current model may be underestimates.

Westward range shifts

Although there are good reasons for doubting the ability of the model to predict reliably range changes in the eastern coastal areas, it seems likely that at least some of the predicted westward range shifts will be real, rather than model artifacts. In particular, the shift of species up the altitudinal gradient (i.e. a gradient of increasing water availability and decreasing temperature) in this region does not seem unlikely. Such shifts already take place on a seasonal basis in a number of local avian migrants (see discussion in Harrison *et al.* 1997; Oatley & Arnott 1998), and have been documented in the Neotropics (Pounds *et al.* 1999). At least amongst the ectotherms, such shifts are likely to be more pronounced for coastal than for interior species. It appears that it is an inability to tolerate low temperatures that prevents the more tropical species from expanding their range up the elevational gradient (Gaston & Chown 1999). Thus, an increase in temperature in this region will undoubtedly lead to an expansion of the ranges of the tropical species.

Range alteration as a whole

Of the 179 species we examined, 30 species expanded their ranges, while 143 species showed range contractions varying between 0 and 98%. Few species showed no response or little response in terms of changes to their absolute range size. Because so many of the species we examined here showed substantial range contractions, it is this facet of range alterations that is of most concern. This concern is especially warranted if it is borne in mind that although the range contractions we have predicted here, are in some instances quite dramatic, they may actually be underestimates because landscape transformation has not been explicitly factored into the model. In practice, theoretical range shifts into transformed landscapes may mean local extinction. On the other hand, it is also possible that some of the dramatic predicted range changes may be mitigated by species immigration from outside the modelled area. Ideally, the geographic extent of the model area should be bounded physiographically, i.e. coastal boundaries for terrestrial animals, and not by political boundaries. In the real world, however, data availability dictated the use of political boundaries for this particular study. The effects of this artificial partition in species' ranges on modelled output have yet to be investigated.

The major reasons for significant concerns regarding range contractions have to do with the negative relationship between range size and extinction probability (Jablonski 1991; Gaston 1994). A reduction in the absolute range size of a species will almost inevitably mean an increase in its risk of extinction. There are several reasons for this. First, a decrease in range size will mean that smaller catastrophic events affect a larger proportion of the species total population. If a species is restricted to just a few sites, then a local catastrophic event could easily cause the extinction of that species (Gaston 1994; see also Lawton & May 1995). Second, the generally positive interspecific relationship between population size and range size is such that population size increases at a disproportionately faster rate with increasing range size, and hence local density also increases with increasing range size (Brown 1995, Gaston et al. 1997). Thus a decrease in range size is likely to effect a reasonably rapid decline in population size, and consequently extinction rate will increase for the usual small population reasons, susceptibility to stochastic variation being amongst the most important (see Gaston 1994; Caughley & Gunn 1996). Contraction of species' ranges towards the centre of their original distributions, as predicted by the model used here, is also unlikely to be entirely realistic given landscape transformation. Channell & Lomolino (2000) recently showed that in many of the 245 species they examined, which have recently shown range contractions, the peripheral, rather than central populations of the species have survived. This means that range fragmentation is likely to be more pronounced than that predicted by our model. Such fragmentation is in itself of considerable conservation concern.

Conclusions

In summary, our model suggests that climate change will have a profound impact on terrestrial animal species in South Africa. Moreover, even this assessment is likely to be conservative given that population growth in South Africa is in the order of 1.9% per year (Anonymous 1998), and that the concomitant increases in population density are likely to result in additional, profound landscape transformation (perhaps exacerbated by high and increasing prevalence of AIDS, Williams *et al.* 2000). Mitigation of the impacts of climate change is ultimately a function of political will to confront difficult issues such as land-use and population planning. However, from a conservation and research perspective several actions can be taken. These include: better communication of findings to politicians, substantial improvements in the quality of information on animal diversity and distribution in South Africa, integration of this information into land-use planning, time-series data for at least some taxa, and the need for substantial improvement of knowledge of the causal links between climate and animal distributions. Of these, the first remains most critical.

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Appendix 1 List of 179 species included in the analyses.

Birds Common name	Scientific name	
Anteating Chat	Myrmecocichla formicivora	
Blackchested Prinia	Prinia flavicans	
Blackheaded Canary	Serinus alario	
Blackthroated Canary	Serinus atrogularis	
Blue Korhaan	Eupodotis caerulescens	
Buffstreaked Chat	Oenanthe bifasciata	
Bully Canary	Serinus sulphuratus	
Cape Bunting	Emberiza capensis	
Cape Canary	Serinus canicollis	
Crimsonbreasted Shrike	Laniarius artococcineus	
Dark Chanting Goshawk	Melierax metabates	
Drakensberg Siskin	Pseudochloroptila symonsi	
Dusky Sunbird	Nectarinia fusca	
Forest Canary	Serinus scotops	
Ground Woodpecker	Geocolaptes olivaceus	
Kalahari Robin	Erythropygia paena	
Karoo Korhaan	Eupodotis vigorsii	
Karoo Robin	Erythropygia coryphaeus	
Larklike Bunting	Emberiza impetuani	
Longtailed Shrike	Corvinella melanoleuca	
Ludwig's Bustard	Neotis ludwigii	
Malachite Sunbird	Nectarinia famosa	
Pale Chanting Goshawk	Melierax canorus	
Redcrested Korhaan	Eupodotis ruficrista	
Redeyed Bulbul	Pycnonotus nigricans	
Rock Bunting	Emberiza tahapisi	
Rock Kestrel	Falco tinnunculus	
South African Cliff Swallow	Hirundo spilodera	
South African Shelduck	Tadorna cana	
Stanley's Bustard	Neotis denhami	
Streakyheaded Canary	Serinus gularis	
Tractrac Chat	Cercomela tractrac	
Whitethroated Canary	Serinus albogularis	
Yellow Canary	Serinus flaviventris	

Mammals	
Common name	Scientific name
Cape pangolin	Manis temminckii
Cape serotine bat	Eptesicus capensis
Caracal	Felis caracal
Giant golden mole	Chrysospalax trevelyani
Grant's rock mouse	Aethomys granti
Gunning's golden mole	Amblysomus gunningi
Hottentot golden mole	Amblysomus hottentotus
Meller's mongoose	Rhynchogale melleri
Namaqua rock mouse	Aethomys namaquensis
Red duiker	Cephalophus natalensis
Riverine rabbit	Bunolagus monticularis
Rock dassie	Procavia capensis

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South African lesser bushbaby
Striped polecat
Suricate
Verreaux's mouse
Wahlberg's epauletted fruit bat
Water mongoose
Yellow mongoose

Galago moholi
Ictonyx striatus
Suricata suricatta
Myomyscus verreauxii
Epomophorus wahlbergi
Atilax paludinosus
Cynictis penicillata

Butterflies

Abantis bicolor Acraea natalica Belenois aurota Catopsilia florella Charaxes jasius Colias electo Colotis danae Danaus chrysippus Dira swanepoeli Eurema brigitta Freyeria trochylus Henotesia perspicua Lepidochrysops bacchus Papilio dardanus Papilio demodocus Papilio euphranor Papilio nireus Pontia helice Zizeeria knysna

Reptiles		77.75 (ate
Lizards	Snakes	Tortoises
Acontias plumbeus	Amblyodipsas concolor	Geochelone pardalis
Bradypodium damaranum	Bitis atropos	Homopus areolatus
Chamaeleo dilepis	Bitis caudalis	Homopus boulengeri
Chamaeleo namaquensis	Bitis cornuta	Homopus femoralis
Chamaesaura aneae	Bitis gabonica	Homopus signatus
Chondrodactylus angulifer	Causus defilippii	Kinixys belliana
Cordylus giganteus	Dasypeltis inornata	Kinixys lobatsiana
Cordylus macrophallus	Dispholidus typus	Kinixys natalensis
Heliobolus lugubris	Hemachatus haemachatus	Kinixys spekii
Homopholis wahlbergi	Homoroselaps dorsalis	Psammobates geometricus
Mabuya capensis	Lamprophis aurora	Psammobates oculifer
Mabuya homalocephala	Lamprophis inornatus	Psammobates tentorius
Nucras lalandei	Lycodonomorphus rufulus	
Nucras livida	Macrelaps microlepidotus	
Pedioplanis laticeps	Naja nivea	
Pedioplanis namaquensis	Psammophis subtaeniatus	
Scelotes mirum	Python sebae	
Tropidosaura montana	Typhlops bibronii	
Varanus albigularis	Typhlops fornasinii	

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Other invertebrates Dung beetles	Jewel beetles	Antlions
Dung beetles Chironitis audens Copris cornifrons Epirinus flagellatus Epirinus gratus Garreta unicolor Gymnopleurus humanus Metacatharsius marani Neosisyphus infuscatus Neosisyphus mirabilis Onitis minutus Onthophagus asperulus	Jewel beetles Acmaeodera albovillosa Acmaeodera grata Julodis cirrosa Julodis viridipes Lampetis gregaria Lampetis viridimarginata Megactenodes reticulata Monosacra lalandei Neojulodis tomentosa Phlocteis exasperata Pseudophlocteis vidua	Antlions Cymothales bouvieri Cymothales illustris Golafrus oneili Palparellus festivus Palpares annulatus Palpares cataractae Palpares lentus Palpares sobrinus Palpares sparsus Palpares speciosus Pamexis karoo
Onthophagus binodis Onthophagus cameloides Onthophagus immundus Onthophagus stigmosis Phalops ardea Scarabaeus gariepinus Scarabaeus rugosus Scrabaeus aesculapius	Scaptelytra aliena Sphenoptera brincki Sternocera orissa	
Termites Stolotermes Porotermes Fulleritermes Apicotermes		
Macrotermes (4 species) Termes Allodontotermes Cryptotermes (2 species) Hodotermes		