

The use of habitat models in conservation of rare and endangered leafhopper species (Hemiptera, Auchenorrhyncha)

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Abstract

For conservation of Auchenorrhyncha species, knowledge of their habitat requirements is essential. However, for most species there is no ‘quantitative’ knowledge that would allow e.g. spatially explicit predictions. Such predictions can be made by habitat models, which quantify the relationship between the environment and the occurrence of species. In two plot-based case studies – the endangered leafhopper *Verdanus bensoni* in mountainous grasslands and four endangered Auchenorrhyncha in urban brownfields – we used habitat models to quantify the habitat requirements of these five species and to exemplify their use for creating habitat suitability maps. In the first case study, the multivariate model showed that occurrence probabilities of the leafhopper *V. bensoni* increase with both decreasing nitrogen indicator values and decreasing tree cover. On urban brownfields, successional age was a driving factor for species’ occurrence. Site age largely determines a range of vegetation characteristics, which, in multivariate models, often replaced the variable age. Internal validation showed the robustness of all models. The models allow predictions of habitat quality under different management regimes (e.g. response to fertilization or abandonment for *V. bensoni* or to different turnover rates on brownfield sites). We discuss the application of habitat models in the conservation of Auchenorrhyncha, especially the use of habitat suitability maps.

Introduction

In cultural landscapes, habitat quality for Auchenorrhyncha is often determined by habitat management. In grasslands, management type and intensity (e.g. mowing, grazing, fertilization) are of great importance (Morris 1981; Sedlacek et al. 1988; Nickel and Hildebrandt 2003). For instance, intensively used grasslands may exhibit different plant species composition and vegetation structure than largely undisturbed ones. The presence of certain host plants is a major habitat requirement of many Auchenorrhyncha species. The actual

quality of host plant patches may be largely determined by the amount, architecture and physiology of the host plant (e.g. Prestidge 1982; Moon et al. 2000). For many Auchenorrhyncha species, additional factors like vegetation structure, microclimate or landscape context may be relevant (e.g. Claridge 1986; Denno and Roderick 1991; Haynes and Cronin 2003).

For conservation of Auchenorrhyncha species, knowledge of their habitat requirements is essential. However, for most species there is no ‘quantitative’ knowledge that would allow e.g. spatially explicit predictions. Nickel (2003) presents a

comprehensive review of habitat requirements and host plants of Auchenorrhyncha species in Germany. However, the information is qualitative and descriptive rather than quantified. For instance, *Neophilaenus minor* is described as preferring ‘sparse cover of vegetation’. Since exact figures are not provided it remains unclear whether the optimum is at 20% vegetation cover, or if 50% is still tolerated. For this reason, data-based predictions of habitat suitability, especially at the landscape level, are not feasible.

Such predictions can be made with habitat models. The aims of habitat models are twofold (e.g. Guisan and Zimmermann 2000; Scott et al. 2002). First, habitat models analyze and quantify the relationship between species’ abundance or occurrence and habitat factors. Second, they yield predictions of species’ abundance or occurrence given certain environmental conditions. The latter makes habitat models a potentially powerful tool in nature conservation: models are able to predict the probability of occurrence for sites or landscapes where species distribution data are not available (Wilson et al. 2005). They can also be used to assess the effects of land use changes or succession on habitat quality (Rudner et al. 2005). Habitat suitability maps which can be obtained from habitat models identify potential core habitats of species and form the basis for the planning of nature reserves (e.g. Cabeza et al. 2004). Such predictions of spatial distribution are essential, since conservation planning has to deal with the whole landscape (Wilson et al. 2005).

Here, we use presence–absence data of Auchenorrhyncha species and environmental data to build habitat models based on logistic regression. In two case studies – the rare leafhopper *Verdanus bensoni* in mountainous grasslands and four endangered Auchenorrhyncha in urban brownfields – we (1) demonstrate the procedure of model building, including variable selection, classification and internal validation, (2) quantify habitat requirements of selected species, (3) exemplify the construction of habitat suitability maps, and (4) discuss the application of habitat models in the conservation of Auchenorrhyncha, especially rare and endangered species. Rare here is understood as locally restricted due to rare habitat; the species can well build up considerable densities in their habitats.

Methods

Study sites

Case study 1: Leafhopper Verdanus bensoni

The first case study investigated habitat requirements of the leafhopper *Verdanus bensoni* (China, 1933). It was conducted in the mountain ranges of Dreisessel (1332 m; 48°47′ N, 13°48′ E) and Arber (1456 m; 49°06′ N, 13°08′ E) in the Bavarian Forest, Germany. The climate is characterized by mean annual temperatures between 5 and 6 °C with annual precipitation between 900 (low altitudes) and 2000 mm (high altitudes). For details on climate, geology and soil types see Hofmann (1984). The area is largely covered by forests. While at altitudes up to approx. 1200 m, mixed forests (mainly beech, fir and spruce) are predominant, above this altitude only spruce forests are found. In the valleys and at lower altitudes land use is characterized by a mix of forests, pastures and fields. At higher altitudes only few patches of grassland are scattered within the forests, mainly small pastures (‘Schachten’, see Hofmann 1984) and ski runs.

Verdanus bensoni has a scattered range and is restricted to European mountain ranges (Nickel 2003). Up to now, it has been recorded from the German Alps, Scottish Highlands, Swiss and French Jura, Bavarian and Bohemian Forest, Giant Mountains, and Ural Mountains. In Germany, *Verdanus bensoni* is a rare species in the Alps and the Bavarian Forest. It is listed in the Red Data Book (Remane et al. 1998). In the study area, the Bavarian Forest, *V. bensoni* was recorded above approx. 800 m a.s.l. (Biedermann unpubl.). *V. bensoni* was found in montane and alpine grasslands, probably feeding on grasses (Biedermann 1998; Nickel 2003). However, the specific habitat requirements have not been studied yet in detail.

Case study 2: Endangered species in urban brown-fields

The second case study was carried out on brownfield sites in the city of Bremen, located in the lowlands of northwest Germany (8°44′ N, 53°05′ E, mean temperature 8.8 °C, mean annual precipitation 694 mm). Urban brownfields, previously-developed land within cities, often support a rich wildlife and house a whole range of rare and

endangered species (Gibson 1998; Eyre et al. 2003). They can provide habitat for stenotopic species from semi-natural habitats like dry sandy grasslands (Eversham et al. 1996). Brownfields form highly dynamic habitats (Gibson 1998; Gilbert 1989) which are continuously being generated, quickly changed by successional processes and destroyed by redevelopment. We assume that, within this cycle, each species finds a limited period of time where its habitat requirements are met. In this study, we investigated four endangered species found on brownfield-sites: the leafhoppers *Rhopalopyx vitripennis* (Flor, 1861) and *Macrosteles quadripunctulatus* (Kirschbaum, 1868), the froghopper *Neophilaenus minor* (Kirschbaum, 1868) and the planthopper *Kelisia sabulicola* (W. Wagner, 1952). They are listed as ‘threatened’ or ‘potentially threatened’ (*N. minor*) in Germany’s Red Data Book (Remane et al. 1998).

Sampling design

Case study 1

For the *Verdanus bensoni* study, 42 plots (5×5 m) were chosen at altitudes between 542 m and 1453 m a.s.l., depending on the availability of grasslands or forests with a grass layer. In each grassland or forest the plots were chosen randomly. In each plot the following parameters were measured: total plant cover and cover of the predominant grass species in the herb layer, tree cover, slope, and altitude. Additionally, the mean Ellenberg indicator values (Ellenberg et al. 1992) for moisture and nitrogen were calculated from the plant species composition. The occurrence of *Verdanus bensoni* was recorded by sweep-netting. At each plot, 20 sweeps were taken covering the entire plot. The sweep-netting was repeated three times.

Case study 2

We investigated urban brownfields within 77 km² in the city of Bremen. On the brownfield sites, 157 sample plots of 225 m² were set up in a random stratified way (Guisan and Zimmermann 2000; Hirzel and Guisan 2002; Maggini et al. 2002). Minimum distance between plots was set to 80 m. To ensure that all characteristic types of brownfields got sampled, the plots covered three gradients: site size, age of brownfields (duration of

abandonment) and soil moisture. In 2003, sweep-net sampling was carried out four times between early June and early September, with 100 sweeps each time.

At each study plot we collected a set of environmental parameters. These included several parameters describing vegetation structure, cover of host plants (as specified by Nickel 2003), soil parameters and landscape context. Site age, as time since demolition of buildings or any other severe disturbance that put succession back to zero, we derived from a time series of aerial photographs. Landscape context was assessed using a map of vegetation types. For examples of these vegetation types see Table 3. Within a GIS, we calculated the proportion of each of these types within a certain distance around every plot (Strauß et al. 2004). We tested radii between 25 and 125 m.

For detailed measurement of vertical vegetation structure, we used a white screen, divided in rectangles, that was erected perpendicular to the ground (see Sundermeier 1999). At six points per plot, vertical cover was estimated for each rectangle looking through a 10 cm wide stand of vegetation. From these estimates, height and density parameters were calculated (Table 3) (Sundermeier 1999; Zehm et al. 2003). 50%-height refers to the height below which 50% of the total vegetation cover is located. 75%-height and 90%-height are defined respectively.

Statistical methods of habitat modeling

Logistic regression

We used species’ presence/absence data for model building. A popular approach for modeling such data is using logistic regression (i.e. generalized linear models (GLM) with a logistic link) (Morrison et al. 1998; Guisan and Zimmermann 2000; Hosmer and Lemeshow 2000; Harrell 2001; Reineking and Schröder 2003). Logistic regression has been successfully used in numerous studies on habitat-occurrence relationships (e.g. Peeters and Gardeniers 1998; Guisan et al. 1999; Manel et al. 1999a). Metric variables can be handled along with nominal ones. The shape of the response curve can be either sigmoid or unimodal (‘bell-shaped’), the latter by including second order terms (Peeters and Gardeniers 1998; Hosmer and Lemeshow 2000). The outcome of a logistic regression model is the

occurrence probability at given parameter values. To distinguish between predicted presence and absence, a threshold probability needs to be defined. Predictions should stay restricted to the range of parameter values that has been covered by the study.

Measures of model performance

Numerous measures assessing performance of logistic regression models are available (Hosmer and Lemeshow 2000; Pearce and Ferrier 2000a; Manel et al. 2001). All of them can only describe certain aspects of model performance. Therefore, we used a set of criteria, threshold-independent as well as threshold-dependent (Manel et al. 1999b).

The difference between predicted and observed values (model calibration) was measured by R^2_N (Nagelkerke 1991). Like R^2 in linear regression, it ranges from 0 to 1. On an univariate level, we used R^2_N to compare the relative influence that single predictor variables had on species' presence. Model discrimination was assessed with AUC (Hanley and McNeil 1982), the Area Under the receiver operating characteristic Curve (AUC). AUC values ≥ 0.7 are regarded as acceptable, ≥ 0.8 as excellent, and ≥ 0.9 as outstanding (Hosmer and Lemeshow 2000).

Sensitivity (proportion of correctly predicted presences), specificity (proportion of correctly predicted absences) and CCR (correct classification rate) are classification threshold dependent measures. CCR is easy to interpret, however largely dependent on the rather arbitrary choice of a threshold (Reineking and Schröder 2003) and should be handled with care. As a threshold, we chose P_{fair} , where specificity and sensitivity are equivalent (Hosmer and Lemeshow 2000). Since the species under study are rare and their prevalence is low, P_{fair} ensures that a reasonable proportion of presences will be predicted correctly. On the other hand, this may result in a lower total number of correct predictions (lower CCR) and, in particular, more predicted presences for observed absences (lower specificity) than with other thresholds. For nature conservation, where often the aim will be to correctly predict as many relevant habitats patches as possible (Morrison et al. 1998), we believe that the advantages of P_{fair} out-run these disadvantages.

Since CCR, sensitivity and specificity are highly dependent on the species' prevalence (Manel et al.

2001), we used Cohen's Kappa κ (Cohen 1960) as another, less sensitive threshold-dependent measure (Fielding and Bell 1997). Kappa ranges from 0 to 1 with values between 0.40 and 0.55 indicating fair agreement and values between 0.55 and 0.70 indicating good agreement between observed and predicted values (Monserud and Leemans 1992). For comparison between models we used the information criterion AIC_c , a version of AIC (Akaike's Information Criterion) modified for small samples (Buckland et al. 1997). AIC indicates how well a model performs the trade-off between model fit and model complexity.

Model building

As recommended by Hosmer and Lemeshow (2000), we performed careful univariate analyses prior to building of multivariate models. For each species, we tested univariate models of all variables. Only significant variables ($p \leq 0.05$) with $R^2_N \geq 0.05$ were considered for further analysis.

A popular approach for building multivariate models uses stepwise procedures for variable selection. Pearce and Ferrier (2000b) recommend the stepwise backward procedure, which we used for the *Verdanus bensoni* study. In general, all stepwise procedures have some disadvantages (Reineking and Schröder 2004). They might not find the best model, or selection is unstable and does not hold for slightly different data. With a large number of predictor variables, like in the brownfield study, stepwise procedures perform poorly. Therefore, in that study, we followed a different approach: we calculated models for all combinations of four, three and two parameters, using Splus 6.1 functions `glm` and `stepAIC` (MASS library). Since the ratio 'number of observations'/'predictor variables' should not fall much below 10 (Morrison et al. 1998; Guisan and Zimmermann 2000), more than four variables per model are not a sound choice for the available data sets.

Strong correlations between predictor variables will lead to abnormally high coefficients and standard errors (Neter et al. 1989). Therefore, maximum spearman rank correlation (r_s) between predictor variables within one model was allowed to be 0.7 (Fielding and Haworth 1995). Since height and density parameters in the brownfield study showed strong correlations, only one of each group was chosen for multivariate modeling.

Model validation

Performance criteria are usually over-optimistic if they are calculated on the same data set that was used for parameter estimation (Reineking and Schröder 2003). Since independent data were not available to correct for this optimism, we used the bootstrap as an internal validation method (Verbyla and Litaitis 1989; Efron and Tibshirani 1993) for evaluating the models. According to Steyerberg et al. (2001) and Harrell (2001), it outperforms other internal validation procedures and allows nearly unbiased estimates of model performance. We performed the bootstrap with Splus 6.1, doing 300 iterations, resulting in corrected measures of model performance.

Habitat suitability maps

Habitat suitability maps can be obtained by applying the regression equations of habitat models to maps of the relevant environmental data within a GIS. These maps spatially explicitly predict the probability of occurrence of the focal species (Osborne et al. 2001; Austin 2002; Joy and Death 2004). Models used for such spatially explicit predictions are restricted to parameters that are available area-wide. In the brownfield-study, these were age of brownfield sites, and all landscape context parameters. For *N. minor*, we calculated a model from these parameters and applied it to part of the study area.

Results

Univariate models

Case study 1

The univariate logistic regression analysis revealed that a number of significant habitat parameters were related to the incidence of *Verdanus bensoni* (Table 1). The occurrence of *V. bensoni* was positively related to altitude and moisture indicator and negatively to nitrogen indicator and tree cover. The habitat parameters slope and total plant cover showed no effect on the occurrence of *V. bensoni*. Likewise, the cover of single grass species had no positive influence on the occurrence of *V. bensoni*.

Case study 2

For the brownfield study, univariate responses for all relevant variables are listed in Table 3. A total

of 29 predictor variables passed the performance criteria. Age was a strong predictor for all four species. *M. quadripunctulatus* showed a sigmoid response, occurrence probability decreasing with increasing age (Figure 2), whereas the other species showed unimodal responses with peaks between 13 and 20 years.

Vegetation height did not play an important role for *M. quadripunctulatus*. *K. sabulicola*, *N. minor* and *R. vitripennis* showed similar, mostly unimodal responses to vegetation height and density parameters. *R. vitripennis* made an exception in preferring high density in the lowest layer. For *M. quadripunctulatus*, high overall density decreased occurrence probability, whereas it preferred moderate densities within the lower vegetation layers.

Most species exhibited strong relationships with moss cover, litter cover and bare ground. As with density and height, *K. sabulicola* was negatively correlated with moss and litter cover, whereas the other species preferred medium to high values for these parameters. In general, high covers of the respective host plants strongly enhanced occurrence probabilities. PH was the most important amongst the soil parameters. *M. quadripunctulatus* preferred high, whilst *N. minor* and *R. vitripennis* preferred medium levels.

Overall, the influence of landscape context was comparatively weak with two exceptions. Occurrence of *N. minor* increased with rising proportions of brownfields with grassy, sparse vegetation. *M. quadripunctulatus* showed an unimodal response to the proportion of open brownfields with < 10% vegetation cover. For all species, R^2_N of landscape context was highest for the 75 m-radius.

Table 1. Case study 1: Univariate responses of the leafhopper *Verdanus bensoni* to various habitat parameters.

| Parameter | Range | Response of <i>V. bensoni</i> | |
|--------------------|------------|-------------------------------|----|
| | | R^2_N | |
| Altitude | 542–1453 m | 0.43 | +S |
| Nitrogen indicator | 2–6 | 0.41 | –S |
| Moisture indicator | 5–7 | 0.34 | +S |
| Tree cover | 0–100% | 0.14 | –S |

–S: sigmoid response, occurrence probability decreases with increasing values of predictor variable; +S: sigmoid, occurrence probability increases with increasing values of predictor variable.

Multivariate models

Case study 1

The multivariate habitat model for *V. bensoni* contained two significant habitat parameters (Table 2). The model showed that with both decreasing nitrogen indicator values and decreasing tree cover the incidence of *V. bensoni* increased (Figure 1). Model discrimination was good (Table 2): in 85% of the plots occurrence of *V. bensoni* was correctly classified.

Case study 2

The final multivariate models for the brownfield species contained three or four (*N. minor*) explanatory variables (Table 3). Model performance measures are given in Table 5, coefficients, standard errors and *p*-values in Table 4.

The model for *R. vitripennis* included the parameters age, moss cover and cover of *Festuca rubra/ovina*. Occurrence probabilities were highest at medium levels of age and moss cover (Figure 3). With increasing cover of *Festuca*, the influence of these parameters became negligible; occurrence probabilities always exceeded the threshold.

Occurrence of *N. minor* could be explained best with a four-parameter model. Occurrence probabilities above the threshold were restricted to low, but non-zero 50%-heights, regardless of the other parameter values (Figure 4). Moderate litter covers were preferred in combination with low cover of *Corynephorus canescens* and low proportions of

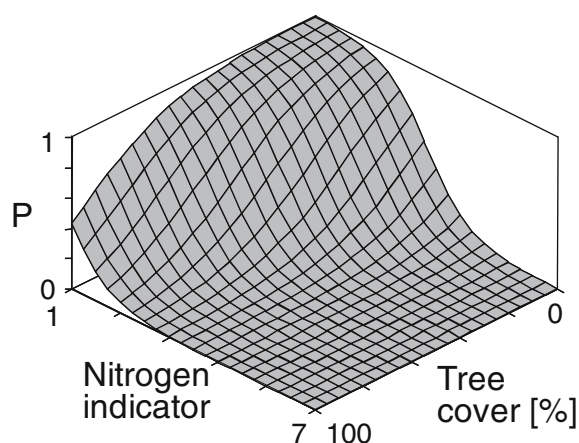


Figure 1. Multiple habitat model for *Verdanus bensoni*. Probability of occurrence (*P*) is plotted against nitrogen indicator and tree cover.

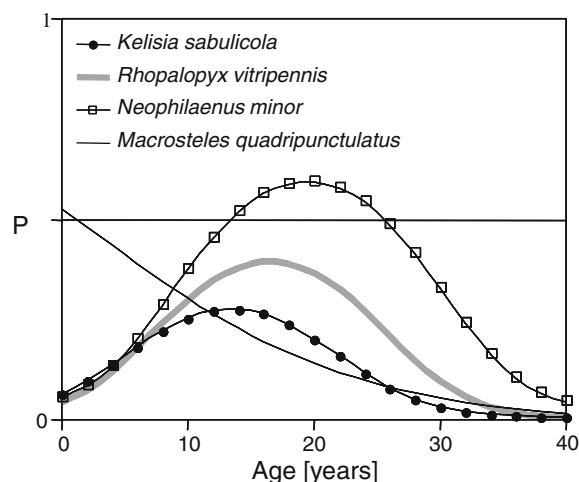


Figure 2. Univariate response curves for the variable 'age'.

Table 2. Case study 1: Multiple habitat model of the leafhopper *Verdanus bensoni*.

| Model parameters & coefficients | |
|---------------------------------|-------------|
| Nitrogen indicator | -1.94218 |
| Tree cover | -0.05667 |
| Intercept | 7.33042 |
| Model performance | |
| Significance | $p < 0.001$ |
| R^2_{Ncorr} | 0.56 |
| AUC_{corr} | 0.88 |
| Classification P_{fair} | |
| P_{fair} | 0.48 |
| κ | 0.72 |
| Sensitivity | 0.88 |
| Specificity | 0.85 |
| CCR | 0.85 |

Model parameters, model performance and classification using the threshold P_{fair} .

BGS75. With increasing values of either or both of these parameters, the modifying influence of litter cover decreased. *M. quadripunctulatus* reached high occurrence probabilities only at moderate to high pH-levels and in combination with both little to no litter cover and medium proportions of BO75 (Figure 6). The shape of the response surface of *Kelisia sabulicola* strongly depended on the cover of *Carex arenaria*. High values of 10% led to occurrence probabilities close to one, regardless of the other two factors (Figure 5). At low cover of *Carex*, presence depended on medium levels of age and vegetation density in the 0–5 cm layer. Model performance was better for *R. vitripennis*, *N. minor* and *M. quadripunctulatus* ($R^2_N > 0.41$, AUC

Table 3. Case study 2: Univariate responses: R^2_N and shape of response curves. R^2_N of variables included in best multiple models printed bold.

| Variable/Parameter | Range | <i>M. quad.</i> | | <i>K. sabul.</i> | | <i>N. minor</i> | | <i>R. vitrip.</i> | |
|------------------------------------------------------------------------------------------------------------------------------|---------|-----------------|----|------------------|----|-----------------|----|-------------------|----|
| | | R^2_N | | R^2_N | | R^2_N | | R^2_N | |
| Age [years] | 0–33 | 0.16 | –S | 0.13 | U | 0.19 | U | 0.21 | U |
| Vegetation height [cm] | | | | | | | | | |
| Veg. height | 0–110 | | | | | | | 0.22 | U |
| Weighted height | 0–24 | | | 0.09 | U | 0.10 | U | 0.18 | U |
| Max. height | 0–117 | | | 0.05 | +S | | | 0.20 | U |
| 50%-height | 0–10 | | | | | 0.14 | U | | |
| 75%-height | 0–28 | | | | | 0.11 | U | 0.08 | U |
| 90%-height | 0–63 | | | 0.06 | U | | | 0.13 | U |
| Vegetation density [%] | | | | | | | | | |
| Veg. cover (horizontal) | 0–90 | 0.05 | –S | 0.14 | U | | | 0.22 | U |
| Veg. density (vertical) | 0–21 | 0.08 | –S | | | | | 0.22 | U |
| Veg. dens. 0–5 cm | 0–92 | 0.16 | U | 0.14 | U | 0.07 | U | 0.18 | +S |
| Veg. dens. 5–15 cm | 0–66 | 0.14 | U | 0.13 | U | 0.07 | U | | |
| Veg. dens. 15–50 cm | 0–28 | 0.06 | –S | | | | | 0.13 | U |
| Veg. dens. 50–100 cm | 0–20 | | | | | | | 0.07 | U |
| Density variation | 0–7 | | | | | | | 0.11 | U |
| Other vegetation parameters [%] | | | | | | | | | |
| Moss cover | 0–100 | 0.23 | –S | 0.05 | +S | 0.05 | +S | 0.23 | U |
| Litter cover | 0–100 | 0.27 | –S | | | 0.08 | U | 0.18 | +S |
| Bare ground | 0–100 | 0.20 | U | 0.12 | –S | 0.06 | –S | 0.18 | –S |
| Cover of host plants [%] | | | | | | | | | |
| <i>Festuca rubra/ovina</i> | 0–88 | | | | | 0.05 | +S | 0.40 | +S |
| <i>Carex arenaria</i> | 0–19 | | | 0.20 | +S | | | | |
| <i>Corynephorus canescens</i> | 0–38 | | | | | 0.29 | +S | | |
| Soil | | | | | | | | | |
| Effective cation exchange capacity | 2–15 | | | 0.09 | U | | | 0.07 | U |
| pH | 3.4–7.7 | 0.24 | +S | | | 0.18 | U | 0.12 | U |
| Stone content (topsoil) | 0–6 | 0.08 | +S | 0.07 | U | 0.17 | –S | | |
| Available water capacity | 4–193 | | | | | 0.08 | +S | 0.05 | +S |
| Landscape context: Proportion of brownfields, covered with a certain structural vegetation type, within a radius of 75 m [%] | | | | | | | | | |
| Open (<10% veg. cover) ('BO75') | 0–100 | 0.14 | U | | | | | 0.05 | –S |
| Grassy, sparse veg. ('BGS75') | 0–100 | | | 0.07 | +S | 0.36 | +S | | |
| Grassy, dense veg. | 0–82 | 0.06 | –S | | | | | 0.07 | U |
| Herbaceous, sparse veg. | 0–98 | 0.09 | U | | | | | 0.06 | –S |
| Bushes/hedges | 0–22 | | | | | | | 0.08 | U |

–S: sigmoid response, occurrence probability decreases with increasing values of predictor variable; +S: sigmoid, occurrence probability increases with increasing values of predictor variable; U: unimodal response.

> 0.84) than for *K. sabulicola* ($R^2_N = 0.29$, AUC = 0.77, Table 3).

Habitat suitability map

The habitat suitability map for *N. minor* (Figure 8) was based on a two-parameter model with age and BGS75. Occurrence probability steeply rose with

increasing proportions of BGS75, in particular in combination with medium age (Figure 7). As the threshold was low (0.13), most of the response surface was above the threshold. Nevertheless, large proportions of the brownfield sites (62%) have low values for BGS75 combined with young age, resulting in occurrence probabilities below the threshold, shown as white regions on the map. The model yielded poorer performance than the best

Table 4. Case study 2: Model performance of multiple models. All performance measures corrected by bootstrapping. (+[^]2) indicates that the second order term is included to model an univariate response.

| Species | pres./abs. | Model parameters | Threshold independent | | Threshold dependent: P _{fair} | | | CCR | |
|--------------------------------------|------------|---------------------------------------------------------------------------------------------|-----------------------------|------|----------------------------------------|------|-------------|------|-------------|
| | | | R ² _N | AUC | P _{fair} | κ | Sensitivity | | Specificity |
| <i>Rhopalopyx viripennis</i> | 33/124 | Age (+ [^] 2) + Moss%(+ [^] 2) + Festuca.rubra/ovina% | 0.42 | 0.85 | 0.21 | 0.49 | 0.77 | 0.81 | 0.80 |
| <i>Neophilaenus minor</i> | 25/132 | 50%-height (+ [^] 2) + Litter%(+ [^] 2) + Corynephorus.canescens% + BGS75 | 0.50 | 0.90 | 0.15 | 0.43 | 0.75 | 0.81 | 0.80 |
| <i>Neophilaenus minor</i> | 25/132 | Age (+ [^] 2) + BGS75 | 0.38 | 0.83 | 0.13 | 0.36 | 0.73 | 0.78 | 0.77 |
| <i>Macrosteles quadripunctulatus</i> | 58/99 | Litter% + pH + BO75 (+ [^] 2) | 0.41 | 0.84 | 0.41 | 0.52 | 0.76 | 0.78 | 0.77 |
| <i>Kelisia sabulicola</i> | 30/127 | Age (+ [^] 2) + Veg.dens.0-5 cm (+ [^] 2) + Carex.arenaria% | 0.29 | 0.77 | 0.17 | 0.28 | 0.67 | 0.72 | 0.70 |

model for the species (Table 5), with R^2_N at 0.38 and AUC at 0.83. Sensitivity and specificity were at 0.76 and 0.77, respectively. Out of the 28 plots within the section shown in the map (Figure 8), all nine presences were predicted correctly, six of the 19 absences were classified incorrectly as presences.

Discussion

Case study 1

For *V. bensoni*, the most important habitat factor was found to be the fertility of the grassland sites. *V. bensoni* was restricted to low productivity sites. Consequently, agricultural intensification and fertilization of the low productivity habitats would pose a threat to *V. bensoni*. Further, the occurrence of *V. bensoni* would decrease if the tree cover of grassland sites increased, for instance, after abandonment of mowing or grazing. There was no relationship between the occurrence of *V. bensoni* and the cover of single grass species. It was known from literature that *V. bensoni* lives on grasses and it has been argued that *V. bensoni* may use several grass species as host plants (Biedermann 1998; Nickel 2003). Our results confirm that *V. bensoni* obviously is not a host plant specialist like, for instance, *Neophilaenus minor*.

Case study 2

Within our dataset, age of brownfield sites was the most driving factor determining species' occurrence. This agrees with the results of Small et al. (2003) for carabid assemblages, who found that time since the last disturbance has a significant influence on species' occurrence. In the study by Brown et al. (1992), successional age had a strong effect on leafhopper assemblages. Characteristic stages of brownfield succession strongly depend on time (Gilbert 1989), but substrate can modify succession rates considerably (Gilbert 1989; Small et al. 2003). The main difference between successional stages lies in their vegetation structures (Hollier et al. 1994). This might be the reason why in two of the 'best' multivariate models, age was substituted by vegetation parameters. They prob-

Table 5. Case study 2: Coefficients and *p*-values of the multiple models.

| | Coeff. | S.E. | <i>p</i> |
|----------------------------------------------------|----------|---------|----------|
| <i>Macrosteles quadripunctulatus</i> | | | |
| Intercept | -5.09507 | 1.32829 | <0.01 |
| BO75 | 0.04916 | 0.03337 | 0.14 |
| BO75 ² | -0.00080 | 0.00040 | 0.04 |
| Litter | -0.03818 | 0.00997 | <0.01 |
| ph | 0.87389 | 0.21201 | <0.01 |
| <i>Rhopalopyx vitripennis</i> | | | |
| Intercept | -3.97124 | 0.74998 | <0.01 |
| Age | 0.20959 | 0.11836 | 0.08 |
| Age ² | -0.00703 | 0.00346 | 0.04 |
| Moss.cover | 0.09384 | 0.03630 | 0.01 |
| Moss.cover ² | -0.00101 | 0.00040 | 0.01 |
| <i>Festuca rubra/ovina</i> | 0.06775 | 0.01595 | <0.01 |
| <i>Neophilaenus minor</i> | | | |
| Intercept | -9.80625 | 3.26620 | <0.01 |
| 50%-height | 3.18859 | 1.65463 | 0.05 |
| 50%-height ² | -0.48593 | 0.22308 | 0.03 |
| Litter cover | 0.14831 | 0.04900 | 0.00 |
| Litter cover ² | -0.00140 | 0.00052 | 0.01 |
| <i>Corynephorus canescens</i> | 0.23070 | 0.11937 | 0.05 |
| BGS75 | 0.06173 | 0.01633 | <0.01 |
| <i>Neophilaenus minor (suitability map)</i> | | | |
| Intercept | -3.40644 | 0.57092 | <0.01 |
| Age | 0.06440 | 0.02664 | 0.02 |
| Age ² | -0.00197 | 0.00122 | 0.11 |
| BGS75 | 0.05627 | 0.01057 | <0.01 |
| <i>Kelisia sabulicola</i> | | | |
| Intercept | -4.61357 | 0.95524 | <0.01 |
| Age | 0.20585 | 0.10274 | 0.05 |
| Age ² | -0.00432 | 0.00292 | 0.14 |
| Veg.dens.0-5 cm | 0.10049 | 0.04486 | 0.03 |
| Veg.dens.0-5 cm ² | -0.00117 | 0.00051 | 0.02 |
| <i>Carex arenaria</i> | 0.64897 | 0.24579 | 0.01 |

ably represent the specific conditions of a particular site more accurately.

Vegetation structure is known to strongly affect species composition of Auchenorrhyncha communities (Murdoch et al. 1972; Denno and Roderick 1991; Achtziger 1995). We assume that vegetation structure is also an indirect measure for a site's microclimate. Sparse vegetation causes more extreme conditions in terms of temperature and moisture (Biedermann 1997, Geiger et al. 2003). Soil conditions influence both plant species composition and food quality of plants (Schoonhoven et al. 1998). The effect of soil conditions on Auchenorrhyncha was shown by Sanderson et al. (1995).

Landscape context we believe to indirectly represent several factors. First, it is a measure for site

isolation. If proportions of favored habitat types are low or those of unsuitable habitat are high, the site is likely to be isolated and thus less likely to be occupied (Haynes and Cronin 2003; Biedermann 2004). Second, landscape context is an indicator for patch size: large proportions of favorable habitat types represent large patch sizes. Large patches have a higher probability of being occupied (e.g. Biedermann 2002). The positive correlation between *N. minor* and proportion of brownfields with grassy, sparse vegetation is probably due to either of these two factors. Third, surrounding vegetation influences a site's microclimate. Bushes and hedges slow down wind and thus provide more balanced, warmer and moister conditions. Sparse vegetation does the opposite. The comparatively small influence of landscape

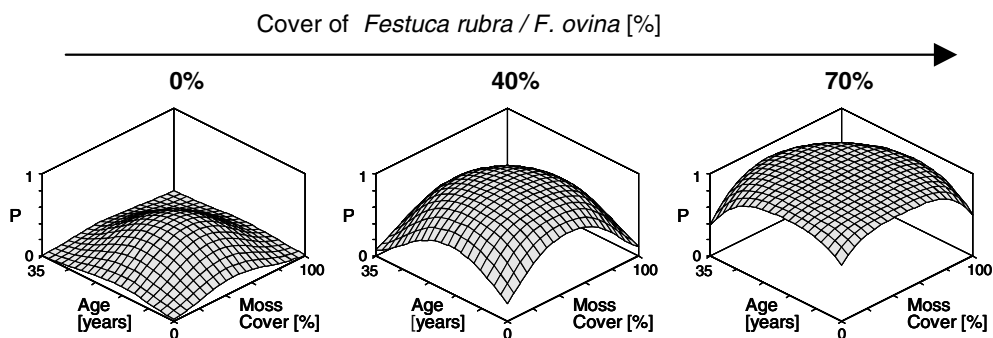


Figure 3. Best habitat model for *Rhopalopyx vitripennis*. Occurrence probability (P) on the z-axis, against age and moss cover. Three levels of *Festuca rubra/ovina*-cover are represented in the three diagrams.

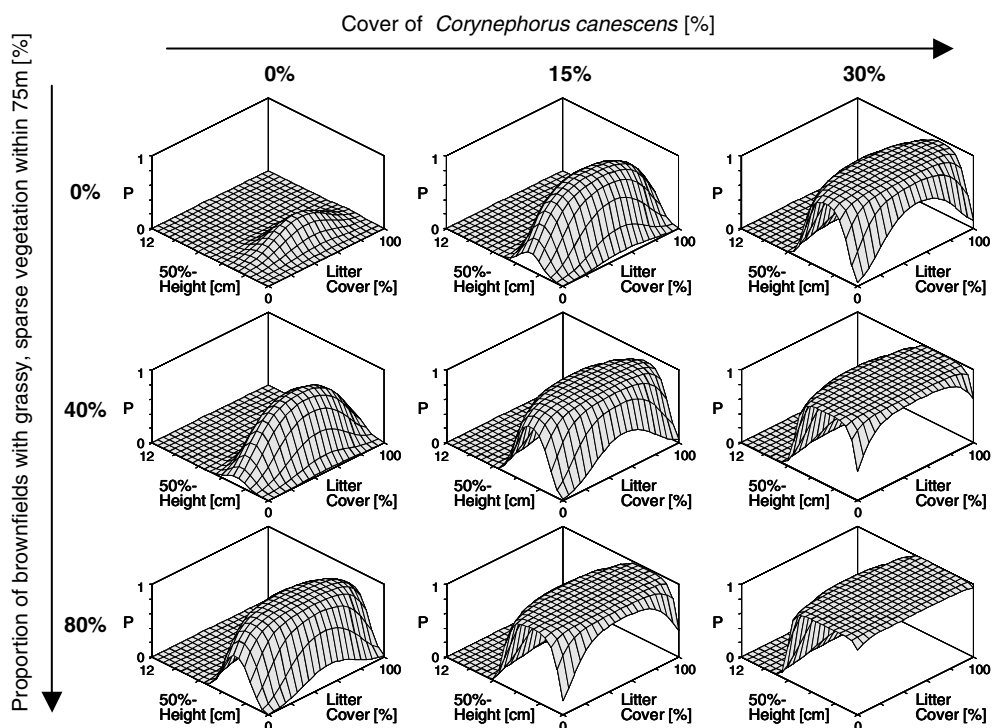


Figure 4. Best habitat model for *Neophilaenus minor*. In each diagram, P is plotted against 50%-Height and litter cover. Columns represent different levels of *Corynephorus canescens*-cover and rows represent different proportions of BGS75.

context might be due to two factors. First, most Auchenorrhyncha species seem to not need large sites to build up viable populations (Biedermann 2002, 2004; Cronin 2004). Second, it is likely that for Auchenorrhyncha, most brownfield sites are not truly isolated. Small patches of potential habitat are found along most roads and tracks and connect the larger sites.

Overall, univariate responses of all four species corresponded well to habitat requirements described in Nickel (2003). For instance, *M. quadripunctulatus* is regarded as a pioneer species preferring sandy, sparsely vegetated and moderately dry to dry sites. This agrees with our results that the species was restricted to young sites with very scarce vegetation.

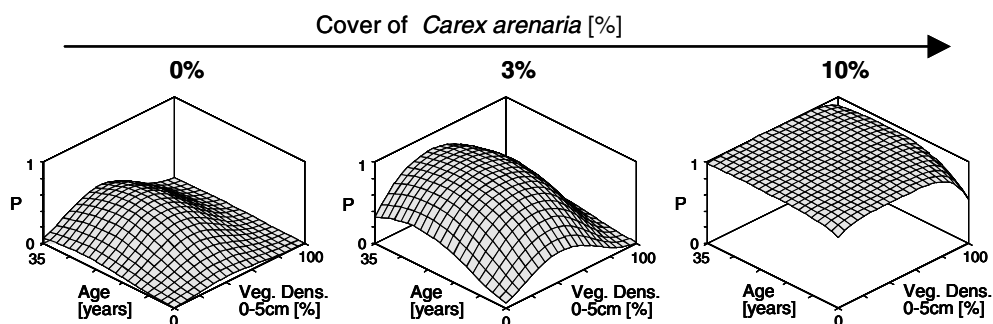


Figure 5. Best habitat model for *Kelisia sabulicola*. P is plotted against age and vegetation density 0–5 cm; diagrams represent different levels of *Carex arenaria* cover.

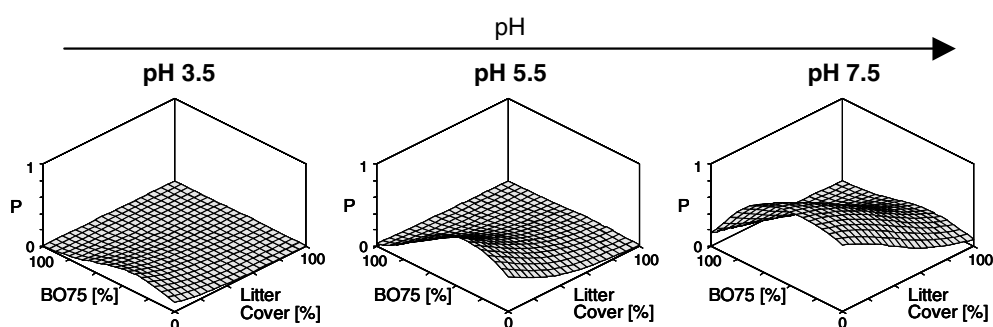


Figure 6. Best habitat model for *Macrosteles quadripunctulatus*. P plotted against BO75 and litter cover; diagrams represent different levels of pH.

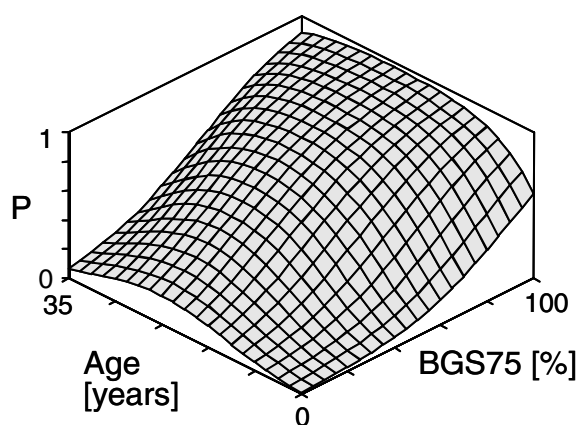


Figure 7. Model used to calculate a habitat suitability map for *Neophilaenus minor*. P plotted against age and BGS75.

The use of habitat models in conservation

Internal validation showed the robustness of habitat models within our studies. However, in perspective, it would be desirable to validate these

models externally, i.e. apply them to independent data sets from other landscapes. In this respect, it would be interesting to test whether the habitat model of *Verdanus bensoni* from the Bavarian Forest is applicable to Alpine populations. The transfer of habitat models has been successfully demonstrated in other insects (Kuhn and Kleyer 1999/2000; Schröder and Richter 1999/2000; Bonn and Schröder 2001; Binzenhöfer et al. 2005). Unfortunately, up till now there have been no attempts with Auchenorrhyncha. However, transferability is regarded as a prerequisite for the broad application of habitat models in the conservation of Auchenorrhyncha.

The habitat models presented here are able to predict the quality of habitats under different management. In the leafhopper *Verdanus bensoni*, the habitat model predicts the response to fertilization or abandonment. In the urban brownfield study, the effects of different turnover rates became obvious. Models are able to predict the occurrence of species along a temporal gradient of

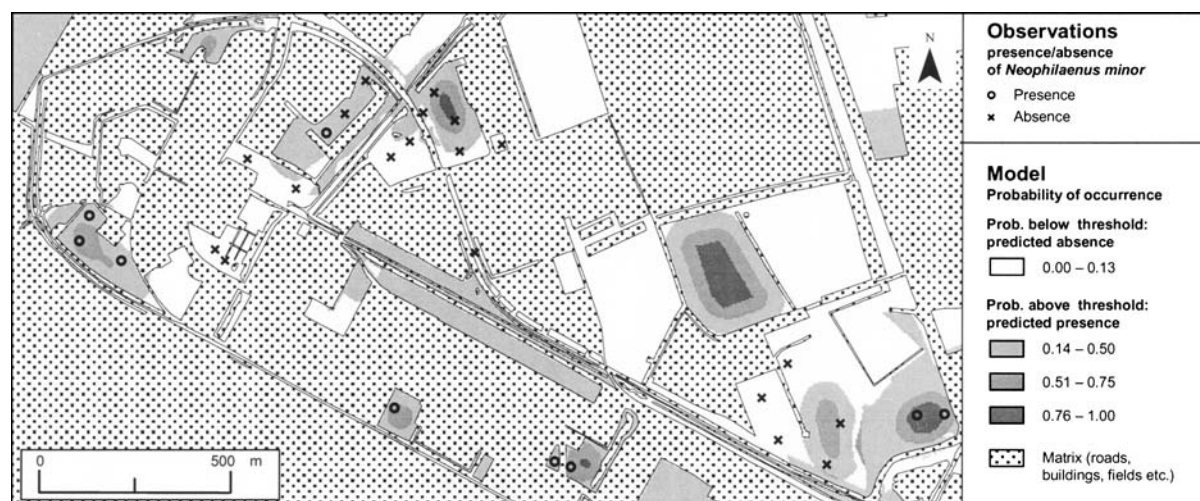


Figure 8. Habitat suitability map for *Neophilaenus minor*. Non-brownfield sites (sealed areas: e.g. roads, buildings, container parks; farmland: mostly wet grassland) are considered *per se* as unsuitable habitats (matrix) and dotted on the map.

succession. This quantitative information may be essential for the conservation of species in brownfields. It was shown that the species under study are restricted to the early or intermediate stages of brownfield succession. Once sites become too old, these species are likely to disappear. Small et al. (2003) found for carabid beetles that the most species rich assemblages are found on early successional sites that can be between 6 and 20 years old. For conservation, this implies that protection of existing brownfield-sites without management will cause many species to disappear over time. In order to preserve high biodiversity, one should focus on the duration of the brownfield stage within the cycle of emergence, succession and demolition of brownfields sites. A constant stock of brownfields of young and intermediate age within an industrial area preserves the typical species assemblage.

The case studies also showed that in some species it is possible to build habitat models with good performance using only a few habitat parameters. For instance, in *Verdanus bensoni* two parameters were sufficient to reach a high correct classification rate. For conservation purpose, those habitat models may be a tool to identify potential habitat relying on only a small number of environmental parameters. Even though a large number of parameters might be necessary to detect the driving forces and build well performing models, once these parameters

are known, models can easily be applied to other regions, assuming the availability of data for the parameters. In the light of increasing availability of area-wide environmental data (e.g. from satellite imagery or public GIS databases) this prerequisite will be easier to meet in future. However, some variables, like the ones describing aspects of vegetation structure in a detailed way, can not be obtained area-wide by these methods. Still, these variables are of great importance when studying a species' ecological needs. Hence, for habitat suitability maps, these variables have to be substituted by ones that are available area-wide. The application of habitat suitability maps in conservation may easily identify and map areas for protection (e.g. Cabeza et al. 2004). However, there are some issues to consider when applying habitat models and habitat suitability maps. First, species' absences can never be recorded with the same certainty as species' presences. Kleyer et al. (1999/2000) suggest regarding presence and absence as a species-specific characteristic. Second, false-positive predictions do not necessarily indicate a poor model fit, since plots recorded as non-use are not always unsuitable habitat (Capen et al. 1986). This is particularly true in declining populations, where many false-positive predictions might result (Wilson et al. 2005): due to an increased extinction rate, suitable habitat might not be inhabited. Thus, habitat suitability maps may help to identify areas for the

reintroduction of endangered or rare species by showing potentially suitable habitat.

In conclusion, this study demonstrated the building and application of habitat models for Auchenorrhyncha. Although further research is needed, especially on the generality of single species habitat models, the value of habitat models for conservation seems obvious. The use of habitat suitability maps could find broad application in future.

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