# Regional and local environmental correlates of native Iberian fish fauna 

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#### Abstract

Knowledge of fishes' habitat requirements is an essential tool for species conservation. We examined the habitat requirements of nine native freshwater fish species using 165 undisturbed or minimally disturbed sites from eight river basins in western Iberia (northern and central Portugal). Six environmental descriptors were considered at the regional level, and six descriptors at a local scale. Statistical associations between local assemblage structure and environmental variables were quantified by canonical correspondence analyses. Logistic regression was performed for each species to identify regional and local factors related to its distribution at the sample sites. The results suggested a reasonable degree of specialisation in the use of local variables and dependence on regional variables. Logistic regression models showed a good adjustment ( $0.75<$ AUC $<0.93$ ). Results of habitat modelling using minimally disturbed sites helped to clarify species preferences for specific abiotic drivers.


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## Introduction

Freshwater ecosystems have frequently been cited as among those that are most altered and threatened by human activities (Mason 1991; Saunders et al. 2002; Clavero et al. 2004). Although initial approaches to decline and extinction phenomena were mostly centred on terrestrial organisms, more recently increasing attention has been paid to the conservation of freshwater fauna (Harding et al. 1998; Ricciardi \& Rasmussen 1999). However, to manage aquatic biodiversity we must understand how fish species are distributed and be able to predict their occurrence in a given area (Ormerod 2003). Knowledge of fishes' habitat requirements is an essential tool for species conservation and fishery management (Heggenes et al. 1991), but the relationship between fish species and fish assemblages and habitat is an extremely complex one (Hubert \& Rahel 1989).

The use of different uni- or multivariate methods to study species distribution and habitat requirements has increased over the last 20 years (Leps \& Smilauer
2003). Regression models in particular have been widely used to predict species distribution, abundance and habitat preference (e.g., Beja \& Alcazar 2003; Filipe et al. 2004; Moreira et al. 2005). As predicted by Margalef (1968), presence/absence data permit the detection of macroscale patterns in community ecological studies. However, most studies either consider sites with different degrees of human disturbance, where populations may thus be in a suboptimal habitat, or else include human-mediated variables to describe the habitat. A valid understanding of natural patterns and processes is an essential prerequisite for viable ecosystem management (Stanford et al. 1996). We have therefore focused our research efforts on investigating patterns and processes in riverine landscapes that are in a near-natural state.

A number of studies indicate that fish assemblage organisation can be determined not only by local processes, but also by large-scale environmental factors (Schlosser 1991; Marsh-Matthews \& Matthews 2000; Magalhães et al. 2002; Hoeinghaus et al. 2006). The importance of the scale of environmental variables
that determine the distribution of individual fish species has not received as much attention (Taylor 1996; Wiley et al. 1997), because most population studies are focused on local conditions, a particular river or a specific region (e.g., Bovee 1982; Gore \& Nestler 1988; Leftwich et al. 1997). In some cases, it has been possible to model the occurrence of fish species at the individual drainage basin scale (Oberdorff et al. 2001; Filipe et al. 2004), and to test the hypothesis of a common response to local and regional environmental variables (Pont et al. 2005). In addition, local habitat characteristics, such as water volume, velocity and thermal regime, are often dependent on regional drivers, such as drainage area, hydrological regime and altitude (Poff \& Allan 1995; Wiley et al. 1997). The relative role and importance of scale factors in shaping and regulating species distribution can also be related to integrity assessment and conservation efforts, because these depend on speciesspecific perturbation drivers and the spatial scale upon which they act.

In this study we aim: (i) to look at the habitat requirements of nine native Iberian fish species by identifying their environmental correlates at undisturbed or minimally disturbed sites; and (ii) to assess the relative importance of these local and regional environmental variables on the occurrence of each species.

## Study area

The study area is located in western Iberia and includes the hydrological network of central and northern Portugal (Fig. 1). These rivers flow NE-SW towards the West Atlantic coast, and the largest basins (Minho, Lima, Douro and Tagus) are shared with Spain. The inland area is dominated by the preMesozoic Hesperic Massif - a complex geological unit that includes granites, schists and quartzites with a various degrees of metamorphism - whereas most of the western coastal fringe is characterised by Tertiary layers under Quaternary deposits. The southern region below the main course of the River Tagus is dominated by the flat platforms of the Hesperic Massif, and in the lower boundary by the Meso-Cenozoic strata, which is composed of sedimentary deposits (e.g., sandstones, limestones, conglomerates and marls), mainly with a calcareous background. Littoral areas form part of the Ceno-Antropozoic Sedimentary Basin, with soft clay and marl deposits.

The rainfall pattern exhibits strong seasonal and inter-annual variability, as well as a discharge regime that is typical of Mediterranean rivers. High floods usually occur in autumn or early winter, with a gradual decline in flow that eventually dries out during late spring and summer in the southernmost areas. Asymmetries in the regional climate are well marked, and

Fig. 1. Map of the study area and location of sampling sites in the different river basins.


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influenced mainly by elevation and the proximity of the Atlantic Ocean, with longer summer droughts in the continental parts and southern regions. Mean annual temperatures in the latter are typically above $17.5^{\circ} \mathrm{C}$, sometimes with less than 400 mm of mean annual rainfall. By contrast, the northern border of the study area presents a more temperate-like climate: the annual rainfall can reach 2200 mm and mean annual temperature is below $10{ }^{\circ} \mathrm{C}$.

The landscape varies greatly across the studied area, with a patchy pattern of land use in central and western basins, including orchards, vineyards, olive groves and irrigation crops. Northern areas and those inland or at higher altitudes are mostly forested, with Pinus and Eucalyptus plantations, or Mediterranean woods, whereas the southern areas are more homogenous and characterised by cork-oak and holm-oak woodlands. The largest human settlements and industries are located in littoral areas.

## Material and methods

## Data collection

Fish sampling was carried out in early spring by electrofishing using European Committee for Normalization standards (norm 14011/2002 for wadeable streams) at 165 sites from eight river basins (Minho, Lima, Cávado, Douro, Vouga, Mondego, Lis and Tagus) in western Iberia (northern and central Portugal). Sample sites were considered undisturbed or minimally disturbed and had previously been extracted from a data set of more than 500 available sampling locations. These were identified using existing European guidelines (Refcond 2003) and the EU-FAME
project approach (http://fame.boku.ac.at), in which human disturbance is ranked semiquantitatively using all field data and available GIS information. In this case, we considered five disturbance variables: integrity of the riparian galleries and naturalness of the banks, longitudinal connectivity of the river segment, sediment load, input of toxic and acidic effluents, and nutrient input. Each variable was scored from 1 (minimum disturbance) to 5 (maximum disturbance), and only sites scoring 1 or 2 for all five variables were retained for the analysis.

Local and regional environmental variables were determined for each site according to their scale of definition and spatial variability (Table 1). The local variables were obtained in situ and regional variables were obtained from arcview GIS and 1:25,000 topographic maps. Local variables included mean width (m), maximum depth (m), instream cover [defined as any submerged structure in which fish could be hidden from overhead view, undercut banks and surface water turbulence and scored subjectively from 0 (absent) to 3 (abundant)], abundance of macrophytes (\%), dominant substrate coarseness (determined visually as $1=$ silt, $2=$ sand, $3=$ gra$\mathrm{vel} / \mathrm{pebble}, 4=$ cobble, $5=$ boulder and $6=$ bedrock), mean depth (m), and overhanging tree shading (scored as 0: absent, $1:<30 \%, 2: 30-60 \%$ and 3 : $>60 \%$ ). Regional variables included mean air temperature $\left({ }^{\circ} \mathrm{C}\right)$, catchment area $\left(\mathrm{km}^{2}\right)$, altitude (m), gradient slope (\%), mean annual rainfall (mm), evapotranspiration (mm), average annual runoff (mm) and distance to source (km). To deal with spatial autocorrelation in the response variable, we included an autocorrelation term (see Augustin et al. 1996). This term was generated for each site as the

|  | Code | Mean/median | Range |
| :--- | :--- | :--- | :--- |
| Regional variables |  |  |  |
| Mean air temperature $\left({ }^{\circ} \mathrm{C}\right)$ | ATEMP | $13.00( \pm 2.10)$ | $7.50-25.0$ |
| Altitude (m) | ALT | $324.40( \pm 243.50)$ | $6.7-1034.0$ |
| Gradient slope (\%) | SLOP | $1.9( \pm 2.4)$ | $0.0-16.7$ |
| Evapotranspiration (mm) | EVA | $601.4( \pm 113.3)$ | $425-900$ |
| Distance to source (m) | DISTSOURC | $27,973.0( \pm 29,338.2)$ | $1000-136,358$ |
| Catchment area (km $\left.{ }^{2}\right)$ | CAREA | $264.61( \pm 475.72)$ | $0.62-2722.50$ |
| Mean annual rainfall (mm) | MAR | $1166.60( \pm 534.58)$ | $450.00-3000.00$ |
| Average annual runoff (mm) | AAR | $8(4)$ |  |
| Local variables |  |  |  |
| Mean depth (m) | MDEP | $0.5( \pm 0.3)$ | $0.1-2.0$ |
| Maximum depth (m) | MAXDEP | $1.10( \pm 0.55)$ | $0.20-2.50$ |
| Mean width (m) | MWIDTH | $9.27( \pm 9.61)$ | $1.00-57.00$ |
| Instream cover (class) | COV | $2(2)$ | $0.00-51.00$ |
| Abundance of macrophytes (\%) | MACRO | $7.65( \pm 9.55)$ |  |
| Substrate coarseness (class) | DOMSUBST | $3(3)$ |  |
| Shading (class) | SHAD | $2(1)$ |  |
| Spatial variable |  |  |  |
| Auto-correlation | ACORR |  |  |

Table 1. Code, mean/median and range of environmental variables measured at 165 sites in western Iberia.

Mean values are given for ATEMP, ALT, SLOP, EVA, DISTSOURC, CAREA, MAR, MAXDEP, MWITH and MACRO followed by standard error (in parentheses), while median values (with interquartil distance in parentheses) are given for MAD, COV, DOMSUBST and SHAD.
number of neighbouring sampling locations within 30 km in the river continuum where a given species occurs.

## Data analysis

A total of 22 species were sampled in western Iberia, but only native fish species with a broad distribution (i.e., occurring at more than $10 \%$ of sampling sites) and presenting a reasonable number of individuals along environmental gradients were considered in the analysis. Altogether, nine species were used in the analysis.

Statistical association between assemblage structure and environmental variables was quantified by canonical correspondence analysis (CCA) (Ter Braak \& Verdonschot 1995), using CANOCO 4 (Ter Braak \& Smilauer 1998) on a 165 sites $\times 9$ species matrix. Separate CCAs were performed using local and regional environmental data sets. A forward stepwise selection procedure was used to obtain the best predictors for the model, at a significance level of 0.05 , using a Monte Carlo permutation test (999 permutations). Fish densities were $\log (x+1)$ to minimise the effect of extreme values (Legendre \& Legendre 1998).

Logistic regression analysis was used to model the influence of different variables (local and regional) on the occurrences of nine fish species. In Mediterranean rivers, many native species are restricted to one or a few basins for palaeo-historical and evolutive reasons (Doadrio 2001). To account for this in the models that we developed for each species, we only considered their natural area of occurrence by analysing only basins where a species is potentially present. Firstly, Spearman's analysis of correlations between the environmental variables was used to detect redundant variables, and those for which correlation values were greater than $|0.8|$ were excluded from the analysis. Twelve variables were considered, plus the variable used to deal with spatial autocorrelation. Secondly, a stepwise selection procedure was applied to each variable with a probability of entry of $P \leqslant 0.05$ and removal of $P>0.05$ (see Hosmer \& Lemeshow 1989). The association between the explanatory variables and interactions with the response variable was tested using the maximum likelihood method (Hosmer \& Lemeshow 1989). This procedure was first applied separately for the variables from the two different scales (local and regional), and subsequently using only the significant environmental correlates obtained in these scales to produce the final model.

Assessing prediction accuracy
The predictive accuracy of the final models was assessed using the area under the curve (AUC) of the
receiver operating characteristic (ROC) curve (Fielding \& Bell 1997; Pearce \& Ferrier 2000). We used this graphical method to represent the relationship between the percentage of presences correctly predicted (sensitivity) and one minus the percentage of the absences correctly predicted (specificity). The area comprised between a $45^{\circ}$ line and the curve measures the discrimination - i.e., the model's ability to correctly classify a species as present or absent at a given plot (AUC). We used a rough guide to classify the accuracy: $0.50-0.60=$ fail; $0.60-0.70=$ poor; $0.70-0.80=$ fair; $\quad 0.80-0.90=$ good; $\quad 0.90-$ $1=$ excellent (Swets 1988).

## Variation partitioning for GLM models

The variation in the data set was partitioned to assess the relative importance of regional and local components in the selected logistic models. The methodology we followed is an adaptation of that proposed by Borcard et al. (1992) and Legendre \& Legendre (1998), but extended to logistic regression (e.g., Reino 2005). The Nagelkerke pseudo- $r^{2}$ (Nagelkerke 1991) was used as a measure of variation explained by each logistic model. The total variation in the expansion model was partitioned into four independent components: (i) a pure local component, $a$; (ii) a shared component - i.e., the local variation that is regionally structured, $b$; (iii) a pure regional component that is probably due to large-scale effects, $c$; and (iv) unexplained variation, $d$. The total variation was partitioned by the expression $\mathrm{Vt}=\sum a+b+c+d$, which was used to extract any single component. This procedure involves calculating the following statistical models: (i) one 'partial' logistic regression using local variables to obtain the significant terms $(a+b)$ using a stepwise selection procedure; (ii) another 'partial' logistic regression for regional descriptors $(b+c)$, again using a stepwise selection procedure; and (iii) one joint local-regional model with all previously significant variables to obtain the shared effects $(a+b+c)$. All data analyses were performed with SPSS v. 12.0.

## Results

## Fish assemblages

The fish species we examined were well distributed throughout the sample sites and presented a mean occurrence of at least $15 \%$ in our data set (Table 2). Barbus bocagei Steindachner, Salmo trutta Linnaeus and Squalius carolitertii Doadrio were the most frequently captured species and occurred at more than $42 \%$ of the sampling sites. Cobitis paludica de Buen and Chondrostoma oligolepis Steindachner were the

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Table 2. Mean, standard error (SE), maximum (Max) and minimum (Min) values of relative abundance for the nine species collected at 165 sites in western Iberia.

| Species |  |  |  |  |  |  | Mean | SE | Max | Min | FO(\%) |
| :--- | ---: | ---: | ---: | ---: | ---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Barbus bocagei (Steindachner) | 50.7 | 13.2 | 1255 | 0 | 45.8 |  |  |  |  |  |  |
| Squalius pyrenaicus (Günther) | 8.8 | 2.2 | 214 | 0 | 21.1 |  |  |  |  |  |  |
| Squalius alburnoides (Steindachner) | 14.5 | 3.5 | 358 | 0 | 24.1 |  |  |  |  |  |  |
| Squalius carolitertiij (Doadrio) | 22.2 | 4.8 | 515 | 0 | 42.2 |  |  |  |  |  |  |
| Chondrostoma polylepis (Steindachner) | 13.9 | 2.8 | 321 | 0 | 34.9 |  |  |  |  |  |  |
| Chondrostoma duriense (Coelho) | 12.8 | 4.1 | 486 | 0 | 18.7 |  |  |  |  |  |  |
| Chondrostoma oligolepis (Steindachner) | 9.0 | 2.2 | 217 | 0 | 15.7 |  |  |  |  |  |  |
| Cobitis paludica (De Buen) | 2.7 | 0.8 | 75 | 0 | 16.3 |  |  |  |  |  |  |
| Salmo trutta (Linnaeus) | 10.0 | 1.5 | 100 | 0 | 44.0 |  |  |  |  |  |  |

Mean frequency of occurrence of each species (FO, \%) is also shown.
least frequent species, and occurred at $16-18 \%$ of the sampling sites - mostly in the Douro ( $53 \%$ ) and Lis River basins ( $100 \%$ ), respectively. The smaller cyprinids Squalius pyrenaicus Günther and Squalius alburnoides Steindachner were mainly restricted to the Tagus River basin, whereas C. paludica was not present in the northern part of the study area, nor was Chondrostoma duriense Coelho found in the southern part of the study area.

## Multivariate response of fish communities

The results obtained in CCA based on regional variables (Fig. 2a) revealed a modest species separation along the two canonical axes ( $\lambda 1=0.380$; $\lambda 2=0.145, P<0.05)$. The explanatory variables accounted for $16 \%$ of the total variation in siteassemblage composition, with the first and second axes displaying $83 \%$ of that variation. The first axis revealed a gradient of fish assemblages going from
those composed mostly of S. trutta, followed by a dominance of small cyprinids, such as $S$. carolitertii and $S$. pyrenaicus, to assemblages characterised by $B$. bocagei and C. duriense. This gradient was associated with altitude ( $t=4.55, P<0.05$ ), mean annual temperature ( $t=6.27, P<0.05$ ) and average annual runoff $(t=3.66, P<0.05)$.

The first two axes in the CCA ordination based on local variables (Fig. 2b) were significant ( $P<0.05$ ), with eigenvalues of 0.259 and 0.187 , respectively. The explanatory variables accounted for $16.1 \%$ of the total variation in site-assemblage composition, with the first and second axes displaying $85.3 \%$ of that variation. As depicted in the biplot, the fish community changed along the first axis from assemblages characterised by the larger sized cyprinids, B. bocagei and C. duriense and also S. trutta, to assemblages characterised by small-sized cyprinids, such as S. pyrenaicus, S. alburnoides and C. oligolepis. This separation was mainly related to instream cover $(t=7.43, P<0.05)$ and abundance of macrophytes $(t=-4.44, P<0.05)$. The second axis revealed a weaker gradient associated with substrate coarseness $(t=-3.89, P<0.05)$ and shading ( $t=-3.94, P<0.05$ ).

## Models for individual species

The variables, evapotranspiration, distance to source and mean depth, were excluded ( $r \geqslant|0.8|$ ) from the analysis. Twelve variables were therefore retained for subsequent analyses. The autocorrelation variable was not significant for any species univariate models and we did not include this term in further analyses.

The models showed that species occurrences were significantly related to both regional and local


Fig. 2. Axes one and two of the canonical correspondence analysis biplot for the 9 species and (a) regional variables, (b) local variables, assessed in western Iberia.
variables, thereby suggesting an interplay of influences between scales (Table 3). Substrate coarseness and overhanging shading were the most frequently selected local variables. Geomorphological variables (gradient slope and altitude) also played an important role in the occurrence of species - namely Chondrostoma polylepis Steindachner C. oligolepis, S. carolitertii, $S$. trutta and B. bocagei. The latter species was mainly associated with larger lowland streams. By contrast, S. trutta displayed a positive association with altitude, depth and shading and a negative association with mean annual temperature. Local habitat variables, such as stream width, cover, substrate coarseness and shading, were significant correlates for the occurrence of S. pyrenaicus, S. alburnoides and C. duriense. Cobitis paludica was found to be mainly confined to warmer areas presenting finer substrates - i.e., coastal and lower courses of main rivers. In the final model, some species related more to regional (C. oligolepis) or local (C. duriense and S. pyrenaicus) variables.

## Accuracy of GLM models

None of the models was classified in the lowaccuracy classes ( $\mathrm{AUC}<0.7$ ) (Table 4). Fit and classification accuracies of logistic regression models were high, indicating a strong predictive power (84\%). The model for S. alburnoides had the lowest model accuracy (75\%). Good accuracy class ( $0.8<\mathrm{AUC}<0.9$ ) was assigned to four species belonging to four genera, with $S$. trutta and C. oligolepis models displaying the highest levels ( $93 \%$ and $90 \%$, respectively).

## Variation partitioning for GLM models

Local, shared and regional components (considering all significant variables obtained in stepwise logistic regression) managed to explain between $70.5 \%$ (S. trutta) and $23.5 \%$ (S. carolitertii) of model variation, but the explanation of the distribution patterns varied from species to species (Fig. 3). Some of the studied distributions are highly associated with pure local attributes which are mainly related to habitat (S. alburnoides, S. pyrenaicus and C. duriense), and which ranged from $17.1 \%$ to $27.9 \%$ of total variation. By contrast, after accounting for local attributes, pure regional covariates are highly represented in three species (S. trutta, C. paludica and C. oligolepis), thus suggesting an underlying large-scale spatial trend in their distribution. Moreover, shared effects also represented a variation in distribution, from 1.9\% (S. alburnoides) to $35.3 \%$ (B. bocagei). We should note that species which are highly associated with regional variables also have strong shared effects, thus suggesting that some of the local attributes may be
 from likelihood ratio statistics (given in parentheses $(P<0.05)$ ).

| Species | $N$ | Regional variables |  |  |  |  |  | Local variables |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  |  | ATEMP | ALT | SLOP | CAREA | MAR | MAD | MAXDEP | MWIDTH | MACRO | COV | DOMSUBST | SHAD |
| Barbus bocagei | 165 |  | -0.004 (0.000) | -0.629 (0.001) | ns |  |  | ns | 0.239 (0.000) |  | ns |  |  |
| Squalius pyrenaicus | 53 |  |  |  |  |  | ns |  |  |  |  | 1.298 (0.000) | 0.037 (0.042) |
| Squalius alburnoides | 143 |  |  | ns |  |  |  |  |  |  |  | 0.681 (0.012) |  |
| Squalius carolitertii | 112 |  |  | -0.276 (0.002) |  |  |  | 1.251 (0.005) |  |  |  |  |  |
| Chondrostoma polylepis | 86 |  |  | -0.723 (0.000) |  |  | 0.001 (0.020) | ns |  |  | ns |  | 0.022 (0.003) |
| Chondrostoma duriense | 79 |  | ns |  | ns |  | -0.002 (0.008) |  | 0.064 (0.009) | -0.102 (0.020) | 1.231 (0.000) |  |  |
| Chondrostoma oligolepis | 163 |  | -0.011 (0.000) | -1.227 (0.005) | -0.002 (0.032) |  |  | ns |  | ns |  | ns | ns |
| Cobitis paludica | 86 | 0.455 (0.001) | ns |  |  |  |  |  |  |  |  | -0.542 (0.001) |  |
| Salmo trutta | 162 | -0.526 (0.001) | 0.003 (0.013) |  |  |  |  | 1.236 (0.011) |  |  |  | ns | 0.030 (0.002) |

ns represents the variables selected in local or regional models that were not selected in the final global model.

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| Species | AUC (\%) | AUC sig. | Sensitivity | Specificity | Nagelkerke $r^{2}$ |
| :--- | :--- | :--- | :--- | :--- | :--- |
| Barbus bocagei | 88 | $P<0.001$ | 0.79 | 0.75 | 0.57 |
| Squalius pyrenaicus | 87 | $P<0.001$ | 0.67 | 0.83 | 0.49 |
| Squalius alburnoides | 75 | $P<0.001$ | 0.65 | 0.7 | 0.28 |
| Squalius carolitertii | 78 | $P<0.001$ | 0.59 | 0.77 | 0.24 |
| Chondrostoma polylepis | 76 | $P<0.001$ | 0.75 | 0.61 | 0.25 |
| Chondrostoma duriense | 88 | $P<0.001$ | 0.91 | 0.66 | 0.49 |
| Chondrostoma oligolepis | 90 | $P<0.001$ | 0.91 | 0.61 | 0.48 |
| Cobitis paludica | 82 | $P<0.001$ | 0.66 | 0.51 | 0.3 |
| Salmo trutta | 93 | $P<0.001$ | 0.84 | 0.8 | 0.7 |

Table 4. Discrimination ability for optimal GLM models for each species including the percentage of the area under the curve (AUC), level of significance, sensitivity and specificity (see Material and methods for explanation) and Nagelkerke's $r^{2}$.


Fig. 3. Variation partitioning applied to the nine GLMs developed for each fish species in western Iberia.
regional or spatially structured. For example, in S. trutta this accounts for $20 \%$ of total variation.

## Discussion

This study focused on fish distribution in western Iberia and on the relative importance of local and regional factors that shape the fish assemblage structure. The most significant way in which our study differed from most others was that the sampling network consisted of undisturbed or minimally disturbed sites.

Canonical correspondence analysis revealed that generic patterns in the fish assemblage composition in western Iberia were primarily associated with watershed features and the hydrological regime, such as altitude, slope, rainfall and run-off. Overall, the fish community went from assemblages dominated by S. trutta, followed by assemblages with a dominance of small cyprinids, such as S. pyrenaicus, S. alburnoides and C. oligolepis, to assemblages characterised by the larger cyprinids, B. bocagei and C. duriense. Our results from the network of minimally disturbed sites
generally corroborated the organisation of freshwater fish communities in western Iberia that had been revealed in some earlier works (Godinho et al. 1997; Filipe et al. 2002; Santos et al. 2004a) and in other geographical areas (Lyons 1989; Rahel \& Hubert 1991). Assemblage composition was also related to local-scale features, such as cover, abundance of macrophytes, substrate coarseness and shading variables that other studies (Godinho et al. 2000; Pires et al. 2004; Santos et al. 2006) have found to be important. Cover may be important for resting and hiding, and can be related to invertebrate production (particularly submerged and emergent species) and structural habitat complexity (Moyle \& Cech 1996). The availability of coarse substrate is important in that it provides protection against widespread mammals (such as the otter Lutra lutra and beech marten Martes foina) and birds (such as kingfisher Alcedo atthis, grey heron Ardea cineria and little egret Egretta garzetta) and prevents fish from being washed away by high flow events. Abundance of macrophytes was also found to be essential. In fact, cyprinids, such as B. bocagei and C. polylepis, eat large amounts of plant material (Magalhães 1992; Collares-Pereira et al. 1996), and S. alburnoides is a phytophil spawner (Fame Consortium 2004). Regional and local-scale factors were thus both significant correlates of the fish assemblage structure. However, there may be other factors influencing species distribution, including population dynamics, species interactions and historical factors, such as isolation arising from the formation and evolution of the basin systems (Carmona et al. 1999). Furthermore, our analyses should be viewed with caution as they represent an ecological snapshot, and interannual or more sampling-intensive studies would help to clarify the environmental correlates that are subject to temporal variability (Lohr \& Fausch 1997; Wiley et al. 1997; Magalhães et al. 2002).

The generally high fit and classification accuracy of the logistic regression models support the basic hypothesis of our modelling approach that significant relationships exist between local habitat features, spatial variables and the occurrence of fish species in western Iberian rivers. However, several factors could
have been responsible for the lower accuracy classification that occurred with some of the models, particularly in the case of S. alburnoides, S. carolitertii and C. polylepis. There may also be habitat measures that are missing from this analysis and could increase predictive power. For example, additional measures often used in predictive models for fish at a local scale include dissolved oxygen, pH and conductivity (Bain \& Robinson 1988; Fausch et al. 1988), and these variables could be important in this system as well. There may be other factors influencing the presence of fishes, including site-based population dynamics, species interactions and historical factors (Guisan \& Zimmermann 2000). Fish are also constrained by their historically defined drainage basins (Williams et al. 2003). Unlike other authors (Hinch et al. 1994), who found that spatial autocorrelation might strongly affect inferences about large-scale distributional patterns, in our study this variable was not a determinant for species models, possibly due to the larger distance between neighbours ( 30 km ) when compared with other studies (Magalhães et al. 2002).

Our models clearly emphasised the importance of examining species-habitat relationships at different spatial scales (local and regional). Altitude remained a major ecological factor for some species: B. bocagei and $C$. oligolepis were negatively related to this variable, whereas $S$. trutta showed a positive association. It is known that altitude influences species occurrence through water temperature (Huet 1949), thus indirectly governing fish population density via growth and fecundity (Lobón-Cerviá et al. 1996). Slope was also a regional key variable selected by the models, as it displayed significant negative associations for four cyprinid species (B. bocagei, S. carolitertii, C. polylepis and C. oligolepis). Slope makes a major contribution to the erosive force acting on substrate and bed scour in a given area (Cobb et al. 1992). Weaker slopes within our study area are associated with higher habitat heterogeneity, thus favouring higher species richness (e.g., coexistence of cyprinid species).

Physical habitat and channel morphology descriptors also played an important role in species occurrence. Stream width was positively related to the probability of the presence of the larger cyprinids, B. bocagei and $C$. duriense. This pattern (i.e., larger fish associated with wider and deeper habitat) has been found elsewhere in other Iberian rivers (Rincon et al. 1990; Godinho et al. 1997; Carmona et al. 1999; Pires et al. 1999; Santos et al. 2004b). Substrate coarseness and shading were also important correlates of fish species occurrence. The availability of coarse substrate for small-sized species, such as S. pyrenaicus and S. alburnoides, is important in that it provides cover and concealment and also relatively
better habitat for macroinvertebrates that provide them with a source of food (Doadrio 2001). In contrast, $C$. paludica's negative response to substrate coarseness is consistent with some studies on this species (Perdices \& Doadrio 1997; Valladolid \& Przybylski 2003; Clavero et al. 2005). Shading was positively related to probability of occurrence of S. pyrenaicus, C. polylepis and S. trutta. The importance of shading, and thus canopy cover, to fishes has been demonstrated in a number of papers (Schlosser 1991; Schiemer et al. 1995; Stauffer et al. 2000). Larger fish, such as C. polylepis and S. trutta, have been found to favour areas with a high canopy cover, and this could be due to a combination of factors including physical stresses, food availability, predation risk or competitive interactions (Schlosser 1991; Collares-Pereira et al. 1995; Mellina et al. 2005; Santos et al. 2006). Shading is also important to small-sized S. pyrenaicus, as it is a major component of instream cover that provides refuge in well-conserved upper reaches and small Iberian streams (Corbacho \& Sánchez 2001; Clavero et al. 2005). But our study also provided insights into the occurrence of less well-known Iberian species. For example, C. oligolepis was found to be negatively associated to large-scale descriptors, such as altitude, slope and catchment area - i.e., it inhabits mainly small coastal rivers and the lower reaches of main tributaries.

Both local and regional attributes were significant correlates of fish species distribution in western Iberia. The purely local attributes accounted for much of the model variation of $C$. duriense and $S$. alburnoides. Previous authors (Godinho et al. 1997; Pires et al. 1999; Santos et al. 2006) had pointed out that the distribution of these species is governed first by the suitability of physical habitat and hydraulic conditions, such as cover, substrate and velocity. On the other hand, factors that operate on a regional scale (temperature, altitude and catchment area) possessed a greater explanatory power for the two species inhabiting the smallest streams (S. trutta and C. oligolepis). Our results seem to demonstrate that the responses of species that inhabit small-order streams differed from those occurring in larger rivers. Also, the largest proportion of explained variation for some of the species (S. pyrenaicus and B. bocagei) was due to the interactions between both sets of attributes. Moreover, species that were highly associated with regional variables (S. trutta, C. oligolepis and C. paludica) also had strong shared effects, thus suggesting that some of the local attributes may be regional or spatially structured. This overlapping could be a consequence of local data acting partially as a synthetic descriptor of unmeasured underlying regional processes. Future research should be extended to the influence of other factors, in an attempt to further elucidate the
mechanisms governing the occurrence of Iberian fish species.

Knowing how fish species respond to habitat heterogeneity on multiple scales can be useful to fishery managers, watershed restoration groups and others with an interest in conservation (Fausch et al. 2002; Filipe et al. 2002; Allan 2004). Species that respond primarily to local factors, such as S. alburnoides, S. pyrenaicus and C. duriense, may have a patchy distribution and hence be more vulnerable to immediate disturbances, but their populations are nonetheless more easily recovered, while species that are principally governed by regional factors, such as S. trutta and C. oligolepis, would be especially affected by broad-scale disturbances, such as climate change and river discontinuities, and therefore less responsive to management actions. Nonetheless, the high rate of endemicity of Iberian native ichthyofauna (Clavero et al. 2004) should deserve special concern for the conservation of inland waters. All studied species, particularly S. alburnoides and S. pyrenaicus, are currently threatened by deterioration or destruction of habitats, both by pollution and intense modifications (i.e., damming and channelisation) and the introduction of exotic species (Cabral et al. 2006). The GLM models provided insights into how both landscape and reach-scale factors affect fish species occurrence, and are thus useful for setting or refining management goals and evaluating the effectiveness of species management and conservation. Subsequent efforts that include ecological attributes of fish species, such as reproductive and trophic guilds (Oberdorff et al. 1993; Belliard et al. 1997) and also colonisation abilities (Grenouillet et al. 2004), and others that focus on variables related to dispersal success (Dunham \& Rieman 1999), may improve predictions of the occurrence of native Iberian fish fauna.

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