Susceptibility of European freshwater fish to climate change: species profiling based on

life-history and environmental characteristics

Running head: European freshwater fish and climate change

Ivan Jarić[1,2,3*], Robert J. Lennox[4], Gregor Kalinkat[2], Gorčin Cvijanović[3] and Johannes

Radinger[2,5]

[1] Biology Centre of the Czech Academy of Sciences, Institute of Hydrobiology, Ceske

Budejovice, Czech Republic

[2] Leibniz-Institute of Freshwater Ecology and Inland Fisheries, Berlin, Germany

[3] Institute for Multidisciplinary Research, University of Belgrade, Serbia

[4] Fish Ecology and Conservation Physiology Laboratory, Department of Biology, Carleton

University, Ottawa, Ontario, Canada

[5] GRECO, Institute of Aquatic Ecology, University of Girona, Spain

[*] Corresponding author: Ivan Jarić, Biology Centre of the Czech Academy of Sciences,

Institute of Hydrobiology, Na Sádkách 702/7, 370 05 České Budějovice, Czech Republic, E-

mail: ivan.jaric@hbu.cas.cz, phone: +420 38 777 5855, fax: +420 385 310 24

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Abstract

Climate change is expected to strongly affect freshwater fish communities. Combined with

other anthropogenic impacts, the impacts will alter species distributions and contribute to

population declines and local extinctions. To provide timely management and conservation of

fishes, it is relevant to identify species that will be most impacted by climate change and those

that will be resilient. Species traits are considered a promising source of information on

characteristics that influence resilience to various environmental conditions and impacts. We

collated life history traits and climatic niches of 443 European freshwater fish species and

compared those identified as susceptible to climate change to those that are considered to be

resilient. Significant differences were observed between the two groups in their distribution,

life-history and climatic niches, with climate-change susceptible species being distributed

more southwardly within Europe, and being characterized by higher threat levels, lower

commercial relevance, lower vulnerability to fishing, smaller body size and warmer thermal

envelopes. We establish a list of species revealed to be of highest priority for further research

and monitoring regarding climate change susceptibility within Europe. The presented

approach represents a promising tool, to quickly assess large groups of species regarding their

susceptibility to climate change and other threats, and to identify research and management

priorities.

Introduction

As ectothermic organisms, fishes are intimately linked to local climatic conditions

through physiological mechanisms that delimit tolerance or resilience (Comte & Olden,

2017a). Zoogeography of fishes is therefore greatly influenced by the average and spread of

temperatures experienced in a given watershed (Pörtner & Farrell, 2008; Isaak & Rieman,

2013). Relative to seas and oceans, freshwater habitats are more drastically impacted by

changes in climate, especially due to changes in temperature and flow, and climate change is

projected to strongly affect freshwater fish communities (O'Reilly et al., 2003; Buisson et al.,

2008; Graham & Harrod, 2009; Harrod, 2016; Radinger et al., 2017). Combined with other

anthropogenic impacts (e.g. land use change and thermal pollution; Radinger et al., 2016;

Raptis et al., 2017), climate change will restrict or redraw thermal envelopes, contribute to

population declines and local extinctions, and overall shifts in the distribution of species.

Riverine fish species losses due to climate change and reduced water discharge are predicted

to reach 75% in some river basins (Xenopoulos et al., 2005). Phenological changes in fish

behaviour (Otero et al., 2014; Dempson et al., 2017; Hovel et al., 2017) have been also

detected and emphasize the powerful changes imposed by a changing climate. In Europe,

there is a broad range of climatic conditions experienced across the landscape and a diverse

ichthyofauna distributed throughout the lakes and rivers (Ficke et al., 2007). Within the IUCN

(International Union for Conservation of Nature and Natural Resources) Red List, as much as

33% of European freshwater fish species are recognized as threatened by climate change

(IUCN, 2017).

Efforts to preserve ecosystem integrity must focus on maintaining species richness

and diversity to ensure that the services provided by freshwater ecosystems are maintained.

Conservation is often limited by funding and therefore must undergo triage to identify

priorities and allocate resources efficiently (McDonald-Madden et al., 2011). To provide

timely management and conservation and allocate resources efficiently, it is important to

identify those species that will be most impacted by climate change and those that might be

rather resilient. Species traits are considered as a promising source of information on

characteristics that influence resilience to various environmental conditions and impacts

(Jiguet et al., 2007; Comte & Olden, 2017b). Species traits represent any morphological,

physiological or phenological feature that is measurable at the individual level of a species

(Floeter et al., 2018). Trait-based evaluation has been demonstrated to be linked to the risk

status of species and can be used to investigate mechanisms that contribute to imperilment,

make predictions about unassessed species, or rank and prioritize species based on their

relative risk (Olden et al., 2007; Bland & Böhm, 2016; Comte & Olden, 2018).

Here we assess various ecological and life-history characteristics of European

freshwater fish species to identify traits that are characteristic for those that are susceptible to

the effects of climate change. Automated scraping of an online trait database and calculation

of climate envelopes using IUCN range maps overlaid on climate maps allowed us to collate

species-specific data on life history, distribution, climatic niches, as well as data on threat and

economic status. This allowed us to compare species identified as susceptible to climate

change with those that are considered to be resilient. Results of the study will contribute to a

better understanding of the expected climate change effects on European freshwater fish

fauna. We also establish a list of European species of highest priority for further research and

monitoring regarding climate change susceptibility. The method allows to extrapolate results

and characterize rare and less studied species, with scarce autecological information.

Materials and methods

Dataset

Our analysis comprised comparisons of in total 443 European freshwater fishes

between those that were identified as threatened by climate change (n=148) within the IUCN

Red List Database (IUCN, 2017) and those without climate change listed as a threat (n=295).

A list of native European freshwater fish species belonging to 25 families, mainly to

Cyprinidae (45%) and Salmonidae (20%), was obtained from the IUCN Red List database

(IUCN, 2017). It included both exclusively freshwater species, as well as those that partly

enter brackish and saltwater. Obtained data also comprised IUCN Red List classification and

maps of their distributional range within Europe. In addition, we obtained information

whether climate change was indicated as one of the threats for each species, which is based on

threat analyses and expert judgement by IUCN species experts. Overall, the dataset comprised

33% species which were categorized as susceptible to climate change.

In addition, for each species we collated trait information related to their life

history, ecology, fishery and threat status, and spatial and bioclimatic data variables (Table 1).

Life-history data were obtained from the FishBase database (Froese & Pauly, 2017) by using

the rfishbase R package (Boettiger et al., 2012, 2017). Traits with low data coverage (i.e.

those that were available for less than 1% of all species) were excluded from the analysis.

Bioclimatic spatial data were obtained from the MERRAclim database (Vega et al., 2017) and

included 19 variables related to temperature and humidity (Table 1). Global Human Footprint

map (map of anthropogenic impacts on the environment) was obtained from WCS and

CIESIN (2005) and the spatial elevation data were obtained from USGS (2010). Based on the

distributional range maps of each species (IUCN, 2017), mean values within each species'

range were estimated for each of the spatial variables using the intersect tool in ArcGIS

(version 10.5) and the *extract* function in the R (version 3.4.3; R Core Team, 2017) package

raster (version 2.6-7; Hijmans, 2017). Range maps were also used to estimate the number of

watersheds covered by each species based on WRI (2006), as well as the area and coordinates

of the range centroid for each species. General descriptive statistics and information on data

sources of all variables used in the analysis is presented in Table 1 and Supplementary

material S1. The complete dataset is presented in Supplementary material S2.

Statistical Analysis

We calculated boosted regression trees (BRT) to evaluate the relationship between the

membership of a species to the group of susceptible vs. non-susceptible species and the 45

explanatory variables. BRT are a statistical learning method that combines and averages

(boosting) many simple single regression trees to form collective models of improved

predictive performance (Elith et al., 2008). BRT can accommodate continuous and categorical

variables, are not affected by missing values or transformation or outliers and are considered

to effectively select relevant variables, identify variable interactions and avoid overfitting

(Elith et al., 2008, Radinger et al., 2015).

Specifically, we first fitted an initial global BRT model (R package *dismo*, *gbm.step*,

version 1.1-4; Hijmans et al., 2017) using the complete set of explanatory variables. An

automatized stepwise backward selection of explanatory variables (*gbm.simplify*) was applied

to eliminate non-informative variables based on model-internal cross-validation of changes in

a models' predictive deviance (Hijmans et al., 2017). Thereafter, we calculated a final BRT

model (*gbm.step*) based on the selected set of explanatory variables. For all BRT modeling

steps, tree complexity and learning rate was set to 3 and 0.001, respectively, to achieve the

recommended number of more than 1000 regression trees (Elith et al., 2008). All other model

settings were set to default or were automatically adjusted by the boosting algorithm. We

calculated a 10-fold cross validation of the BRT model as already implemented in the

algorithm. In addition, we extracted the mean AUC (area under the receiver operating

characteristic curve) as a measure of the model's predictive quality. The AUC is a threshold-

independent rank-correlation coefficient with high values typically indicating a strong

agreement between the model prediction and the membership of species to the susceptible vs.

non-susceptible group (Hijmans & Elith, 2017).

The relative importance (%) of each explanatory variable in the final BRT model was

quantified based on the number of times each variable was used for splitting, weighted by the

squared improvement at each split and averaged over all trees (Elith et al., 2008). For BRT

models with Gaussian distribution the relative variable importance equals the reduction of

squared error attributable to a given variable. Differences between groups were also assessed

by bootstrapping, by sampling each group independently and estimating the difference based

on confidence intervals (functions *two.boot* and *boot.ci*, R package *simpleboot*, version 1.1-3;

Peng 2008). Differences were considered to be significant if 95% confidence intervals did not

overlap with zero.

Subsequently, species were ranked based on the subset of variables selected by the

BRT analysis (i.e., those with >1% variable importance score), and weighed by the

importance of each variable, in order to estimate their position along the climate change

susceptibility continuum. For each species, the value of each variable was standardized based

on its position between the minimum (t_{min}) and maximum values (t_{max}) observed in the

dataset, with 0-1 possible range, and multiplied by the importance score (I_x) of the given

variable:

$$R_{tx} = \frac{t_x - t_{min}}{t_{max} - t_{min}} \times I_x \qquad\qquad (1)$$

where R_{tx} represents the rank value of variable t in species x, and t_x is the value of variable t

for species x. For variables where the lower endpoint (t_{min}) is associated with the climate

change susceptibility, equation should be adjusted as follows:

$$R_{tx} = \frac{t_{max} - t_x}{t_{max} - t_{min}} \times I_x \qquad\qquad (2)$$

Summing of all ranking scores across all variables yielded the total species ranking score,

which can range from 0 to 100, with higher values indicating stronger climate change

susceptibility.

Results

Our analyses indicated substantial differences between the two groups, climate

change susceptible versus non-susceptible species. The BRT analysis selected 35 most

relevant variables, which were subsequently assessed for their relative importance to

discriminate between the two groups (Figure 1 and Supplementary material S3). The BRT

model with the selected set of explanatory variables was successfully modeled

(Supplementary material S3) with a good cross-validated AUC of 0.87 (standard error =

0.014). Out of all explanatory variables, latitude of the species range centroid was selected as

by far the most relevant variable (31% variable importance), followed by the IUCN Red List

classification (8%), commercial relevance (6%) and vulnerability to fishing (6%). Climate

susceptible species were characterized by more southwardly positioned distribution range

centroids (41.6º vs. 47.8º N as a mean value in susceptible and non-susceptible species,

respectively), smaller range sizes (175×10^3 km^2 vs. 1686×10^3 km^2), and lower elevations

within their ranges (717.7 m vs. 892.2 m a.s.l.), with a higher proportion of exclusively

freshwater species (93% vs. 66%; Figure 2). Susceptible species were also characterized by a

smaller maximum body length (23.4 cm vs. 41.0 cm), higher proportion of threatened species

(63% vs. 31%), lower proportion of commercially relevant species (25% vs. 74% of highly

commercial, commercial and minor commercial species), and lower vulnerability to

overfishing (32.6 vs. 38.5 vulnerability index; Figure 3), as well as by higher temperature-

related values (Supplementary material S4). Bootstrapping indicated significant differences

between the groups in each of the variables. Species that are susceptible to climate change are

mainly distributed within the Mediterranean region, while the non-susceptible species

distribution mainly covers central and northern European regions, as well as the Carpathian

region (Figure 4).

Species ranking based on the association of their traits with climate change

susceptibility characteristics is presented in Table 2 and Supplementary material S5. The five

top-ranked climate susceptible species were Acheron spring goby (*Knipowitschia milleri*),

Corfu toothcarp (*Valencia letourneuxi*), *Iberochondrostoma almacai,* Evia barbel (*Barbus*

euboicus) and Malaga chub (*Squalius malacitanus*). Most of the species with the high climate-

susceptibility ranks are also classified as highly threatened according to the IUCN

classification (Table 2). Interestingly, the highest ranked species, *K. milleri*, was not classified

within the IUCN Red List as threatened by climate change. Other high-ranking species that

were not recognized as threatened by climate change were Malaga chub (*Squalius*

malacitanus), Almiri toothcarp (*Aphanius almiriensis*), and Trichonis dwarf goby

(*Economidichthys trichonis*). Species with the lowest ranking scores, i.e. with low climate

change susceptibility, were humpback whitefish (*Coregonus pidschian*), Arctic flounder

(*Liopsetta glacialis*), northern pike (*Esox lucius*), burbot (*Lota lota*), and European perch

(*Perca fluviatilis*). A complete list of all species' rankings is presented in Supplementary

material S5.

Discussion

In the present study, significant differences in life-history and climatic niches were

observed between the European freshwater species susceptible to climate change and those

that are not, such as species body size, range size, distribution and thermal envelopes. The

latitude of the species range centroid was by far the most influential trait. Overall, southern

regions with the warmer, Mediterranean climate comprised a higher proportion of species

susceptible to climate change (Figure 4). These results support recent findings that the species

from lower latitudes and tropical, warm-water habitats, are at greater risk from climate change

and warming (Payne & Smith, 2016; Payne et al., 2016; Comte & Olden, 2017b). In such

species, evolved towards higher upper thermal tolerances, specialization to thermal extremes

is accompanied by a reduced physiological flexibility and adaptation capacity to respond to

changing environmental conditions (Payne & Smith, 2016; Payne et al., 2016; Comte &

Olden, 2017b). Such heat-tolerant species are also adapted to temperatures close to their

physiological thermal limits, with a narrow safety margin for further temperature increases

(Sinclair et al., 2016; Comte & Olden, 2017b). Freshwater basins in Southern Europe are also

of particular conservation concern due to an elevated pressure by a range of anthropogenic

impacts that further exacerbate effects of climate change, such as pollution, water resource

development and consumption, and biological invasions (Xenopoulos et al., 2005; Clavero &

García-Berthou, 2005; Walther et al., 2009; Vörösmarty et al., 2010; Comte & Olden, 2017a).

Climate-susceptible species were also characterized by a smaller body and range

size (Figures 2, 3). These traits, which are also related to a lower dispersal ability (Radinger &

Wolter, 2014), are well recognized as predictors of climate change susceptibility in fish (e.g.

Ficke et al., 2007; Isaak & Rieman, 2013; Chessman, 2013; Pearson et al., 2014; Radinger et

al., 2017). Smaller-bodied fish are facing elevated overall extinction risk in freshwater habitat

due to multiple threats, such as habitat loss and fragmentation (Olden et al., 2007; Kalinkat et

al., 2017; Kopf et al., 2017), which explains higher threat level observed in climate-

susceptible species in the present study. Observed lower commercial relevance and lower

vulnerability to fishing of climate-susceptible species both stem from a lower body size and

related faster life history of such species.

It is important to acknowledge certain limitations of the data sources used in this

study, such as species and trait coverage, reliability of methods applied for threat and

extinction risk classification, and potential assessors' biases (Clavero & García-Berthou, 2005;

Keith et al., 2014; Trull et al., 2018). Furthermore, species that are not classified within IUCN

Red List as threatened by climate change can comprise also those that are not yet assessed for

their major threats. Nevertheless, the focus of our study on a well-studied group such as

European species ensured that the basic life history data and IUCN Red List assessments were

mostly available (Kopf et al., 2017). IUCN Red List is sometimes considered to understate or

improperly account for climate change as a threat, mostly due to ambiguous definitions and

criteria (Trull et al., 2018). However, recent studies have indicated that the IUCN

classification is more efficient in detecting species vulnerable to climate change than

anticipated (Keith et al., 2014; Pearson et al., 2014). Notwithstanding all the caveats, the

databases used in the present study still represent the most comprehensive sources of data and

the best available knowledge (Olden et al., 2007; Vega et al., 2017).

Trait-based risk assessments are increasingly used for species profiling (Pacifici et

al., 2015; Liu et al., 2017; MacLean & Beissinger, 2017). The approach presented in this study

might be considered a valid and promising approach to be used as a screening tool, i.e. to

quickly assess large groups of species regarding their susceptibility to climate change and

other threats based on species traits, and to identify research and management priorities. Our

results indicate that the European environmental policy related to climate change mitigation

and adaptation (EEA, 2012, 2017) should be mainly focused on Mediterranean region. This is

especially important since this region is also predicted to have the highest frequency of

droughts and extreme high temperatures, strongest reduction in precipitation and river

discharges, the highest aggregate climate change impact and the lowest adaptation capacity

(Milly et al., 2005; Dankers & Feyen, 2008; Fischer & Schär, 2010; ESPON Climate, 2011;

Stagge et al., 2011; Rojas et al., 2012; Jacob et al., 2014; Russo et al., 2014). Moreover,

Mediterranean region was also identified as a European priority area regarding climate change

impacts for other species groups. A similar distributional pattern of species susceptible to

climate change was previously also reported for aquatic insects such as Plecoptera,

Ephemeroptera and Trichoptera (Hering et al., 2009; Conti et al., 2014), mammals (Levinsky

et al., 2007), as well as for terrestrial species in general (Pacifici et al., 2015).

271   Species ranking conducted here indicated priority species for further research and

monitoring regarding climate change (e.g. *V. letourneuxi*, *I. almacai* and *B. euboicus*; Table 2).

Moreover, it also identified species whose IUCN Red List status potentially needs to be

reconsidered or updated, such as highly ranked but apparently non-susceptible species (e.g. *K.*

milleri), or highly ranked species without a proper threat category (e.g. *K. goerneri*, classified

as Data Deficient species). As such, it has a potential to be used as a "Robin Hood Approach"

(Punt et al., 2011), where assessments based on information-rich species are used to evaluate

and categorize those that are information-poor. There is a need for climate change

vulnerability assessments that would be based on quantitative approaches and consistent set of

criteria, such as trait-based approaches advocated by IUCN (Foden et al., 2013; Trull et al.,

2018). The approach presented here could be a good step in that direction.

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Figure legends:

Figure 1. Variables selected by the boosted regression tree (BRT) model as the most relevant

descriptors of climate change susceptibility in European freshwater fish species; 20 most

relevant variables are presented, which together account for 90% of the total relative variable

influence.

Figure 2. Violin-boxplots and barplots of the most relevant spatial variables in European

freshwater fish species indicated as either susceptible (n = 148) or non-susceptible (n = 295)

to climate change. Habitat preference: blue - exclusively freshwater species, green - species

that also enter saltwater.

Figure 3. Violin-boxplots and barplots of the most relevant life history traits and variables

related to threat and commercial status in European freshwater fish species indicated as either

susceptible (n = 148) or non-susceptible (n = 295) to climate change.

Figure 4. Richness of freshwater fish species across Europe indicated as either susceptible

(middle panel) or non-susceptible (left panel) to climate change, and the relative share of

susceptible species in the local total species richness (right panel).

Number of non-susceptible species

≤5   25   50   100   ≥150

Number of susceptible species

≤2   5   10   20   ≥30

Share (%) of susceptible species

0   25   50   75   100