Conservation networks do not match ecological requirements of amphibians

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Abstract

1. Amphibians are among the most threatened taxa as they are highly sensitive to habitat degradation and fragmentation. They are considered as model species to evaluate habitats quality in agricultural landscapes. In France, all amphibian species have a protected status requiring recovery plans for their conservation. Conservation networks combining protected areas and green infrastructure can help the maintenance of their habitats while favouring their movement in fragmented landscape such as farmlands. Yet, assessing the effectiveness of conservation networks is challenging.

2. Here, we compared ecological requirements of amphibian species with existing conservation network coverage in a human-dominated region of western France. First, we mapped suitable habitat distributions for nine species of amphibian with varying ecological requirements and mobility. Second, we used stacking species distribution modelling (SSDM) to produce multi-species habitat suitability maps. Then, we defined species and multi-species core habitats and performed a connectivity analysis using Circuitscape theory to identify spatial continuity in suitable habitats at regional scale. Finally, we compared different suitability maps with existing conservation networks to assess conservation coverage and efficiency.

3. We highlighted a mismatch between the most suitable amphibian habitats at regional scale and the conservation network, both for common species and for species of high conservation concern. We also found two bottlenecks between areas of suitable habitat which might be crucial for population movements induced by global change, especially for species associated with hedgerow mosaic landscapes. These bottlenecks were not covered by any form of protection and are located in an intensive farmland context.

4. *Synthesis and applications* - We advocate the need to better integrate agricultural landscape mosaics into species conservation planning as well as to protect and to promote agroecological practices suitable for biodiversity, including mixed and extensive livestock farming. We also emphasize the importance of interacting landscape elements of green infrastructure for amphibian conservation and the need for these to be effectively considered in land-use planning policies.

Keywords

Green infrastructure, protected area, multi-habitat network, ecological network, stacked species distribution models, gap analysis, other effective area-based conservation measure citizen science data

1 Introduction

The biodiversity decline is still accelerating across the world affecting both scarce and common species (Dirzo et al., 2014; Gaston & Fuller, 2008) especially in agroecosystems (Kleijn et al., 2011; Seibold et al., 2019). To palliate this decline, the establishment of protected areas (PAs) is a relevant conservation tool in a human dominated world (Rodrigues & Cazalis, 2020). However, PAs alone are not sufficient to protect biodiversity and they should be part of an ensemble of management and planning strategies aiming to create functional conservation networks. Green infrastructure (GI) (also called Green and Blue infrastructure in France) has been proposed as a way to address this issue (Chatzimentor et al., 2020; Salomaa et al., 2017). Such initiatives aim to maintain a coherent and functional network of suitable habitats and to improve biodiversity integration into land-use planning. GI design is based on the *ecological network* concept (Opdam et al., 2006) emphasizing complementarity of core habitat patches and ecological corridors at different spatial and temporal scales (Opdam et al., 2006; Salomaa et al., 2017).

Evaluating the efficiency of conservation networks (here PAs and GI) is a real challenge (e.g., Rodrigues & Cazalis, 2020). In France, GI has mostly been defined based on spatial analysis of land-cover data without considering species data. It is based on a simplistic classification of ecosystems reduced to broad categories of land cover data (i.e. different layers for forest or wetlands separately) with a poor consideration of habitat complementation. In contrast, PAs have usually been defined according to the needs of focal endangered species and/or habitats, often charismatic species such as birds and endangered plants, which might be poorly representative of species-habitat relations (Rodrigues et al., 2004). Gap analysis and conservation prioritization are common approaches for a first-step assessment of the effectiveness of a conservation network and to inform decisions for conservation measures (Jennings 2000; Moilanen et al. 2005). To conduct a gap analysis, outputs of spatial modelling are particularly useful (Ahmadi et al., 2020). Species distribution models (SDMs) are commonly used to make spatially explicit predictions of the suitable environment of a given species from biological data (Guisan & Zimmermann, 2000) and multi-specific approaches have recently been developed (Scherrer et al., 2018; Thuiller et al., 2015). In addition, recent methods integrate both graph theory and SDMs for ecological network modelling (Duflot et al., 2018; Godet & Clauzel, 2020). These multi-species and integrative approaches could be relevant for gap analyses and spatial prioritization but remain underused (Foltête et al., 2020).

In western Europe, traditional hedgerow landscapes are of high conservation concern for numerous taxa such as plants, amphibians, birds, bats and arthropods (Baudry et al., 2000; Boissinot et al., 2019). In France, these rural landscapes are associated with dense hedgerow networks interconnected with a mosaic of small interlocked patches of pastures, cultivated fields and bushes with high pond density (Burel & Baudry, 1995). They have been greatly impacted by agricultural intensification and have declined from 40 % to 80 % in Europe (Bazin & Schmutz, 1994). Amphibians are good candidates to assess the ecological quality of human-dominated landscape mosaics and to question the efficiency of existing conservation networks. Indeed, They are among the most threatened taxa with rapid and widespread population declines in both natural and modified habitats (Stuart et

al., 2004). They have a bi-phasic life cycle with different ecological requirements inducing seasonal migrations between habitat patches (Sinsch, 2014). Their poor mobility and their permeable skin increases their sensitivity to anthropogenic disturbances, pollution and ecological fragmentation induced by urbanization and intensive agriculture (Cushman, 2006; Hamer & McDonnell, 2008; Stuart et al., 2004). Hence, some authors consider amphibians as good ecological indicators of general environmental health (Collins and Storfer 2003; Díaz-García et al. 2017 but see Sewell & Griffiths, 2009). In addition, amphibians have been the target of several citizen science programs in different countries around the world (De Solla et al., 2005; Petrovan & Schmidt, 2016) leading to a large amount of available data over wide areas which could be relevant for conservation tools assessment (Snäll et al., 2016).

The aim of this study is to propose a method to assess the ecological quality of agricultural mosaics landscapes from species data and to conduct a gap analysis on existing conservation networks. In a human-dominated region of western France with traditional hedgerow landscapes of high conservation concern, we compared the predicted habitat requirements of amphibian species with existing conservation network coverage (of both protected areas and green infrastructure). Specifically, we: (1) took single habitat suitability maps from Matutini, Baudry, Fortin, et al., 2021 and produced multi-species habitat suitability maps for amphibians with differing ecological requirements and mobility (2) compared these habitat suitability maps with existing conservation networks (PAs and GI), and (3) identified gaps in conservation coverage which could be a priority for new potential conservation areas. Additional analyses on all the species in the region were carried out to discuss the potential generalization of the results obtained on rare species.

2   Material and Methods

2.1   Regional context and studied species

Ecological context

Our study was performed in a region of western France (Pays-de-la-Loire) covering 32,082 km². Intensification of agricultural practices has greatly transformed the landscape over the last century from a complex matrix of small fields, meadows and woody elements (i.e. traditional hedgerow landscapes) to more homogeneous landscapes with larger fields. With 20 native species of amphibian (out of 36 native species recorded in metropolitan France), of which 12 are priority species for conservation, the region has considerable responsibility for the preservation of amphibian species and their habitats, including traditional hedgerow landscapes and wetlands.

Studied species

Nine amphibian species in the Pays-de-la-Loire region were studied (see Table 1 for species list and conservation issues). Three form a "FOR" group of forest species or species associated with dense hedgerow mosaic landscape (*Triturus marmoratus, Rana temporaria* and *Salamandra* s*alamandra*) and five form a "GEN" group of more generalist species (*Lissotriton helveticus*, *Bufo spinosus, Hyla arborea, Rana dalmatina* and *Triturus cristatus). Pelodytes punctatus* is a pioneer species mainly associated with disturbed open environments, emblematic of alluvial valleys. Finally, the "ALL" group contained all nine

studied species. Species were selected according to data availability (i.e. at least 500 presence after filtering operations according to the number of predictors used for species distribution modelling, see Matutini, Baudry, Fortin, et al., 2021). Supplementary information about data, species selection and other species of the region are presented Appendix 1.

Biodiversity conservation policies

In France, there are different conservation tools including protected areas, green infrastructures, and inventoried sites with recognised biodiversity value but that are not protected (Categories "PA","GI" and "INV" respectively in Table 2). We defined five groups of conservation tool based on IUCN categories and protection levels (Table 2 and Fig. 1). For two types of PAs (SCEN and ENS in PA-Group 2), mapped site boundaries were not available unlike other PAs and GI where we have more precise GIS data. As a result, only a point location according to the site centroid was considered in the analysis (see Appendix 2). If one conservation area was overlapped by another, we classed the area into the group with the strongest protection level and therefore all PAs were excluded from GI. French GI is divided into different land cover or landscape classes (Table 2). Each GI layer includes two types of elements: habitat nodes, considered as "biodiversity reservoirs", and "corridors" ensuring potential continuity between nodes based on geographical criteria.

2.2 Single and multi-species habitat suitability maps

We used 100 m-resolution habitat suitability maps from Matutini, Baudry, Fortin, et al., 2021. This used heterogeneous data from citizen sciences completed by professional surveys to perform and evaluate (with independent dataset) presence-only habitat suitability modelling with functional friction-based and multi-scale predictors (see Matutini, Baudry, Fortin, et al., 2021 and Matutini, Baudry, Pain, et al., 2021 for full details and method tests). Model calibration sets were opportunistic presence-only data from a regional Atlas project with distance-based filtering to reduce spatial autocorrelation coupled with a weighted pseudo-absence selection to reduce sampling bias. SDM were performed combining Random Forest and General additive models. An independent and a standardised detection-nondetection dataset, stratified by model predictions for each species, were used for a robust model evaluation. This dataset included 576 ponds monitored by experts at least 2 nights with a 5-min acoustic survey followed by a visual inspection using halogen light and direct sampling using a fishing net. Data are described Appendix 1.

We produced multi-species suitability maps by compiling individual species maps (Stack-SDM, e.g. Ferrier & Guisan, 2006) using a simple probabilistic stacking method (pSSDM, see Zurell et al., 2019). We summed continuous individual maps without binary "presence-absence" conversion (Calabrese et al., 2014; Scherrer et al., 2020) for three focal species groups: (1) forest species (FOR), (2) generalist species (GEN), and (3) all species (ALL). According to the ecological context of our region (i.e. a hedgerow landscape with small interlocked landscape elements) for a best integration of mosaic effect for landscape evaluation, we summed species maps at 500 m-resolution after 100 m-pixels aggregation

using maximum suitability values (see Appendix 3 for additional information and selection
tests). Hence, we compiled single species standard deviation maps to obtain a general
representation of levels of confidence in our mapped distributions. To assess the accuracy
of the three pSSDMs, we compared multi-species suitability index (i.e., the sum of single
species suitability values from 0 to 1) with species richness from independent pond
monitoring (Matutini, Baudry, Pain, et al., 2021). We calculated a community-AUC
(Scherrer et al., 2020) and R^2 value for each species group with 500 iterations. For each
iteration, the evaluation-set was built by randomly selecting the pixels (500m) containing
evaluation data respecting a distance of 1km between each pixel (independence) and
stratifying the pixels along the suitability gradient of the predicted map (methods tested
and validated in Matutini, Baudry, Pain, et al., 2021).
207
208 ## 2.3   Reclassification and core habitat node definition
209 For each species and stacked species group, we reclassified the final suitability map into
210 five categories of potential habitat: not suitable (below the ten-percentile value P_{10th}
211 method in Huang et al 2020), Less suitable (P_{10th}-55%), moderately suitable (55%-65%),
212 suitable (65%-80%), and highly suitable (>80%).
213
2.4   Network patterns
To map potential connections between each highly suitable patch (i.e., patch with a
suitability index above 80%) at regional scale, we used circuit theory with Circuitscape
software (McRae et al., 2016). Circuit theory considered each cell as an electric node
connected to neighbouring cells by resistors and defined by a landscape resistance (or
conductance) value (McRae et al., 2008). We used highly suitable core habitats as source
patches, and a friction map as a measure of conductance (i.e., conductance of each raster
100 m cell for movement). This method provides an exploration of potential links between
nodes according to landscape permeability combining habitat suitability and connectivity
modelling (Ahmadi et al., 2020; Koen et al., 2014).
Habitat suitability could however be a poor proxy of permeability as species may
move in unsuitable habitats (Keeley et al., 2017). Hence, the friction map was computed
using different friction cost values derived from the multi-species habitat map and
landcover data (Keeley et al., 2016, 2017). Therefore, we considered moderately to highly
suitable habitat classes (i.e., from 65% to 100%) as permeable features except for pixels
crossed by a linear barrier. In less suitable habitats (i.e., under the 65% threshold), we
calculated a friction cost based on a 100m-grid analysis performed over 10 m-resolution
landscape layers followed by a classification procedure (see Appendix 4). As a precaution,
the friction values were defined according to the least mobile species of each group. 100m-
resolution were selected to reduce computational power needed for Circuitscape analysis
and to consider landscape complementation of fine-scale landscape elements (e.g., ditches,
hedgerows and field margins, Pope et al., 2000; Mazerolle, 2005). The cost values of
impermeable barriers (i.e., high-density urban area, highspeed train line, highway and dual
carriageways) were considered as infinite. Permeable features facilitating movement across
barriers (e.g., viaduct, wildlife pass and others, see Table 2) were digitized from aerial
photographs. To reduce border effects, we allocated resistance values to a 5km-buffer

around the studied region which were randomly selected if land-cover data were not
available.
242
2.5 **Gap analysis and prioritizing for conservation**
We based the gap analysis framework on a method used by Ahmadi et al. (2020) that
compared reclassified suitability maps with areas covered by PA. We calculated "total
conservation coverage" (i.e., the proportion of suitable habitat overlapping a PA or GI),
"conservation efficiency" (i.e., the proportion of PA or GI overlapping suitable habitats)
and plotted histograms showing the total distribution of suitability values covered by
different types of conservation area. To complete the analysis, we overlaid the existing
conservation network with the final regional maps obtained using Circuitscape for each
species group to discuss network configuration in relation to potential regional continuities.
All spatial analyses were performed using R environment v. 3.5.3 (R Development Core
Team, 2019).
254
3 Results
3.1 **Multi-species maps and core habitat definition**
For each group, the AUC varied from 0.76 (forest species) to 0.87 (generalist species) and
R^2 from 0.23 (forest species) to 0.31 (nine studied species). The suitable maps are shown in
Fig. 1 while associated confidence maps are available Appendix 5. For the nine studied
species, highly suitable habitats covered 12% of the region and suitable habitats covered
49% of the region (Fig. 1). Highly suitable habitat for the forest species, covered 11% and
suitable habitat covered 17% of the region. Larger amounts of highly suitable and suitable
habitats were available to generalist species, covering 33% and 40% of the region
respectively.
265
266 3.2 **Protected areas (PAs)**
267 The distribution of protected areas along the habitat suitability gradients modelled for each
268 species or species group (FOR, GEN, ALL) is shown Fig. 2.
269
Conservation coverage
PAs covered a relatively low proportion of highly suitable habitat: 15%, 19% and 8% for
all species, forest species and generalist species respectively (Table 4). The most strictly
protected areas protection (PA group 1) covered even lower proportions of suitable habitat:
1% of highly suitable habitat and 2% of suitable habitat for each species group. Similarly,
for two of the three species of high conservation concern, PA covered only a relatively low
17% or 26% of highly suitable habitat, for *T. marmoratus* and *R. temporaria* respectively.
Only for *P. punctatus* was conservation coverage high: 90% of highly suitable habitat was
included in PA. We also found (see Appendix 1) low conservation coverage of PAs for
rarer species based on available presence-only data, except for *Pelobates cultripes* with
72% of presence data located in protected areas (of which 37% in PA Group 1).
281
Conservation efficiency

Among PA-Group 1, no sites with the highest level of protection (names "RNN", see Table
2) contained highly suitable area for any species group or individual forest species except
P. punctatus (12% of RNN areas contain high suitable habitat) (Table 4). Other PA-
Group 1 protection levels were covered by 7 to 24% of multi-species core area and by 3 to
24% of species-specific highly suitable habitat, with a higher proportion for *P. punctatus*.
Regional Park (PA-Group 2) and Natura 2000 network (PA-Group 3) included less than
16% multi-species core area and lower proportions for the two-forest species individually.
For SCEN and ENS (in PA-Group 2) 26%, 12% and 8% of these sites were located in
highly suitable habitats for generalists, forest species and all nine species respectively.
Overall, PAs contain a higher proportion of highly suitable habitat for *P. punctatus* than for
other species.
294

3.3   **Inventoried sites (INV)**
Inventoried sites with known ecological value but no legal protection (INV-Group 5)
covered 13% of the region. 20% to 34% of their areas were covered by multi-species
highly suitable habitat (24 to 57% for suitable area). 11%, 21% and 1% are covered by
highly suitable habitat of *T. marmoratus*, *R. temporaria* and *P. Punctatus* respectively
(Table 4). INV-Group 5 covered the most suitable and highly suitable habitat of all studied
groups and species except for *P. punctatus* with a poor overlapping between this species
issues and areas of INV-Group 5 (i.e., 2%).
303

3.4   **Green infrastructure (GI)**
GI included and covered a larger area of highly suitable and suitable habitat for amphibian
species than PA particularly hedgerow networks BOCAGE followed by WOOD and WET
(see Table 2). Globally, GI covered around 50% of potential habitat for amphibian species.
Among the five GI's classes (Table 4), GI relating to traditional hedgerow landscapes
(BOCAGE) were the most widespread and covered from 16% to 33% of highly suitable
habitat of species or species groups (especially *T. marmoratus* with 33% and excluding *P.*
punctatus with only 2%). In addition, the GI BOCAGE contained 42% highly suitable
habitat for generalist species and between 12-21% for other species except *P. punctatus*.
GI relating to woodlands (WOOD) and wetlands (WET) covered more than 10% of highly
suitable habitat except for *T. marmoratus* and *P. punctatus*. GI relating to coastland
(COAST) and natural open grasslands (OPEN) covered small areas compared with other
GI and also a low area of suitable habitat for studied species.
317

Potential multi-habitat continuities patterns
The distribution patterns associated with highly suitable and suitable habitats are shown
Fig. 1 and supplemented in Fig. 3 by a more detailed analysis of the spatial organisation
and potential links between habitat patches. The central and the southern parts of the
region have a few patches of highly suitable habitat, especially for forest species.
Generalist species have a more homogeneous distribution of suitable habitats spreads over
the region. Suitable and highly suitable habitat distribution patterns associated with forest
species are mostly common with other species, with the exception of large suitable forest
patches especially at the north of the region. Indeed, 92% and 81% of highly and suitable

habitat of FOR species (i.e., suitability index > 65%) are include in GEN and ALL species'
high and suitable habitat respectively.
Regarding potential multi-specific regional continuities, two major sectors form
"bottlenecks" to north-south (sector B, Fig. 3) or east-west continuity axis and potential
movements (sector A, Fig. 3). This is particularly marked for forest species. These two
sectors contain few protected areas and show a cover mismatch with GI (Fig. 3).
333

334   **4   Discussion**

We highlighted a mismatch between the most suitable habitats for studied species at
regional scale and the existing conservation network, in particular protected areas (in
coverage and efficiency). Indeed, habitat patches with a high suitability index associated
with common or slightly rarer species were poorly covered by PA except for one species of
high conservation concern: *P. punctatus*. Analysis of other rarer species distributions
suggest that the observed mismatch concerns other species present in the studied region
(Appendix 1). However, most suitable areas were better covered by GI (especially GI
layers associated with wood, hedgerow landscapes and wetlands) and by INV sites
(inventoried sites with ecological value but no regulatory protection). Difference in
coverage and efficiency obtained for PAs and INV highlights that some sites with
amphibian conservation issues are known but not protected. In addition, we identified two
narrow "bottlenecks" in suitable habitat continuities at regional scale which are only very
sparsely covered by the conservation network (PAs and GI).
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349   4.1   **General limits and interpretation framework**

In addition to the limits related to the presence-only SDM detailed in Matutini, Baudry,
Fortin, et al. (2021) recently developed stacking-SDM methods require further
development to improve their evaluation. SDM are useful tools to predict the distribution
of habitat suitability but rarely predict the biology of populations (Lee☐Yaw et al., 2022).
The multi-species networks identified should be considered simply as maps of highly
suitable habitat (habitat perspective) - measured with highly sensitive and biphasic
amphibian species – rather than as a real multilayer network of amphibian communities
(species perspective). In addition, resistance values based on literature and expert opinions
or derived from SDM values are still debated (see Godet & Clauzel, 2020 for methods
comparison and discussion) and information about species movement and dispersion
processes are still very scarce in the literature (Pittman et al. 2014).
Amphibians also have poor movement capacities, rarely exceeding a few kilometres
during dispersal movements (Sinsch, 2014). In this work, continuities were identified from
habitat distributions at regional scale and species level reflecting long-term ecological
processes and population movements across decades. This study takes a first step towards
assessing potential barriers to species distribution shifts in response to global change, but
does not directly include landscape dynamics and climate change scenarios. Rather it
provides elements at regional scale to guide regional policies and further studies to
improve multi-scale integration and efficiency of measures for biodiversity conservation.
Collect and integrate independent genetic data to assess our final maps, particularly in the

two bottleneck sectors, would be important to evaluate connectivity and complete the
investigation in view to propose relevant conservation or restoration measures.
372

4.2 **A mismatch for the conservation of amphibians**
Two main reasons may explain the mismatch between location of PAs and amphibian
habitat requirements. Firstly, PAs are generally defined according to the needs of rare
and/or emblematic species or habitats). In the study region, most PA focus on birds or
plants species, large open wetlands near the western coast or the Loire River valley but
only a few target species of amphibian and these tend to be rare species (e.g., *P.*
cultripes,see Appendix 1). Most local, rare species of high conservation concern are
pioneer species and/or associated with loose or sandy soils with poor vegetation that are
found in particular on the coast, along the Loire River and in some quarries. Their ecology
contrasts with the other studied species, except for *P. punctatus* which shares many of the
same ecological requirements, explaining the better conservation coverage and efficiency
of PA for this species.
Secondly, PA definition in the region presents a "location bias". Indeed, PAs are
defined primarily in areas of low economic interest, less used for human activities
(Rodrigues et al., 2004). This bias is common worldwide and widely described in the
literature (Ahmadi et al., 2020; Joppa & Pfaff, 2009). It can be especially problematic for
species living in human-dominated landscapes, like most amphibian species. For example,
Rodrigues et al. (2004) demonstrated that amphibians contain a large number of "gap
species" worldwide, i.e., they are the biological group least covered by the global PA
network. In addition, our results show a potential "network bias" similar to Ahmadi et al.
(2020), i.e., the extension of PA is established without sufficiently considering their
position within the network. Indeed, the current PA sites (and the sites intended for the
future extension of PAs – see Appendix 5) are not located in the two areas with highest
potential defined by our primary analysis of the regional continuity of suitable habitats.
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4.3 **Green infrastructure (GI) as a complementary tool for biodiversity**
conservation
Our results show GI offers better coverage of highly suitable habitats than PAs for
amphibian species groups, but this remains partial, in particular in the two identified
"bottleneck" sectors. Indeed, agricultural areas are poorly covered by PAs, generally due to
a conflict of economic interest (Rodrigues et al. 2004) while GI provides complementary
coverage in areas dominated by human activities. Agricultural landscapes with high
conservation value for biodiversity such as hedgerow mosaic landscapes depend on the
maintenance of traditional farming practices that may even be inhibited by PAs
conservation (Schmitz et al., 2017).
In addition, we underline the importance of GI layers associated with wetlands,
woodlands and traditional hedgerow mosaic landscapes for amphibians. Consequently, our
results highlight the importance of considering several GI layers with potential
interactions. GI tends to compartmentalize conservation issues by splitting them into
distinct layers of habitat types considered as independent "subnetworks", where the
"habitat" concept is limited to landcover categories (see Pilosof et al., 2017 for general

perspective on multi-layer nature of ecological networks). Species survival is however conditioned by different ecological processes which generally go beyond a single type of land cover class. Amphibian habitat network studies supporting GI definition mainly focus on pond networks but natural or semi-natural forest patches, especially mature forest (e.g. >70 years) and even small woodlots (Boissinot et al. 2015), are essential for amphibian population persistence, in particular in agricultural regions (Hartel et al., 2010; Collins & Fahrig 2017). Species other than amphibians may also have very contrasting requirements during their life cycle (e.g., many holometabolous insects, odonates and bats).

Lastly, we expected a higher overlap of the GI layer called Bocage (traditional hedgerow mosaics in agricultural) based on knowledge of amphibian ecology (Boissinot et al., 2019). This GI layer was mapped using combined land use data (i.e., hedges, ponds and meadows only) at 1km-resolution with a poor consideration of the interactions between these elements (i.e., simple metrics as density or area related to hedges, ponds and meadows were combined independently). Of course, other elements of the landscape are important for amphibians, such as small streams and wooded patches, as well as more complex processes related to landscape complementation and structural heterogeneity (Boissinot et al., 2019; Collins & Fahrig 2017).

Therefore, an important weakness of GI is its exclusive reliance on land use data without including biological data and landscape complementation as well as the multi-layered nature of ecological networks. Ecological niche modelling tools such as SDM, coupled with ecological connectivity models, are of particular interest for improving the ecological realism of identified networks and supporting related decisions (Clauzel & Godet 2020). These modelling tools need to be used and interpreted with care (e.g. within the framework specified in 4.1.).

4.4 **Implications for conservation**

First, we identified "bottleneck" sectors not covered by any conservation tools such as PA or GI despite high potential conservation value. These areas might be especially important for amphibian populations in a context of global change requiring distribution shifts to maintain suitable conditions for population viability. Indeed, according to Préau et al. (2019), important distribution shifts at regional scale are expected for certain amphibian species based on global change predictions. These two "bottlenecks" are located in a context of open fields and intensive farming which are considered by nature protection organisations to have limited potential for biodiversity. In these areas, some semi-natural elements such as small woods associated with wetlands may nonetheless play a key role in maintaining functionality and connectivity of the network at regional scale and should be strengthened. In these two areas, habitat continuities are maintained by riparian woods (sometime very thin and degraded) and associated river landscapes with a higher proportion of pastures and hedgerows, as well as small wood patches close to wetlands including ponds. Considering the general limits of this studies (see 4.1), these two areas need further investigation - especially connectivity evaluation by integrating genetic data – in order to defined appropriate conservation and restoration measures.

Overall, the protected areas network needs to be strengthened and supplemented by conservation tools more adapted to landscapes with strong interactions between biodiversity and human activities. Green infrastructure (GI) has better efficiency and coverage in particular because it concerns landscapes where natural and semi-natural habitats and human activity are closely intertwined. However, GI is not regulatory. Other effective area-based conservation measures (OECMs) are new conservation tools intended to complement protected areas (PAs) and might strengthen GI at local scale by sharing the common objective of restoring large scale connectivity (Convention on Biological Diversity, 2018). OECMs might be the complementary regulatory brick to PAs and GI with, in a European context, a better consideration and recognition of management practices and certain agroecological farming practices that are more biodiversity-friendly.

The different maps obtained from this study provide support for prioritizing certain regional policies with different objectives: (1) strengthening PA network, GI and future OECM (Appendix 6 provides more specific information to prioritize new protected or conservation area from a regional preselected set of sites according to our results); (2) identifying potential conflict points with transport networks (i.e., roads and railways); (3) spatially prioritizing actions supporting the agroecological transition, particularly in the two bottleneck areas considered as "degraded" for biodiversity. Conservation actions for amphibian conservation often focus on ponds and pond networks. The complementarity of ponds (or wetlands) and forest patches should be considered to define conservation actions (Cushman, 2006; Denoël & Lehmann, 2006; Pope et al., 2000) as well as certain associated extensive and traditional farming practices in western Europe (Boissinot et al., 2019; Hartel et al., 2010). In our region, amphibians have interactions with traditional agricultural practices and with related landscape dynamics. As multi-habitat species, amphibians have potential as biodiversity indicators of ecological quality of hedgerow network landscapes, composed of a mosaic of trees, grasslands and ponds upon which they depend. The success of conservation measures depends on our ability to integrate biodiversity issues into farming practices for the benefit of amphibians but also many other forms of biodiversity benefiting from such landscape mosaics.

Acknowledgements

This work would not have been possible without the support of the Pays de la Loire Herpetological Group, the CPIE Regional Union and the French BirdLife partner (LPO). We are especially grateful to Morgane Sineau and Benoit Marchadour who coordinate regional naturalists' databases. We also acknowledge the many naturalists involved, for access to data and for additional fieldwork, especially Dorian Angot, Baptiste Gaboriau and Martin Bonhomme. We thank Jean Secondi and Aurélien Besnard for providing helpful discussion. Our work was supported by funding from L'Ecole Supérieure des Agricultures d'Angers, Angers Loire Metropole, The French Society for Ecology and Evolution (SFE[2]) and "Humanité et Biodiversité".

Authors' contributions

Florence Matutini: Conceptualization (Equal), Data curation (Lead), Formal analysis (Lead), Funding acquisition (Supporting), Methodology (Lead), Investigation (Lead), Visualization (Lead), Writing-original draft (Lead), Writing-review & editing (Lead); **Marie-Josée Fortin**: Conceptualization (Equal), Methodology (Supporting), Supervision (Supporting), Writing-original draft (Supporting), Writing-review & editing (Supporting); **Jacques Baudry, Guillaume Pain & Josephine Pithon**: Conceptualization (Equal), Supervision (Equal), Writing-original draft (Supporting), Writing-review & editing (Supporting), Funding acquisition (Lead).

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Fig. 1. Identification of highly suitable and suitable multi-specific patches of habitat using stacked habitat suitability map for three species groups.

Legend

■ Group 1 - high conservation level (UICN class from I to IV)
■ Group 2 (UICN class V)
■ Group 3 (Natura 2000)
□ All region area (no protected area)

Fig. 2. Distribution of pixels along the regional habitat suitability gradient covered or not by protected area. Suitability index (from the habitat suitability map) is calculated for each species and varies from 0 to 1. Suitability index of the three groups (ALL, FOR and GEN) is the sum of the individual species index related to each group (pSSDM) divided by the number of species in order to have a suitability index on the same scale. The frequency corresponds to a number of pixels covered by protected areas from the Group 1, 2, 3 in relation to the whole region area.

Fig. 3. Large scale permeability between high suitability areas and associated continuities patterns confronted with regional green infrastructure. A and B show potential conflict area and issues for continuity conservation at regional scale with a mismatch with the green infrastructure network.

Table 1. Description of the studied species. RL FR: France red list (2015); RL Region: regional red list of Pays-de-la-Loire (2021); Priority: regional priority level for species conservation from the regional red list from 0 (low) to 3 (very high) defined by Marchadour et al. (2021); Hab. Dir. N2000: habitats directive classification (Natura 2000); Nb presence data: Number of presence-only data were calculated after a 100 m-resolution rasterization and correspond to the number of pixels with at least one observation of the species. In bold, vulnerable or near threatened species at the regional scale. See appendix 1 for species selection and presentation of other species present in the studied area.

	Species	RL FR	RL Region	Hab. Dir. N2000	Nb presence data (Atlas)
Species used for SSDM	*Pelodytes punctatus*	**LC**	**NT**		**1714**
	Triturus marmoratus	**NT**	**NT**	IV	**906**
	Rana temporaria	**LC**	**VU**	V	**848**
	Lissotriton helveticus	LC	LC		4580
	Bufo spinosus	/	LC		6071
	Hyla arborea	NT	LC	IV	4555
	Rana dalmatina	LC	LC	IV	6118
	Triturus cristatus	**NT**	**NT**	II & IV	**1040**
	Salamandra salamandra	LC	LC		3514

Table 2. Classification of main conservation areas according to protection levels used in Pays-de-la-Loire (France). PA: protected area; GI: Green infrastructure; INV: area with an inventory of biodiversity and having a strong natural heritage interest but not protected by regulatory tools (ZNIEFF, type I and II). The level of protection varies from 0 (null) to +++ (strong).

Cat.	Group	Protect. level	Code name	UICN class	Protection approaches	Level	French denomination
PA	1	+++	RNN	I-IV	Regulatory	National	National Nature Reserve
		++	SIN,SCL	III	Regulatory	National	Classified site and Registred site
		++	RNR,RB, SCL,APB	I-V	Regulatory essentially	National or Regional	Biological Reserve, Conservatoire du littoral Site, Regional Nature Reserve, Biotope Protection Order
	2	+/++	SCEN, ENS	I-VI	Regulatory, Contractual management or Land purchase and management	Regional	Area managed by Natural Area Conservation Societies (CENs)
		+	PNR	V	Contractual management	Regional	Regional Nature Park
	3	+/++	N2000	/	Contractual management	European	Natura 2000 sites (N2000)
GI	4	0	BOCAGE	/	Non-regulatory Advisory / informative	National to local	Traditional hedgerow mosaic landscapes Infrastructure
		0	WOOD	/			Woodlands infrastructure
		0	WET	/			Wetlands infrastructure
		0	COAST	/			Coastlands infrastructure
		0	OPEN	/			Natural open fields infrastructure
		0	Corridor	/			Ecological cooridors
INV	5	0	ZNIEFF	/	Biodiversity inventories only	National	Natural areas of special ecological, faunal and floral value

Table 3. Accuracy of stacked species distribution models for three species groups according to the nine studied amphibian species. Results obtained with external data with 500 permutations. AUC were calculated from predicted multi-species suitability values compared to observed species diversity. Community AUC (AUCc) were obtained with predicted suitability index for each species compared to the presence/absence observations in the assemblage (Scherrer et al. 2020). Individual AUC and SEDI are evaluation metrics obtained by Matutini, Baudry, Fortin, et al. (2021) for each species specific map.

Species group	R^2	AUC	AUCc	Min individual AUC / SEDI	Max individual AUC / SEDI
All nine species (*n*=9)	0.31	0.86	0.79	0.73 / 0.67	0.86 / 0.89
Forest species (*n*=3)	0.23	0.76	0.96	0.78 / 0.68	0.86 / 0.84
Generalist species (*n*=5)	0.30	0.87	0.86	0.73 / 0.67	0.81 / 0.89

Table 4. Proportion of the conservation area covered by different habitat suitability classes (1. conservation efficiency) and proportion of different habitat suitability classes covered by the conservation area (2. conservation coverage) for different species and species groups. Classes of conservation area: Protected (groups 1 to 3) – classes I to V derived from the UICN classification; EURO: European conservation network from Natura 2000; Non-protected: GI (Green infrastructure) and INV (inventoried sites define as "high naturalist interest"). Suitability class for each groups or species: "+++" = highly suitable (>0.80) ; "++" = suitable (>0.65) ; +/- = "moderately or not suitable" (<=0.55). See Table 2 for "code names" details. Names with * show sites with point data only (surface area is not available).

1) AnB / A — CONSERVATION EFFICIENCY

| | | | | SSDM 500m resolution | | | | | | | | | SDM 100m resolution | | | | | | | | |
| | | | | All studied species (n=9) | | | Forest species (n=3) | | | Generalist species (n=5) | | | Triturus marmoratus | | | Rana temporaria | | | Pelodytes punctatus | | |
Group	Name	Class	Area (ha)	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-
1	RNN	I-IV	3183	0	23	26	0	19	26	0	21	24	0	3	21	0	4	12	12	1	4
	S N,SCL	III	80912	7	33	36	7	11	18	12	35	29	8	13	19	4	2	3	24	18	20
	RNR,RB,SCL,APB	I-V	4902	17	30	28	20	12	8	24	32	28	3	12	17	22	4	2	14	11	15
2	SCEN,ENS*	I-VI	/	12	50	26	12	24	23	26	44	19	8	21	20	6	7	7	6	10	15
	PNR	V	358663	11	28	27	15	15	10	13	32	24	5	10	13	15	8	4	10	12	24
3	N2000	EURO	144070	10	33	36	10	16	12	19	34	26	2	10	23	10	8	6	24	14	13
4.1	BOCAGE	GI	424568	20	63	17	21	26	29	42	43	13	12	27	30	15	11	9	0	3	10
	WOOD	GI	176157	26	55	18	38	25	23	38	38	20	6	20	22	31	12	11	0	1	4
	WET	GI	270266	19	56	22	21	25	25	38	44	15	5	18	26	15	11	11	1	3	8
	COAST	GI	9376	2	51	35	0	12	30	4	59	26	12	34	25	0	0	0	13	20	26
	OPEN	GI	19445	15	53	28	11	22	20	42	41	13	1	5	11	13	11	12	1	5	17
	Total core hab.	GI	650572	20	60	19	24	25	27	40	43	15	9	24	26	18	10	9	0	3	9
4.2	Corridor	GI	378391	17	52	25	13	21	26	39	41	16	5	14	23	9	8	9	0	4	14
5	ZNIEFF	INV	279266	20	57	20	29	24	25	34	44	18	11	28	25	21	9	10	1	4	10
	REGION		3236407	12	49	28	11	17	23	33	40	17	5	13	19	9	7	7	3	6	15

Protected (PA): groups 1–3. Non protected (GI & INV): groups 4.1, 4.2, 5.

2) AnB / B — CONSERVATION COVERAGE

| | | | SSDM 500m resolution | | | | | | | | | SDM 100m resolution | | | | | | | | |
| | | | All studied species (n=9) | | | Forest species (n=3) | | | Generalist species (n=5) | | | Triturus marmoratus | | | Rana temporaria | | | Pelodytes punctatus | | |
Group	Name	Class	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-	+++	++	+/-
1	RNN	I-IV	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	S N,SCL	III	1	2	3	1	2	2	1	2	4	4	2	2	1	1	1	19	8	3
	RNR,RB,SCL,APB	I-V	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
2	SCEN,ENS*	I-VI	/	/	/	/	/	/	/	/	/	/	/	/	/	/	/	/	/	/
	PNR	V	10	6	10	14	10	5	4	9	15	11	9	8	20	13	6	35	23	18
3	N2000	EURO	4	3	6	4	4	2	3	4	7	2	3	5	5	5	4	35	11	4
4.1	BOCAGE	GI	21	17	8	24	20	17	16	14	10	33	27	20	23	21	17	2	8	9
	WOOD	GI	11	6	3	18	8	5	6	5	6	7	8	6	20	9	9	1	1	2
	WET	GI	13	10	7	16	12	9	10	9	7	8	12	11	15	14	14	2	4	5
	COAST	GI	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	1	1	1
	OPEN	GI	1	1	1	1	1	1	1	1	0	0	0	0	1	1	1	0	0	1
	Total core hab.	GI	33	25	13	42	30	24	24	21	17	39	36	28	43	32	28	3	11	13
4.2	Corridor	GI	16	13	10	14	15	13	14	12	10	12	12	12	12	15	16	2	9	11
5	ZNIEFF	INV	14	10	6	22	12	10	9	9	9	21	18	12	21	12	12	2	6	6

Protected: groups 1–3. Non protected: groups 4.1, 4.2, 5.