

# **Riparian habitat connectivity restoration in an anthropized landscape: A multi-species approach based on landscape graph and soil bioengineering structures**

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#### **Abstract**

29 In urbanized areas, rivers and riparian ecosystems are often the only ecological corridors available for wildlife movement. There, riverbanks are often protected by civil engineering structures (dykes, riprap). This can lead to habitat degradation and loss of landscape connectivity. Fascines (willow bundles tied together) could be an alternative to riprap, since they maintain the quality of the natural ecosystems by using native vegetal species instead of rocks, but their potential positive impact needs to be assessed.

 We proposed a landscape-scale decision-making method for river managers who want to restore banks by transforming riprap into fascines to improve landscape connectivity. We applied our methodology to a case study involving a 25km-stretch of the Arve river, France. We selected four target vertebrate species based on biological traits to cover a wide range of dispersal capacities. For each species, we used landscape graphs to assess habitat connectivity under different contrasted riverbank scenarios. Scenarios included replacing all natural banks with ripraps or replacing all ripraps with fascines. In addition, we systematically tested the effect of replacing individual 100 or 500m sections of ripraps by fascines, to locate where riverbank restoration would maximize connectivity gain.

 The four species selected responded very differently to the scenarios (up to +14% and +46% change in Probability of Connectivity for common toads and Eurasian beavers, respectively, 0% for common sandpipers and barred grass snakes). The restoration of specific riverbank sections could result in important gains in PC (up to +33% for one single section for one species) but no section maximized connectivity gain for all the target species.

#### **Keywords**

 Ecological connectivity restoration, graph theory, riverbank, riverine landscape management, vertebrates

#### **Introduction**

 Riparian habitats are transition areas, or ecotones, at the interface between terrestrial and aquatic habitats. They offer habitat to a wide variety of animalspecies, whether aquatic (e.g. for spawning and nursery of juvenile fish [\(Mouton et al., 2012\)\)](https://www.zotero.org/google-docs/?2yF9Hy), terrestrial (such as forest species), or dependent on these specific environments (e.g. beaver). Riparian habitats are also crucial for connectivity across these different types of environments for amphibious species, whose life cycles require both terrestrial and aquatic habitats (Wells, 2007). These habitats are therefore of disproportionate ecological importance in relation to their size as they are often only a few dozen metres wide (Sabo et al., 2005; González et al., 2017; Hunter, 2017). The specific linear nature of rivers makes them potential biological corridors, particularly in human-modified landscapes (Naiman et al., 1993; Erős et al., 2011; de la Fuente et al., 2018). Biological corridors are areas that allow organisms to move through a landscape. In particular, they enable gene flow, which is of great importance for biodiversity conservation. Their conservation is often integrated into public environmental policies (Rosenberg et al., 1997; Van Der Windt & Swart, 2007).

 Under natural conditions, rivers and riparian zones are highly dynamic systems changing their position and extensions in the landscape following hydrogeomorphological processes at various temporal scales (Tabacchi et al., 1990; Benda et al., 2004). Riverbank erosion along rivers is one of these natural processes and is necessary to create new pioneer habitat and feed the river's sediment transport flow (Florsheim et al 2008). To limit this erosion in urban areas, riverbanks are often stabilized using bank stabilization structures like dikes, riprap (civil engineering structures made of rocks or concrete) or other civil engineering works.

 Therefore, riverbeds and riparian habitats are subject to numerous pressures linked to anthropization, in particular through changes in hydrological regime, installation of dams, and channelization of watercourses. Damming and channelization can lead to a loss of habitat and functional connectivity – i.e. the way the landscape structure affects ecological processes like gene flow across landscapes

 [\(Wainwright et al., 2011; Baguette et al., 2013\).](https://www.zotero.org/google-docs/?iiMlUU) This has been observed in many species such as aquatic macroinvertebrates and fish [\(Smith et al 2009; Van Looy et al., 2014\),](https://www.zotero.org/google-docs/?gHWWT7) but also in riparian plants (Merritt & Wohl, 2006) and mammals, such as otters (Leoncini et al., 2023), whose dispersal is impeded by dams. River embankments with civil engineering structures are usually linked to a loss and/or degradation of natural riparian habitats as they consist in replacing native woody and herbaceous riparian plants to bare ground or rock [\(Ward & Stanford, 1995; Van Looy et al., 2003; Dudgeon et al.,](https://www.zotero.org/google-docs/?j4RJo0)  [2006\).](https://www.zotero.org/google-docs/?j4RJo0) They also lead to a loss of lateral connectivity by inducing differences between upstream and downstream plant communities [\(Van Looy et al., 2003; Martin et al., 2021\).](https://www.zotero.org/google-docs/?j7Pisf) Loss of connectivity is one of the main causes of biodiversity loss (Haddad et al., 2015; Thomson et al., 2017; Pardini, 2018).

 Soil water bioengineering (SWBE) bank protection structures are an alternative to 'hard' civil 87 engineering structures such as riprap. SWBE structures make it possible to maintain habitats with higher quality and functional diversity [\(Cavaillé et al., 2015; Janssen et al., 2019\),](https://www.zotero.org/google-docs/?zIGNz7) and suitable for more terrestrial and aquatic species (Cavaillé et al., 2013, 2018). For instance, fascine toes (small-diameter willow bundles tied together to form a solid structure) show a higher vegetation cover, a higher abundance of plants, a higher species richness and a lower abundance of invasive alien species than riprap [\(Cavaillé et al., 2013; Martin et al., 2021\).](https://www.zotero.org/google-docs/?jDTFEm) Soil water bioengineering structures can help protect riverbank, while restoring habitats [\(Rey et al., 2019\)](https://www.zotero.org/google-docs/?OFXdB1) and improving habitat connectivity (Martin et al 94 2021). In general, restoring riparian habitats can greatly improve landscape connectivity (Rojas et al., 2020). River managers may therefore wish to restore riparian habitats by replacing civil engineering structures with SWBE structures. However, in highly urbanized areas, hosting a large number of civil engineering bank protection structures, it is unrealistic to replace all of them simply because of financial and technical constraints.

 Therefore, efforts to identify where restoration would improve habitat connectivity the most can facilitate implementation of SWBE structure. To maximize the positive impact of new restoration projects on landscape connectivity, we need to identify key areas for the dispersal of organisms.

 Restoration ecology and landscape ecology have provided a wealth of literature on how to prioritize areas for restoration to maximize landscape connectivity (e.g. Hodgson et al., 2011; Tarabon et al., 2019; Bergès et al. 2020). These studies focus in particular on habitat patch creation (Hodgson et al., 2011; Clauzel et al., 2015) or the detection of barriers to movement to remove in priority (McRae et al., 2012; Roy & Le Pichon 2017; Kraft et al., 2019). How to optimize the location of the terrestrial habitats to be restored has been extensively addressed in the literature, but not specifically for riparian habitats. The applied techniques range from least-cost distance calculations based on estimated resistance surfaces (McRae et al., 2012 ; Roy & Le Pichon 2017) to more complex spatially explicit models, e.g. ecological network modeling based on landscape graphs [\(Clauzel et al., 2015; Avon &](https://www.zotero.org/google-docs/?1Sb4Gl)  [Bergès, 2016; Kraft et al., 2019; Clauzel & Godet, 2020\).](https://www.zotero.org/google-docs/?1Sb4Gl) Landscape graphs are models derived from graph theory, a mathematical discipline dedicated to the study of networks. Graph theory applied to landscape ecology borrows some of its concepts from the theory of island metapopulation models. In this conceptual framework, target organisms frequent patches of adapted habitats located in a matrix of non-adapted habitats (Urban et al. 2001). The ease with which organisms can move through the matrix from one habitat patch to another depends on their movement capabilities and the characteristics of the elements composing the matrix. Resistance surfaces are used to model such movements of animals between habitat patches across heterogeneous landscapes. Resistance surfaces are raster layers in which all the elements of the landscape are assigned values indicating their propensity to movement of target organisms. Even if resistance surfaces are largely used for connectivity analyses (Zeller et al., 2012), there is no consensus on how to parameterize them. Resistance values can ideally be derived from empirical data collected in the study area, such as genetic data (landscape genetics), movement data (telemetry), species habitat model [\(Duflot et al., 2018;](https://www.zotero.org/google-docs/?Mt75nn)  [Lalechère & Bergès, 2021\)](https://www.zotero.org/google-docs/?Mt75nn) or from a combination of data from the literature and expert opinion (Dutta et al., 2022). The use of landscape graphs and resistance surfaces in landscape ecology makes it possible to calculate specific metrics for quantifying the connectivity of habitats in a spatial network. In particular, the concept of amount of reachable habitat (ARH) focuses on habitat accessibility,

 including the surface and quality of the habitat patches themselves and their degree of habitat connectivity (Saura et al., 2011). This ARH can be assessed in the form of a single metric (Saura and Rubio, 2010).

 Connectivity has to be assessed taxon by taxon as species differ in their response to habitat availability and configuration (Vos et al., 2001) and it is rarely possible to carry out detailed analyses for every species. How to select target species to best quantify multi-species landscape connectivity is still under debate in the literature (e.g. [Albert et al., 2017; Meurant et al., 2018; Dutta et al., 2023\)](https://www.zotero.org/google-docs/?2ijCZE). Meurant et al. (2018) showed that a small number of surrogate species covering different biological traits related to dispersal might adequately represent the needs of the full list of species found in a given habitat type, and provide more informative results than simply maximizing taxonomic diversity of surrogate species. Working with surrogate species is a good way to reconcile conservation objectives with stakeholder constraints and field data availability (Opdam et al., 2008; Wiens et al., 2008).

 In this study, we proposed a method for prioritizing sections of the riparian habitat for restoration to maximize gains in habitat connectivity restoration to a peri-urban river located in the South East of France. First, based on a multivariate analysis of species traits, we selected several surrogate species, which allowed us to characterize the ability of species to reach available habitats within the riparian ecosystem. For each surrogate species, we used landscape graphs to quantify the importance of riparian habitats to total connectivity. Then, we assessed the contribution of existing and potential SWBE structures, using scenarios. Scenarios were chosen to assess the maximum potential gain in connectivity by replacing all riprap with SWBE structures, or to assess the contribution of existing bank protection structures to connectivity. Finally, in order to prioritize restoration areas, we systematically tested the ability of each potentially restorable bank section along the river to improve habitat connectivity for our surrogate species.

 We hypothesize that, as each species has different needs and dispersal capacities, the sections that maximize habitat connectivity will differ from one taxon to another. However, the aim of the study is

 to identify optimal sections that will enable a trade-off to be found to increase habitat connectivity for as many species as possible.

#### **Materials and methods**

*Study area*

 The study area is located in the Haute-Savoie department, France. The section of interest is a 25 km long stretch of the Arve river between the towns of Magland (upstream) and Bonneville (downstream). As the spatial analysis methods used require the use of raster layers, the study area considered is a 160 157 km<sup>2</sup> rectangle encompassing the 25 km stretch of river studied (Fig. 1). The Arve has a mean annual 161 discharge of 27.4 m<sup>3</sup>/s (measured at the hydrological station of Sallanches). The elevation in the study site ranges between 440m (Bonneville) and 500m (Magland). It is mainly (54% of the linear) located in an urbanized area (Fig. 1). It is characterized by number of bank protections along the river corridor to control erosion. These protections consist of riprap (46.1% of the total length, see sample photo in the supplementary material), mixed technique (riprap toe with vegetation on the upper part of the bank, 2.2%) or soil water bioengineering structures (live toe willow fascines, 0.5%, see sample photo in the supplementary material). Natural banks represent 51.2% of the total length. The detailed cartography and typology of the bank protections were provided by the river manager (Syndicat mixte d'aménagement de l'Arve et de ses affluents hereafter named SM3A) and supplemented by a field observation campaign along the entire length to verify the information and assess the vegetation height.



 Fig. 1 Land cover map of the study area. Data from France OSO soil mappin[g \(Inglada et al., 2017\).](https://www.zotero.org/google-docs/?CCrfCP) The names of the main towns along the river are shown in white boxes.

## *Selection of surrogate species*

 Species differ in their response to landscape configuration, i.e. the size and position of habitat patches and the composition of the landscape matrix (Vos et al., 2001). In order to avoid performing a connectivity analysis for all the species surveyed in the area, we developed a method to select a few surrogate species that cover a diversity of biological traits.

 First, we listed the 54 vertebrate species present in our study area by compiling georeferenced data from inventories carried out by the SM3A, the natural heritage information system SINP (INPN, 2023) as well as the data present in the inventories of the zone of floristic and faunistic ecological interest (INPN, 2023b). As these three sources were not exhaustive in terms of invertebrate inventories, we limited our analysis to the vertebrates found in the area. The use of vertebrates as umbrella species in conservation studies is relevant and very common (Yang et al., 2023). The detailed species list is provided in the Appendix 1 in the supplementary material. Then, for mammals, reptiles and

 amphibians, we selected only species using watercourses as habitats according to IUCN (environment 5.1 in IUCN nomenclature, see IUCN (2022) for detailed information). For birds, we only selected species whose main breeding habitats are moving riverbeds, riparian habitats and torrents, according to the habitat preference classification of Roché et al. (2016), and removed generalist species. Our final list encompassed 18 species that depend on riparian habitats: five amphibians, 10 birds, two mammals and one reptile (Table 1).

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 Table 1. The 18 candidate species surveyed in the area after retaining only those strictly associated with watercourses. The four variables (biological traits) used to perform the factor analysis are indicated, as is the cluster resulting from the hierarchical clustering. Id = unique identifier used for each species. The paragon species selected for each cluster are in bold.





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201 We based our species selection on several biological traits that are representative of species ability to reach distant habitats. As we restricted the analysis to those species dependent on a single habitat type (riparian habitats), we did not include habitat preferences in the factorial analysis. Following Henle et al. (2004), we selected traits linked to vulnerability to habitat change: two functional traits related to dispersal capacity (maximum dispersion distance and dispersal mode) and two life history traits related to population dynamics (maximum longevity and fecundity per year). For each species, we collected information on these traits from the literature, using either existing databases and generalist papers (Sutherland et al., 2000; Grimm et al., 2014; Trochet et al., 2014; Myhrvold et al., 2015; Oliveira et al., 2017) or articles more specific to each taxon (Rouland & Migot, 1990; Smith & Green, 2005; Decout et al., 2010; Meister et al., 2010; Quaglietta et al., 2013). Depending on data precision and accessibility in the literature, we coded the traits either as categorical or as continuous variables. The categorical variables were the maximum dispersion distance (coded into the following

213 categories: 0.1-1 km; 1-10 km; >10 km) and the dispersal mode (categories: aerial; terrestrial; terrestrial and aquatic). The continuous variables were the maximum longevity (in years) and the fecundity per year (in number of offspring) (Table1).

 To select surrogate species, we performed a k-means clustering on these four variables using the 217 Hierarchical clustering on principal components (HCPC), based on a factor analysis for mixed data (FAMD). To do so, we used the functions FAMD and HCPC of the package FactoMineR (Lê et al., 2008) in R environment (R Core Team, 2022). We kept the number of clusters with the highest relative loss of inertia as recommended by Lê et al. (2008). For each cluster, we selected the species located closest to the barycenter as surrogate (thereafter called paragon species).

*Habitat characterization*

 SM3A provided us with a detailed mapping of the habitats along the rivers in the study area. For more remote areas, we used Occupation des Sols (OSO) soil mapping [\(Inglada et al., 2017\),](https://www.zotero.org/google-docs/?AZ49gI) which provides 225 a 23-class land cover map at 10m resolution produced from satellite data. For the aquatic environment, the location of obstacles to flow (weirs) was provided by the SM3A.

227 We treated the riverbanks separately from the rest of the study area. We mapped them using a 10m 228 buffer zone along the Arve River. We determined three characteristics of the banks: type of protection structure, vegetation height and slope. The type of protection structure (5 categories: riprap, lower- bank ripraps with upper-bank plantings, bare rock, natural bank, live toe fascine) and the vegetation height (3 categories: low vegetation cover <25%, mainly herbaceous or shrubby vegetation, mainly tree cover) were mapped in a field survey (2017). The slope was determined using a 1m-resolution digital elevation model obtained by LIDAR and supplied by SM3A. We used two categories: low slope (<60%) and steep slope (>60%). Crossing these three characteristics allowed us to obtain a classification of the banks in 5\*3\*2=30 categories. 68.1% of the riverbanks were categorized as steep (vs. 32.9% as shallow). Vegetation cover on the banks was sparse (<25%), herbaceous/shrubby and woody on 4.2%, 14.1% and 81.7% of the linear length, respectively.

238 We carried out the cartographic work using QGIS (QGIS.org, 2023). We combined the collected 239 information in a land use raster map adopting a resolution of 2m, the finest resolution usable by 240 Graphab software (Foltête et al., 2012, 2021) in our computer configuration (Intel Core i7, 32GB RAM).

241 *Resistance maps*

242 We first determined the main habitat types for each target species, based on literature and discussions with experts. All patches of potential habitats were mapped within the study area. Then, we classified 244 the different elements of the matrix according to their resistance to species movement based on literature and expert opinion. We used the literature to select prior resistance values, which were then corrected and validated by local naturalist experts. The resistance values were attributed on a 247 logarithmic scale according to previous papers that demonstrated the importance of using a contrasted resistance scale in landscape connectivity models (Verbeylen et al., 2003; Clauzel et al., 2013; Clauzel & Godet, 2020). Our resistance values thus ranged from 1 (habitat patch) to 10,000 (barrier), with a value of 10 assigned to elements favorable to movement, a value of 100 to elements neutral to movement and a value of 1000 to elements unfavorable to movement (Table 2 and Table 252 3). The resistance map obtained covered the entire study area (157km<sup>2</sup>).

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254 Table 2. Resistance values assigned to the various landscape elements based on literature and expert 255 opinion, for the 4 surrogate species.





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257 Table 3. Resistance values assigned to different riverbanks based on literature and expert opinion, for

258 the 4 surrogate species, depending on their protection structures, vegetation cover and slope.





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#### 260 *Assessment of connectivity using landscape graphs*

 We constructed landscape graphs for each species/scenario combination using Graphab 2.4 software (Foltête et al., 2012, 2021). To avoid modeling patches too small to constitute real habitats for 263 vertebrate species, we kept habitat patches larger than 500m<sup>2</sup> as nodes for the graphs. Graph links were constructed between all habitat patches, and we only retained the ones that were shorter than the maximum dispersion distance of each species based on literature. The link impedance between patches was calculated in least-cost distances based on the previously constructed resistance map.

 We assessed the landscape connectivity by computing the Probability of Connectivity (PC) index for each graph. The PC accounts for the amount of reachable habitat for the target species (Saura & Pascual-Hortal, 2007). It is defined as the probability that two individuals randomly selected within the landscape are located in interconnected habitat patches. The PC takes into account 1) the total amount of habitat available, 2) the spatial configuration of habitat patches, 3) the resistance of the landscape matrix to species movement and 4) the dispersal capacity of the target species modeled:

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$$
PC = \frac{\sum_{i=1}^{n} \sum_{j=1}^{n} a_i a_j p_{ij}^*}{A_L^2}
$$

274 with  $a_i$  and  $a_j$  the respective capacities of habitat patches *i* and *j and A<sub>L</sub>* the total landscape capacity. 275 We considered habitat patch capacity to be equal to habitat patch area.  $p_{ij}^*$  is the maximum value of 276 the probability of dispersal  $p_{ij}$  calculated on all the possible paths connecting patches i and j.  $p_{ij}$  is the 277 direct probability of dispersal between the two habitat patches *i* and *j* and is a negative exponential 278 function of the distance between them  $d_{ij}$  in km. :

$$
p_{ij} = e^{-kd_{ij}}
$$

 k is a constant expressing the intensity of the decrease in the probability of movement (p) with distance 281 (d). The value of k was fixed such that  $p_{ii} = 0.05$  when  $d_{ii}$  equals the maximum dispersion distance of the target species

*Contribution of each habitat type to landscape connectivity*

 To focus on the importance of each habitat type in the ecological network, we classified habitats into three categories: riverbanks, other river-related habitats (e.g. gravel banks in the middle of the watercourse) and offstream habitats (e.g. pond banks, wetlands). To test each species' level of dependence on riverbank versus offstream or other river-related habitats, we created new graphs, each time removing all habitat patches in a given category. We replaced the removed habitats with matrix elements to which we assigned a resistance value equal to the average resistance of the study area. For each graph, we calculated the PC and the loss of PC compared with the current situation (in %).

292 We scaled this analysis down to the habitat patch level. We also evaluated the importance of each patch of habitats, using the dPC (for delta PC) index calculated with Graphab. dPC is calculated by removing each patch one by one from the network and calculating the associated percentage of PC change. This quantifies the importance of each habitat patch in the spatial ecological network [\(Saura](https://www.zotero.org/google-docs/?hbGby1)  [& Rubio, 2010\).](https://www.zotero.org/google-docs/?hbGby1) To control for habitat patch area, we also calculated the dPC/area index, which quantifies the connectivity importance of each patch per surface unit, which seems appropriate in a context where riverbanks often represent very small surfaces and neighbor larger habitats (e.g. lakes and reservoirs).

*Effect of soil water bioengineering structures on habitat connectivity*

 To investigate the potential for improvement and further degradation in the studied landscape, we drew up three contrasting scenarios of land use change and compared them with the baseline scenario, reflecting the current landscape configuration: 1) the 'riprap to fascines' scenario, or a best-case scenario, in which all riprap structures would be replaced in the study area with low-slope, live toe  fascines and plantings above, considered *a priori* to be the most favorable situation for connectivity; 2) the 'fascines to riprap' scenario, in which the existing live toe fascines would be transformed into riprap; and 3) the 'fascines and natural banks to riprap' scenario, or worst-case scenario, in which the live toe fascines and all natural banks would be replaced by riprap. The first scenario was designed to assess the potential for improving habitat connectivity through SWBE structures. The second would quantify the contribution of existing live toe fascines to the observed connectivity. The third was designed to assess the contribution of existing natural banks to connectivity. We computed the PC for each scenario and each target species. The PC of the baseline scenario was used to compare the overall connectivity of the ecological network between target species.

## *Prioritization of bank sections to restore*

 As the 'riprap to fascines' scenario is not a realistic short-term development scenario, we tested intermediate scenarios where the manager would only have the means to restore one section of the bank. The aim of this approach was to identify the sections that would offer the highest gain in connectivity for the target species. We first isolated the riprap embankments. Using the QGIS plugin Polygon divider, we divided them into sections of around 100 m in length. For each target species, we then systematically changed the resistance of each section one by one, switching from a riprap resistance value to a live toe fascine resistance value. For each stage (one species and one section), we recalculated the least-cost paths and recreated a graph. For each graph, we calculated PC change, expressed as a percentage of change in connectivity (positive or negative) relative to the baseline scenario. We obtained a change in connectivity for each surrogate species. This made it possible to calculate the impacts of the restoration of any 100 m section, and thus to select those that would maximize connectivity gains once restored. Finally, to test for the sensitivity of this analysis to section size considered, we repeated the operation using 500 m sections (an equally plausible landscape change scenario). A total of 214 sections of 100 m and 77 sections of 500 m were tested one by one.

**Results**

## *Selection of surrogate species*

 The first two components of the HCPC captured 73% (46.6% and 26.4% respectively) of the variation of the four variables reflecting traits related to dispersal capacity in 18 species (Fig. 2). The variables 'maximum longevity' and 'fertility per year' were negatively correlated. At the same time, dispersal mode and maximum dispersion distance appeared to be related (Fig. 2A). In particular, species that fly can disperse over longer distances.



 Fig. 2 Factor analysis for mixed data (FAMD) performed on the 18 species and projected into the factorial map formed by the first two components. A: Qualitative and quantitative variables (traits). The continuous variables appear in black while the categorical variables appear in red. B: Species clustered according to their traits. Each point represents a species, and numbers correspond to species identifiers mentioned in Table 2. The hierarchical principal component clustering (HCPC) resulted in four clusters. The silhouettes of some emblematic animals are shown for illustrative purposes

 Hierarchical clustering identified four clusters, each comprising between two and 10 species (Fig. 2B, Table 1). Clusters 1 and 4 were taxonomically homogeneous, being composed of birds and amphibians, respectively. Cluster 2 included one reptile species and one amphibian, while cluster 3 contained two mammals and two amphibians (Table 1). We labeled the clusters according to their main ecological characteristics: "flying animals" (Main dispersal mode = 'Aerial'), "amphibious weak dispersers" (Main dispersal mode = 'Terrestrial and aquatic' and Maximum dispersion distance = "100m–1km'), "amphibious strong dispersers" (Main dispersal mode = 'Terrestrial and aquatic' and Maximum dispersion distance = "+10km') and "walkers" (Main dispersal mode = 'Terrestrial). The paragon species of these four clusters were the common sandpiper (*Actitis hypoleucos*), the barred grass snake (*Natrix helvetica*), the Eurasian beaver (*Castor fiber*) and the common toad (*Bufo bufo*), respectively. These four species were used as target species thereafter.

## *Contribution of each habitat type to landscape connectivity*

357 For the Eurasian beaver, the main habitats identified (0.29 $km^2$ , 0.2% of the total study area) were wooded banks with no ripraps, with a quiet zone of at least 10m around them, free of any infrastructure (roads, tracks, buildings), as well as water bodies less than 30m from the Arve riverbed. For the common sandpiper, the main habitats (0.29km², 0.2% of the total study area) were gravel banks, riparian vegetation with small cattail and riverbed with herbaceous riparian vegetation. For the barred grass snake, the main habitats (0.52km², 0.3% of the total study area) were lakes and reservoirs banks and low-slope herbaceous stream banks. Finally, for the common toads, main habitats (0.57km², 0.4% of the total study area) were wooded riverbanks with low slopes, lake banks and reservoirs.

 The highest probability of connectivity (PC) in the study area was found for the common sandpiper (PC 366 = 3.0\*10-6), followed by the Eurasian beaver (PC = 2.1\*10-6), the common toad (PC = 8.1\*10-7) and 367 the barred grass snake (PC =  $6.3*10-7$ ).

 Calculating PC with each habitat type removed one by one, we observed that the Eurasian beaver was the species most dependent on riverbank habitats, which contribute to almost all habitat connectivity  for this species (Table 4). The common sandpiper main habitats were gravel banks, which were classified as 'other river related habitats'. Habitat connectivity for the common toad depended on both riverbank and offstream habitats. Connectivity for the barred grass snake was mainly linked to offstream habitats (Table 4). Note that it is normal for the sum of PC losses for each species to be >100%, as several habitat patches of different types may be redundant in terms of connectivity.

375 Table 4. Variation in PC obtained by removing each habitat type one by one. It is expressed as a 376 percentage loss compared with the current situation.



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 Calculating the dPC for each habitat patch showed that for the Eurasian beaver and the barred grass snake, each offstream habitat patch contributed more in absolute values to total network connectivity than riverbank habitat patches. To a lesser extent, this was also the case for the common toad. In our landscape, the only habitats of the common sandpiper were gravel banks (classified as 'other river related habitats'), so no comparison across habitat types was possible. When controlling the contribution of habitat patch to overall connectivity by habitat patch area (dPC/area), riverbank habitats appeared to contribute more to overall network connectivity than offstream habitats for the



3).



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 Fig. 3 Boxplot of the distribution of the delta Probability of connectivity (dPC) computed for each habitat patch, differentiating between riverbank habitats, other river-related habitats (mainly gravel banks) and offstream habitats. Top: dPC is shown as a percentage of PC change. Bottom: dPC is divided by patch area to eliminate the 'area effect' associated with the PC calculation method. We separated species into two groups for easier reading

## *Impact of landscape change scenarios*

395 The live toe fascines were identified as very favorable environments (resistance  $= 1$ ) for the Eurasian beaver and common toad, and neutral (resistance = 100) for the common sandpiper and barred grass snake (Table 3). The 'riprap to fascines' scenario responded in an average increase in PC of 15% for the four target species. The 'fascines to riprap' scenario had no effect (+0%), while the 'fascines and natural  banks to riprap' scenario corresponded to a PC decrease of 2%. However, these values masked strong differences among species. The barred grass snake (amphibious low disperser) and the common sandpiper (aerial disperser by flight) were insensitive to any bank modifications. In contrast, the 'riprap to fascines' scenario increased connectivity by 46% and 14% for the Eurasian beaver (amphibious high disperser) and the common toad (walker), respectively. For these two species, the 'fascines and natural banks to riprap' scenario led to a decrease in PC of 4% and 5%, respectively (Table 5).

405

- 406 Table 5. Percentage change in probability of connectivity (PC) relative to the baseline scenario, for each
- 407 species and each scenario.



408

## 409 *Prioritization of bank sections to restore*

 As the analysis of the different scenarios showed that two of the four species (the common sandpiper 411 and the barred grass snake) were insensitive to bank modifications, the prioritization of the sections to restore was only made for the other two species (Fig. 4). Whatever the section length (100m or 500m), the vast majority of the sections tested had little or no effect on connectivity if they were the only ones to be restored. Thus 98.1% (210/214) of the 100m sections and 94.8% of the 500m sections  (73/77) increase the PC of any species by less than 3%. Only four 100m sections and four 500m sections offered greater potential for connectivity improvement. Over the 100m sections, two of them would improve PC for common toad by more than 7.5% and two would improve PC for beaver by more than 12%. For the 500m sections, two significantly improved connectivity (gain above 5%) for common toad (+13.1% and +14.2% respectively), and two for Eurasian beaver (+12.4% and +32.7% respectively). 420 However, the sections that best improved connectivity could only do so for a single target species at a time. In other words, the most favorable sections for the Eurasian beaver did not at all intersect with the most favorable sections for the common toad.



424 Fig. 4 Gains in habitat connectivity for two target species achieved by replacing riprap sections with 425 live toe fascines. The orange dots represent 100 m sections, while the blue triangles represent 500 m sections. The gain in connectivity for each species(Eurasian beaver *Castor fiber* and common toad *Bufo bufo*) is expressed as the PC gain relative to the PC of the baseline scenario. Rectangles highlight sections with a significant gain in connectivity (above 5%) for a given species

 Fig. 5 confirmed that the sections favorable to the two target species were different and distant. The high-priority sections for the restoration of Eurasian beaver riparian corridors were located in the town of Cluses, on either side of a weir (B). The high-priority sections for the restoration of common toad riparian corridors were located upstream (east) of the study area, under the bridge over which the highway crosses the Arve river (C). The location of the sections to restore did not change when we used 500m sections, which suggests that at this spatial scale the prioritization does not depend on the length of the section considered (see Appendix 3 in the supplementary material).



 Fig. 5 PC gain obtained by replacing one by one 100m riprap section with a live toe fascine. Sections are represented by circles for visibility. As each section is tested for two species, the circles are divided in two. The gain in connectivity is represented by a gradient of blue (Eurasian beaver) or red (common toad). We provided detailed inserts B, C for greater readability

## **Discussion**

 We showed that using soil water bioengineering (SWBE) structures instead of civil engineering for riverbank protection could greatly change landscape connectivity for some species, while showing no effect for others. Our results showed that not all tested vertebrate species responded in a similar way to changes in riverbank structures, some of them not responding at all (common sandpiper and barred grass snake). Restoring small, well-positioned bank sections could lead to significant gains in connectivity for certain species like the Eurasian beaver or the common toad. However, sections that simultaneously favor all taxa were not detected.

### *Selection of surrogate species*

 We used a method for selecting surrogate species based on the use of species traits linked to sensitivity to landscape fragmentation. This method was mainly based on data from generalist databases covering a wide number of species (Grimm et al., 2014; Trochet et al., 2014; Myrvolt et al., 2015) and is therefore easily transposable to other case studies. However, for some species, more specific literature would be required for conducting this analysis. In our study, this was particularly the case for certain reptile and amphibian species, for which the literature is less extensive than for birds and mammals.

 Our method enabled us to highlight four clusters of species, and therefore four surrogate species, to represent the diversity of traits related to dispersal capacities and life cycle present in our pool of candidate species. This number of species was slightly lower than the results o[f Meurant](https://www.zotero.org/google-docs/?YEdu9Y) et al. [\(2018\)](https://www.zotero.org/google-docs/?YEdu9Y), who used an initial list of 296 vertebrate species – compared with 18 in our study, which probably corresponds to a much greater diversity of biological traits. In this configuration, they demonstrated that selecting 5 to 7 species based on traits was generally sufficient to retain the majority of 84 conservation criteria. The use of surrogate species representative of their cluster, rather than virtual species in connectivity analysis, seems also better suited to communicate with river managers and  decision-makers. Generalist species that use riparian habitats may also be sensitive to bank modifications. Nevertheless, we have chosen to include only specialist species in the analysis, which are particularly sensitive to landscape modifications [\(Devictor et al., 2008; Keinath et al., 2017\).](https://www.zotero.org/google-docs/?dVMfes) We 470 believe this conservative approach is relevant to environmental management research.

 Unsurprisingly, the factorial analysis showed that the variables 'maximum longevity' and 'fertility per year' were negatively correlated. This is a good example of the r/K dichotomy [\(Pianka, 1970\),](https://www.zotero.org/google-docs/?Oo5XYA) which explains why there's always a trade-off between quality and quantity of offspring. The so-called r 474 species (mainly represented by the amphibians in our study) have a reproductive strategy that enables 475 them to deal with major variations in the availability of vital resources. They are characterized by short life spans and high annual fecundity. K species invest more in the survival of their young in a context 477 of predictable access to resources. They are characterized by a long lifespan and low annual fecundity. On the other hand, there were a statistical relationship between dispersal mode and maximum dispersal distance. In particular, species capable of flight are able to disperse further, while the walking group disperses over shorter distances. The four clusters identified, "flying animals", "amphibious weak dispersers", "amphibious strong dispersers" and "walkers", appeared to be ecologically coherent. The fact that they were not all taxonomically homogeneous is consistent with the results of Meurant et al. (2018) who showed that grouping the species by taxon was not necessarily the most appropriate approach.

#### *Connectivity analysis and riverbank change scenarios*

 PC ranking mainly reflected the target species's dispersal capacities in the studied landscape. It was logical to find that connectivity of the landscape was highest for the common sandpiper first, as birds are naturally less sensitive to landscape fragmentation and barriers compared to other taxa such as reptiles [\(Keinath et al., 2017\).](https://www.zotero.org/google-docs/?OYTz1s) Next came the Eurasian beaver, which disperses mainly along watercourses (Table 2) followed by the common toad and barred grass snake, which disperse mainly

 along terrestrial habitats and are therefore more sensitive to the quality of the landscape matrix [\(Keinath et al., 2017\).](https://www.zotero.org/google-docs/?spJHk7)

 Interestingly, scenario analysis revealed significant disparities between species (Table 5). The common sandpiper was not sensitive to changes in riverbank structures, which can be explained by its ability to fly and high dispersal distances. Changing from riprap to vegetated banks and vice versa had little impact on it, particularly as its main habitat is gravel banks in the middle of watercourses. The importance of gravel banks in landscape connectivity for this species was therefore significant (Fig. 3).

 The barred grass snake (*Natrix helvetica*) had some dependence on riverbanks (Table 4). However, it did not appear to be sensitive to riverbank change scenarios at all (Table 5). This can be explained by the fact that riprap banks and fascines were categorized neither as favorable habitats, nor as elements that particularly impede this species movements. Experts therefore assigned them an equal resistance value (100) (Table 3). Consequently, switching from one type of bank to another as part of the scenario did not affect landscape connectivity for this species. This result is interesting from a management perspective, as it shows that ecological restoration is not necessarily favorable for all species, particularly species that are well adapted to human infrastructures or degraded habitat (Walts & Wallace Covington, 2004; Noreika et al., 2015).

 The analysis of connectivity for Eurasian beaver (*Castor fiber*) and common toad (*Bufo bufo*) showed that both species seemed to be highly dependent on riverbank habitats for their connectivity (Table 4). The 'riprap to fascines' scenario greatly improved connectivity, while the 'fascines and natural banks to riprap' scenario reduced it (Table 5). The 'fascines to riprap' scenario only concerned the replacement of the single live toe fascine in the landscape (0.5% of the total bank linear), so its impact on connectivity was expected to be low. The dPC analysis showed that offstream habitats could be very important for habitat connectivity in absolute value (Fig. 4). This was particularly the case for the Eurasian beaver, which occurs in offstream water bodies that can cover large areas of the study zone (Table 2). When divided by surface area (dPC/area), the importance of riverbank habitats for ecological

 connectivity appeared to be greater, highlighting their disproportionate ecological importance in relation to their surface. Their importance had already been demonstrated by Sabo et al. (2005), who showed that riparian habitats increased regional species richness by >50% on average worldwide, despite their narrow surface area. Riparian habitats are therefore what Hunter (2017) calls small natural features with large ecological roles. Nevertheless, maintaining favorable offstream habitats in addition to riverbank habitats for semi-aquatic species seems to be a good solution for maintaining or improving landscape connectivity. This is particularly the case for species that are not strictly river- dwelling, such as the common toad and the barred grass snake (Table 3). This result is consistent with previous studies showing the importance of the offstream wetland network in ecological connectivity, particularly when riverbed habitats are subject to change as a result of hydromorphological variations like flooding (Karim et al., 2012). Our study shows that this is also the case in urbanized areas, which are much less subject to hydromorphological variations due to the channelization of rivers. This study is based on resistance surfaces parameterized based on expert opinions and the literature. It could be useful to supplement it with field studies on the actual movements of species, for example using genetics or telemetry.

#### *Prioritization of bank sections to restore*

 Our prioritization analysis showed that at the tested scale (100 and 500m-long sections) there was no single section where restoration would maximize connectivity for all the target species. On the contrary, the majority of sections tested had only a limited effect on connectivity when considered separately (PC gain < 3%). The few sections with a strong effect were only effective for one species at a time, either the Eurasian beaver or the common toad (Fig. 4). Multi-species connectivity studies often reveal inter-taxon disparities in response to habitat change (e.g. Albert et al., 2017; Meurant et al., 2018). It is noteworthy that a few sections have a very high potential for increasing connectivity, and their restoration alone would allow to achieve or approach the level of connectivity of the best case 540 ('riprap to fascines') scenario. For example, one 500m section would increase connectivity for common  toad by 14%, which is the potential total gain of the 'riprap to fascines' scenario in the studied riverine landscape. For Eurasian beavers, a single section can increase PC by more than 32%, close to the total gain of 46% for the 'riprap to fascines' scenario.

 A cartographic analysis of the location of the sections tested showed that there were key sectors of the river where the priority sections to restore were concentrated. These were areas whose position in the habitat network made the reduction of their resistance to movement particularly interesting for connectivity. For example, panel B (Fig. 5) showed that the most interesting sectors to restore in terms of habitat connectivity for Eurasian beavers were located on either side of a weir in an urban environment. The weirs were identified as potential barriers to beaver longitudinal movement (Table 2). As the concerned weirs were located in urban areas, the environments above the banks were not crossable by beavers (built-up areas). The fact that the beavers could cross the banks laterally (10m wide) were not sufficient to enable them to reach areas favorable to movement. They therefore needed to travel a longer distance along the banks to bypass the weirs and move from one section of the river to another. This cost of movement would be reduced by replacing riprap with SWBE structures, improving the crossability of the banks to improve habitat connectivity. Panel C (Fig. 5) contained the main sectors to restore for the common toad. These were located under the highway bridge. As the highway itself is a strong barrier to toad movement, being able to pass underneath it would increase habitat connectivity between downstream and upstream. This could be made possible by replacing the bank riprap under the freeway bridge with a fascine, which is more permeable to toad movement than riprap.

 Our analysis therefore showed that the restoration of small, well-chosen areas of riverbanks could enable semi-aquatic animals to avoid ecological barriers present in urbanized landscapes (buildings, roads) or in riverbeds (weirs). Detecting and removing barriers to movement to improve connectivity is a well-known issue in landscape ecology, in both terrestrial (McRae et al., 2012) and aquatic contexts

 (Kraft et al., 2019). In the case of the Eurasian beaver, it is likely that the same result could be achieved by removing the barrier in the riverbed (weir) as by restoring the surrounding banks.

 The prioritization analysis was not very sensitive to the size of the sections tested. We could expect that switching from 100m to 500m sections should reveal new favorable zones, but the most favorable zones to restore did not change (Appendix 3). The switch just revealed low-priority areas (potential PC gain <3%) for the common toad and the Eurasian beaver. However, high-priority areas were immediately apparent when testing 100m sections. The switch to 500m only confirmed the potential PC gain (Fig. 4, Appendix 3).

#### *Conclusion: insight for management*

 Our study proposes a valuable decision-making method to help river managers determine where to restore bank protection structures using soil water bioengineering. As many species with variable dispersal capacities may be present in the study area, we recommend that managers rely on accurate, up-to-date species inventories to guide management choices. We showed that it can be difficult to satisfy the needs of all riparian habitat species when selecting a section to restore. This result supports previous studies showing that different riparian species can respond differently to deleterious changes in their environment (e.g. Van Looy et al., 2013). It seems logical to obtain similar results for a priori more favorable changes in the landscape, such as ecological restoration. To improve habitat connectivity in a multi-species context, it may be worthwhile to restore several small, appropriately located sections according to the ecological requirements of several target species. It may also be worthwhile to focus on one or two species of particular concern (e.g., a species with high conservation value) to limit the number of target species with different ecological needs. As the restoration of connectivity is not necessarily at the heart of all restoration projects, this choice of target species can be made as part of a more general framework for defining the objectives of the restoration project (Hallett et al., 2013; Coutinho et al., 2023). We believe that our method will find its place as a decision-making tool for planning restoration projects, alongside others from other fieldsthat may include other

- areas of ecology as well as governance or economic benefit criteria that may be crucial to the chances
- of success of ecological restoration projects (Hallett et al., 2013).

## **Authors' contribution**

 AE, LB and MA designed the study. BD and TH performed the field work. EB performed the analysis and led the writing of the manuscript. AE, LB, MA and EB contributed critically to the drafts and gave final approval for publication.

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