

Connectivity among wetlands matters for vulnerable amphibian populations in wetlandscapes

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1 Connectivity among wetlands matters for vulnerable amphibian

2 populations in wetlandscapes

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Abstract

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- 18 Wetlands have been degraded and destroyed, resulting in the decline of many
- wetland-dependent species populations. Many conservation efforts are based on
- 20 protection of individual wetlands; however, fluxes of energy, materials and organisms

between wetlands create important structural and functional connections upon which several species depend. We investigated the role of individual wetlands within a wetlandscape in sustaining an amphibian population. Wetlandscapes were represented as networks, where nodes were wetlands and links were flows of organisms described by an amphibian population model. Relationships between a wetland's connectivity to the other wetlands and the abundance of amphibians under different wetland management strategies were examined. The first finding was that wetlands within a network can be classified into sinks (where local mortality exceeds birth rate), sources (where local birth rate exceeds mortality), and pseudo-sinks (where excessive immigration maintains the population above the carrying capacity). These three wetland classes have low, medium, and high *indegree* (a parameter reflecting a wetland's connectivity), respectively. The second finding was that management interventions in wetlands have different consequences according to the wetland's *indegree*: wetland removal has the worst impact on amphibian populations if the wetland is a source, and wetland restoration has the best impact if the wetland is a pseudo-sink. These findings provide support for policies that managing wetlands not as independent objects but as integral parts of the wetlandscape.

Key Words

Connectivity, Ecological network, Wetland configuration, Amphibian model, Graph theory, Population dynamics.

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1. Introduction

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Wetlands are important ecosystems as they provide several functions and services (Calhoun et al. 2016, Yao et al. 2016, Creed et al. 2017) and constitute an important source of biodiversity (Costanza et al. 1997, Gibbs 2000). In recent decades, many wetlands have been drained because of urban or agricultural expansion (Davidson 2014, Dixon et al. 2016, Golden et al. 2017). Wetland loss impacts on biodiversity both directly, by removing habitat (Gibbs 2000), and indirectly, by increasing the distance among remaining wetlands and resulting in functional isolation and increasing mortality of organisms migrating from one wetland to another (Baguette et al. 2013). Wetlands are not isolated features; on the contrary, they are dynamic, complex ecosystem with biotic and abiotic connections to other wetlands on the wetlandscape (Cohen et al. 2016, Thorslund et al. 2017). Understanding the ecological dynamics of wetlandscapes is important to sustaining biodiversity (Semlitsch and Bodie 1998, Gibbs 2000). In particular, amphibians' survival is influenced by both wetland habitat and wetland connections to other wetlands (Dudgeon et al., 2006). In fact, these wetland qualities determine the success of amphibians' breeding (Mushet et al., 2012): wetland habitat is used by adults for mating and by offspring to complete their metamorphism from egg. Factors such as availability of resources and dispersal capabilities influence the amphibian population in wetlands (Pechmann et al. 1989, Semlitsch et al. 1996). Availability of resources depends on wetland habitat properties (e.g., area, vegetation) and on the number of amphibians competing for available resources. Amphibian dispersal relies on wetland distribution within the surrounding terrestrial habitat. Every

year at the end of the summer, amphibians start their migration through the terrestrial

habitat and the following spring they reach a new aquatic breeding habitat (Pittman et al. 2014). Alteration of wetland habitat and distribution within the landscape, such as wetland loss, negatively influences both breeding and dispersal success by decreasing wetland density and increasing travel distances for amphibians (Gibbs, 1993)."

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Management strategies have been implemented to protect biodiversity promoted by wetlands. Many of these management strategies focus on wetlands of special importance (Amezaga et al. 2002). Policy goals vary from "no net loss" to "net gain" (Accating et al., 2018) to general statements about the need to address adverse impacts to these wetlands (e.g., Calhoun et al. 2016). Few of these management strategies focus on the physical, chemical, or biological connections among wetlands (e.g., Cohen et al. 2016). Although it is widely recognized that wetland connectivity is important for biodiversity (Semlitsch 1996, Semlitsch and Bodie 1998, Skelly et al. 1999, Marsh and Trenham 2001, Cushman 2006), concrete strategies in policies are still not well formulated. The lack in wetland policies of clear operational strategic schemes based on wetland connectivity is at least in part due to the lack or rarity of quantitative assessments of the role of wetland connectivity in sustaining wetland biodiversity. Important steps forward would be to determine if and how wetland connectivity plays a role in sustaining biodiversity in a wetlandscape (Fortuna et al. 2006, Albanese and Haukos 2017), and to explore if management interventions on the wetland itself (i.e., wetland removal or restoration) are influenced by wetland connectivity.

The "sink-source" framework describes the distribution of species in the variety of interconnected habitat patches within a region (Pulliam 1988, Watkinson and Sutherland 1995). According to this framework, a productive patch serves as a source of

individuals, which are dispersed to less productive patches called sinks (Pulliam 1988, Dunning et al. 1992). Pulliam (1988) argued that in sink habitat patches reproduction is insufficient to balance local mortality, whereas in source habitat patches reproduction balances local mortality; the population in sinks is thus maintained by immigration from sources. Most studies classify sinks and sources only by demographic measures (i.e., birth and death rate) (Watkinson and Sutherland 1995). The role of connectivity in the classification of sinks and sources has not been explored.

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Models are useful to test the "sink-source" framework for exploring organism dispersal through wetlandscapes, especially when empirical data are lacking or extremely difficult and costly to collect (Pittman et al. 2014). Patch-based models (e.g., Skelly and Meir 1997, Trenham 1998) focus on population dynamics within patches, which are important to describe fundamental ecological processes such as breeding (Marsh and Trenham 2001), interspecific competition, and predation (Wilbur 1997, Beebee et al. 1996). Patch-based models were successfully applied to wetlandscape (Marsh and Trenham 2001, Wilbur 1997). However, an exclusively patch-based approach does not consider the role of wetland isolation and the mobility of individuals to other wetlands (Cushman 2006). In contrast, network-based models focus on connectivity within network's node and they can be applied to wetlandscape, too (e.g., Albanese and Haukos 2017). They allow to quantify changes to the connectivity of wetlands and identify wetlands critical to the maintenance of system connectivity. Network-based models are useful tools for combining both the within-wetland population dynamics and the dispersal of individuals among wetlands (Estrada and Bodin 2008). Network-based models can be used to identify keystone patches that are integral to the persistence of populations

(Urban & Keitt 2001, Keitt 2003) and to quantify the robustness of populations to wetland loss (Bunn et al. 2000, Hanski 2001, Jordán et al. 2003).

In this paper, we addressed the role of wetland connectivity in determining the role of different wetlands to sustain amphibian populations. We focused on amphibian species characterized by a bi-phasic life-cycle, migrating into different wetlands during the course of their life. We built a model of amphibian population dynamics in a wetland network and we formulated scenarios to address two research questions: how does the connectivity of a wetland influence the abundance of the local population in the wetland itself? And, how does a management intervention on a single wetland (e.g., wetland removal or wetland restoration) influence the total landscape population by changing connectivity within the wetlandscape?

2. Methods

We focused on amphibian species with life history traits characterized by a terrestrial and an aquatic phase, but the approach could be adapted to amphibian species with other life history traits. In summer, amphibians congregate in wetland for mating. At the end of the summer, amphibians leave wetlands and migrate through the terrestrial habitat searching food and refuges for overwintering until the next spring, when they disperse again, looking for aquatic breeding habitat (Pittman et al. 2014). Examples of species having such as life cycle are the northern leopard frog (Rana pipiens) having a dispersal distance ranging between 2 km and 10 km (Kendell 2002), and the great plains

toad (Anaxyrus cognatus) having a dispersal distance ranging between 300 m and 1300 m (COSEWIC 2010).

2.1 Model description

We built a theoretical model for simulating the dynamics of a biphasic life-cycle amphibian population within a wetlandscape. The population dynamics of amphibians consist of a continuous repetition of reproduction phase and migration phase. In the reproduction phase, new offspring are produced in each wetland as a function of the abundance of the local population. In the migration phase, amphibians migrate from a wetland to another, according to how wetlands are connected and their mortality. Our model was conceived in the Medawar zone (Loehle 1990, Grimm et al. 2005) (i.e., we included only the necessary elements for addressing our research questions and to avoid unnecessary details, such as predation phenomena or climate variables like air and surface water temperature of wetlands).

2.2 Wetlandscape

A network consists of a set of nodes that are defined as spatial elements, and links that represent linkages between nodes (Urban and Keitt 2001). We represented a wetlandscape as a network, where the nodes are wetlands, whereas the links are flows of amphibians between wetlands. We connected two wetlands by a link if the distance between a given pair of wetlands was less than or equal to the maximum distance walkable by the amphibians. Links were assigned a weight equal to the potential connectivity, \tilde{c}_{ij} , which is defined as the probability of an amphibian leaving wetland i to choose wetland j as breeding site the following year. The potential connectivity was

determined by the Euclidean distance and the adjacency between two wetlands. To determine the connectivity between two wetlands either a boundary approach or a distance approach could be used. The boundary approach consists of methods that determine the weight based on the presence or the length of a shared boundary (Ermagun and Levinson 2018). The distance approach consists of a spectrum of methods that determine the weight using the geographic distance (Ermagun and Levinson 2018). Both the boundary approach and the distance approach or a combination of them have been used to construct a theory-driven spatial weighting matrix (Dray 2006), as well as biological considerations such as propagation process (Sokal and Oden 1978), patch size (Hanski 1994) and dispersion capability (Knapp et al. 2003). In our work, the potential connectivity represented only the spatial interactions among wetlands and combined the boundary approach and distance approach using a weighted average, with the parameter β as the weight to balance the relative importance of adjacency over the inverse distance (Figure 1). The underlying assumption is that amphibians will move from wetland i to wetland j if the two wetlands are adjacent (see below for definition) or if two wetlands are close each other. The potential connectivity has this expression:

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$$\tilde{c}_{ij} = \beta \cdot a_{ij} + (1 - \beta) \cdot \frac{1}{x_{ij}}$$
 Eqn 1

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The term a_{ij} was the adjacency term and was computed with the Voronoi tessellation (Galpern et al. 2012). This method consisted of a subdivision of the landscape into polygons, each polygon associated with a wetland. The polygon was formed by all the points of the landscape nearer to that wetland than to any other (Kenkel et al. 1989). Since each wetland was the centroid of a polygonal region, for each one we assigned an

adjacency value of $a_{ij} = 1$ if the polygons of the two wetlands were adjacent, and $a_{ij} = 0$ otherwise. The adjacent polygons in the tessellation described hypothesized regions of proximity for the organism within which there is high connectivity (Galpern et al. 2012). The term x_{ij} was the Euclidean distance between wetlands i and j. This term took into account that an amphibian was more likely to go to a wetland at a shorter distance. We set $\beta = 0.5$ as a constant value assuming an equal effect played by adjacency and distance.

2.3 Amphibian population dynamics

The amphibian population dynamics was described using a time-discrete model with a time step t of one year, divided in two sub-steps, the first for breeding in wetlands and the second for migration from one wetland to another (Fig. 1c). In the notation of our model, N_t^i represented the number of amphibians in wetland i at the beginning of year t, $N_{t+\frac{1}{2}}^i$ represented the number of amphibians in wetland i after the breeding phase, and N_{t+1}^i represented the number of amphibians in wetland i after the dispersal phase of year t.

In the breeding phase, the adults in each wetland produced offspring that went through the metamorphic cycle and became juveniles ready to move to reproduce in another wetland the following year. The population $N_{t+\frac{1}{2}}^i$ consisted of the population N_t^i increased by the number of new individuals generated that successfully completed the metamorphic cycle (here after called *newborns*) in wetland *i*.

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$$N_{t+\frac{1}{2}}^{i} = N_{t}^{i} + max\left(0, rN_{t}^{i}\left(1 - \frac{N_{t}^{i}}{K}\right)\right)$$
 Eqn 2

The survival of the new individuals generated was density-dependent: at low density, individuals did not interfere with each other, whereas at higher densities, resources became less available for each individual. We characterized each wetland by its carrying capacity, K, which is the maximum size of the population that each wetland can sustain. Parameter r represented the intrinsic growth rate. The max operator assured that the number of new individuals generated was not negative.

In the migration phase, each amphibian in wetland i moved to wetland j with a certain probability. Such a probability corresponded to the *effective connectivity* w_{ij} , which consisted of the *potential connectivity* \tilde{c}_{ij} modified for taking into account the dispersal distance of amphibians p_{ij} , and the probability of mortality due to distance travelled d_{ij} , and the natural mortality μ :

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$$w_{ij} = \tilde{c}_{ij} \cdot p_{ij} \cdot (1 - d_{ij}) \cdot (1 - \mu)$$
 Eqn 3

The dispersal distance was computed with a negative exponential function of the distance travelled x_{ij} : $p_{ij} = e^{-\delta x_{ij}}$, where δ is the dispersal capacity evaluated as a function of the amphibian maximum walkable distance x_{max} : $\delta = \frac{1}{x_{max}}$. The same functional shape was used in other similar models (e.g., Bunn et al. 2000, Hanski and Ovaskainen 2000, Saura and Pascual-Hortal 2007).

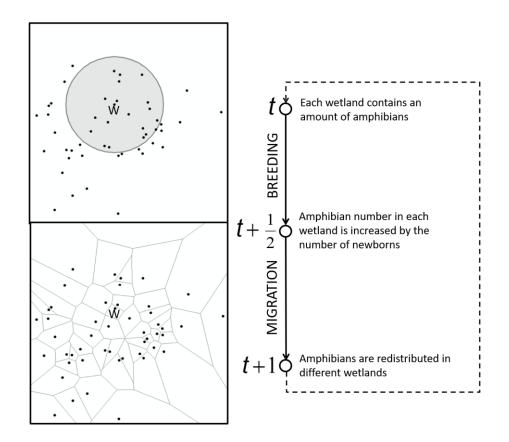


Figure 1. Spatial (panels (a) and (b)) and temporal (panel (c)) schematization of the model. Wetlands are represented by points on a plane. Panel (a): The gray circle around the wetland marked with "W" represents all wetlands within the maximum walking distance that can be reached by the adults in wetland W, but the connectivity probability decreases with the Euclidean distance. Panel (b): Thiessen polygons are represented around each wetland. In case the two polygons have a common boundary, the relative wetlands are considered as adjacent. Panel (c): The cyclic succession of the fundamental events in the model is represented along with notation about time.

The term d_{ij} described the mortality during the winter for migration and it is proportional to the distance x_{ij} between the two wetlands $d_{ij} = \gamma \cdot x_{ij}$. The term γ was a coefficient representing the probability of mortality per distance unit, implicitly including the probability of dying in the search of a wintering place. The natural mortality μ was considered a constant parameter of the amphibian species.

The effective connectivity w_{ij} is the fraction of amphibians in pond j, which choose wetland i as destination and arrive there safely in the following time step. The number of amphibians at the time t+1 that reached wetland i was calculated as:

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$$N_{t+1}^i = \sum_{i=1}^N w_{ij} N_{t+\frac{1}{2}}^j$$
 Eqn 4

The sum $\sum_{i=1}^{N} w_{ij}$ (i.e., the sum of weights of the ingoing links of wetland i) represented the fraction of the amphibians $N_{t+\frac{1}{2}}^{i}$ incoming from the other wetlands of the network at time t+1. N_{t+1}^{i} are defined as Immigrants. The sum $\sum_{i=1}^{N} w_{ij}$ gave a measure of how much the wetland i was reachable from other wetlands in the network and was called indegree (Jordán et al. 2003, Estrada and Bodin 2008): we used indegree for quantifying wetland connectivity.

2.4 Scenarios for different wetlandscapes

We used the population model to run simulations over different randomly generated wetland networks. To investigate the role of wetland indegree on the local amphibian populations we first analyzed the behavior of the system on a single randomly generated wetland network (Single-network scenario) and then analyzed the relationship between wetlands indegree and population on a set of randomly generated wetland networks using a Monte Carlo simulation approach (Multi-networks scenario). To investigate the impact of wetland removal and restoration on the population, we conducted wetland removal and restoration scenarios also using a Monte Carlo simulation approach (Wetland removal and restoration scenarios).

The single-network scenario was used to explore the overall population dynamics and the local population dynamics in single wetlands of the network, with each wetland characterized by a different value of *indegree*. We randomly generated a network of 50 wetlands and ran the population dynamics model on it. For generating the random network we used the algorithm developed by Fachada et al. (2014). This algorithm generates a random network configuration starting from deterministic topological parameters (e.g., Slope, Number of clusters, Total points) and stochastic parameters (e.g., Average separation of line centers along the X axis, Average separation of line centers along the Y axis, "Cluster fatness"). Stochastic parameters varied for each network and were randomly extracted from a uniform distribution of values. Stochastic parameters made it possible to define a network configuration that was not deterministically assigned but had some random variations. The model was initialized to 20 amphibians per wetland and parameterized as in Table 1 with a fixed simulation time horizon. All the wetlands in the network were characterized by the same carrying capacity, K; this allowed us to explore the relationship between wetlands indegree and population, since wetlands only differed for their indegree value, without other confounding factors due to habitat difference. We classified the wetlands of the network into three classes of indegree (low, medium or high) using the Jenks natural breaks classification, a data clustering method that minimizes each class's average deviation from the class mean while maximizing each class's deviation from the means of the other cluster (Jenks 1967). We then explored the local population dynamics in three wetlands, each one belonging to a different class of indegree. Specifically, we investigated 1) the number of newborns produced in the wetlands, and 2) the number of immigrants.

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Parameter	Range of values	Description	Source*
Breeding			_
r(/)	6 - 8	Growth rate without density effect	Dray et. al., 2006
<i>K</i> (individuals)	400 - 600	Carrying Capacity	-
Mortality			_
μ (/)	0.4 - 0.6	Natural probability of dead	Vonesh and De la Cruz, 2002
Dispersal			_
MWD (Km)	3	Maximum Walkable Distance	Ray et al., 2002
β (/)	0.5	Connectivity matrix weight	-

* The reference in which the parameter has a similar range.

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The multi-network scenario consisted of running simulations for several randomly generated networks (all having the same topologic characteristic). This scenario served to assess if the observations made from the single-network scenario could be considered general and to explore with more detail the relationship between *indegree* of the wetland and its local population. We generated 100 networks of 50 wetlands each and ran the population dynamics model for each one of the networks. For each network configuration, the values of r, K and μ were extracted from a uniform distribution within a fixed range. Within each experiment, the parameters were kept constant and K was equal for all the wetlands. The model was initialized to 20 amphibians per wetland with the range of values of parameters provided in Table 1, i.e., the same demographic parameters of the single-network scenario. As in the single network scenario, wetlands carrying capacity was set equal for all the wetlands for all the networks. We ran the simulations over a period of 60 years, checking that this time horizon was long enough to reach a steady state. We compared the population size in each wetland with its *indegree* value, pulling together all the wetlands in all the simulations. We then explored how the population dynamics evolved within the wetland network analyzing the different contribution played by Immigrants, Newborns and Survivals at each time step. The

wetlands within the random networks were divided into three classes (low, medium, high) according to their *indegree*, using the Jenks method. We made pairwise comparison between the distribution of amphibian population abundance within classes with the Kolmogorov-Smirnov and Student's t-test (significance level = 0.05). Kolmogorov-Smirnov test was used to detect the significance of the difference between the distributions of the amphibian population abundance at the steady states in the different classes of *indegree*. Student's t-test was used to detect the difference between the means of the amphibian population abundance at the steady states in the different classes of *indegree*.

Wetland-removal-and-restoration-scenarios were used to estimate the role of wetland indegree on amphibian population when a wetland was removed or restored. The wetland removal scenario consisted of removing a wetland from the network, simulating the amphibian population dynamics in the altered network, and comparing the result with the non-altered network. For a given network, the experiment was repeated by iteratively removing one wetland at a time. For all the other wetlands, we set an equal value for carrying capacity, *K*. We repeated the procedure with 100 randomly generated wetlands and then examined the relationship between the *indegree* of the wetland removed and the impact on the total population. This relationship was independent from the carrying capacity of the removed wetland, since all the wetlands have the same K. The wetland restoration scenario consisted of increasing the carrying capacity of a wetland in the network by duplicating its original value while keeping constant the carrying capacity of the other wetlands in the network. The population dynamics of the altered network was simulated and the results compared to the non-altered network. For a given network, the

experiment was repeated by iteratively enhancing one wetland at a time. We repeated the procedure with 100 randomly generated wetlands and examined the relationship between the *indegree* of the wetland improved and the impact on the total population. We generated 100 wetland networks and repeated the wetland restoration scenario for each network. We then compared the population of the altered and non-altered networks to analyze the impact of management strategies. As in the *multi-network scenario*, the wetlands within the random networks were divided into three classes (low, medium, high) according to their *indegree*, using the Jenks method. We made pairwise comparison between the distribution of amphibian population abundance within classes with the Kolmogorov-Smirnov and Student's t-test (significance level = 0.05) to detect statistically significant differences in the distribution and in the mean value respectively.

The methodology and all the simulations are implemented with Matlab R2016a.

3. Results

3.1 Single-network scenario

The simulation on the single-network scenario showed that the total population abundance, after a transient phase, reached a steady state. The local population in each wetland behaved in the same way. This shows that the length of the time horizon (60 years) was sufficient to represent the population dynamics of the wetland network. Hereafter, we present the results referred to the steady state. For the wetland network generated (Fig. 2A), we calculated the frequency distribution of the *indegree* and divided the wetlands into three classes (low, medium, and high) (Fig. 2B). We investigated the

abundance of the population in three wetlands (marked in Fig. 2A), each one belonging to a different class: Wetland A (low indegree) showed the smallest population, wetland B (medium indegree) showed the largest population, and wetland C (high indegree) showed an intermediate population. Wetland A had a relatively small proportion of newborns, wetland B had a larger proportion of newborns compared to wetland A and C, and wetland C had no newborns (Fig. 3).

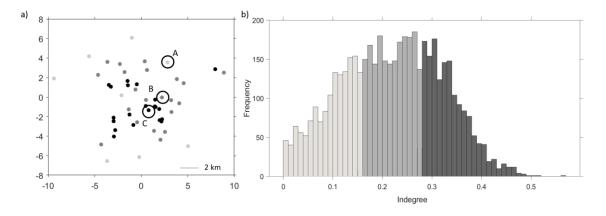


Figure 2. a) Spatial configuration of the 50 wetlands that form a single network; the total population of three wetlands (A, B and C) are given in Figure 3. b) Frequency distribution of indegree in the 100 simulated wetland networks. Indegree was computed for each wetland, and the wetland was then classified into one of three classes (i.e., low (light grey), medium (grey) or high (black) indegree), with the classes defined using the Jenks natural breaks classification method.

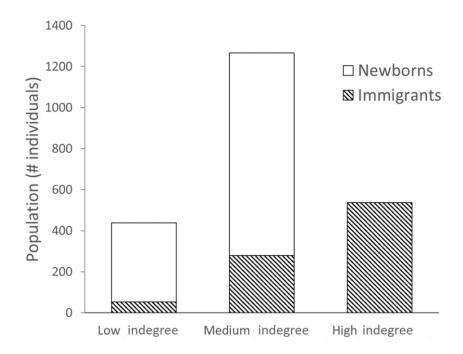


Figure 3. The wetland population, with relative contribution of *Immigrants* and *Newborns*, for three wetlands (A, B, C, see Figure 2a) for the wetland network depicted in Figure 2.

3.2 Multi-network scenario

The behavior of the single-network scenario (Fig. 3) was generalizable (Fig. 4). The low indegree class had the smallest population, the medium indegree class had the largest population, and the high indegree class had an intermediate population (Fig. 4A). The Kolmogorov-Smirnov test revealed that the data belonging to different indegree classes had significantly different distribution (p-value < 0.001). The Student's t-test revealed that data belonging to different indegree classes had significantly different mean values (p-value < 0.02). The contribution of immigrants and newborns to the population for the different classes is shown in the pie charts in Figure 4B. Wetlands in all indegree classes showed that the majority of the population was composed of newborns, but that the proportion of newborns in the population of the high indegree class was not as high as in the low or medium indegree classes. This was because the pie chart could not capture

the natural oscillation given by the model of the population dynamics, but only displayed a median value.

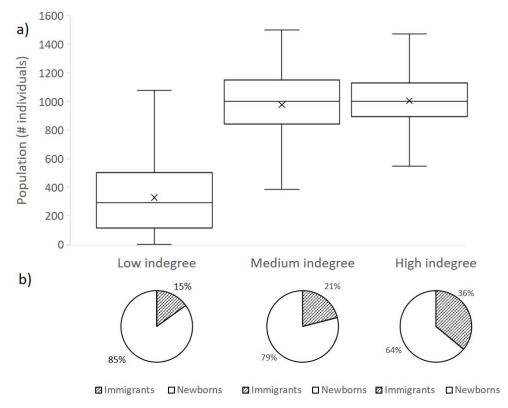


Figure 4. (a) The wetland population in the three classes of indegree, resulting from the Monte Carlo experiment, and (b) the contribution of *Immigrants* and *Newborns* to the total population. The number of spatial configurations of wetlands' network is equal to 100 and each network is made up of 50 wetlands. For each spatial configuration, the population dynamics were simulated, and for each wetland the total population, the number of *Newborns* and the number of *Immigrants* were computed and related to the indegree of each wetland. The three classes of indegree were defined by the Natural Jenks algorithm.

The multi-network scenario showed a non-linear relationship (Figure 5A) between wetland indegree and wetland population; as indegree increased from low to medium the population increased, at medium indegree the population reached a peak, and

as indegree increased from median to high the population decreased. The number of immigrants showed a linear increase as indegree increased (Fig. 5B), and the number of newborns showed a non-linear relationship, similar to the total population (Fig. 5C).

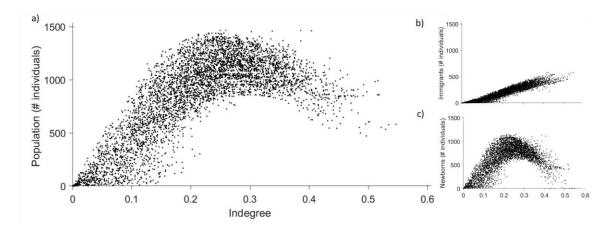


Figure 5. *Indegree* vs. (a) Population in each wetland for each network, which is the sum of (b) the number of *Immigrants* and (c) the number of *Newborns*.

3.3 Wetland removal and restoration scenarios

The wetland removal scenario showed greater negative effects on the amphibian population when the removed wetland belonged to a medium indegree class (Fig. 6B). In contrast, the wetland restoration scenario showed greater positive effects on the amphibian population when the restored wetland belonged to the high indegree class (Fig. 6A). Fig. 7 directly compares the wetland removal and restoration scenarios by showing the deviation in the population abundance of the entire wetland network between altered and non-altered networks, based on the class of indegree of the altered wetland within the network. As before, the Kolmogorov-Smirnov test revealed that the data belonging to different indegree classes had significantly different distribution (p-value < 0.001). The Student's t-test revealed that the data belonging to different indegree classes had significantly different mean values (p-value < 0.001).

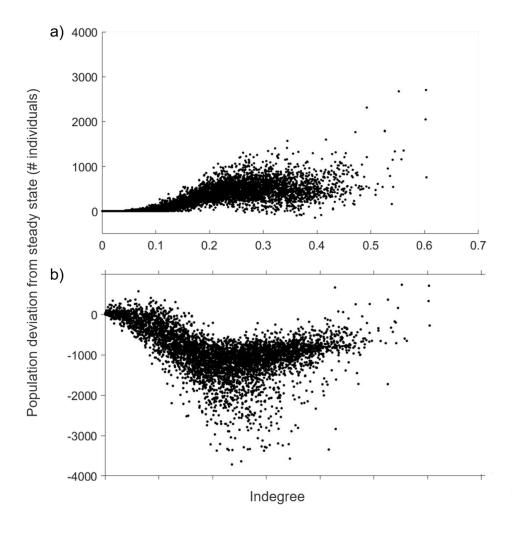


Figure 6. Changes in the population (expressed as deviation from steady state of the altered with respect to the non-altered networks) for different spatial configurations of wetlands' network under the (a) wetland restoration and (b) wetland removal scenario. The deviation from steady state of the altered with respect to the non-altered networks is plotted against the indegree of the altered wetland.



Figure 7. Changes in the total population (expressed as deviation from steady state) within the wetlands' network for different spatial configurations of wetlands classified as low, medium or high indegree and under scenarios of wetland restoration and removal.

4. Discussion

Wetlandscapes sustain sink-source population dynamics with individual wetlands playing different roles (sources or sinks) according to their indegree. We identified source wetlands (where local birth rate is greater than mortality) and sink wetlands (where local mortality is greater than the birth rate, and the population is maintained by continued immigration from source wetlands nearby). Among sink wetlands, we identified two types: those in which the population goes extinct in the absence of immigration, and those in which immigration maintains the population above the carrying capacity. Such a distinction corresponds to true sinks and pseudo-sinks, respectively, according to Watkinson and Sutherland's (1995) terminology. Mortality in pseudo-sinks

is caused by excessive immigration, which provokes excessive mortality due to "density effect", the decline in population growth caused by crowding, predators and competition (Hixon and Johnson 2009). If immigration stops in a pseudo-sink, the population would decline, but only to the carrying capacity. Both true sinks and pseudo-sinks have been observed in nature (Watkinson et al. 1989, Engelhard et al. 2017, Erickson et al. 2017).

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Indegree is a useful proxy for classifying wetlands into sources, sinks, and pseudo-sinks. Wetlands with a low indegree – representing relatively isolated wetlands – could be classified as sinks because few amphibians migrated to these wetlands and consequently they produced few newborns. Wetlands with medium indegree could be classified as sources because they had the highest population. In fact, the number of immigrants arriving in the wetlands and the generated offspring were fully sustained by wetlands resources (expressed by the carrying capacity, K). Specifically, the population within each wetland did not exceed carrying capacity and a large number of amphibians was able to successfully emigrate from these wetlands. These wetlands were highly productive, in fact the number of Newborns exceeded the number of Immigrants and contributed positively to the total wetland population growth. Wetlands with a high indegree – representing relatively connected wetlands – could be classified as pseudosinks because the largest number of *Immigrants* arrived at these wetlands, but the available resources in these wetlands were insufficient to sustain both the immigrant population and the generated offspring. The number of Newborns declined until all newborns died due to a "density effect", that is the population of these wetlands exceeded carrying capacity. Most of the amphibians in these wetlands were *Immigrants*, since the breeding success within the wetland was low due to the competition for resources during

the breeding phase (Fortuna et al. 2006). It should be noted that all the wetlands had the same carrying capacity value, K, within each simulation; therefore, the difference between non-competition for resources in source wetlands and competition for resources in *pseudo-sink* was due to differences in the wetland's indegree.

Management strategies have different impacts on amphibian populations depending on the *indegree* of the wetland. For example, the largest negative impact to the amphibian population was produced when the removed wetland belonged to the medium *indegree* class since a *source* wetland was removed (Fig. 6B). Removing wetlands with high amphibian productivity results in a large loss to the amphibian population of the wetlandscape. Furthermore, the largest positive impact to the amphibian population was produced when the enhance wetland bellowed to the high *indegree* class since the enhanced wetland was originally limited by its carrying capacity (i.e., *pseudo-sink* wetlands) (Fig. 6A). In fact, restoring wetlands with high indegree means that a high number of immigrants can now come to the wetland and reproduce successfully, potentially shifting the wetland from a *pseudo-sink* to a *source*. In contrast, restoring wetlands with low or medium indegree (i.e., *source* or *sinks*) did not lead to an increase in the amphibian population because they were limited by low indegree and not by low carrying capacity.

Wetland removal and restoration strategies produced impacts on amphibian population dynamics that varied with wetland indegree (Fig. 6). The impacts were not "symmetrical" (Fig. 7). A loss in the amphibian population caused by wetland removal could not be compensated equally by wetland restoration. In fact, wetland removal had "cascading" consequences to the remaining wetlands because of: (1) reducing the number

or density of individuals dispersing; and (2) increasing the dispersal distance among the remaining wetlands and thus reducing effective indegree in the network (Semlitsch and Bodie 1998). In contrast, wetland restoration had "cascading" consequences to the remaining wetlands because of reducing "density effect" of individuals breeding.

The modelled amphibian population dynamics are supported by empirical studies (Eggert et al. 2015, Arntzen et al. 2017, Lannoo et al. 2017). Amphibians occur on a continuum of *source-pesudo-sink* wetlands and contribute differentially to the population of the wetlandscape (Eggert et al. 2015). Knowledge of the *sink-source-pseudo-sink* behavior can improve wetland management strategies designed for conservation of species-at-risk (Pulliam 1988, Lawton 1993, Eggert et al. 2015). For example, reintroducing a species into presently unoccupied parts of a species' range is more likely to succeed if the reintroduction occurs into a *source* wetland rather than a *sink* wetland (Pulliam 1988). Furthermore, wetland removal could focus on wetlands with low indegree, and wetland restoration could focus on wetlands with high indegree to increase species population. With these insights, the conservation strategies can include design of wetland networks to optimize *sink-source* dynamics.

In the future, our model can be improved by adding a two-step migration system, which would enable us to simulate the migration from breeding habitat to winter habitat at the end of summer and the dispersal from winter habitat to breeding habitat at the beginning of spring. In addition, simulation scenarios where wetlands are added to the network could be analyzed to assess the wetland positions leading to the highest positive impact on the population.

5. Conclusion

It is important to recognize the role of individual wetlands as defined by their connectivity within wetlandscapes. The adoption of a single-object perspective for wetland management is incomplete given the flux of energy, materials, and organisms that occur among wetlands and in light of our results. Wetlands can have different roles (sink, source, pseudo-sinks) in sustaining an amphibian population within wetlandscapes. Wetland indegree is a key property for 1) quantifying wetland connectivity and 2) classifying wetlands into sources, sinks and pseudo-sinks. Wetland removal and restoration are not compensatory. Rather, these management strategies have different impacts on species population dynamics depending on the indegree among wetlands; for example, wetland removal can result in population loss that is not compensated for by wetland restoration. Understanding the role of indegree of wetlands within wetlandscapes, and the differential impacts of wetland management strategies on these connections, is important for sustainable management of wetland dependent species-at-risk.

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Supporting information

Appendix S1

Networks of wetlands were generated with the algorithm by Fachada et al. (2014) using the software MATLAB (The MathWorks 2016). The algorithm generates a network on a two-dimensional plane through the coordinates of nodes; the coordinates are the centroids of wetlands, and a spatial unit corresponds to 1 km. The algorithm's parameters are listed in Table S1, with their values and their ranges, and a brief description of their meaning. Deterministic parameters were fixed for all the networks (i.e. *Slope, Slope standard deviation, Number of clusters, Length average, Length standard deviation, Total points*). Stochastic parameters (i.e., *Cluster separation on X axis, Cluster separation on Y axis* and "*Cluster fatness*") varied for each network and were randomly extracted from uniform distributions according to the range of variation of each parameter. When the coordinates of nodes were generated, the distances between each node were used to compute the *indegree* index.

The parameter values were selected in order to simulate a wetlandscape where amphibians could move according to their dispersal capabilities. We focused on anuran species that were reported to be able to move up to 10 km during their dispersal stage (Kendell 2002). Thus, we created wetlandscapes where distances are generated in order to have different degree of connectivity.

Table S1. Parameters of the network algorithm used in the simulations.

691	Parameter	Value*	Description
692	Slope (/)	1	Growth rate without density effect
693 694 695 696 697	Slope Standard deviation (/)	1	Standard deviation of the slope; used to obtain a random slope variation from the normal distribution, which is added to the base slope in order to obtain the final slope of each cluster.
698	Number of clusters (/)	1	Number of clusters to generate
699 700	Cluster separation on X axis (km	a) 2.5 – 20	Average separation of line centers along the X axis.
701 702	Cluster separation on Y axis (km	2.5-20	Average separation of line centers along the Y axis.
703 704	Length Average (km)	1	The base length of lines on which clusters are based.
705 706 707 708 709 710	Length standard deviation (km)	1	Standard deviation of line length; used to obtain a random length variation from the normal distribution, which is added to the base length in order to obtain the final length of each line.
711 712 713 714 715	"Cluster fatness" (km)	0.5 – 6	The standard deviation of the distance from each point to the respective line, in both x and y directions; this distance is obtained from the normal distribution.
716 717 718	Total points	50	Total points in generated data (will be randomly divided among clusters).

^{*} Where a range is given, the parameter is randomly selected from a uniform distribution within the range (bounds included).