



HAL
open science

Integrating opportunistic and structured non-invasive surveys with spatial capture-recapture models to map connectivity of the Pyrenean brown bear population

Maëlis Kervellec, Cyril Milleret, Cécile Vanpé, Pierre-Yves Quenette, Jérôme Sentilles, Santiago Palazón, Ivan Afonso Jordana, Ramón Jato, Miguel Mari Elósegui Irurtia, Olivier Gimenez

► **To cite this version:**

Maëlis Kervellec, Cyril Milleret, Cécile Vanpé, Pierre-Yves Quenette, Jérôme Sentilles, et al.. Integrating opportunistic and structured non-invasive surveys with spatial capture-recapture models to map connectivity of the Pyrenean brown bear population. *Biological Conservation*, 2023, 278, pp.109875. 10.1016/j.biocon.2022.109875 . hal-04234608

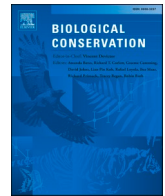
HAL Id: hal-04234608

<https://hal.science/hal-04234608>

Submitted on 22 Feb 2024

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Integrating opportunistic and structured non-invasive surveys with spatial capture-recapture models to map connectivity of the Pyrenean brown bear population

Maëlis Kervellec^{a,*}, Cyril Milleret^b, Cécile Vanpé^c, Pierre-Yves Quenette^c, Jérôme Sentilles^c, Santiago Palazón^d, Ivan Afonso Jordana^e, Ramón Jato^f, Miguel Mari Elósegui Iurrtia^g, Olivier Gimenez^a

^a CEFE, University of Montpellier, CNRS, EPHE, IRD, Montpellier, France

^b Faculty of Environmental Sciences and Natural Resource Management, Norwegian University of Life Sciences, NO-1432 Ås, Norway

^c Équipe Ours, Office Français de la Biodiversité, Impasse de la Chapelle, 31800 Villeneuve-de Rivière, France

^d Fauna and Flora Service, Department of Territory and Sustainability, Dr. Roux, 80, 08017 Barcelona, Spain

^e Departament de Territori, Paisatge e Gestió Ambiental, Conselh Generau d'Aran, Plaça d'Aran 1-2, 25530 Vielha, Spain

^f Departamento de Producción y Asistencias Técnicas, Sociedad Aragonesa de Gestión Agroambiental, S. L. U., Ctra HUV-Fornillos, 22195 Fornillos de Apiés, Huesca, Spain

^g Servicio de Biodiversidad, Sección de Espacios Naturales y Especies Amenazadas, Gobierno de Navarra-Nafarroako Gobernua, Pamplona, Iruñea, Spain

ARTICLE INFO

Keywords:

Fragmentation
Functional connectivity
Large carnivore
Non-invasive genetic sampling
Road network
Ursus arctos

ABSTRACT

Connectivity, in the sense of the persistence of movements between habitat patches, is key to maintain endangered populations and has to be evaluated in management plans. In practice, connectivity is difficult to quantify especially for rare and elusive species. Here, we use spatial capture-recapture (SCR) models with an ecological detection distance to identify barriers to movement. We focused on the transnational critically endangered Pyrenean brown bear (*Ursus arctos*) population, which is distributed over Spain, France and Andorra and is divided into two main cores areas following translocations. We integrate structured monitoring from camera traps and hair snags with opportunistic data gathered after depredation events. While structured monitoring focuses on areas of regular bear presence, the integration of opportunistic data allows us to obtain information in a wider range of habitat, which is especially important for ecological inference. By estimating a resistance parameter from encounter data, we show that the road network impedes movements, leading to smaller home ranges with increasing road density. Although the quantitative effect of roads is context-dependent (i.e. varying according to landscape configuration), our model predicts that a brown bear with a home range located in an area with relatively high road density (8.29 km/km²) has a home range size reduced by 1.4-fold for males and 1.6-fold for females compared to a brown bear with a home range located in an area with low road density (1.38 km/km²). When assessing connectivity, spatial capture-recapture modeling offers an alternative to the use of experts' opinion when telemetry data are not available.

1. Introduction

Habitat loss and fragmentation are a major concern for the conservation of animal populations (Fardila et al., 2017). Landscape structure can constraint the movement of individuals, their ability to find resources, to disperse, and to establish a home range (Fahrig, 2003). More generally, the presence of barriers can isolate populations and reduce their size, which can lead to the reduction of genetic diversity and

ultimately affect population viability (Jackson and Fahrig, 2011). Landscape connectivity – “the degree to which the landscape facilitates or impedes movement among resource patches” (Taylor et al., 1993) – is increasingly included in conservation plans (Keeley et al., 2021).

Carnivores are usually the first species affected by the loss of connectivity because they live at low density and over large areas (Correa Ayram et al., 2016; Zeller et al., 2012). They are also flagship species, which increases stakeholder involvement in corridor projects (Beier

* Corresponding author at: CEFE 1919 Route de Mende, 34090 Montpellier, France.

E-mail address: maelis.kervellec@cefe.cnrs.fr (M. Kervellec).

<https://doi.org/10.1016/j.biocon.2022.109875>

Received 13 May 2022; Received in revised form 13 December 2022; Accepted 17 December 2022

Available online 10 January 2023

0006-3207/© 2023 The Authors. Published by Elsevier Ltd. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

et al., 2008). In particular, the transport infrastructure (e.g. road networks) has been a primary focus, because it causes direct mortality (i.e. roadkills), behavioral modifications (attraction or avoidance), or can act as physical barriers to movement (Coffin, 2007; Forman and Alexander, 1998).

A common approach to measure landscape connectivity is to build resistance maps. Resistance surfaces quantify the degree of potential flow through each cell given land cover types and using expert opinion or empirical data (Fletcher and Fortin, 2018). In practice, there is a trade-off between the information needed and the data available. Expert opinion is more subjective and less predictive of connectivity than biological data but it is often the only information available (Zeller et al., 2012). Telemetry and GPS data are the most informative data to study connectivity, because the process studied is movement (Zeller et al., 2018). However, these data are expensive to collect, can sometimes only be acquired at a coarse temporal resolution and often constitute a small proportion of the population monitored (Zeller et al., 2012, 2018).

Capture-recapture methods and, increasingly spatial capture-recapture (SCR) models, are the standard framework to monitor elusive populations from individual encounter data, like large carnivores (Royle et al., 2018). SCR models integrate a latent ecological process modeling the distribution of individuals and their activity centers, and a detection process accounting for heterogeneity in detectability by explicitly considering the distance between the traps and individual activity centers (Efford, 2004). The detection process of standard SCR models assumes circular home ranges, and therefore that the movement of individuals is homogeneous around their activity centers and not affected by landscape structure. When these assumptions are unlikely to be met, e.g. when habitat features restrict within home range movement, more complex SCR models can accommodate an ecological distance detection model. These models allow quantifying the impact of the landscape characteristics and estimating a resistance surface, as well as improving the estimation of space use patterns (Royle et al., 2013; Sutherland et al., 2015). Combined with the estimation of density, such SCR models allow an empirical quantification of connectivity based on spatial encounter histories from non-invasive detection data (Morin et al., 2017; Zeller et al., 2012). They model the distribution of activity centers in space (i.e. second-order selection defined by Johnson, 1980) and estimate how landscape structure affects within home range movement and space use (i.e. third-order selection (Johnson, 1980)). Connectivity is measured at the scale of the home range in the sense of the “area traversed by the individual in its normal activities of food gathering, mating and caring for young” (Burt, 1943). Since SCR models can make use of monitoring data collected at the population level, and can account for imperfect detection, they can make population-level inferences (Royle et al., 2018).

In this study, we focused on the critically endangered Pyrenean brown bear (*Ursus arctos*) population, which is distributed over Spain, France and Andorra. Due to human persecution, the population was almost extinct in 1995 with only five individuals remaining in the western Pyrenees (Aspe and Ossau valleys) (Piédallu et al., 2019). Since 1996, eleven bears have been translocated from Slovenia in the western and the central Pyrenees to reinforce the population and avoid its extinction. The Pyrenean brown bear population is currently recovering with a minimum of 70 individuals detected in 2021 (Vanpé et al., 2022). The population is divided in two main core areas isolated with regard to exchange of females and located in areas with moderate human disturbance (Martin et al., 2012; Parres et al., 2020). Our objectives are to identify barriers that limits home range movement and to evaluate how road fragmentation affects brown bear space use. The road network is assumed to impede brown bear movements, like it has been shown for other European populations (e.g. Slovakia: Skuban et al., 2017, Cantabrian: Mateo-Sánchez et al., 2014). However, the degree of avoidance to roads and the population-level consequences have never been quantified. We used transnational non-invasive genetic sampling data across France and Spain to draw population-level inferences about

connectivity. To do so, we accounted for imperfect and heterogeneous detections by building a SCR model with non-invasive genetic sampling data obtained from the structured and opportunistic monitoring of the population.

2. Material and methods

2.1. Study area

The transnational study area (36,600km²) is located at the border between Southwestern France (6 counties), Northeastern Spain (3 autonomous regions) and Andorra, and encompasses the entire range of the Pyrenean brown bear population (Fig. 1). Elevation ranges from 0 to 3400 m and is characterized by large valleys and steep mountains. The study area is composed at 41 % of forests with deciduous (dominant beech *Fagus sylvatica*) and coniferous trees (dominant silver fir *Abies alba*) (Martin et al., 2012). Between forest patches there are large open areas with shrubs such as rhododendron (*Rhododendron ferrugineum*) and wild blueberry (*Vaccinium myrtillus*) above 1800 m. The landscape is also shaped by human activities. Traditional pastoral activity, mostly sheep, occupies pastures at higher altitudes from June to October. Other human activities are forestry and recreational activities (e.g., hiking, hunting, skiing). Human population (mean = 67 inhabitants/km²), and the road network (mean = 3.8 km/km²) are mostly concentrated in the valleys. The Pyrenean Mountains are framed by highways, and few primary roads cross perpendicularly the massif to link France and Spain (Fig. A2).

2.2. Data collection

2.2.1. Structured monitoring

Three surveys were conducted in 2017, 2018 and 2019 from May to November. We assumed that the population was demographically closed since births take place in winter and the Pyrenean population is isolated from other populations. We used a structured sampling design, and we set up two types of traps, DNA hair snags alone or combined with camera traps. These traps were distributed in the areas known to be regularly occupied by bears in France and in Catalonia. Camera traps were essentially set up to detect reproductions, while hair snags allowed the genetic identification of individuals. Some individuals with distinctive marks (GPS collar, ear mark, distinctive spot) could also be identified on camera traps. Hair snags were baited with beech tar. We visited the stations at least once a month, and twice for some traps in France, in May, June and September. More details about the monitoring protocols can be found in Vanpé et al. (2022).

2.2.2. Opportunistic monitoring

We also used opportunistic detection data consisting in all the genetic samples collected following a depredation event on livestock or beehives. We did not consider other opportunistic signs of presence, because we were not able to define a corresponding search effort. In order to match the study period defined for the structured monitoring, we only considered those collected from May to November. The trap array can be assimilated to the distribution of livestock in the Pyrenees. It is recommended that aggregated traps be spaced 1.5 σ apart (σ is the scale parameter estimated in spatial capture-recapture model), or approximately 5 km for bears in the Pyrenees (Milleret et al., 2018). To match opportunistic detection data to a flock, we defined a 5 × 5 km grid and consider a cell to be active if at least one depredation event occurred between 2010 and 2020. Then, every sample collected was attributed to the center of the cell, making all centers of active cells opportunistic traps.

2.2.3. Brown bears individual identification

The individual identification of bears by genetic analysis of the samples (hair) consisted in the amplification and identification of

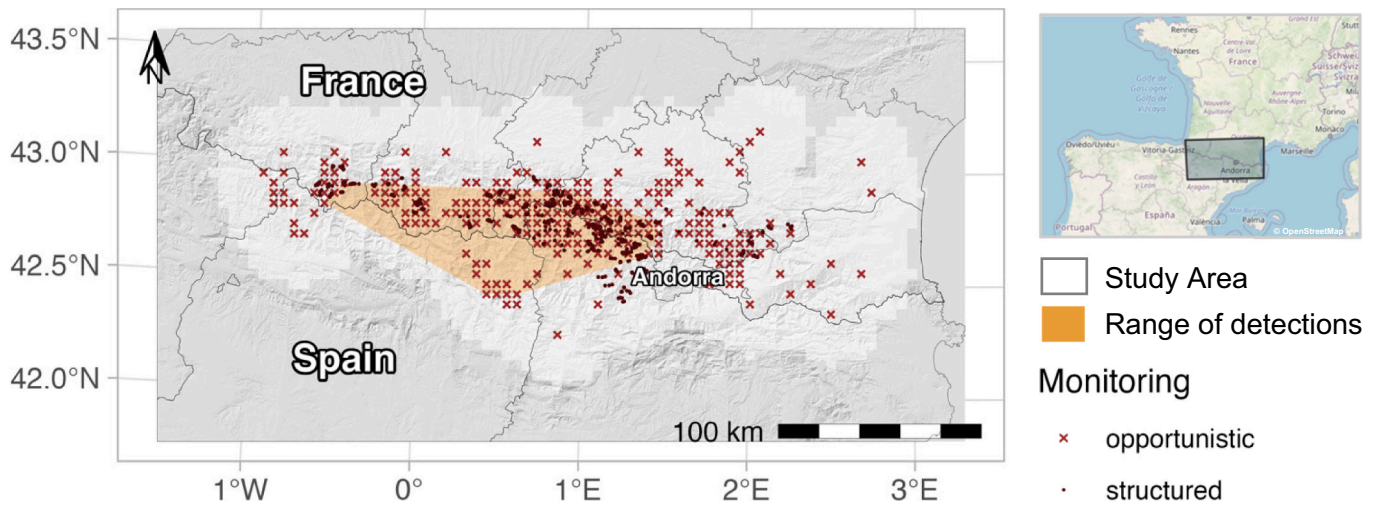


Fig. 1. Study area, minimum convex polygon to represent range of detections and locations of traps for the monitoring of the brown bear population in the Pyrenees in 2019. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

genetic markers. We obtained species and genetic lineage by mitochondrial DNA analysis. We determined individual identity using a multiple tubes Polymerase Chain Reaction (PCR) approach with 13 microsatellite markers (Vanpé et al., 2022). Sex was identified by combining three markers present on the sex chromosomes (Vanpé et al., 2022). When a sample could not be attributed to an individual, it was discarded from the analyses. In total, 156 samples were genetically identified in 2017, 110 in 2018, and 200 in 2019, which corresponded to 31, 31, and 33 different individuals, respectively (Table A1).

2.3. Statistical modeling: Spatial capture-recapture model

We estimated bear density and landscape connectivity simultaneously with spatial capture-recapture (SCR) models. These models combine 1) a detection process to account for the imperfect detection of individuals from individual encounter history and 2) a spatial point process to model the distribution of individuals in space through the estimation of their activity center’s location (Efford, 2004; Royle et al., 2013). Here, we used a multi-session sex-structured SCR model (Sutherland et al., 2019). This model provides a framework to account for variation in densities between males and females and between different monitoring sessions. First, we fitted the SCR model to the structured survey data and accounted for heterogeneity in detection. Second, we integrated the opportunistic data, and evaluated the added value of data integration to improve inference about spatial heterogeneity in density, detection and resistance. Finally, we quantified the impact of roads on the shape of brown bear home range, using non-Euclidean distance SCR models (Sutherland et al., 2015). Parameters were estimated by maximum likelihood implemented in R using the package oSCR (Sutherland et al., 2019).

2.3.1. Baseline detection probability

SCR models are based on encounter history data y_{ijk} of an individual i at trap j during capture occasion k :

$$y_{ijk} \sim \text{Bernoulli}(p_{ijk}) \tag{1}$$

where the detection probability, p_{ijk} describes the probability to detect an individual i at trap j during sampling occasion k . We assumed that this probability decreases with the distance between the activity center and the trap. We modeled this probability with a half-normal detection function:

$$p_{ijk} = p_{0ijk} \times \exp\left(-\frac{d_{euc}(x_j, s_i)^2}{2\sigma^2}\right) \tag{2}$$

where $d_{euc}(x_j, s_i)$ is the Euclidean distance between trap x_j and activity center s_i . The baseline detection probability p_{0ijk} is the detection probability when an individual’s activity center is exactly on the trap (i.e. $d_{euc}(x_j, s_i) = 0$).

We considered six variables to explain variation in p_{0ijk} (Table 1):

- (1) The behavioral response (**b**) is a binary individual covariate that differentiates the first detection of an individual at a given trap ($b = 0$) from the following ones. This persistent response accounts for trap happiness, because hair snags are baited and individuals detected once seems more likely to be detected afterwards.
- (2) The sampling occasion (**m**) is defined as a month from May to November ($m = 1, \dots, 7$). To acknowledge that bears are more likely to be detected in the summer, we considered a quadratic effect of the sampling occasion ($m + m^2$).
- (3) The search effort (**effort**) is a three-level factor. It is defined according to the country where the trap is placed (France or Spain) and the number of visits performed per month to collect genetic material. Our hypothesis is that the greater the number of visits at a station, the greater are the chance to obtain DNA material on

Table 1
Description of the SCR model parameters and variables considered.

Parameter	Notation	Meaning
p_0	b	Behavioral response to traps
	$m + m^2$	Quadratic effect of the sampling occasion
	Session	Year (2017, 2018, 2019)
	Sex	Sex (female, male)
Effort		Search effort: country and number of visits in the station per month (1 visit in France, 2 visits in France, 1 visit in Spain)
	Trap	Type of traps: structured (hair snag alone or hair snag combined with camera trap), or opportunistic (center of 5 km grid surrounding depredeations since 2010)
Sigma	Sex	Sex
	Elevation	Mean elevation over 200 km ²
	Ruggedness	Mean terrain ruggedness index over 200 km ²
	Human density	Log mean human density over 200 km ²
Movement	Forest cover	Mean forest cover over 200 km ²
	Road density	Length of all types of roads (in km) by cells of 6.25 km ²

the trap of sufficient quality to allow identification of the individual (De Barba et al., 2010). The search effort at opportunistic traps was assumed to be equivalent to traps in the structured monitoring that were visited once per month, and depended of the country where the depredation occurs.

- (4) The type of trap (**trap**) describes whether the trap consists of a hair snag alone, a hair snag combined with a camera trap, or an opportunistic trap. Hair snag combined with a camera trap is supposed to have higher detection probability, because hair founded in front of a camera trap are identified in priority.
- (5) The session (**session**), here define by the year (2017, 2018, 2019), accounts for annual variations in detection probability.
- (6) The sex covariate (**sex**) is considered to model behavioral differences (e.g., females with cubs avoid areas where males are present, and males disperse over large distances) (Swenson et al., 2000).

2.3.2. Scale parameter

The scale parameter σ controls the shape of the detection function. The larger σ is, the slower the detection probability decreases with distance from the activity center. We assumed that σ varied with sex because males are known to have larger home range than females (Swenson et al., 2000). In preliminary analyses, we noticed that two males, Néré and Goiat, were detected over half of the study area. As we could not assume that they had the same home range size as the other males and to avoid bias in σ estimate, we dropped these individuals from our analyses. From the estimation of σ in the case of a half-normal detection function, it was possible to estimate the size of the home range according to the relation: $\pi(\sigma\sqrt{5.99})^2$ (Royle et al., 2014).

2.3.3. Spatial variation in density

Activity centers s_i can be uniformly and independently distributed in the spatial domain S (set of potential positions of the activity centers of detectable individuals) with constant intensity, or the intensity can vary according to landscape variables. This spatial domain is defined from the traps around which we add a buffer zone (i.e. a zone large enough to encompass all activity centers of individuals which can be detected by the trap array) (Sutherland et al., 2019). We defined a buffer of 25 km around the stations (approx. 3 and 4σ) and used a 5 km resolution of the spatial domain.

We selected four habitat variables (Fig. A1) to model spatial variation in density of bears in the Pyrenees:

- (1) Forest cover is obtained from the Corine Land Cover (European Environment Agency, 2018) forest data (deciduous, coniferous, and mixed forests) for France and Spain at 100 m and the Andorra habitat map of 2012 (Institut d'Estudis Andorrans, 2012) at 250 m resolution.
- (2) Elevation is defined at 90 m resolution (Shuttle Radar Topographic Mission, 2018). High elevation generally correlates with areas of low human presence and low food resource availability.
- (3) Ruggedness is the average of the absolute values of elevation differences between the focal cell and the eight surrounding cells.
- (4) Human density (Columbia University, 2018) is intended to inform the model about human activities that bears seek to avoid (Swenson et al., 2000).

To ensure that each variable described the habitat at the scale of the bear home range and not just at a given point, we averaged each habitat variable with a sliding window of 200km². The size of the window corresponds to the average home range size of an adult female bear estimated in preliminary analyses. We scaled the three first variables and log-transformed and scaled human density to obtain four rasters of habitat at 5 × 5 km² resolution.

2.4. Estimation of road resistance on connectivity from detection data

The shape of individuals' home ranges depends on the distribution of resources in the landscape as well as constraints on their movements, like the presence of roads (Dahle and Swenson, 2003; Proctor et al., 2019). Home ranges can be irregular, asymmetric, and non-stationary (i.e. varying with location) (Royle et al., 2013). We accounted for the impact of landscape structure on movement using the ecological distance. This distance, involved in the calculation of p_{ijk} (Eq. 2), is based on the least cost path distance instead of the Euclidean distance. Given this discrete landscape ν , the distance between two points can be represented by a sequence of m steps connecting cells denoted $\nu_1, \nu_2, \dots, \nu_p$. We computed the cost of joining two points through this path, and through all possible paths $\mathcal{L}_1, \dots, \mathcal{L}_w$. The least cost path is defined as the minimum of these paths:

$$d_{lcp}(\nu, \nu') = \min_{\mathcal{L}_1, \dots, \mathcal{L}_w} \sum_{g=1}^{m-1} cost(\nu_g, \nu_{g+1}) \times d_{euc}(\nu_g, \nu_{g+1}) \quad (3)$$

$$\text{where } cost(\nu_g, \nu_{g+1}) = \frac{\exp(\alpha_2 z(\nu_g)) + \exp(\alpha_2 z(\nu_{g+1}))}{2}$$

The cost of using a cell is estimated by the resistance parameter α_2 , which is estimated with the maximum likelihood method and by using the value taken by the variable of interest $z(\nu_p)$ in the cell ν_p . The estimated resistance parameter reflects the extent to which a given habitat variable facilitates ($\alpha_2 < 0$) or impedes ($\alpha_2 > 0$) the movement of individuals across the landscape (Royle et al., 2013; Sutherland et al., 2015). When $\alpha_2 = 0$ (i.e. the resistance is null) we have $cost(0) = 1$ and the distance is exactly Euclidean.

The probability that an individual with its activity center s_i uses a cell s_u in S , noted $Pr(g[s_u, s_i])$, can be calculated from the estimated parameters and $p_0 = 1$. We distinguished cell use (g) from imperfect detection (y) (Sutherland et al., 2015). Given an activity center location, we computed the 95 % kernel of the utilization distribution to estimate the home range size in a particular landscape context. The potential connectivity, denoted PC, represents the total utilization assuming that one individual activity center is located in each cell of the habitat. In other word, PC illustrates the number of individuals reaching a cell given that individuals are evenly distributed in the landscape. In the oSCR package, the matrix recording the least-cost path between all cells ν , is computed by the Dijkstra algorithm implemented in the *gdistance* package (van Etten, 2012). Realized densities and potential connectivity can be combined to obtain a realized map of connectivity called density weighted connectivity (DWC) for males and females since their home range size are estimated separately (Morin et al., 2017).

To determine the impact of roads on connectivity, we built a landscape variable considering the length of all roads defined in OpenStreetMap from motorway to track at a resolution of 2.5 km (Fig. A2). This covariate is not correlated (Pearson $r < 0.7$) to the four covariates considered as related to density (Table A2).

2.5. Model selection

To avoid having to test too many models we conducted a hierarchical selection. The best model was selected according to the Akaike information criterion (AIC; Burnham and Anderson, 2002).

2.5.1. Model heterogeneity in detection probability

Firstly, we focused on the detection probability and we tested all combinations of behavioral, sampling occasion, session and sex on p_0 only with data from the structured monitoring. We included type of trap and search effort as additive linear effects on p_0 and a sex effect on σ in each model. At the same time, we also tested whether density was constant between 2017 and 2019 or varied between sessions. In this step, we compared 32 models.

2.5.2. Model spatial variation in density with habitat and integration of opportunistic data

Secondly, we selected the variables explaining best spatial variation in density according to the AIC. In case several models had a better AIC than the model with uniform density, we also tested their additive effect if they were not correlated (Pearson $r < 0.7$). We conducted the same model selection procedure on models considering only structured detection data and on models with both structured and opportunistic detection data, resulting in 10 additional models to compare. We hypothesized that the integration of opportunistic data could improve inferences of spatial variation in density (Tenan et al., 2017). In addition to providing more detections, it also improved the spatial coverage of the survey as opportunistic traps area also located in areas with low density or absence of bears, contrary to the structured survey that was located only in areas regularly occupied by bears (Fig. 1).

The model best supported by the data then served as the basis for quantifying connectivity. Finally, we used AIC to evaluate whether the Euclidean model performed better than the non-Euclidean one (Sutherland et al., 2015).

3. Results

3.1. Heterogeneity in detection probability in the structured monitoring data

The eight best models (AIC differences with the best model: $\Delta AIC < 6$) modeling heterogeneity in detection probability included a behavioral response and a quadratic effect of time. Because estimates were close, we used in the next step the model with the lowest AIC (Table A3). In this model, density was constant in the three monitoring sessions. The baseline detection probability, p_0 , was maximum in June and July, close to zero in October and November and was higher if the bear had already

been detected once. The baseline detection probability was also higher if the trap was composed of a camera trap and hair snags and was visited twice per month in France (Fig. A3). The estimated scale parameter (σ) was larger for males, 7.31 km ($CI_{95} = [6.56, 8.13]$), than for females, 4.89 km ($CI_{95} = [4.35, 5.49]$), as we anticipated (Fig. A4). According to the relationship $\pi(\sigma\sqrt{5.99})^2$ and the assumption of circular home range we found that the estimated 95 % home range size was 252.1km² ($CI_{95} = [179.0, 355.1]$) for females, and 1433.3km² ($CI_{95} = [1056.5, 1944.5]$) for males.

3.2. Spatial variation in density

When considering only detection data from the structured monitoring, none of the models that accommodated spatial covariates on density performed better than the null model (Table A4. A). In contrast, after integrating the opportunistic detection data, the model with an additive effect of ruggedness and human density was considered as the best model (Table A4. B). Bear density was negatively correlated with human density (Fig. 2D), and positively correlated with ruggedness (Fig. 2C). When there was on average one inhabitant/km² over the 200km² surrounding, brown bears density was estimated maximum at 0.015 ($CI_{95} = [0.007, 0.029]$) and 0.009 ($CI_{95} = [0.005, 0.018]$) respectively for females and males (Fig. A5).

3.3. The influence of roads on connectivity

The use of non-Euclidean distance SCR model, with the density of roads as a spatial covariate, strongly improved the AIC of the model ($AIC = 3995.5$). The estimated resistance parameter $\alpha_2 = 0.428$ ($CI_{95} = [0.12, 0.74]$) was positive (Table A5), meaning that the density of roads impeded movement of brown bear. We selected two representative points of high and low road density within the brown bear range. In an

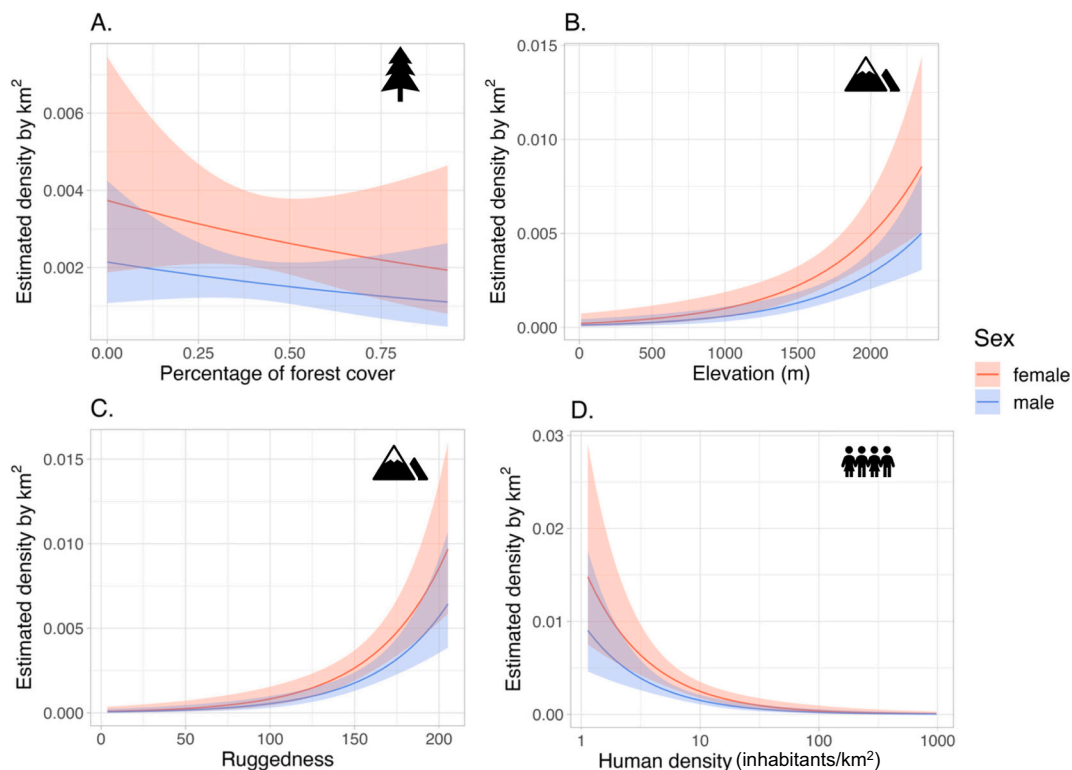


Fig. 2. Estimated density of male (blue line) and female (red line) Pyrenean brown bears per km² as a function of four habitat covariates: (A) percentage of forest cover, (B) elevation (m), (C) ruggedness and (D) human density (mean number of inhabitants per km² over 200 km²). The curves represent the estimated values and the shaded zone the confidence interval at 95 %. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

area with high density of roads (i.e. 8.29 km/km², near Vielha in Val d'Aran, Spain), the model predicted 95 % home range size to be 736.5km² for males and 321.1km² for females. Conversely, in an area with low density of roads (i.e. 1.38 km/km², near Coufflens in Ariège, France), the model predicted 95 % home range size to be 1052.7km² for males and 549.9km² for females (Details about the variation of home range size according to road density are available in the supplementary material (Fig. A6 & A7)).

4. Discussion

Using a non-Euclidean SCR model (Morin et al., 2017; Sutherland et al., 2015) and multiple sources of non-invasive sampling data, our study provided evidence that roads acted as a barrier to the movement of brown bears in the critically endangered Pyrenean population. The home range size of brown bears is smaller with increasing density of roads, which can impede the connectivity and ultimately limit access to resources patches. In addition, we showed the importance of collecting and integrating opportunistic sampling data in SCR models to reveal ecological patterns, such as spatial variation in density and connectivity.

4.1. Spatial variation in density

Bear density was not uniformly distributed over the study area and decreased with human density and increased with ruggedness. Rugged terrain can be considered as refuge areas as they are generally remote and characterized by terrain with high elevation. These results are consistent with previous studies of this population (Martin et al., 2012; Piédallu et al., 2019), and also on other bear populations (e.g. Cantabrian: Mateo-Sánchez et al., 2014; Andean: Morrell et al., 2021). These selected variables are habitat descriptors that correlate relatively well with the characteristics of the two cores areas where the bears from Slovenia were translocated and established their home range, but are not necessarily synonymous of high-quality habitat (Parres et al., 2020). These results cannot therefore be used to infer the future distribution of the population. This also means that other uncolonized habitat which is not characterized by high ruggedness and low human density may support the presence of bears.

4.2. Integration of opportunistic monitoring

In the Pyrenees, the structured monitoring is restricted to the area of regular bear presence, while the opportunistic sampling can, in theory, occur anywhere where domestic animals are located and where bear depredation can occur. The integration of opportunistic detection data using a detector grid located almost continuously through the population range (Fig. 1) allowed us to have traps with no detection of any individual in areas with low or sporadic bear presence. Obtaining information in such areas was very important to detect spatial variation in density and likely the reason why we could only detect association between density and habitat variables after integrating opportunistic samples (Sun et al., 2019). By integrating multiple data sources into SCR models, we provided density maps of the Pyrenean brown bear population across its entire distribution range, which was not possible by considering the structured monitoring only.

Brown bear is an elusive species living at low density over large areas (Swenson et al., 2000). SCR models account for imperfect and heterogeneous detection probability, which varies across time, space and individual characteristics (Efford, 2004; Sutherland et al., 2019). The baseline detection probability varied during the year according to the ecological characteristics of the species, with higher detectability in June during the mating season, and in the summer when their frequency of depredation events usually picks, before it decreased to be close to zero in October and November during hyperphagia when bears can stay on the same feeding area for several days before hibernation (Swenson et al., 2000). The baseline detection probability was also higher when a

bear was at least detected once. This behavioral effect accounted for the trap happiness, because traps are baited, and also because some individuals seem to have a higher detection rate than others. Considering heterogeneity in detectability was important in order to obtain unbiased abundance estimates.

Usually, the monitoring of large carnivores is composed of different types of survey to maximize the number of individuals detected (e.g. Sollmann et al., 2013). Here we considered two types of surveys: the structured monitoring (i.e. hair snags and camera traps) and the opportunistic monitoring (i.e. biological samples gathered after a depredation). Camera traps alone do not permit the identification of bears. They inform on the presence of cubs, the date to which a bear left hair on the traps, and whether several bears use the same hair snag between two visits. Usually, opportunistic detection data are not considered even though they allow the capture of individuals missed or the spatial recapture of bears already captured by structured monitoring, and can improve the spatiotemporal extent of inferences (Sun et al., 2019).

The main obstacle to the use of opportunistic detection data is the difficulty to obtain robust data on the spatial variation in search effort because the detection process is opportunistic and therefore not recorded (Zipkin et al., 2021). Consequently, we had to consider a spatially homogeneous opportunistic search effort. In the spirit of Tenan et al. (2017), we checked for data consistency by comparing parameter estimates in the model integrating or not opportunistic data (Figs. A3 & A4). In our case, the baseline detection probability at structured monitoring traps (hair snags alone and hair snags combined with a camera trap), as well as the scale parameter, did not change after the integration of opportunistic detection data (Fig. A3 & A4). In the integrated model, the baseline detection probability increased with the number of visits (i.e., the search effort). When visits are too sparse, the DNA may be degraded, in too small quantity, or the sample may be composed of DNA originated from several bears (De Barba et al., 2010), leading to genotyping issues. Moreover, some genetic material could not be analyzed for financial reasons. Failure to identify the bear associated with a detection may lead to overestimation of densities and underestimation of p_0 (Royle et al., 2014). Here the population is well known and individuals are recaptured frequently. This process is unlikely to happen because we estimated a population size close to actual population counts (Vanpé et al., 2022).

4.3. The influence of roads on connectivity

We found that road density impeded the movement of brown bear. In an area with low road density (1.38 km/km²), home range is 1.4 times larger for males and 1.6 times larger for females than in an area with high road density (8.29 km/km²). This shows that landscape fragmentation and roads can act as barrier and as a limiting factor for the distribution, and space use of individuals (Coffin, 2007). This situation is similar to the case of the Cantabrian brown bear population which were divided into two cores areas and suffered from low genetic diversity (Pérez et al., 2009). Spanish managers focused on restoring connectivity by creating a corridor between the two nuclei (tree plantations, crossings) and these actions now show encouraging results (i.e. improve demographic and genetic exchanges) (Gonzalez et al., 2016). The area near Andorra, the Mediterranean coast between Narbonne and Gerona and the north-west part of the study area near Pau and Tarbes have a low potential connectivity, suggesting that connectivity might be challenging to achieve in this part of the Pyrenees. Roads from Saint-Gaudens to Vielha limit connectivity in the west of the centro-oriental core, which may limit connectivity between the two cores (Fig. 3). Note that we consider that all roads have the same weight even though forestry roads do not have the same ecological effect as highways, that is roads with a high traffic volume or which are fenced are less crossed (Skuban et al., 2017). During preliminary analyses relying only on French data different characterizations of the road network were explored (distance to main roads (i.e. motorway, trunk and primary

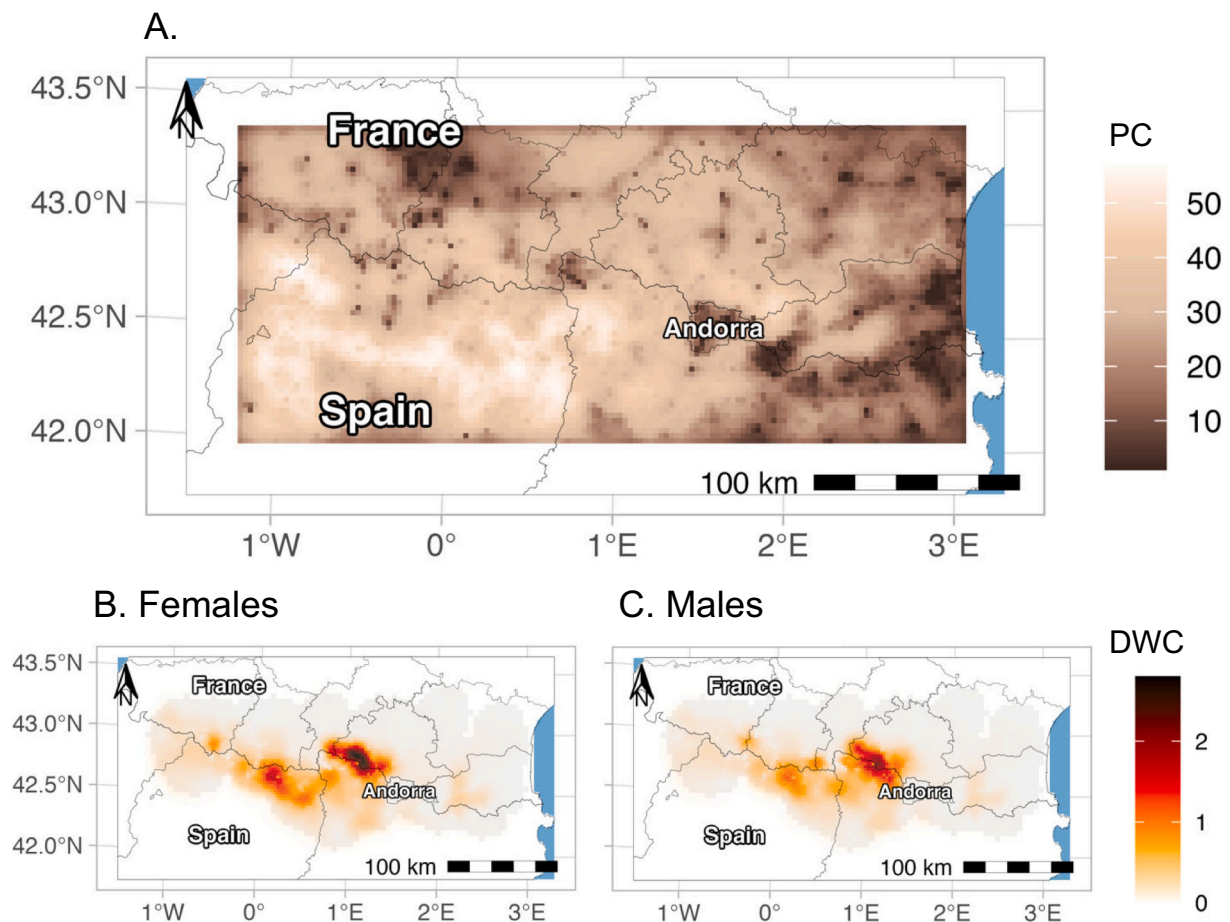


Fig. 3. Maps of metrics of connectivity for the Pyrenean brown bear population in 2019. (A) Potential connectivity (PC) for both sex and density weighted connectivity (DWC) for (B) females only and (C) males only. The grey shaded zone in (B) and (C) represents the study area. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

roads) at 2.5 km, density of paved roads (i.e. motorway, trunk, primary, secondary and tertiary roads) at 1 km, density of all roads at 1 km). The model considering the density of all roads had the lowest AIC and was retained for further analyses (Table A6). However, this choice of implementation gives more importance to urban areas, which have higher road density. Moreover, our model remains correlative and other variables correlated with road density, might be ecologically responsible of the resistance to movement (Royle et al., 2013). In the Pyrenees, roads are concentrated in valleys, where there are villages, train track, semi-canalized rivers, human activities, higher traffic volume and lower food availability (Martin et al., 2012). These factors can also shape brown bear movements (Proctor et al., 2019; Skuban et al., 2017; Swenson et al., 2000).

Density weighted connectivity (DWC) provides a realized measure of connectivity at the population scale (Morin et al., 2017). DWC map showed two cores (Fig. 3). The larger one was in the centro-oriental part of the Pyrenees at the boundary between Ariège and Catalonia and the second is in the western Pyrenees (Piédallu et al., 2019). However, we also estimated a high DWC in the Ordesa y Monte Perdido National Park, because the model predicted the presence of undetected individuals. The model likely predicted the presence of bears since the habitat characteristics in this area was associated with high bear density (very low human density (between 0 and 3 inhabitants/km²) and steep slopes (Fig. 3)), but with a low recorded search effort.

The spatial capture-recapture framework including an ecological distance allows the estimation of a resistance surface based on encounter history data (Sutherland et al., 2015). Compared to occupancy models with unknown individual identity (Howell et al., 2018; Vasudev et al.,

2021), SCR models that use known individual identity, encounter data offer the advantage to mechanistically model the influence of covariates on individual movement. This framework offers an efficient and cost-effective alternative to the use of expert opinion or telemetry data to quantify connectivity (Royle et al., 2018; Zeller et al., 2012). However, if GPS or telemetry data are available, they can be integrated with capture-recapture data to provide more precise estimate of movement and inform the estimation of the scale parameter (Dupont et al., 2021).

Euclidean SCR models assume a half-normal detection function, meaning that the estimated home range is circular, regular and symmetric, which is biologically unrealistic and lead to biased estimates of density (Royle et al., 2013). Also, the consideration of landscape characteristics explaining variation in the size and the shape of home ranges provided us with a more realistic measure of space use. Our estimates of home range size are comparable to what was found for other European populations (e.g. Southern Scandinavia: males [314, 8264] km², and females without cubs [81, 999] km² (Dahle and Swenson, 2003) or in Trento province in Italia [34,1813] km² (Preatoni et al., 2005)). However, we supposed that all males and all females had a similar home range size, which can be a source of bias if some individuals display higher movement abilities. Here, we discarded such individuals from the analysis, even though their contribution to connectivity could be important. Although the ecological reason for this outlier behavior remains unclear, finite mixture models associated with spatial capture-recapture framework could be used to capture this heterogeneity and the contribution of these individuals to connectivity be quantified.

The SCR framework does not explicitly integrate movement like other connectivity models based on GPS and telemetry data, which

precludes from modeling trajectories between captures (Zeller et al., 2018). The SCR framework models the habitat used by an individual during a year around its activity center, therefore integrating concepts (e.g. resistance surface) consistent with the landscape connectivity theory (Royle et al., 2013). Moreover, the density weighted connectivity provides a functional metric of connectivity which is not simply the inverse of the habitat suitability (Keeley et al., 2016; Morin et al., 2017).

4.4. Conservation implication

Spatially explicit estimator of connectivity and density are key to the conservation of recovering populations. However, the biological movement data needed to obtain such estimator are often lacking or difficult to collect in sufficient quantity (Zeller et al., 2012). SCR models with ecological distance provide a framework to integrate structured and opportunistic detection data often collected in monitoring programs on carnivores. SCR models also permit avoiding double counting of individuals that live on both sides of a boundary, as it is often the case in transnational populations (Bischof et al., 2016), and therefore population size overestimation. Overall, SCR models enable ecologists without GPS data to identify problematic areas that limit movement and to estimate a resistance surface from encounter data. These maps based on the probability of space usage and population density provide spatial information for managers to place wildlife crossing for example (Morin et al., 2017; Royle et al., 2013) and corridors, although we acknowledge that the implementation of corridors should consider other species requirements (e.g. dispersal abilities, or migratory behavior) (Beier et al., 2008).

CRediT authorship contribution statement

Maëlis Kervellec: Conceptualization, Methodology, Software, Formal analysis, Data curation, Writing – original draft, Visualization. **Cyril Milleret:** Conceptualization, Methodology, Writing – review & editing. **Cécile Vanpé:** Conceptualization, Investigation, Writing – review & editing. **Pierre-Yves Quenette:** Investigation, Funding acquisition, Writing – review & editing. **Jérôme Sentilles:** Investigation, Data curation, Writing – review & editing. **Santiago Palazón:** Investigation, Funding acquisition, Writing – review & editing. **Ivan Afonso Jordana:** Investigation, Funding acquisition, Writing – review & editing. **Ramón Jato:** Investigation, Funding acquisition, Writing – review & editing. **Miguel Mari Elósegui Iruña:** Investigation, Funding acquisition, Writing – review & editing. **Olivier Gimenez:** Conceptualization, Methodology, Supervision, Project administration, Writing – review & editing.

Funding

The work was partially supported by the French National Research Agency grant ANR-16-CE02-0007. Financial and logistical support for this study was provided by the ONCFS in France. Genetic analyses were performed by the ANTAGENE Company.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Data availability

The authors do not have permission to share data.

Acknowledgements

We gratefully acknowledge all field co-workers from the Brown Bear

Network (ROB) and from Spanish and Andorran teams. We thank the anonymous referee and the associate editor, Varun R. Goswami, for their helpful and constructive comments.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.biocon.2022.109875>.

References

- Beier, P., Majka, D.R., Spencer, W.D., 2008. Forks in the road: choices in procedures for designing wildland linkages: design of wildlife linkages. *Conserv. Biol.* 22, 836–851.
- Bischof, R., Brøseth, H., Gimenez, O., 2016. Wildlife in a politically divided world: insularism inflates estimates of brown bear abundance. *Conserv. Lett.* 9, 122–130.
- Burnham, K.P., Anderson, D.R., 2002. *Model Selection and Multimodel Inference - A Practical Information - Theoretic Approach*, 2nd ed. Springer-Verlag, New York.
- Burt, W.H., 1943. Territoriality and home range concepts as applied to mammals. *J. Mammal.* 24, 346–352.
- Coffin, A.W., 2007. From roadkill to road ecology: a review of the ecological effects of roads. *J. Transp. Geogr.* 15, 396–406.
- Correa Ayram, C.A., Mendoza, M.E., Etter, A., Salicrup, D.R.P., 2016. Habitat connectivity in biodiversity conservation: a review of recent studies and applications. *Prog. Phys. Geogr. Earth Environ.* 40, 7–37.
- Dahle, B., Swenson, J.E., 2003. Home ranges in adult scandinavian brown bears (*Ursus arctos*): effect of mass, sex, reproductive category, population density and habitat type. *J. Zool.* 260, 329–335.
- De Barba, M., Waits, L.P., Genovesi, P., Randi, E., Chirichella, R., Cetto, E., 2010. Comparing opportunistic and systematic sampling methods for non-invasive genetic monitoring of a small translocated brown bear population. *J. Appl. Ecol.* 47, 172–181.
- Dupont, G., Linden, D.W., Sutherland, C., 2021. Improved inferences about landscape connectivity from spatial capture–recapture by integration of a movement model. *Ecology* 1–11.
- Efford, M., 2004. Density estimation in live-trapping studies. *Oikos* 106, 598–610.
- Fahrig, L., 2003. Effects of habitat fragmentation on biodiversity. *Annu. Rev. Ecol. Evol. Syst.* 34, 487–515.
- Fardila, D., Kelly, L.T., Moore, J.L., McCarthy, M.A., 2017. A systematic review reveals changes in where and how we have studied habitat loss and fragmentation over 20 years. *Biol. Conserv.* 212, 130–138.
- Fletcher, R., Fortin, M.-J., 2018. *Spatial Ecology and Conservation Modeling: Applications With R*. Springer.
- Forman, R.T.T., Alexander, L.E., 1998. Roads and their major ecological effects. *Annu. Rev. Ecol. Syst.* 29, 207–231.
- Gonzalez, E.G., Blanco, J.C., Ballesteros, F., Alcaraz, L., Palomero, G., Doadrio, I., 2016. Genetic and demographic recovery of an isolated population of brown bear *Ursus arctos* L., 1758. *PeerJ* 4, e1928.
- Howell, P.E., Muths, E., Hossack, B.R., Sigafus, B.H., Chandler, R.B., 2018. Increasing connectivity between metapopulation ecology and landscape ecology. *Ecology* 99, 1119–1128.
- Jackson, N.D., Fahrig, L., 2011. Relative effects of road mortality and decreased connectivity on population genetic diversity. *Biol. Conserv.* 144, 3143–3148.
- Johnson, D.H., 1980. The comparison of usage and availability measurements for evaluating resource preference. *Ecology* 61, 65–71.
- Keeley, A.T.H., Beier, P., Gagnon, J.W., 2016. Estimating landscape resistance from habitat suitability: effects of data source and nonlinearities. *Landscape Ecol.* 31, 2151–2162.
- Keeley, A.T.H., Beier, P., Jenness, J.S., 2021. Connectivity metrics for conservation planning and monitoring. *Biol. Conserv.* 14.
- Martin, J., Revilla, E., Quenette, P.-Y., Naves, J., Allainé, D., Swenson, J.E., 2012. Brown bear habitat suitability in the Pyrenees: transferability across sites and linking scales to make the most of scarce data: pyrenean brown bear habitat suitability. *J. Appl. Ecol.* 49, 621–631.
- Mateo-Sánchez, M.C., Cushman, S.A., Saura, S., 2014. Connecting endangered brown bear subpopulations in the Cantabrian Range (north-western Spain): enhancing connectivity to allow brown bear persistence. *Anim. Conserv.* 17, 430–440.
- Milleret, C., Dupont, P., Brøseth, H., Kindberg, J., Royle, J.A., Bischof, R., 2018. Using partial aggregation in spatial capture recapture. *Methods Ecol. Evol.* 9, 1896–1907.
- Morin, D.J., Fuller, A.K., Royle, J.A., Sutherland, C., 2017. Model-based estimators of density and connectivity to inform conservation of spatially structured populations. *Ecosphere* 8, 1–16.
- Morrell, N., Appleton, R.A., Arcese, P., 2021. Roads, forest cover, and topography as factors affecting the occurrence of large carnivores: the case of the Andean bear (*Tremarctos ornatus*). *Glob. Ecol. Conserv.* e01473.
- Parres, A., Palazón, S., Afonso, I., Quenette, P.-Y., Batet, A., Camarra, J.-J., Garreta, X., Gonçalves, S., Guillén, J., Mir, S., Jato, R., Rodríguez, J., Sentilles, J., Xicola, L., Melero, Y., 2020. Activity patterns in the reintroduced pyrenean brown bear population. *Mammal Res.* 65, 435–444.
- Pérez, T., Vázquez, F., Naves, J., Fernández, A., Corao, A., Albornoz, J., Domínguez, A., 2009. Non-invasive genetic study of the endangered Cantabrian brown bear (*Ursus arctos*). *Conserv. Genet.* 10, 291–301.

- Piédallu, B., Quenette, P.-Y., Bombillon, N., 2019. Determinants and patterns of habitat use by the brown bear *Ursus arctos* in the French Pyrenees revealed by occupancy modelling. *Oryx* 53, 334–343.
- Preatoni, D., Mustoni, A., Martinoli, A., Carlini, E., Chiarenzi, B., Chiozzini, S., Van Dongen, S., Wauters, L.A., Tosi, G., 2005. Conservation of brown bear in the Alps: space use and settlement behavior of reintroduced bears. *Acta Oecol.* 28, 189–197.
- Proctor, M.F., McLellan, B.N., Stenhouse, G.B., Mowat, G., Lamb, C.T., Boyce, M.S., 2019. Effects of roads and motorized human access on grizzly bear populations in British Columbia and Alberta, Canada. *Ursus* 16–39.
- Royle, J.A., Chandler, R.B., Gazenski, K.D., Graves, T.A., 2013. Spatial capture–recapture models for jointly estimating population density and landscape connectivity. *Ecology* 94, 287–294.
- Royle, J.A., Chandler, R.B., Sollmann, R., Gardner, B., 2014. *Spatial Capture-recapture*. Elsevier, Amsterdam.
- Royle, J.A., Fuller, A.K., Sutherland, C., 2018. Unifying population and landscape ecology with spatial capture-recapture. *Ecography* 41, 444–456.
- Skuban, M., Findo, S., Kajba, M., Koreň, M., Chamers, J., Antal, V., 2017. Effects of roads on brown bear movements and mortality in Slovakia. *Eur. J. Wildl. Res.* 63, 82.
- Sollmann, R., Torres, N.M., Furtado, M.M., de Almeida Jácomo, A.T., Palomares, F., Roques, S., Silveira, L., 2013. Combining camera-trapping and noninvasive genetic data in a spatial capture–recapture framework improves density estimates for the jaguar. *Biol. Conserv.* 167, 242–247.
- Sun, C.C., Royle, J.A., Fuller, A.K., 2019. Incorporating citizen science data in spatially explicit integrated population models. *Ecology* 100.
- Sutherland, C., Fuller, A.K., Royle, J.A., 2015. Modelling non-Euclidean movement and landscape connectivity in highly structured ecological networks. *Methods Ecol. Evol.* 6, 169–177.
- Sutherland, C., Royle, J.A., Linden, D.W., 2019. oSCR: a spatial capture–recapture R package for inference about spatial ecological processes. *Ecography* 42, 1459–1469.
- Swenson, J.E., Gerstl, N., Dahle, B., Zedrosser, A., 2000. Action Plan for the conservation of the Brown Bear (*Ursus arctos*) in Europe (No. 114), Nature and Environment. Council of Europe Publishing.
- Taylor, P.D., Fahrig, L., Henein, K., Merriam, G., 1993. Connectivity is a vital element of landscape structure. *Oikos* 68, 571–573.
- Tenan, S., Pedrini, P., Bragalanti, N., Groff, C., Sutherland, C., 2017. Data integration for inference about spatial processes: a model-based approach to test and account for data inconsistency. *PLOS ONE* 12, e0185588.
- Van Etten, J., 2012. R Package gdistance: Distances and Routes on Geographical Grids.
- Vanpé, C., Piédallu, B., Quenette, P.-Y., Sentilles, J., Queney, G., Palazón, S., Jordana, I. A., Jato, R., Elósegui Irurtia, M.M., Solà de la Torre, J., Gimenez, O., 2022. Estimating abundance of a recovering transboundary brown bear population with capture-recapture models. *Peer Community J.* 2, e71.
- Vasudev, D., Goswami, V.R., Oli, M.K., 2021. Detecting dispersal: a spatial dynamic occupancy model to reliably quantify connectivity across heterogeneous conservation landscapes. *Biol. Conserv.* 253, 108874.
- Zeller, K.A., McGarigal, K., Whiteley, A.R., 2012. Estimating landscape resistance to movement: a review. *Landsc. Ecol.* 27, 777–797.
- Zeller, K.A., Jennings, M.K., Vickers, T.W., Ernest, H.B., Cushman, S.A., Boyce, W.M., 2018. Are all data types and connectivity models created equal? Validating common connectivity approaches with dispersal data. *Divers. Distrib.* 24, 868–879.
- Zipkin, E.F., Zylstra, E.R., Wright, A.D., Saunders, S.P., Finley, A.O., Dietze, M.C., Itter, M.S., Tingley, M.W., 2021. Addressing data integration challenges to link ecological processes across scales. *Front. Ecol. Environ.* 19, 30–38.