Improving abundance estimation by combining capture–recapture and occupancy data: example with a large carnivore

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Summary

1. Abundance is a key quantity for conservation and management strategies but remains challenging to assess in the field. Capture–recapture (CR) methods are often used to estimate abundance while correcting for imperfect detection, but these methods are costly. Occupancy, sometimes considered as a surrogate for abundance, is estimated through the collection of presence/absence data and is less costly while allowing gathering of information at a large spatial scale.

2. Building on the recent pieces of work on the combination of different data sources, we showed how abundance data can be complemented by presence/absence data and can be analysed conjointly to improve abundance estimates. Our approach relies on a hierarchical model that makes explicit the link between the abundance and occupancy state variables while formally accounting for imperfect detection.

3. We used a population of Eurasian lynx in France monitored via camera traps and a collection of presence signs as an illustration of our approach.

4. Synthesis and applications. We combined capture–recapture and occupancy data and demonstrated that we can efficiently improve abundance estimates. Our method can be used by managers when estimates of trends in abundance lack power due to sparse data collected during an intensive survey, by simply integrating data collected during non-systematic survey. Furthermore, combining these two sampling procedures makes full use of all available data and allows the development of conservation and management strategies based on precise abundance estimates. Overall, the combination of different data sources in an integrated statistical framework has great potential, especially for elusive species.

Key-words: abundance, Bayesian approach, camera-trapping, Lynx lynx, population size, presence signs, site-occupancy

Introduction

Management and conservation strategies require information on abundance, a key quantity that is notoriously difficult to assess in the field. Due to the impossibility of carrying out an exhaustive census for free-ranging animal populations, sampling methods only yield a count of some portion of the population. As a consequence, researchers have to estimate in some ways the probability that an individual present and exposed to sampling is actually detected and, therefore, that it appears in a count (Williams, Nichols & Conroy 2002).

Capture–recapture (CR) methods are often used to estimate abundance while correcting for imperfect detection (Otis et al. 1978; Williams, Nichols & Conroy 2002). However, CR methods with physical capture may be challenging when species are cryptic and individuals range over large distances. Non-invasive techniques have recently been developed to promote advancement in ecological research on such species (MacKay et al. 2008). However, gathering CR data requires individual identification that usually involves a great commitment of time and money.
In parallel to the estimation of abundance, there has been increasing interest in alternative state variables such as occupancy, that is the proportion of area occupied by a species. Similar to CR models, site-occupancy models have been developed to infer occupancy while explicitly modelling detection probabilities (MacKenzie et al. 2002). This approach relies on replicate observations of ‘detection/non-detection’ (‘presence/absence’) from repeated surveys of a generic site, therefore allowing the separation of true absences from non-detections. Site-occupancy sampling is attractive to ecologists as it tends to be less costly than CR methods and enables the gathering of information at a relatively large spatial scale. As a consequence, such surveys are increasingly used in many areas of investigation in ecology (e.g. macroecology: Royle, Nichols & Kéry (2005)).

In general, abundance and occupancy are considered as two distinct variables answering different biological questions: abundance is often used to address questions dealing with population dynamics, while occupancy is used in biogeography and macroecology. Although apparently disconnected, occupancy and abundance are two linked state variables. By considering the sites at which a species is detected or not as individuals, the link between site-occupancy and CR models is straightforward. Occupancy data alone may inform estimates of abundance (Royle & Nichols 2003) since when a site is known to be occupied by a species, there is at least one individual of this species on this site (Mackenzie et al. 2005). This has led some authors to see occupancy as truncated abundance data (Mackenzie & Nichols 2004; Sólymos, Lele & Bayne 2012). Royle & Nichols (2003) exploited the relationship between occupancy and abundance to estimate population size from presence/absence data and occupancy techniques. Nevertheless, the relationship between the two variables is not explicit in their approach, in that they assumed the sources of variation in detection were mainly driven by the variation of abundance and used the detection probability to extract information on abundance.

Recognizing the interest in combining capture-recapture and occupancy protocols, Freeman & Besbeas (2012) developed an integrated approach in which they combined count survey and ring-recovery data to estimate abundance. Based on simulations and the analysis of a real case study, they demonstrated that combining different sources of data provided a better precision allowing the detection of change in abundance and other demographic parameters that would not be possible otherwise. Nevertheless, Freeman & Besbeas (2012) assumed that detectability was equal to 1 and did not explicitly consider the observation process.

Here, we developed an approach to combine site-occupancy models and CR models to estimate abundance while formally accounting for species and individual imperfect detection. We implemented our approach in a Bayesian framework and used a case study as an illustration to estimate the local abundance of Eurasian lynx *Lynx lynx* in an area of the Jura Mountains, France. We used data collected through intensive monitoring protocols analysed with CR models and presence signs through volunteers monitoring programme analysed with site-occupancy models. Large carnivores are an ideal case study to demonstrate the improvement in estimating abundance by combining capture-recapture and occupancy data as they have low detection probability and large distribution areas leading to sparse data.

Materials and methods

In this section, we introduce the general procedure for implementing our hierarchical approach beginning with (i) the presentation of abundance and occupancy and the relationship between these two state variables that permits combining CR with presence/absence data then (ii) developing the observation processes leading to the measurement of these state variables using CR modelling for estimating abundance and site-occupancy modelling for estimating occupancy. We adopt a Bayesian approach (McCarthy 2007) to properly accommodate estimation uncertainty at every level of our hierarchical model. The Bayesian approach provides a convenient framework to combine information on important ecological parameters from multiple sources of data (Ellison 2004; Brooks et al. 2008). It uses powerful computer-intensive methods for handling complex models and combines the likelihood with prior probability distributions of the parameters to obtain the posterior distribution of the parameters of interest based on the Bayes’ theorem. We use Markov chain Monte Carlo (MCMC) methods to simulate observations from the posterior distributions.

State variables and the relationship between abundance and occupancy

Our objective is to estimate the abundance state variable *N*. We assume that *N* is a realization of a Poisson random variable with rate λ. In parallel, we assume that we have access to another state variable, namely occupancy, defined as the proportion of occupied spatial units (e.g. patches or sites). We assume that occupancy *Z* is a realization of a Bernoulli process with index ψ.

Both state variables are inferred from monitoring protocols and modelling approaches that specifically deal with detectability less than one (see below) (Fig. 1). The link between abundance and occupancy is made by using the fact that when the probability of a site to be occupied is >0, the abundance at this site is >0 (Fig. 1). More formally, the probability of having at least an individual is:

\[ P(N > 0) = 1 - P(N = 0) = 1 - e^{-\lambda} \]

Eqn 1

which is exactly the probability of having a true presence, that is the probability of occurrence ψ in patch occupancy models. Here, Eqn 1 gives information on the probability that abundance is nonzero if one individual is detected on one site; we do not consider that two presence signs collected nearby necessarily come from two distinct individuals. For occupancy to bring information about abundance, the link between these two state variables needs to be made explicit. This is achieved by expressing the rate
Combining capture–recapture and occupancy data

Fig. 1. Diagram representing the capture–recapture and occupancy models used to estimate, respectively, abundance and occupancy. Both models deal with imperfect detection, that is not all the individuals are detected with camera traps and some sites were falsely considered as unoccupied while collecting presence/absence data. Red crosses represent individuals detected as occupied. The link between abundance and occupancy is made by using the fact that when a site is occupied (orange circles), the abundance at this site is >0 (blue crosses).

From the area of interest, R spatial units are selected to be surveyed for the species using appropriate methods (e.g. collection of presence signs). These sites are surveyed multiple times each season for T years. In each survey, detections are considered as true, meaning that we cannot falsely recognize the species, but the non-detections can arise from both (1) true absence and (2) false absence (i.e. presence but not detected). Within a season, we assume that sites were closed to changes in occupancy. So a unit must be either occupied or not for all visits in this season (MacKenzie et al. 2002). We denote \( Y_{s,t} \) a binary random variable corresponding to the observation of an individual on site \( s \) at time \( t \), which takes value 1 if the individual is observed and 0 otherwise. Given a realization \( z_{s,t} \) of the state variable occupancy, \( Y_{s,t} \) is distributed as:

\[
[Y_{s,t}|z_{s,t}] \sim \text{Bernoulli}(z_{s,t} \times p_{Os})
\]

where \( p_{Os} \) is the probability of detecting the species at a site \( s \).

APPLICATION

Material

For many elusive species such as large carnivores, it is easier to detect evidence of their presence than to actually see the animal. To overcome these issues, monitoring a carnivore population may be performed at various levels of resolution (Gese 2001). Biologists often resort to large-scale sign survey to provide information on species distribution or relative abundance at a lesser cost (mustelids: Ruiz-Olmo, Saavedra & Jiménez (2001); felids: Rolland, Basille & Marboutin (2011); canids: Llaneza et al. (2005)). If the financial support is substantial enough, intensive monitoring (camera trapping or DNA sampling) is conducted to provide absolute abundance estimates or density but is often restricted to small areas [mustelids: Graham (2002); felids: Karanth et al. (2004); canids: Cubaynes et al. (2010)].

In the French Jura Mountains (Fig. 2), this two-scale survey method is used for monitoring a population of Eurasian lynx. The monitoring is organized at two levels: an extensive sign survey carried out by a network of observers (state employees, hunters, naturalists, farmers and mountain guides; Duchamp et al. (2012) who collect evidences of presence in the field (i.e. tracks, scat, hair, sightings, livestock and wildlife killed) and an intensive

OBSERVATION PROCESS FOR ABUNDANCE AND OCCUPANCY

Abundance and occupancy cannot be directly measured in the field. The observation process needs to be made explicit to reliably estimate these state variables. Regarding abundance, we resort to CR closed population model (Otis et al. 1978) to estimate population size. Because \( N \) is the key parameter that makes the link between occupancy and abundance (see above), the challenge was to find a flexible approach that would treat population size as an explicit parameter, in contrast with standard CR models in which it is often calculated as a derived parameter. We use the complete data likelihood approach proposed by Schofield & Barker (2014) to fit a CR model with individual heterogeneity on the detection process \( p_i \) for individual \( i \). The originality of their approach lies in the fact that they consider \( N \) explicitly in the likelihood via the trans-dimensional approach (e.g. King & Brooks 2008). We denote \( X \) the complete \( N \times k \) matrix of capture histories over \( k \) capture occasions where \( x_{ij} = 1 \) when individual \( i \) is captured at occasion \( j \) and 0 otherwise. Given \( N \) and the mean \( p_i \) and variance \( \tau_i \) governing the normal distribution that is used to deal with heterogeneity in the detection process, it can be shown that:

\[
[X,p|N,p_i,\tau_i] = \frac{N!}{\sum_{i=1}^{N!} \prod_{j=1}^{N} \left( p_i \right)^{x_{ij}} \left( 1 - p_i \right)^{1 - x_{ij}} \prod_{j=1}^{N} \text{Normal}(p_i,\tau_i)}
\]

eqn 3

where \( z_h \) is the number of individuals observed with capture history \( h \). The unobserved \( p_i \) are dealt with a trans-dimensional algorithm (Carlin & Chib 1995). We refer to Schofield & Barker (2014) for the technical details of the implementation of eqn 3.

Now we define the occupancy as the probability that a random selected site or sampling unit in an area of interest is occupied by a species (at least one individual of the species). Obviously, we cannot have access to the genuine proportion of occupied sites since we do not see all the individuals, and species imperfect detection needs to be accounted for.

\[
\lambda = -\log(1 - \psi)
\]

eqn 2

\[
\text{Normal}(\mu,\tau)
\]

\[
\text{Bernoulli}(p)
\]
ual attributes (e.g. age and sex classes) and home range features
under interest, that is abundance and occupancy.

We fit three models, starting with the capture–recapture model to
the camera-trapping data, then we model the presence/absence
data with a single-season occupancy model, and finally, we imple-
ment the model to combine the two sources of data, that is CR
data and presence/absence data, by combining the two state vari-
able occurring at the scale of species’ distribution range. Only
the cells within the camera-trapping zone are considered. We
assume that the entire camera-trapping zone was surveyed by the
observers’ network that collected the presence signs, thus prevent-
ing false absence. Duchamp et al. 2012 estimated that 15 km was
the mean distance at which signs were recorded relatively to the
observers’ living place. This information provides good support
for our assumption that the camera-trapping area was covered by
the network of lynx observers. To model the occurrence of lynx
in the camera-trapping zone, we use a 3-year temporal window in the
occupancy model (Stahl & Vandel 2001).

Due to the lack of an appropriate integrated framework, the
data collected during extensive and intensive surveys have so far
been separately analysed. Here, we illustrate how our new
approach can be used to improve abundance estimates by com-
bining these two sources of data.

Models

We then fit a site-occupancy model (MacKenzie et al. 2002)
with site-dependent \( \psi_i \) and site-specific detection probability \( p_{O} \),
as suggested by a previous analysis of the data (Rolland, Basille &
Marboutin 2011). The site-dependent occupancy probability is
defined by the mean occupancy \( \mu_s \) and a site random effect \( \beta_s \),
normally distributed with mean 0 and variance \( \sigma^2_w \). Since we do
not have any information on the occurrence of the species, we
assign a Normal (0, 5) prior for \( \mu_s \) on a log scale and a Uniform
(0, 5) prior for \( \sigma^2_w \). Regarding the detection probability, the logit
of \( p_{O} \) is normally distributed with mean \( \mu_p \) and a variance \( \sigma^2_p \).
We set a Normal (0, 10) prior for the logit of \( \mu_p \) and a Uniform
(0, 10) prior for \( \sigma^2_p \).

Finally, we consider the model that combined the two state
variables of interest, that is abundance and occupancy (see eqn 1).
We use a site-dependent occupancy \( \psi_i \) and the mean occupancy \( \mu_s \)
in the function of the index \( l \) for the Poisson dis-
bution of \( N \). Regarding the observation process, we consider a
site-specific probability of detecting occupancy \( p_{O} \), for the site-
occupancy likelihood and a detection probability varying among
individuals \( p_{i} \) in the capture–recapture likelihood. We keep simi-
lar priors that were used in the previous models.

STATISTICAL INFERENCE IN A BAYESIAN FRAMEWORK

We fitted the three models to the data by Bayesian statistical
inference using MCMC sampling. Three chains were generated
using 15 000 iterations each with a burn-in phase of length 2500.
Convergence was diagnosed both visually (checking the traces
and mixing of the chains) and numerically using the Gelman–
Rubin statistic (Gelman & Rubin 1992). We implemented these
analyses in JAGS (Plummer 2003) called from R using the rjags
package. The code for the combined model is available in
Appendix S1.

Results

A total of 172 presence signs were collected by the volun-
teers of the Réseau Loup-Lynx from 2009 to 2011 across the
camera-trapping area. Regarding the camera-trapping
protocol, lynx photographs were collected from February
to April 2011 from 33 trap sites. We removed one trap
site because of technical problems. The effective effort
was reduced to 1816 effective trapping nights. We
obtained 39 detections of lynx during the sampling period
and identified nine individuals that were photographed on
14 of the 32 trap sites. Individuals were captured at up to
six different trap sites. When the sole CR model with het-
erogeneous detection probability was fitted to these data,
we found that the mean estimated detection probability $\mu_p$ was 0.11 ± 0.07 and that the estimated abundance was 14.46 ± 7.50 individuals [95% CI: 9.0; 35.0] (Fig. 3). Using the single-season occupancy model, the mean occupancy was estimated at 0.96 ± 0.04 [95% CI: 0.83; 0.99] and the mean detection probability was quite high: 0.71 ± 0.16 [95% CI: 0.30; 0.97]. Using our new approach to combine CR and presence/absence data, abundance was estimated at 9.96 ± 1.18 individuals [95% CI: 9.0; 13.0] (Fig. 3) while detection probability was on average estimated at 0.16 ± 0.06 in the CR model likelihood. The probability of detecting occupancy was estimated on average at 0.71 ± 0.15, while the occurrence probability was estimated at 0.99 ± 0.01. All the parameters estimated in the three models converged towards their stationary distribution.

**Discussion**

The choice of the state variable to monitor is challenging when dealing with rare and elusive species. Abundance data have commonly been assumed to be more informative on a per site basis, but presence/absence data allow for more sites to be visited and allow carrying out inference across a much larger study area. Moreover, estimating animal abundance requires consistent and standardized application of a technique to be able to detect changes or differences with some degree of accuracy (Macdonald, Mace & Rushton 1998). These techniques are often costly and time-consuming, to a point that systematic surveys are in some situations unaffordable.

Indeed, while studying wild populations, practitioners frequently encounter difficulties in collecting good quality data in sufficient quantity to estimate abundance. Most practitioners have to cope with limited financial resources and data sampling jeopardy. In the case study we developed, we raised the issue of scarce data due to the ecology of the species. Large carnivores have secretive and nocturnal behaviour, are distributed over large areas and live at low density, which means that even standardized and consistent protocols do not guarantee precise estimates. In this context, there is clear need to develop optimal methods to analyse data resulting from such population monitoring schemes, in particular in regions where resources for monitoring programmes are limited (Balmford, Green & Jenkins 2003; Roberts, Donald & Green 2007), or for species with low detectability. Additional presence/absence data, coming from citizen science programmes for instance, may profoundly increase our confidence in abundance estimates, enabling to develop more sound management and conservation strategies. Our approach is a first step in that direction.

Indeed, we showed how abundance data can be complemented by presence/absence data and be analysed conjointly to improve abundance estimates. Our approach is similar to the Freeman and Besbeas approach in that we combine different sources of data but differs from it in the way that we extended their model by adding the detection process to account for imperfect detection. By combining camera-trapping data and presence signs through an explicit link between the state variables of abundance and occupancy, we significantly reduced the width of the credible interval on lynx abundance estimates. The reduced effort associated with the collection of presence/absence data means that occupancy surveys have already proved to be very useful in large-scale surveys for various species. They now are shown to be very useful in small-scale surveys to estimate population size. There is an increasing number of participative programmes on birds (Greenwood 2007, Christmas Bird Count, eBird) and amphibians (FrogwatchUSA, Global Amphibian BioBlitz) that could be used to complement individual-level data, collected occasionally, when financial resources are sufficient. Our model is easily applicable to such taxa since the spatial unit to which occupancy is estimated is well defined: closed units, such as nests or ponds, fulfil the closure assumption underlying single-season occupancy models, although application to mammals is more complicated.

Precautions must be taken when using our new approach regarding the assumptions underlying the model and the sampling design. First, we used presence signs that were collected during 3 years prior to the camera-trapping session and so, we assumed that the population of interest was geographically closed during the sampling period. Thus, the presence/absence data collected were assumed to be presence signs of individuals that were part of the population monitored via the camera-trapping protocol. The geographical assumption is valid for the Eurasian lynx, but it would not be the case for a non-territorial species. Secondly, false positives in presence data may lead to abundance overestimation, and care

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**Fig. 3.** Posterior distribution of the abundance of Eurasian lynx in the camera-trapping area in the Jura Mountains while using only capture-recapture data (salmon) or combining the presence/absence data and the capture-recapture data (blue). The overlap between the two distributions is displayed in purple.
must be taken to ensure the quality of the presence signs which are collected, possibly with a validation step (Molinari-Jobin et al. 2011) or using models that explicitly account for identification errors (Miller et al. 2011). In our case study, objective and standardized identification and reliability assessment criteria have been defined for each category of field signs (Vandel & Stahl 1998), which guarantees a negligible false-positive rate. Thirdly, we assumed that abundance \( N \) was a realization of a homogeneous Poisson random variable. The violation of this assumption might lead to underestimating the variance of \( N \). To relax this assumption, we would need to resort to spatially explicit capture-recapture (SECR) surveys that were formulated by Borchers and Efford (2008) using a general non-homogeneous Poisson process to model animal home range locations. However, to combine the SECR model likelihood with the occupancy model likelihood, \( N \) has to appear explicitly as a parameter in the former likelihood, which is not the case since \( N \) is obtained as a by-product. Further work is needed to extend the (Schofield & Barker 2014) approach to SECR surveys.

Presence/absence surveys in studies with small budgets or targeting elusive species remain useful in an attempt to successfully assign IUCN Red List categories of threat (Joseph et al. 2006) and can accurately represent trends in population size. For these reasons, Royle & Nichols (2003) suggested the idea of using presence/absence data to infer abundance in a proper way. Assuming that the most important source of heterogeneity in detection was in animal abundance, they defined the probability of detecting occupancy as a function of \( N \) and used heterogeneity in detection to extract information about abundance in the occupancy surveys. As for our study, we focused on the direct relationship between abundance and occupancy and directly connected abundance and occupancy in the model. It results in more flexibility when combining various sources of information as our approach can accommodate several sources of data. For example, recent studies made use of replicated counts on spatial units that do not rely on individual identification. As an alternative to CR models, these so-called N-mixture models (Royle 2004) are used to estimate population size while adjusting for detectability <1 (Kéry, Royle & Schmid 2005; Royle, Nichols & Kéry 2005). These models are widely used to estimate bird abundance (Royle 2004; Kéry, Royle & Schmid 2005). Various bird surveys have two levels of resolution as it is the case for carnivores: a standardized survey with repeated visits on a sampling area provides count data used to estimate abundance, while some observers distributed over the area all year long collect presence of the bird species of interest. The first kind of data can be analysed in the N-mixture framework, while the presence/absence data collected by can be analysed in a site-occupancy framework. Our approach can also be used with data arising from distance sampling protocols (Buckland et al. 2001) pending some modifications of the abundance component of the model. We assumed the lynx population was closed during the CR survey since the sampling period was out of the breeding (Breitenmoser-Würsten et al. 2007) and the dispersal period (Zimmermann, Breitenmoser-Würsten & Breitenmoser 2005), and the species being territorial, migration in or out of the study area that would have biased the abundance estimates was avoided. We used the same closure assumption for the occupancy survey. We assumed that one site that was occupied once has been occupied during all survey and any non-detection at this site is considered as a 'false negative' (Rota et al. 2009). Moreover, here, we clearly deal with what Efford and Dawson (2012) referred to as 'asymptotic occupancy'. The authors made the distinction between instantaneous occupancy referring to the probability of a cell being occupied at a particular time, whereas the overall occupancy of a given cell refers to the probability that at least one individual will enter the cell during the study period. Since large carnivores are territorial, we can safely assume that if a cell is detected as occupied once, it will remain occupied during all the survey (provided that the size of their home range does not go through major fluctuations). For large carnivores and other organisms, the generalization of our approach to geographically open population system involving local extinction and colonization processes deserves further exploration. Since our case study involved a real data set and the true abundance was unknown, simulations will be used in future to test the robustness and accuracy of our estimates.

Acknowledgements

L. Blanc was supported by Labex Centre Méditerranéen de l’Environnement et de la Biodiversité (CeMEB) and University of Montpellier 2. We especially thank the staff from the French National Game and Wildlife Agency, the Forest National Agency and the Departmental Federation of Hunters of Jura department, who collected the photographs during the camera-trapping session. We also thank Matthew Schofield for his technical support for modelling explicitly capture-recapture data. We finally thank all the volunteers that are members of the Réseau Loup-Lynx that collect every year precious presence signs of lynx all over its distribution area.

References


Combining capture–recapture and occupancy data


Received 12 February 2014; accepted 10 July 2014
Handling Editor: Hamish McCallum

**Supporting Information**

Additional Supporting Information may be found in the online version of this article.

**Appendix S1.** R code for the combined model.