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Evaluating camera trap methods for monitoring population trends in ungulates: insights from simulation

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Abstract

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2	Camera traps have been widely used in the last decade to monitor abundance of
3	unmarked animal populations. Most estimation methods rely either on the number of
4	times animals pass through the detection zones, like random encounter models (REM)
5	or on the number of capture occasions in a time-lapse program when animals were seen
6	on the pictures, like the instantaneous sampling approach (IS). We simulated a setup of
7	either 100 or 25 camera traps randomly distributed on a 2600-ha area (respectively ≈ 4
8	and 1 trap/km ²), along with the movements of a fictional population of 300 roe deer
9	$(\it Capreolus \ capreolus).$ We assessed the ability of these two classes of popular methods
10	to estimate population size and detect a 20% decline over five years. Simulations were
11	informed by field data on habitat, habitat selection and activity patterns of
12	GPS-monitored roe deer. Both IS and REM estimated population size without bias,
13	with a coefficient of variation only equal to about $15\%~(4~{\rm traps/km^2})~{\rm or}~30\%~(1$
14	$\rm trap/km^2).$ Despite a huge sampling effort and simplified assumptions (perfectly known
15	day range, constant sensor sensitivity), both methods failed to detect the strong
16	population decline in $2/3$ to $3/4$ of simulations (4 traps/km ²), and in about $4/5$ of
17	simulations (1 $\mathrm{trap/km^2}$). We tested other sampling strategies to improve this
18	sensitivity, which either led to an unchanged population size estimation precision
19	(stratified sampling) or to biased estimated trends (sampling only in high-quality
20	habitats). Simulating animals with a 10 times larger home-range, like red deer (${\it Cervus}$
21	$elaphus),$ allowed to detect the decline more frequently (60% to 95% with 4 $\rm traps/km^2,$
22	and 1/3 to 2/3 of the simulations with 1 trap/km ²). These results suggest that the key
23	metric for camera trap use is the average number of different traps visited per animal,
24	which in turn depends on trap density, home-range size and space use heterogeneity.
25	We provide a R package allowing the reader to reproduce these simulations, and carry
26	out their own.

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- ¹ Keywords: random encounter models, instantaneous sampling, simulations, animal
- ² movement, habitat selection, sampling design

1 Introduction

Camera traps are used to reach many goals in wildlife studies, including estimating 2 occupancy, studying animal behaviour, and investigating species interactions, in particular 3 predator-prey interactions (see the review by Burton et al. 2015). Estimating the size of a 4 population with capture-recapture methods, taking advantage of the possibility to identify 5 animals based on phenotypic patterns, has also been among the first use of camera traps, in 6 particular concerning elusive carnivores (Karanth 1995). More recently, a large amount of 7 literature has been devoted to the use of camera traps to estimate the size of the 8 population of non-identifiable animals (Rowcliffe et al. 2008, Nakashima et al. 2017, Howe 9 et al. 2017, Moeller et al. 2018). The basic idea of these methods is that the number of 10 animals detected by the traps can be used to infer the number of animals present in the 11 study area. 12

Two main families of methods have been proposed to achieve this aim: those based on 13 animal-trap encounters, which quantify the number of animals passages through the 14 detection zones of traps motion sensors, and those based on animal-trap associations, which 15 count animals at capture occasions within time-lapse programs (Campos-Candela et al. 16 2018). The random encounter model (REM, Rowcliffe et al. 2008) is a prominent example 17 of the first family, while camera trap distance sampling (Howe et al. 2017), instantaneous 18 sampling (IS, Moeller et al. 2018) or the similar "association model" of Campos-Candela 19 et al. (2018) exemplify the second. 20

These estimation methods are often seen as a grail, in that they allow the estimation of population abundance, a key population parameter (Williams et al. 2002) without resorting to the labor-intensive capture-recapture approach. Many practitioners are attracted by such

approaches (Gilbert et al. 2020), and some even want to replace their well-established
monitoring methods with these new, easier to implement ones (pers. obs.). This might be a
problem, as the properties of these new methods are not yet fully understood, and
numerous factors must be considered when using them.

Thus, one major issue when estimating a population size is the imperfect detectability 5 of the animals in the detection zone (Gilbert et al. 2020). Most camera trap studies rely on 6 the use of motion sensors to trigger the traps when an animal enters the detection zone. 7 However, such sensors are less sensitive when the animals are small or far from the trap. 8 The camera traps characteristics (brand, model, etc.) may also have an effect on the 9 detection probability of the animals in the detection zone. Obstacles may further limit its 10 effective surface area. Numerous authors have insisted on the need to account for this 11 imperfect detection in population size estimation (e.g., Howe et al. 2017, Moeller et al. 12 2023). However, most practitioners continue to ignore this issue, leading to potential biases 13 in their estimates (Burton et al. 2015). Nevertheless, population trend estimates may still 14 be unbiased, unless habitat structure changes over time (e.g. vegetation growth or human 15 disturbance, Guillera-Arroita 2017). 16

A related issue is the presence of resting periods, during which animals are inactive and 17 less detectable by motion sensors. Such periods do not necessarily affect estimation 18 accuracy if the camera traps are randomly distributed over the study area, and if the 19 resting areas have the same probability to be monitored as other areas (e.g., no burrows). 20 However, even when these assumptions are satisfied, the decreased detectability of inactive 21 animals by motion sensors in the trap detection zone can lead to density underestimation. 22 Numerous methods have been proposed to account for such activity patterns in population 23 size estimation (e.g. Nakashima et al. 2017), though a common approach is to focus the 24

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monitoring on periods when it is reasonable to suppose that all animals are active (Howe
et al. 2017).

Empirical validation of camera trap-based methods for estimating population size 3 remains limited. Most comparative studies employ reference density values estimated using established methods such as conventional distance sampling (Rovero and Marshall 2009, 5 Caravaggi et al. 2016) or capture-recapture (Twining et al. 2022). However, these 6 comparisons, carried out over a few years, often involve only a limited number of imprecise 7 estimates from both camera-trap estimation methods and "reference" methods, often 8 leading to conclusions that no difference can be detected between the two families of 9 methods, which does not constitute proof of estimate equivalence. On the other hand, 10 simulations offer a valuable alternative, enabling numerous replications of a specific camera 11 trap study setup under fully known and user-controlled demographic and spatial conditions 12 (Santini et al. 2022).13

In this paper, we assess the feasibility of using camera traps to monitor roe deer 14 Capreolus capreolus) abundance through simulation, while examining how camera and 15 habitat characteristics, along with animal behaviour, influence detection probabilities. 16 Specifically, we explore how the differential use of habitats with varying detection 17 properties, combined with the animals' activity rhythms, affects their detectability. To do 18 so, we simulated realistic movements of animals, incorporating habitat selection and 19 activity patterns (resting, foraging and exploring) informed from the GPS-monitoring in a 20 roe deer population. We simulated a decreasing trend in population size across years, and a 21 correlated increase in home-range size (Kjellander et al. 2004). We assessed the estimation 22 precision and the ability to detect the simulated declining trend of two population size 23 estimation methods: one method relying on encounters (random encounter model, Rowcliffe 24

et al. 2008) and one method relying on associations (instantaneous sampling, Moeller et al. 1 2018). We also assessed the effect of placing all camera traps in the most selected habitat 2 type, as well as the effect of stratified sampling, on the precision of the trend estimates. 3 Finally, we explored the possibility to expand these results to other species, so that we also simulated the monitoring of a species with a larger home range, which is more 5 characteristic of larger ungulates like the red deer (*Cervus elaphus*). 6 We provide a companion R package named simCTChize, available at 7 https://github.com/ClementCalenge/simCTChize, containing all the code and data used 8 for these simulations. The supplementary material corresponds to the package PDF 9 vignette, and describes how the user can install this package and easily reproduce the 10 calculations carried out in this paper. This vignette also includes numerous additional 11 details on the simulations that we omit in this paper to keep it concise (specific values of 12 the parameters used for the movement processes or the detection functions of the camera 13 traps, density of trees simulated in the detection zone, etc.). Finally, this package also 14 contains a tutorial allowing the user to design their own simulations if they want to design 15 a monitoring for another forest species. 16

¹⁷ **1** Material and Methods

1.1 Simulating realistic roe deer movements and their detection by camera 19 traps

We first simulated a population of moving animals in the Chizé reserve, France (46.083° N, 0.417° W), during five years. We chose this enclosed 2614 ha forested study area because its roe deer population has been monitored for over 30 years (Pellerin et al. 2017), and

space use by the roe deer was extensively studied (Saïd et al. 2005, Pellerin et al. 2008, 1 2010, Gaudry et al. 2018). This population monitoring provides us with a wealth of data on 2 demographic parameters (survival, reproductive rates), enabling the development of 3 realistic population simulations. The northern part of the study area (1385 ha) is mainly composed of an oak stand (Quercus spp.) characterized by a high quality forage source, 5 whereas the southern part (1228 ha) is mainly composed of a beech stand (Faque sylvatica) 6 with poor resource quality Pettorelli et al. (2001). Consequently, roe deer density is higher 7 in the northern part of the area, a spatial heterogeneity that we incorporated into our 8 subsequent simulations. 9 We simulated animal movements using a model that combined multiple 10 Ornstein-Uhlenbeck with foraging (OUf) processes (Fleming et al. 2014). Each process was 11 characterized by distinct parameters – attraction points, speed and relocation 12 autocorrelations, and patch sizes – to represent various animal movement behaviours, such 13 as foraging and inter-patch movements, analogous to the mixed models proposed by 14

Blackwell (1997) and Michelot et al. (2019). The model also accounted for the presence of
resting periods.

We estimated 25 utilization distributions for March using GPS data from 15 roe deer 17 monitored in Chizé from 2003 to 2008 (Note: some roe deer were monitored during multiple 18 years). The kernel method was applied, and we subsequently identified the modes of these 19 distributions (see Pellerin et al. 2008, for more details on the monitoring of these animals). 20 This resulted in 25 sets of points (between 1 and 17 points per set), which we used to 21 simulate the presence of attraction points for our simulated animals. To simulate the 22 movement of an animal, we randomly selected one of these sets and randomly placed it on 23 our study area, giving its centroid a probability equal to 0.64 to be located in the northern 24

part to reflect the higher habitat quality in this region, as reported by Pettorelli et al.
(2001). We made sure that all the simulated movements were located within the limits of
the study area when sampling a centroid position, to make sure that all animals were
present within the study area during the whole study period and to avoid edge effects on
our results. When an animal survived across multiple years, we used the same centroid for
this animal during all years, thereby simulating its sedentarity.

We then simulated our movement model. We first supposed two types of movements for 7 each animal: (i) Patch-level movements characterized by OUf processes with small 8 variances and centred on one attraction point (see below); (ii) Between-patch movements 9 characterized by OUf processes with larger variances, also centred on one attraction point. 10 The supplementary material gives more details on the precise parameterization of these 11 OUf processes. Analysis of the GPS data showed that we could roughly describe animal 12 movements as randomly switching between three behaviours: resting (immobility), 13 patch-level movements (movement concentrated within a very small portion of the home 14 range) and between-patch movements (larger movements within the home range). From 15 this dataset, we derived rough estimates of both the daily frequency and timing of these 16 movements types (unpublished results). This information was used to parameterize the 17 simulated movements in our simulations. Fig. 1 gives a schematic description of the 18 approach used to simulate this stochastic switching, and we describe this approach more in 19 detail below. 20

We simulated the same activity cycle for all animals and all days of the study period. We simulated the animal movement during each day between the 1st and 31st of March, as follows. For each day (starting at 17:00 and ending à 16:59), we first simulated the presence of a main resting period with a probability equal to 0.85, its starting time *m* being

randomly drawn between 8:00 and 15:00. The duration (in hours) of this resting period was 1 equal to 15-m + e, with e a residual randomly drawn from an Gaussian distribution $\mathcal{N}(0, 1)$, 2 and truncated so that this resting period never ends after 16:59. We then randomly placed 3 patch-level movements during the remaining periods of the 24h sequence. We simulated the start time of these patch-level movements by simulating a Poisson process with constant 5 intensity ensuring the presence of 0.66 patches per night on average, with a duration 6 randomly drawn between 0h and 5h. Finally, we placed resting patches in the remaining 7 periods (except between 6:00 and 8:00, when all deer are considered active). We simulated 8 the start time of these resting patches by simulating a Poisson process with constant 9 intensity ensuring the presence of 0.33 patches per night on average, with a duration also 10 randomly drawn between 0h and 5h. Between-patch movements were simulated in the 11 remaining periods. 12

When the animal switched from one movement type to another (e.g., from between-patches to patch-level, from patch-level to between-patch, or from resting to between-patch), a new attraction point was randomly drawn from the simulated set. When the animal initiated a resting period, it stopped moving immediately and remained stationary until the period ended. An example of simulated movements is given in Fig. 2A).

We randomly placed camera traps on the study area (see next sections), and simulated for each trap a motion sensor (i.e., camera traps triggered by the presence of animals, taking at most one picture per second). We defined a detection zone for each trap consisting of a 20 metres circular sector with an angle of 10 degrees. We chose this angle based on Rowcliffe et al. (2008), but the results of the simulations were not affected by this choice (see section 5.5 of the supplementary material for a set of simulations with an angle

of 42 degrees). When the movement of animals crossed the detection zone, we simulated 1 imperfect detection by the sensors (Fig. 2B). We simulated a detection probability 2 decreasing with the distance to the camera trap, and depending on the habitat type where 3 the trap was located. Three habitat types are present in our area: (i) Coppice (76% of the study area), (ii) Open habitat (6% of the study area), (iii) Regeneration (18% of the study 5 area). In Coppice and Open habitat types, we chose the parameters of the detection 6 probability to ensure that the detection probability was equal to 0.99 at 5 metres from the 7 trap and to 0.01 at 18 metres from the trap (calibrated visually from the results of Howe 8 et al. 2017). We simulated the presence of trees obscuring the view in the Coppice habitat 9 type (see supplementary material for more details on the distribution of tree diameters). In 10 the regeneration habitat type, we did not simulate explicitly the presence of such obstacles, 11 but we subjectively chose parameters ensuring a detection probability of 0.99 at 0.5 metres 12 and 0.01 at 3 metres. Finally, we supposed that an animal resting in the detection zone 13 could not be detected by the sensors. More details on the parameters of the detection 14 process are given in supplementary material. 15

16 1.2 Three types of simulations

17 1.2.1 Roe deer population, no habitat selection

We simulated a simple population dynamics process. We used the data collected on the Chizé study area for the last 30 years to derive rough estimates of the key demographic parameters in this are, which were used to calibrate the simulations (based on the estimates of a capture-recapture model fitted every year to field data by the last author, but see also Gaillard et al. 1992, 1993, 1997, for other sources). We established an initial population size

of 300 roe deer with a balanced sex ratio. Each year, 90% of females were assumed to
reproduce, with an average of 1.7 fawns per reproducing female. We simulated a fawn
survival probability of 0.3 and an adult survival probability of 0.7. These parameters,
selected from realistic ranges observed on the area, were chosen to induce a gradual
population decline from 300 animals during year 1, to 239 animals during year 5 (see
supplementary material for exact numbers of animals simulated every year).

We simulated an influence of the decreasing population density on animals' home-range 7 by gradually increasing mean home-range size over time. Kjellander et al. (2004) indeed 8 observed this negative correlation between home-range size and population density in two 9 populations. In reality this relationship can be complex (see discussion), but our aim here 10 was to assess the effect of the opposite effects of the density decline and increasing 11 home-range size on the number of detections by the traps. More precisely, we gradually 12 increased the variance of the OUf process controlling for the between-patch movements. We 13 also varied the list of sets of attraction points used to define the patches, considering only 14 the 19 "small" sets with less than 10 attraction points during the first year, and 15 progressively adding an increasingly larger sample of the 6 "large" sets with more than 10 16 attraction points in year 2 to 5. This led to a progressive increase of the home-range size, 17 from an average of 13.1 ha during year 1 (SD = 4.4 ha) to 25.8 ha during year 5 (SD = 18 12.9 ha). Note that this increase in home-range size also affected the average movement 19 speed of the animals, which may be important given that REM estimators require the 20 knowledge of average speed. 21

Thus, we simulated the movements of all animals of the population during the month of March (i.e., after reproduction when animals are no longer territorial, and before the births of fawns) for every year of the 5-year period. To simulate camera trap monitoring, we

randomly placed either 100 or 25 camera traps over the study area, corresponding to a
density of approximately 4 traps/km² and 1 trap/km² respectively. We then used the REM
and IS to estimate the population size from the data collected by these traps (see below).
For each density of camera traps (4 or 1/km²), we carried out 1000 simulations of a camera
trap monitoring.

6 1.2.2 Roe deer population, selection of paths neighborhood

The previous simulations did not account for habitat selection by the target species. In 7 practice, researchers utilize knowledge of habitat use by the species to enhance their 8 monitoring design. This can involve stratification, where pre-specified proportions of 9 camera traps are placed in the different habitats, with overall density calculated as a 10 weighted average of the habitat densities (Rowcliffe et al. 2008). Alternatively, some studies 11 abandon unbiased population size estimates in favor of unbiased trends estimates. To 12 achieve this, they maximize detections by deploying camera traps only in highly selected 13 habitats or in locations with high detection probabilities (e.g., open habitat), assuming a 14 constant bias in size estimates. This strategy relies on increased detections to enhance 15 trend precision. We assessed these two strategies with a new set of simulations, by 16 simulating a strong habitat selection by the roe deer. We limited these simulations to a 17 trap density of 4 traps/km² (100 traps). 18

We simulated a strong selection of the human paths and roads by the roe deer on our study area. This specific selection of linear features by the roe deer is not expected in reality from a biological point of view, as such features are generally found at the home-range periphery (Seigle-Ferrand et al. 2021). However, this artificial scenario allows us to illustrate the ideal situation where the camera traps are placed in habitats with both

¹ high target species abundance and maximum detection probability.

We defined "paths neighborhood" as the 20-metres buffer around each human path or 2 road present in our study area. We simulated a selection of this habitat type by the roe 3 deer: we first selected the location of the home-range centroid in the northern part with a probability of 0.64, as before. However, for this set of simulations, we ensured that: (i) the 5 centroid was located in the paths neighborhood, (ii) it was located at more than 600 m 6 from the border of the study area (to make sure that all the movements were located within 7 the area). Then a random number of attraction points comprised between 3 and 10 was 8 selected within a random distance comprised between 100m and 300m from this centroid, 9 making sure that all these points were located within the path neighborhood. We then 10 carried out again the simulations described in the previous section, using these randomly 11 generated attraction points instead of a randomly selected set of attraction points as before. 12 As for the previous set of simulations, we varied the parameter σ^2 of Ornstein-Uhlenbeck 13 processes to simulate an increase in home-range size across years (see supplementary 14 material). 15

We then simulated two types of camera trap monitoring, corresponding to the two strategies described previously. In the first simulation type, we assessed the impact of placing all camera traps within highly selected habitat, specifically by randomly distributing them in the paths neighborhood.

In the second type of simulations, we evaluated the impact of stratified sampling on the precision of population size estimates. We simulated various stratified sampling designs, allocating 5% to 95% of the 100 traps to path neighborhoods in 5% increments. In practice, we did not simulate a single study with X% of traps in the path neighborhoods. Instead, we simulated two separate studies: one with all 100 traps placed in path neighborhoods and

another with all 100 traps placed outside of path neighborhoods. To simulate stratified 1 sampling, we combined the detection data from these two scenarios by randomly selecting 2 the detections from X% of the traps from the first study (all traps in path neighborhoods) 3 and the detections from 100-X% of the traps from the second study (all traps outside path neighborhoods). This allowed us to assess the impact of different proportions of traps in 5 path neighborhoods on population size estimates without directly simulating a stratified 6 sampling design within the population itself (which was less computer intensive). Although 7 the animals simulated in the two scenarios (100% and 0% in paths neighborhood) are not 8 the same, we showed in the supplementary material that the results obtained with this 9 approach do not differ from an approach where we simulated an actual stratified monitoring 10 of the population where all 100 traps study the same animals. 11

¹² We simulated each simulation type 500 times.

13 1.2.3 "Red deer" population, no habitat selection

Finally, we aimed to assess the effect of the species home-range size on the results. To do 14 so, we considered an animal with a home-range size similar to that of the red deer (*Cervus* 15 *elaphus*). Although the simulated animal remains the same as before (e.g. same activity 16 rhythm), with the only difference being a significantly larger home range, we will refer to 17 this animal as the "red deer" in the following. We used the same movement algorithm as 18 for the roe deer, with the following differences in parameterization: (i) we increased 19 strongly the value of the variance of the between-patch and patch-level movement, (ii) the 20 attraction points are randomly drawn in a rectangular box of 4 kilometres wide centred on 21 the home range centroid. We did not simulate a home-range size varying with population 22 decline over the 5 years. The simulated home-ranges covered 469 ha on average 23

1 (interquartile range: 383 to 548 ha).

As for the roe deer, we simulated a population size of 300 red deer, which corresponds to a high density for this species though not unrealistic (Borowik and Jwdrzejewska 2018). This choice allowed a comparison with the results obtained for the roe deer. We assumed a sex-ratio of 0.5, an adult annual survival of 0.76, and a reproduction equal to 0.5 fawns per female. The simulated number of red deer was equal to 240 during the last year, corresponding to a 20% decrease in 5 years.

We then placed randomly either 4 traps/km² or 1 trap/km² over the study area. For
each number of traps, we simulated 1000 times the camera trap monitoring.

10 1.3 Population size estimates

For each simulation of each simulation set, we estimated the population size using two methods, REM and IS. To use the REM population size estimator, we need to consider the total number Q of animal-trap encounters. The estimator is:

$$\hat{N}_{\text{rem}} = \frac{S \times Q \times \pi}{A \times v \times r \times (2+\theta)} \tag{1}$$

¹⁴ With \hat{N} the population size estimate, S the study surface area, A the the cumulated ¹⁵ activation duration of the traps, v the mean travel speed of the animals, and r, θ the depth ¹⁶ and angle of the detection zone. Note that one of the biggest difficulties with the REM is ¹⁷ that the mean travel speed of the animals is to be estimated, which can be an important ¹⁸ source of imprecision of the population size estimation. In our simulations, we have ¹⁹ considered that this mean speed is known. Although this will never be the case in practice, ²⁰ our aim was not to devise means to estimate travel speed.

To use the IS population size estimator, we needed to reshape our data. We discretized 1 our study period (the month of March of every year during five years) into one-second 2 intervals. We define "capture occasions" as J discrete moments marking the start of these 3 intervals. For each capture occasion and each animal-camera trap pair, we determined whether a detection occurred (i.e., if the simulated motion sensor of the camera had 5 detected the animal at that time). For each trap i and each capture occasion j, we define a 6 new variable n_{ij} corresponding to the number of animal-trap associations (number of animals present in the detection zone at that time and detected by the trap). Let s_i be the 8 surface area of the detection zone of the camera trap i. The IS population size estimator is: 9

$$\hat{N}_{is} = S \cdot \frac{1}{J} \cdot \frac{1}{M} \sum_{j=1}^{J} \sum_{i=1}^{M} \frac{n_{ij}}{s_i}$$
(2)

¹⁰ where M is the number of traps placed on the area.

The two estimation methods were applied each with three modalities: (i) we used the 11 above estimators without accounting for the limited detectability of animals due to 12 imperfect sensors, lack of visibility or activity rythm (hereafter, these estimators are simply 13 called REM and IS), (ii) we accounted for the limited visibility in the detection zones by 14 modifying these estimators (hereafter called REM d and IS d, see below), (ii) we 15 accounted for both this limited visibility and the reduced detectability during resting 16 (hereafter called REM_da and IS_da). For the REM estimator, the mean travel speed was 17 computed exactly from the simulated movements, using all data for the REM and REM_d, 18 and using only data collected during active periods for the REM_da. 19

To account for the imperfect visibility in the detection zone in the REM estimator, we replaced Q in equation 1 by Q/\bar{p} , where \bar{p} is the proportion of all encounters that were

detected by the traps. This average detection probability of encounters was estimated from 1 the simulated data, where we knew the exact times when animals crossed trap detection 2 zones. To account for this imperfect detectability of animals in the detection zone in the IS 3 estimator, we replaced s_i by $s_i \times \overline{d}$ in equation 2 where \overline{d} is the mean detection probability of associations. This average detection probability of associations was also estimated from 5 simulated data, where we knew the exact detection probability for all traps (as in fig. 6 2(B)). Finally, to account for the limited detectability of animals in the detection zone 7 during resting, we changed our study period to remove all the data collected between 8:00 8 and 17:00, when the main resting period occurs. 9

For each estimator and each estimation modality, we estimated the population size for each year of the 5 years period, as well as the change rate: $CR = (N_5 - N_1)/N_1$. We also calculated a mean estimated trend over the 5-year period:

$$\hat{\lambda} = \exp\left\{\frac{\operatorname{Cov}(\log \hat{N}_t, t)}{\operatorname{Var}(t)}\right\} - 1$$

where t is the year and \hat{N}_t the estimated population size during year t. This parameter $\hat{\lambda}$ estimates the proportion of the population disappearing in one year.

¹² When the simulations implied a stratified sample, we calculated one population size ¹³ estimate for each strata and summed the two estimates. We used the bootstrap to estimate ¹⁴ the variances and 90% confidence intervals on population size, change rate and trend ¹⁵ estimates, by resampling for each estimation 1000 times with replacement the camera traps ¹⁶ and recomputing the estimates to obtain a distribution of 1000 bootstrap estimates. When ¹⁷ the simulations implied stratified samples, we bootstrapped the two strata separately. ¹⁸ In total, all these simulations required ≈ 20 days of calculation on a Dell T5610

¹ workstation with a processor Intel(R) Xeon(R) CPU E54-2650 (2.6GHz).

2 2 Results

³ 2.1 First simulation set: roe deer with no habitat selection

We present the results of the simulations in Tables 1 (4 traps/km^2) and 2 (1 trap/km^2). In 4 the two cases, it is clear that the two estimators strongly underestimate the population size 5 when the imperfect visibility in the detection zone is not accounted for, and more strongly 6 so for the IS. Accounting for this imperfect visibility leads to an unbiased estimate of the 7 population size with the REM, but IS still results in a biased estimate. Accounting for the 8 limited detectability of the animals in the detection zone when they rest results in an 9 unbiased estimate for both IS and REM. The bootstrap allowed us to correctly estimate the 10 standard error of the sampling distribution for all estimates. The trends and change rate 11 were correctly estimated for all modalities and methods. 12

¹³ With 4 traps/km² on the study area, the population size estimation was rather

¹⁴ imprecise (with a coefficient of variation of about 15% to 20%, depending on the method).

¹⁵ A population size decrease was correctly identified in more than 80% of the simulations

with all methods; however, this decrease was significant only in 1/4 to 1/3 of the

¹⁷ simulations, depending on the estimation method.

With only 1 trap/km² on the study area, the population size estimation was even more imprecise (with a coefficient of variation of about 30% to 40%). A population size decrease was estimated correctly in about 2/3 to 3/4 of the cases, but this decrease was rarely significant (only in about 1/6 simulations).

1 2.2 Habitat selection by the roe deer

When we simulated habitat selection and placed all the camera traps in the most selected habitat type, the population decrease was strongly overestimated (Table 3). Indeed, the estimated population decline was approximately 8% per year, leading to a 28% decrease in year 5 (i.e., $1 - (1 - 0.08)^4 = 0.28$), instead of the simulated 20% decline (corresponding to a yearly decline of 5.5%).

Using a stratified sample allowed an unbiased population size estimation. The highest
precision of the estimation was obtained when half of the traps were placed in the paths
neighborhood and half out of this habitat (Fig. 3). However, this stratified sampling did
not improve noticeably over a simple random sampling (the smallest coefficients of variation
on this graph were similar to those obtained in Table 1).

12 2.3 Simulations of a red deer population

The simulations of a red deer population led to the same conclusions regarding the bias and 13 precision of population size estimates: accounting for imperfect visibility in the detection 14 zone led to less biased estimates by the REM and the IS, and accounting for the reduced 15 detectability of resting animals by sensors led to unbiased estimates for the IS. The 16 precision of population size estimates was comparable to that obtained for roe deer, for 17 both 4 traps/km² (Table 4) and 1 trap/km² (Table 5). However, the trend and change rate 18 estimations were much more precise. With 4 traps/km², the simulated decrease was nearly 19 always identified, and was significant in more than 95% of the cases for the REM, and in 20 1/2 to 3/4 of the cases for the IS. With 1 trap/km², the decrease was also frequently 21 identified in most cases with the REM, and in more than 80% of the cases for the IS. The 22

¹ decrease was significant in 1/3 (IS) to 2/3 (REM) of the cases.

We also compared various descriptive statistics between the roe deer and the red deer 2 population (Table 6). The mean number of encounters with red deer was more than twice 3 that of roe deer. However, the mean speed of red deer was also more than twice bigger than that of roe deer, so that the resulting number of animal-trap associations (and its standard 5 deviation) was similar for the two species. As a result, the population size estimates were 6 comparable between the two populations, whether estimated with REM or by IS. However, 7 since the home-range size was much larger for the red deer than for the roe deer (36 times 8 larger during year 1), the average red deer visited 7 times more traps than the average roe 9 deer. As a result, each camera trap, on average, detected 7 times more red deer individuals 10 than roe deer individuals. Therefore, when a red deer died between two years, the number 11 of encounters and associations decreased in a larger number of traps than for the roe deer 12 (and conversely when a new animal was born). As a result, by bootstrapping the camera 13 traps to calculate the standard deviation for the trends, it was much easier to detect red 14 deer population changes than roe deer population changes. 15

16 **3** Discussion

We have simulated different settings of camera trap monitoring of a roe deer population.
Our simulations revealed the importance of accounting for imperfect visibility in the
detection zone to estimate the population size correctly with the REM method and even
more with the IS method. Moreover, it was also important to account for the limited
detectability of animals due to the activity rhythm of the animals for the IS method only.
This was less important for the REM, as long as the mean speed was correctly estimated by

including inactivity periods. Although the population trend or the change rate were 1 accurately estimated, even without accounting for imperfect visibility or activity rhythm 2 (Table 1), the estimates were very imprecise and often not significantly different from zero. 3 With 1 camera $trap/km^2$, the roe deer population size and trend estimates were so imprecise that they were virtually useless (Table 2). When we simulated animals with a 5 much larger home range, the trend estimates were much more precise, even with only 1 6 trap/km². We also showed that placing the traps preferentially in the habitat selected by 7 the species led to a strong overestimate of the population decrease. Finally, we showed that 8 using a stratified sampling did not improve the precision of population size estimation 9 compared to random sampling. 10

We used the REM and IS to estimate the population size of the roe deer and red deer 11 populations. We did not compare the two methods to determine which one was the more 12 interesting for the monitoring. Such a comparison would be difficult, as the two methods do 13 not require the same elements to be used. Thus, our results seem to indicate that the two 14 methods give comparable results, though the REM seems more precise than the IS in our 15 simulations. However, as Nakashima (2022) noted, the mean movement speed, which we 16 supposed known in our simulations, is never known in practical studies and has to be 17 estimated. The uncertainty of this parameter estimation results in a larger uncertainty of 18 the population size estimates, so that comparing IS and REM is not fair without 19 accounting for the speed estimation. There are numerous other methods available to 20 estimate the population size based on camera trap data collected on unmarked populations 21 (Nakashima et al. 2017, Moeller et al. 2018), and we chose IS and REM because they were 22 the most easily automated (easily computed estimator that does not rely on the complex fit 23 of a model) and because they belonged to the two main families of estimation methods: 24

methods relying on the encounters and methods relying on associations. We think that our 1 simulation results, that are similar for REM and IS, also generalize to other methods. 2 Our results are consistent with previous research highlighting the importance of 3 accounting for the imperfect visibility in the detection zone to obtain unbiased population size estimates (Moeller et al. 2018, 2023). While we assumed perfect knowledge of detection 5 probabilities in our simulations, real-world applications require estimating these 6 probabilities, which can introduce additional uncertainty into population size estimates. 7 Accounting for this imperfect detectability of animals in the detection zones with methods 8 relying on detection is basically the rationale behind the camera-trap distance sampling 9 (Howe et al. 2017). Thus, IS without accounting for detectability is very similar to 10 camera-trap distance sampling where the detection probability is set equal to 1. 11 In our study, the roe deer population was characterized by a home-range size increasing 12 with time, as a result of the decreasing animal density. This correlation can occur under 13 various ecological conditions. For instance, in the hypothetical case where reduced habitat 14 productivity (e.g., caused by shifts in land management practices) triggers a population 15 decline due to diminished foraging resources, the habitat productivity hypothesis predicts 16 that resource scarcity should also drive an increase in home-range sizes, as individuals 17 compensate for limited resources (Harestad and Bunnel 1979). Alternatively, this 18 correlation might emerge from density-dependent behavioral mechanisms: declining 19 populations could reduce competition among males, potentially allowing for larger 20 individual territories. Such an inverse relationship between male home-range size and 21 density has been identified in the roe deer by Vincent et al. (1995). Note that the habitat 22 productivity hypothesis may also predict a positive correlation between home-range size 23 and population density under different conditions. For example, if population reductions 24

stem from non-habitat factors (e.g., hunting pressure), lower densities could increase 1 resource availability per capita, thereby shrinking home ranges. Regardless of the 2 underlying mechanism, such correlations do exist in nature, and may bias density 3 estimation with the REM when the increase in home-range size is also correlated with increased mean animal travel speed: in our simulations, the mean speed was 30% smaller in 5 year 1 than in year 5. As the population size estimator is inversely related to this mean 6 speed, using a single value of mean speed for all years, as commonly done in studies using REM, would result in a substantially biased estimation (e.g., using the mean speed of year 8 1 for all years in equation 1 would result to a 30% underestimation in population size 9 during year 5). 10

Accounting for activity patterns had a minimal impact on REM population size 11 estimates. The random camera placement meant resting areas were sampled as frequently 12 as active ones. While resting animals are undetectable by motion sensors, their entry and 13 exit from a trap's detection zone, which requires movement, allow encounter detection. 14 Therefore, resting periods within encounters did not affect REM results. On the other 15 hand, since association-based methods strongly rely on the detection of animals at any 16 moment of the encounters, the presence of a resting period during an encounter may lead to 17 a large number of associations missed by a motion-triggered sensor, which explains why 18 association-based methods are more sensitive to activity patterns. Avoiding to include the 19 monitoring periods during which a substantial fraction of the animals are resting allows to 20 remove this small bias, which is also consistent with the findings of previous authors (Howe 21 et al. 2017). This is the simplest approach to account for the activity rhythm of animals, 22 and although other approaches are possible (e.g., by estimating the proportion of the day 23 during which the animal is active and by modifying the duration of the study period in the 24

equations Nakashima et al. 2017), our approach led to unbiased population size estimates. Our results raise the question of whether motion-sensors or time-lapse programming of 2 camera traps are preferable for camera trap studies. Association-based methods require 3 time-lapse data, which can be obtained by direct time-lapse programming or post-processing of motion-triggered trap data. Direct time-lapse programming eliminates 5 sensor sensitivity bias, but image analysis, often AI-assisted (Rigoudy et al. 2023), may still 6 introduce detectability issues related to distance and visibility. Nevertheless, contrarily to 7 motion sensors, direct time-lapse programming is not affected by the activity of the animal 8 and detects both active and resting animals equally. However, the short time-lapses needed 9 for association-based methods (Howe et al. 2017) raise storage challenges: trap's memory 10 fills up quickly. Therefore, direct time-lapse programming might not be adequate with 11 association-based methods. 12

The fact that it was not necessary to account for the activity rhythm of animals and 13 imperfect visibility in the detection zones in our trends estimate is probably a result of our 14 simulation design. Indeed, as long as the simulated detection probability and activity 15 rhythms of animals were the same across years, the resulting bias in the population size 16 estimate was constant, allowing for unbiased estimation of the population trends. However, 17 we do not expect this result to hold if the detectability in the detection zones or the 18 activity rhythm changes across years. For example, the encroachment of shrubs into the 19 study area would lead to an average detection probability decreasing with time. 20 Considering constant detection, or not considering detection at all, would result in an 21 increasing underestimation of population size with time and, consequently in our example 22 of declining population, in an overestimation of the decreasing trend. Not accounting for 23 the activity rhythm or the imperfect detection in trend estimation is only possible under 24

¹ the assumption that these phenomena do not vary with time.

This variation of animal behaviour across years does explain our overestimation of the 2 decline when the traps are placed in highly-selected habitats only. Indeed, since we knew 3 that the roe deer selected preferentially the paths neighborhood, it seemed intuitive to place all the traps in this habitat to maximize the number of detections, and thereby 5 enhance the precision of the estimated trend. However, this placement led to a strong 6 overestimation of the decreasing trend (the percentage of the population of year 1 that had disappeared in year 5 was 20%, but was estimated at 28%). This was caused by the 8 simulated change in home-range size. Indeed, the simulated negative correlation between 9 population size and home-range size led to larger home ranges during year 5. Even if the 10 habitat preferences of the animals were the same during the 5 years of the study, 11 individuals spent more time travelling between selected patches in later years. Therefore, in 12 average the animals spent less time in paths neighborhood in year 5 than in year 1 (Fig. 4). 13 The mean proportion of the population located in the paths neighborhood at a given 14 moment was therefore not constant across years, which resulted in an overestimation of the 15 decrease of the population. The supplementary material demonstrates mathematically that 16 combining the actual population decrease with this decrease in the proportion of the 17 population in the paths neighborhood results in an annual decrease of 8% of the number of 18 encounters/associations in this habitat. 19

While altering the sampling strategy could potentially enhance estimate precision, we found no suitable approach; specifically, stratification did not improve precision compared to simple random sampling. Another common strategy involves placing camera traps at the nodes of a regular grid, thereby ensuring a good spatial coverage of the population. We did not consider this grid-based strategy, as it would have yielded results similar to random

1 sampling: given our random distribution of home ranges across the study area, the spatial
2 relationship between animals and traps would have been equally random for both
3 grid-based and simple random sampling. The estimates would therefore have been
4 equivalent. Furthermore, density estimation theory assumes random trap placement (e.g.
5 Rowcliffe et al. 2008), making grid-based sampling potentially unsuitable.

6 Several aspects of our simulations were oversimplistic: our simulated animals

 $_{7}\,$ corresponded to a point moving in a 2D space, and not to a volume moving in a 3D space.

8 This simplification may affect some of our results. For example, when the point

representing the animal was located behind a tree, it was supposed to be completely hidden 9 by the tree, whereas in reality animals can be detected by traps when they are located 10 behind small trees. We think that this did not affect strongly our results, as we simulated 11 the presence of trees only for the coppice habitat type (which represents 76% of the study 12 area), and always at a low density (see supplementary material for details): there was in 13 average 0.37 trees simulated in the detection zone of a camera trap located in the coppice 14 habitat. Simulations showed that the trees obscured in average only 1.2% of the detection 15 zone of a trap in the Coppice habitat, and these trees had more chance to be located far 16 from the traps, i.e., where the detection probability is already low, since this is where the 17 area of the detection zone is the largest. 18

More generally, simplifying assumptions in our simulations led to an overestimation of precision in our results: Our camera traps did not deteriorate with time. We did not simulate any camera trap effect on the detectability of the encounters/associations. Although our simulated animals were characterized by a variable home range size, they all had the same activity rhythm. There were no differences between male and female, or young and adults. Animals were not gregarious (simulating gregarious animals would have

led to a greater variance of the encounter rate across camera traps). In other words, it is 1 expected that the uncertainty characterizing the population size and trends estimates in 2 actual studies will be much more important than those obtained in our simulations. And 3 yet our results concerning the roe deer show that the estimation of population size and trends are very imprecise. Even with 4 camera $traps/km^2$, the coefficient of variation of the 5 population size estimate was about 15 to 20% and increased to 30 to 40% when only 1 6 $trap/km^2$ were used. Similar coefficients of variations were obtained in previous studies 7 (e.g., Palencia et al. 2022, obtained CV comprised between 34% to 75% when estimating 8 the population size of various species with the REM, on areas comprised between 1400 to 9 6600 ha with 7 to 37 camera traps during periods covering 15 to 138 days). Our study 10 showed that a strong population decrease of 20% in 5 years was not significant in 2/3 to 11 3/4 of the simulations when 4 traps/km² were used, and this proportion rose to 5/6 when a 12 density of only 1 $trap/km^2$ was used. As our simulations were very optimistic, this 13 proportion is expected to be much smaller in real studies. This shows that the use of 14 camera traps for the monitoring of roe deer population trends is probably of limited value. 15 The results obtained for the red deer were more encouraging. Even with only 1 camera 16 $trap/km^2$, we were unable to detect a 20% decrease of the population size in 1/3 to 2/3 of 17 the simulations, depending on the methods used. This is still a low sensitivity, given the 18 strong simulated population decrease and the numerous simplifying assumptions of our 19 simulations. However, these results indicate that camera traps monitoring might not be 20 immediately dismissed for larger species in areas of size similar to the Chizé forest. An 21 important quantity governing the choice to use or not camera traps for population 22 monitoring is the number of different camera traps that can capture a given animal in a 23 given study area. This metric depends on multiple factors, including the number of traps 24

deployed, the species' home-range size, heterogeneity of space use within the home range, 1 and the size of the study area. For example, considering the red deer in our study, 2 randomly placing 100 camera traps over the 2600 ha study area resulted in a density of 3 approximately 4 traps/km², meaning that roughly 18 traps were included within the average 4.5 km^2 home-range. Because red deer home-range use is highly heterogeneous, 5 with activity concentrated around a limited number of attraction points (see supplementary 6 material for an illustration), the actual number of traps detecting a given red deer is 7 considerably reduced, averaging 3.5 traps per animal (Table 4). Consequently, the death of 8 red deer between two years is detected by an average of 3.5 traps, resulting in a clear 9 decrease in the number of encounters/associations detected by the traps. This explains the 10 effectiveness of camera trap monitoring for this species under these conditions. In contrast, 11 with a larger study area of 20000 ha and the same 100 traps, the trap density would 12 decrease to 0.5 traps/km². This would result in approximately $0.5 \times 3.5/4 = 0.44$ different 13 traps encountered by a given red deer in average, similar to the value obtained for the roe 14 deer in our simulations (Table 4). Indeed, for roe deer, with only 0.45 traps visited per 15 animal in Chizé, most individuals remain undetected, resulting in low power to detect 16 population declines. 17

We advise the reader to proceed to simulations to determine whether or not camera traps are a suitable choice to design a monitoring of a given population. Our R package simCTChize can be useful to carry out such simulations. The supplementary material of the present paper, also available as a vignette of the package, contains a tutorial explaining how it can be used to reach this aim.

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Table 1: Results of 1000 simulations of a roe deer population monitored for 5 years using 4 camera traps/ km^2 (see text). We use two methods to estimate population size, the random encounter model (REM) and the instantaneous sampling (IS). For each method, we estimate the population size: (i) without accounting for imperfect visibility in the detection zone or the reduced detectability of animals by the motion sensors in this zone during their resting periods (REM and IS), (ii) accounting for imperfect visibility but not for the limited detectability of animals during their resting periods (REM d, IS d), (iii) accounting for both imperfect visibility and limited detectability of animals during the resting periods (REM da and IS da). For each method, we present the results for the population size during year 1 (N1) and year 5 (N5), as well as the trend λ over five years (see text; lam, theoretical value = -0.0547) and the change rate CR (= [N5-N1]/N1; theoretical value = -0.2). For each method and each parameter, we present the simulated value of the parameter (TrueValue), the mean of estimated values (MeanEst), the standard deviation of the distribution of estimated values (SE), the mean of the bootstrap estimates of this standard deviation (SEEst), the standard deviation of the distribution of the bootstrap estimates of this standard deviation (SE.SEEst), and the proportion of simulations for which the 90% confidence interval estimated by the bootstrap includes the true value of the parameter (Pcov). For lam and CR, we also present the proportion of simulations for which a decrease is estimated (negative value of lam or CR; PDec), as well as the proportion of simulations for which the upper limit of the confidence interval is smaller than 0 (PsigD).

$\begin{array}{c c c c c c c c c c c c c c c c c c c $	Method	Param	TrueValue	MeanEst	SE	SEEst	SE.SEEst	Pcov	PDec	PsigD
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM	N1	300	237.31	37.875	38.687	6.215	0.511		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM	N5	239	181.767	33.792	31.553	6.114	0.444		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM	lam	-0.0547	-0.063	0.049	0.048	0.007	0.869	0.906	0.364
$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM	CR	-0.2033	-0.22	0.165	0.165	0.043	0.859	0.906	0.378
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$										
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_d	N1	300	303.86	45.68	49.51	7.853	0.923		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_d	N5	239	232.533	40.045	40.352	7.572	0.882		
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	REM_d	lam	-0.0547	-0.063	0.049	0.048	0.007	0.867	0.906	0.367
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_d	CR	-0.2033	-0.22	0.165	0.169	0.042	0.9	0.906	0.327
$\begin{array}{cccccccccccccccccccccccccccccccccccc$										
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_da	N1	300	303.224	47.154	50.959	8.427	0.928		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_da	N5	239	232.706	41.018	41.146	7.83	0.882		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_da	lam	-0.0547	-0.062	0.051	0.05	0.007	0.867	0.889	0.344
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	REM_da	CR	-0.2033	-0.217	0.172	0.176	0.045	0.904	0.891	0.311
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$										
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	IS	N1	300	54.903	9.955	10.084	1.897	0		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	IS	N5	239	42.983	8.975	8.3	1.844	0		
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IS	lam	-0.0547	-0.058	0.055	0.055	0.008	0.88	0.843	0.289
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IS	CR	-0.2033	-0.198	0.195	0.198	0.057	0.87	0.848	0.279
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$										
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	IS_d	N1	300	262.512	47.812	46.688	11.689	0.737		
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	IS_d	N5	239	205.374	40.169	37.611	9.195	0.682		
IS_d CR -0.2033 -0.197 0.196 0.197 0.056 0.882 0.841 0.276 IS_da N1 300 298.869 57.784 56.794 15.407 0.881 IS_da N5 239 233.97 49.233 45.015 12.228 0.849 IS_da lam -0.0547 -0.057 0.059 0.057 0.009 0.885 0.833 0.272 IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257	IS_d	lam	-0.0547	-0.057	0.056	0.054	0.008	0.884	0.854	0.279
IS_da N1 300 298.869 57.784 56.794 15.407 0.881 IS_da N5 239 233.97 49.233 45.015 12.228 0.849 IS_da lam -0.0547 -0.057 0.059 0.057 0.009 0.885 0.833 0.272 IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257	IS_d	CR	-0.2033	-0.197	0.196	0.197	0.056	0.882	0.841	0.276
IS_da N1 300 298.869 57.784 56.794 15.407 0.881 IS_da N5 239 233.97 49.233 45.015 12.228 0.849 IS_da lam -0.0547 -0.057 0.059 0.057 0.009 0.885 0.833 0.272 IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257										
IS_da N5 239 233.97 49.233 45.015 12.228 0.849 IS_da lam -0.0547 -0.057 0.059 0.057 0.009 0.885 0.833 0.272 IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257	IS_da	N1	300	298.869	57.784	56.794	15.407	0.881		
IS_da lam -0.0547 -0.057 0.059 0.057 0.009 0.885 0.833 0.272 IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257	IS_da	N5	239	233.97	49.233	45.015	12.228	0.849		
IS_da CR -0.2033 -0.193 0.211 0.214 0.067 0.891 0.834 0.257	IS_da	lam	-0.0547	-0.057	0.059	0.057	0.009	0.885	0.833	0.272
	IS_da	CR	-0.2033	-0.193	0.211	0.214	0.067	0.891	0.834	0.257

Method	Param	TrueValue	MeanEst	SE	SEEst	SE.SEEst	Pcov	PDec	PsigD
REM	N1	300	235.668	78.199	72.339	23.699	0.688		
REM	N5	239	181.565	64.232	59.381	21.176	0.638		
REM	lam	-0.0547	-0.057	0.099	0.097	0.026	0.87	0.732	0.186
REM	CR	-0.2033	-0.161	0.383	0.417	0.358	0.871	0.738	0.193
REM_d	N1	300	301.727	93.669	92.96	29.521	0.868		
REM_d	N5	239	232.242	76.207	76.196	26.018	0.864		
REM_d	lam	-0.0547	-0.057	0.099	0.097	0.026	0.872	0.732	0.192
REM_d	CR	-0.2033	-0.161	0.383	0.417	0.289	0.887	0.738	0.161
REM_da	N1	300	301.069	97.229	95.555	31.466	0.871		
REM_da	N5	239	231.893	78.276	77.683	26.871	0.861		
REM_da	lam	-0.0547	-0.056	0.104	0.1	0.027	0.861	0.722	0.187
REM_da	CR	-0.2033	-0.15	0.413	0.436	0.276	0.877	0.734	0.159
IS	N1	300	54.785	20.805	18.818	7.119	0		
IS	N5	239	42.953	16.803	15.415	6.117	0		
IS	lam	-0.0547	-0.053	0.115	0.111	0.034	0.856	0.701	0.159
IS	CR	-0.2033	-0.107	0.503	0.628	0.908	0.848	0.701	0.166
IS_d	N1	300	263.562	97.551	87.342	36.81	0.768		
IS_d	N5	239	204.538	75.535	69.373	28.575	0.745		
IS_d	lam	-0.0547	-0.053	0.113	0.108	0.03	0.851	0.699	0.157
IS_d	CR	-0.2033	-0.119	0.482	0.605	1.06	0.856	0.703	0.167
IS_da	N1	300	301.313	120.668	105.91	50.082	0.816		
IS_da	N5	239	233.349	91.489	83.106	36.841	0.826		
IS_da	lam	-0.0547	-0.051	0.124	0.115	0.033	0.856	0.699	0.147
IS_da	\mathbf{CR}	-0.2033	-0.091	0.586	0.813	3.492	0.85	0.688	0.155

Table 2: Results of 1000 simulations of a roe deer population monitored for 5 years using 1 camera trap/km² (see text). We present the same estimates and the same methods as in Table 1.

Table 3: Results of 500 simulations of a roe deer population monitored for 5 years using 4 camera $traps/km^2$, where all animals display a preference for the neighborhood of paths and where all the traps have been placed in this neighborhood (see text). We present the same statistics for the estimated trend as Table 1, for the same estimation methods.

Method	MeanEst	SE	SEEst	SE.SEEst	Pcov	PDec	PsigD
REM	-0.08	0.03	0.03	0.00	0.73	1.00	0.87
REM_d	-0.08	0.03	0.03	0.00	0.73	1.00	0.86
REM_da	-0.08	0.03	0.03	0.00	0.75	1.00	0.83
IS	-0.08	0.04	0.04	0.01	0.77	0.99	0.70
IS_d	-0.08	0.04	0.04	0.01	0.79	0.99	0.71
IS_da	-0.08	0.04	0.04	0.01	0.81	0.99	0.69

Method	Param	TrueValue	MeanEst	SE	SEEst	SE.SEEst	Pcov	PDec	PsigD
REM	N1	300	207.511	40.895	40.456	4.87	0.325		
REM	N5	240	165.729	33.028	32.409	4.006	0.315		
REM	lam	-0.0543	-0.055	0.013	0.012	0.002	0.875	1	0.989
REM	CR	-0.2	-0.2	0.047	0.046	0.009	0.885	1	0.976
REM_d	N1	300	296.257	52.281	57.931	6.602	0.916		
REM_d	N5	240	236.537	41.961	46.383	5.297	0.916		
REM_d	lam	-0.0543	-0.055	0.013	0.012	0.002	0.874	1	0.991
REM_d	CR	-0.2	-0.2	0.047	0.046	0.009	0.937	1	0.993
REM_da	N1	300	296.516	52.561	57.848	6.605	0.921		
REM_da	N5	240	236.556	42.166	46.411	5.377	0.916		
REM_da	lam	-0.0543	-0.055	0.015	0.015	0.003	0.879	1	0.968
REM_da	\mathbf{CR}	-0.2	-0.2	0.057	0.054	0.011	0.931	0.999	0.964
IS	N1	300	51.406	10.502	10.236	1.301	0		
IS	N5	240	41.187	8.642	8.3	1.119	0		
IS	lam	-0.0543	-0.054	0.022	0.022	0.005	0.888	0.988	0.761
IS	CR	-0.2	-0.195	0.086	0.084	0.02	0.872	0.982	0.704
IS_d	N1	300	259.127	48.484	47.458	7.392	0.748		
IS_d	N5	240	208.346	39.598	39.027	6.823	0.76		
IS_d	lam	-0.0543	-0.053	0.026	0.026	0.007	0.892	0.974	0.672
IS_d	CR	-0.2	-0.19	0.101	0.099	0.034	0.889	0.959	0.611
IS_da	N1	300	295.255	57.31	55.822	10.413	0.879		
IS_da	N5	240	237.832	47.503	46.435	10.191	0.874		
IS_da	lam	-0.0543	-0.053	0.031	0.03	0.008	0.893	0.957	0.583
IS da	CR	-0.2	-0.187	0.12	0.117	0.045	0.897	0.936	0.51

Table 4: Results of 500 simulations of a red deer population monitored for 5 years using 4 camera traps/km² (see text). We present the same statistics for the same estimation methods as in Table 1.

Method	Param	TrueValue	MeanEst	SE	SEEst	SE.SEEst	Pcov	PDec	PsigD
REM	N1	300	210.484	80.766	78.826	19.945	0.655		
REM	N5	240	167.931	65.534	62.97	15.982	0.658		
REM	lam	-0.0543	-0.055	0.028	0.025	0.009	0.835	0.979	0.679
REM	CR	-0.2	-0.197	0.103	0.105	0.058	0.846	0.967	0.63
REM_d	N1	300	299.684	104.803	113.481	27.342	0.902		
REM_d	N5	240	239.059	84.642	90.663	21.937	0.895		
REM_d	lam	-0.0543	-0.055	0.028	0.025	0.01	0.835	0.979	0.673
REM_d	CR	-0.2	-0.197	0.103	0.105	0.059	0.88	0.967	0.645
REM_da	N1	300	299.633	104.979	112.892	27.098	0.902		
REM_da	N5	240	238.704	84.816	90.233	22.085	0.903		
REM_da	lam	-0.0543	-0.055	0.034	0.028	0.01	0.817	0.95	0.582
REM_da	CR	-0.2	-0.194	0.134	0.123	0.069	0.88	0.936	0.561
IS	N1	300	51.964	20.603	19.837	5.205	0		
IS	N5	240	41.302	16.859	15.898	4.35	0		
IS	lam	-0.0543	-0.056	0.05	0.043	0.017	0.828	0.878	0.423
IS	CR	-0.2	-0.185	0.199	0.227	0.348	0.816	0.85	0.376
IS_d	N1	300	260.332	95.011	91.322	25.821	0.823		
IS_d	N5	240	208.851	80.911	74.454	22.685	0.808		
IS_d	lam	-0.0543	-0.055	0.055	0.047	0.018	0.844	0.858	0.365
IS_d	CR	-0.2	-0.178	0.224	0.249	0.382	0.841	0.834	0.34
IS_da	N1	300	295.397	110.777	106.145	33.628	0.862		
IS_da	N5	240	238.746	98.29	87.876	31.69	0.854		
IS_da	lam	-0.0543	-0.054	0.065	0.054	0.02	0.824	0.819	0.328
IS_da	CR	-0.2	-0.164	0.277	0.325	0.726	0.813	0.791	0.289

Table 5: Results of 500 simulations of a red deer population monitored for 5 years using 4 camera traps/km² (see text). We present the same statistics for the same estimation methods as in Table 1.

Table 6: Mean value (SD in parentheses) for various statistics calculated for each simulation of a camera trap monitoring of a roe deer and red a deer population of 300 individuals with 100 camera traps.

	Roe Deer	Red Deer
Number of associations	19172 (SD = 3476)	17951 (SD = 3667)
Number of encounters	684 (SD = 101)	1584 (SD = 279)
Travel speed $(m.s^{-1})$	0.0165 (SD = 0.0001)	0.0387 (SD = 0.0002)
Number of traps visited by an animal	0.459 (SD = 0.056)	3.492 (SD = 0.62)
Number of animals seen by a trap	1.376 (SD = 0.167)	10.475 (SD = 1.86)



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Figure 1: Simulation of the movement of a roe deer during a typical day, starting at 17:00 and ending at 16:59 the following day. The sequence 1-5 describes the simulation of the activity cycle. The point 6 illustrates the simulation of the movement process itself, zooming on the beginning of the day.



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Figure 2: Simulation process used : (A) simulation of the movements of a roe deer using a model combining multiple Ornstein-Uhlenbeck process, including resting patches (green points) within the movement (when the animals stay at the same place without moving at all, (B) simulation of the detection process by a camera trap. The detection zone is modelled as a circular sector with a radius of 20 m and an angle equal to 10 degrees. Detection probability decreases with increasing distance from the trap, represented by varying shades of grey (darker = lower detection probability, and black = no detection at all). We also simulate the presence of trees that may limit the visibility within the detection zone. The movements of the animal simulated on panel (A) are displayed in red on this figure. The camera traps can capture the animals every second. The points correspond to the detections of the animals (more numerous close to the trap).



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Figure 3: Precision of the estimation of population size in a monitoring of a roe deer population with 300 individuals with 4 camera traps/km², using stratified sampling. The animals display a strong preference for the paths neighborhood on our study area (which represents 38% of the study area), and we vary the proportion of traps in this habitat type. We use two methods to estimate population size, the random encounter model (REM) and the instantaneous sampling (IS). For each method, we estimate the population size: (i) without accounting for imperfect visibility in the detection zones or the limited detectability of the animals in the detection zones during their resting periods (REM and IS), (ii) accounting for imperfect detectability of animals in the detection zones but not for these resting periods (REM_d, IS_d), (iii) accounting for both imperfect detectability and these resting periods (REM_da and IS_da)



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Figure 4: Estimated mean time spent by the roe deer in the paths neighborhood every year, in our simulations of a camera trap monitoring over 5 years. Since the home-range size of the animals increases with time, animals spend less time in average in "good" habitats.