SUPPLEMENTARY INFORMATION: A GENERALISED RANDOM ENCOUNTER MODEL FOR ESTIMATING ANIMAL DENSITY WITH REMOTE SENSOR DATA

Authors:

Tim C.D. Lucas^{1,2,3}†, Elizabeth A. Moorcroft^{1,4,5}†, Robin Freeman⁵, Marcus J. Rowcliffe⁵, Kate E. Iones^{2,5}

Addresses:

- 1 CoMPLEX, University College London, Physics Building, Gower Street, London, WC1E 6BT, UK
- 2 Centre for Biodiversity and Environment Research, Department of Genetics, Evolution and Environment, University College London, Gower Street, London, WC1E 6BT, UK
- 3 Department of Statistical Science, University College London, Gower Street, London, WC1E 6BT, UK
- 4 Department of Computer Science, University College London, Gower Street, London, WC1E 6BT, UK
- 5 Institute of Zoology, Zoological Society of London, Regents Park, London, NW1 4RY, UK † First authorship shared.

1

S1. Table of symbols

Symbol	Description	Units
θ	Sensor width	rad
α	Animal signal width	rad
x_i	Focal angle, $i \in \{1, 2, 3, 4\}$	rad
r	Detection distance	m
\bar{p}	Average profile width	m
p	A specific profile width	m
v	Velocity	$\mathrm{m}\mathrm{s}^{-1}$
t	Time	s
z	Number of detections	-
D	Animal density	m^{-2}
T	Step length	s
N	Number of steps per simulation	-
d	Distance moved in a time step	m
S	Probability of remaining stationary	-
\boldsymbol{A}	Maximum turning angle	rad

Table S1. List of symbols used to describe the gREM and simulations. '-' means the quantity has no units.

S2. SUPPLEMENTARY METHODS

S2.1. **Introduction.** These supplementary methods derive all the models used. For continuity, the gas model derivation is included here as well as in the main text. The calculation of all integrals use in the gREM is included in the Python script S3.

S2.2. **Gas model.** Following **?**, we derive the gas model where sensors can capture animals in any direction and animal signals are detectable from any direction ($\theta = 2\pi$ and $\alpha = 2\pi$). We assume that animals are in a homogeneous environment, and move in straight lines of random direction with velocity v. We allow that our stationary sensor can capture animals at a detection distance r and that if an animal moves within this detection zone they are captured with a probability of one, while animals outside the zone are never captured.

In order to derive animal density, we need to consider relative velocity from the reference frame of the animals. Conceptually, this requires us to imagine that all animals are stationary and randomly distributed in space, while the sensor moves with velocity v. If we calculate the area covered by the sensor during the survey period we can estimate the number of animals the sensor should capture. As a circle moving across a plane, the area covered by the sensor per unit time is 2rv. The number of expected captures, z, for a survey period of t, with an animal density of D is z = 2rvtD. To estimate the density, we rearrange to get D = z/2rvt.

S2.2.1. gREM derivations for different detection and signal widths. Different combinations of θ and α would be expected to occur (e.g., sensors have different detection widths and animals have different signal widths). For different combinations θ and α , the area covered per unit time is no longer given by 2rv. Instead of the size of the sensor detection zone having a diameter of 2r, the size changes with the approach angle between the sensor and the animal. For any given signal width and detector width and depending on the angle that the animal approaches the sensor, the width of the area within which an animal can be detected is called the profile, p. The size of the profile (averaged across all approach angles) is defined as the average profile \bar{p} . However, different combinations of θ and α need different equations to calculate \bar{p} . This \bar{p} is the only thing that changes

We have identified the parameter space for the combinations of θ and α for which the derivation of the equations are the same (defined as sub-models in the gREM) (Fig. S2.1). For example, the gas

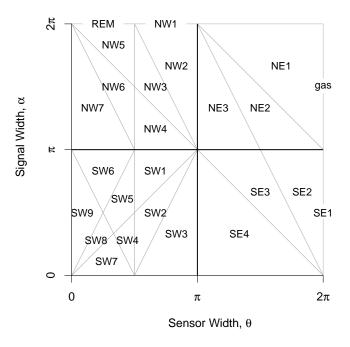


Figure S2.1. Locations where derivation of the average profile \bar{p} is the same for different combinations of sensor detection and animal signal widths. Symbols within each polygon refer to each gREM submodel named after their compass point, except for Gas and REM which highlight the position of these previously derived models within the gREM. Symbols on the edge of the plot are for submodels where $\alpha, \theta = 2\pi$

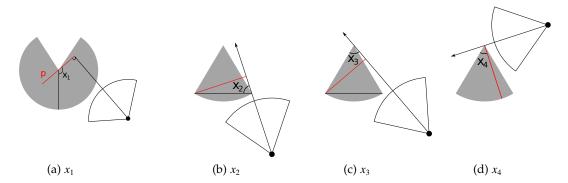


Figure S2.2. The location of the focal angles $x_{i\in[1,4]}$. x_1 is used in SE and NE models (including the gas model). $x_2 - x_4$ are used in NW and SW models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile p is shown with a red line. (a) Animal is directly approaching the sensor at $x_1 = \frac{\pi}{2}$. (b) Animal is directly approaching the sensor at $x_2 = \frac{\pi}{2}$. x_2 then decreases until the profile is perpendicular to the edge of the detection region. (c) When the profile is perpendicular to the edge of the detection region and the profile.

model becomes the simplest gREM sub-model (upper right in Fig. S2.1) and the REM from (?) is another gREM sub-model where $\theta < \pi/2$ and $\alpha = 2\pi$.

Models with $\theta = 2\pi$ are described first (the gas model described above and SE1). Then models with $\theta > \pi$ are described (NE then SE). Finally models with $\theta < \pi$ (NW then SW) are described.

S2.3. **Model SE1.** SE1 is very similar to the gas model except that because $\alpha \le \pi$ the profile width is no longer 2r but is instead limited by the width of the animal signal. We therefore get a profile width of $2r\sin(\alpha/2)$ instead.

$$\bar{p}_{\text{SE1}} = \frac{1}{\pi} \int_{\frac{\pi}{2}}^{\frac{3\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 \qquad \text{eqn S1}$$

$$\bar{p}_{\text{SE1}} = 2r \sin\left(\frac{\alpha}{2}\right)$$
 eqn S2

This profile is integrated over the interval $\left[\frac{\pi}{2}, \frac{3\pi}{2}\right]$ which is π radians of rotation starting with the animal moving directly towards the sensor (Fig. S2.2a).

S2.4. **Models NE1–3.** When the detection zone is not a circle, we have more complex profiles and need to explicitly write functions for the width of the profile for every approach angle. We then use these functions to find the average profile width \bar{p} for all approach angles by integrating across all 2π angles of approach and dividing by 2π .

There are three submodels within quadrant NE (Fig. S2.1). Note that NE1 covers the area $\alpha = 2\pi$ as well as the triangle below it as these two models are specified exactly the same, rather than happening to have equal results.

These models have up to five profiles.

- (1) The profile width starts, from $x_1 = \frac{\pi}{2}$ as 2r.
- (2) At $x_1 = \theta/2$, the right hand side of the profile cannot be r wide as the corner of the 'blind spot' limits its size to being $r \cos(x_1 \theta/2)$ wide (Fig. S2.3a).
- (3) The third profile is only found in NE3. If $\alpha < 4\pi 2\theta$, then at $x_1 = \theta/2 + \pi/2$, when the profile is perpendicular to the edge of the blind spot, the whole right side of the profile is invisible to the sensor (Fig. S2.3b). This gives a profile size of just r.
- (4) At some point, the sensor can detect animals once they have passed the blind spot giving a profile width of $r+r\cos(x_1+\theta/2)$ (Fig. S2.3c). From $x_1=\pi$, if the animal signal is wide enough to be detected in this area, this is the wider profile. This then defines the split between NE1

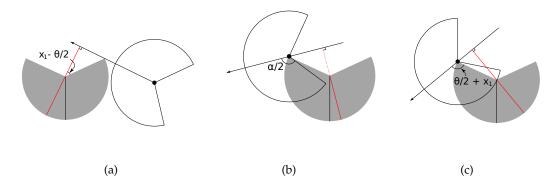


Figure S2.3. Three of the integrals in NE models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile p is shown with a red line. Dashed red lines indicate areas where animals cannot be detected. (a) The second integral in NE with width $r + r\cos(x_1 - \theta/2)$ (b) The third integral in NE3. $\alpha/2$ is labelled. As it is small, animals to the right of the detector cannot be detected. (c) After further rotation, $\alpha/2$ is now bigger than the angle shown and animals to the right of the detector can again be detected.

and NE2. In NE1, with $\alpha > 3\pi - \theta$, the animal signal is wide enough that at $x_1 = \pi$ the animal can immediately be detected past the blind spot and so this profile is used. In NE2, with $\alpha < 3\pi - \theta$, the latter profile is reached at $5\pi/2 - \theta/2 - \alpha/2$.

(5) Finally, common to all three models, at $x_1 = 2\pi - \theta/2$ the profile becomes a full 2*r* once again.

S2.4.1. *Model NE1*. Submodel NE1 exists within the area bounded by $\alpha \le 2\pi$, $\theta \le 2\pi$ and $\alpha \ge 3\pi - \theta$ (Fig. S2.1). It has four profiles; it does not include the r profile at $x_1 = \pi$ (profile described in point (3) in Section S2.4). Furthermore, θ is wide enough that the $r + r\cos(x_1 + \theta/2)$ profile starts at π . This then gives us

$$\bar{p}_{\text{NE1}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, \mathrm{d}x_1 + \int_{\frac{\theta}{2}}^{\pi} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, \mathrm{d}x_1 \right)$$

$$+ \int_{\pi}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, \mathrm{d}x_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, \mathrm{d}x_1$$

$$= \frac{r}{\pi} \left(\theta + 2 \sin\left(\frac{\theta}{2}\right)\right)$$
eqn S4

S2.4.2. Model NE2. Model NE2 is bounded by $\alpha \le 3\pi - \theta$, $\alpha \ge 4\pi - 2\theta$ and $\alpha \ge \pi$ (Fig. S2.1). It is the same as NE1 except that the third profile starts at $5\pi/2 - \theta/2 - \alpha/2$ instead of at π which is reflected in the different bounds in the second and third integral.

$$\bar{p}_{\text{NE2}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, \mathrm{d}x_1 + \int_{\frac{\theta}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, \mathrm{d}x_1 \right)$$

$$+ \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, \mathrm{d}x_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, \mathrm{d}x_1$$

$$= \exp \text{S5}$$

$$\bar{p}_{\text{NE2}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + \cos\left(\frac{\alpha}{2} + \theta\right)\right)$$
eqn S6

S2.4.3. *Model NE3*. Model NE3 is bound by $\alpha \le 4\pi - 2\theta$, $\alpha \ge \pi$ and $\theta \ge \pi$ (Fig. S2.1). It is the same as NE2 except that it contains the extra profile with width r (third integral).

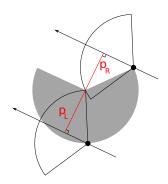


Figure S2.4. The second integral in SE. The right side of the profile (p_R) is limited by the size of the sensor region while the left side of the profile (p_L) is limited by the size of the signal width. The full profile has width $p = r \sin(\alpha/2) + r \cos(\theta/2 - x_1)$. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile p is shown with a red line.

$$\bar{p}_{\text{NE3}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, dx_1 + \int_{\frac{\theta}{2}}^{\frac{\theta}{2} + \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, dx_1 \right)$$

$$+ \int_{\frac{\theta}{2} + \frac{\pi}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \, dx_1 + \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, dx_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, dx_1$$

$$\bar{p}_{\text{NE3}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S8

- S2.5. **Models SE2-4.** Quadrant SE contains three submodels excluding SE1 (Fig. S2.1). The differences between these three models are similar to the differences between the models in NE. There are four possible profiles.
 - (1) As α is less than π the profile is smaller than 2r, even when the sensor width is a full diameter. The profile width starts as $2r \sin(\alpha/2)$.
 - (2) Similar to NE, at a certain point the blind spot of the sensor area limits the profile width on one side. This gives a profile width of $r \sin(\alpha/2) + r \cos(x_1 \theta/2)$ (Fig. S2.4).
 - (3) Also similar to NE, there can be a point where the right side of the profile is 0 giving a profile width of $r \sin(\alpha/2)$.
 - (4) If $\alpha \le 2\pi \theta$, then at $x_1 = \theta/2 + \pi/2 + \alpha/2$ the profile width becomes 0. This inequality distinguishes between SE3 and SE4.
 - (5) The third profile $r \sin(\alpha/2)$ starts at $\theta/2 + \pi/2$ while at $5\pi/2 \alpha/2 \theta/2$ the profile returns to size $2r \sin(\alpha/2)$. If $\theta/2 + \pi/2 \ge 5\pi/2 \alpha/2 \theta/2$ we go straight into the $2r \sin(\alpha/2)$ profile and miss the $r \sin(\alpha/2)$ profile. SE2 and SE3 are separated by this inequality which simplifies to $\alpha \le 4\pi 2\theta$.
- S2.5.1. *Model SE2*. SE2 is bounded by $\alpha \ge 4\pi 2\theta$, $\alpha \le \pi$ and $\theta \le 2\pi$ (Fig. S2.1). As $\alpha \ge 4\pi 2\theta$, there is no $r \sin(\alpha/2)$ profile. As $\alpha \le 4\pi 2\theta$, the profile returns to $2r \sin(\alpha/2)$ rather than going to 0. These integrals relate to profiles (1), (2) and (5) in Section S2.5.

$$\bar{p}_{SE2} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 + \int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) + r \cos\left(\frac{\theta}{2} - x_1\right) dx_1 + \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{3\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 \right)$$
 eqn S9
$$\bar{p}_{SE2} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + \cos\left(\frac{\alpha}{2} + \theta\right)\right)$$
 eqn S10

S2.5.2. *Model SE3.* SE3 is bounded by $4\pi - 2\theta \le \alpha \le 4\pi - 2\theta$ and $\alpha \le \pi$ (Fig. S2.1). Therefore there is a $r \sin(\alpha/2)$ profile but no 0r profile. This relates to profiles (1), (2), (3) and (5) above.

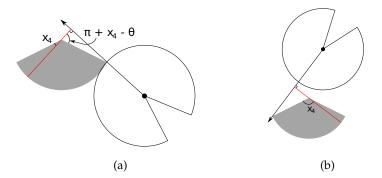


Figure S2.5. The second and fourth profiles of NW1. The left side of both profiles is of width r while the right side differs. (a) The right side of the profile is $r\cos(\pi + x_4 - \theta) = -r\cos(\theta - x_4)$ (b) The right side is $r\cos(\pi - x_4) = -r\cos x_4$ respectively. In both images the sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile p is shown with a red line.

$$\begin{split} \bar{p}_{\text{SE3}} &= \frac{1}{\pi} \left(\int\limits_{\frac{\pi}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) \mathrm{d}x_1 + \int\limits_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) + r \cos\left(\frac{\theta}{2} - x_1\right) \mathrm{d}x_1 \\ &+ \int\limits_{\frac{\theta}{2} + \frac{\pi}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) \mathrm{d}x_1 + \int\limits_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{3\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) \mathrm{d}x_1 \right) \end{aligned} \quad \text{eqn S11}$$

$$\bar{p}_{\text{SE3}} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right) \quad \text{eqn S12}$$

S2.5.3. *Model SE4*. Finally SE4 is bounded by $\alpha \le 4\pi - 2\theta$, $\alpha \le \pi$ and $\theta \le \pi$ (Fig. S2.1). It is the same as SE3 except that the profile becomes 0 rather than returning to $2r\sin(\alpha/2)$. This relates to profiles (1), (2), (3) and (4) above though profile (4) with width 0 is not shown.

$$\bar{p}_{\text{SE4}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 + \int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) + r \cos\left(\frac{\theta}{2} - x_1\right) dx_1 + \int_{\frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\alpha}{2} + \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_1 \right) \quad \text{eqn S13}$$

$$\bar{p}_{\text{SE4}} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right) \quad \text{eqn S14}$$

S2.6. **Model NW1.** NW1 is the first model with $\theta < \pi$. Whereas previously the focal angle has always been x_1 , we now use different focal angles. x_2 and x_3 correspond to y_1 and y_2 in (?) while x_4 is new. They are described in Fig. S2.2b–d.

There are five different profiles in NW1.

- (1) x_2 has an interval of $[\pi/2, \theta/2]$ which is from the angle of approach being directly towards the sensor until the profile is parallel to the left hand radius of the sensor sector (Fig. S2.2b). During this interval the profile width is $2r\sin(\theta/2)\sin(x_2)$ which is calculated using the equation for the length of a chord . Note that while rotating anti-clockwise (as usual) x_2 decreases in size.
- (2) From here, we examine focal angle x_4 (note that x_3 is used in later models, but is not relevant here.) The left side of the profile is a full radius while the right side is limited to $-r\cos(x_4 \theta)$ (Fig. S2.5a).
- (3) At $x_4 = \theta \pi/2$, the profile is perpendicular to the edge of the sensor area. Here, the right side of the profile is 0r giving a profile size of r.

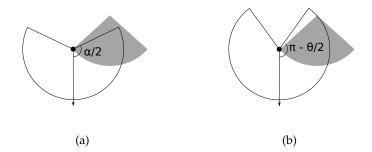


Figure S2.6. Profile sizes when an animal approaches from behind in models NW2–4. If α is relatively large, animals can be detected when approaching from behind. Otherwise animals cannot be detected. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. (a) If $\alpha/2$ is less than $\pi - \theta/2$, as is the case here, then the width of the profile when an animal approaches directly from behind is zero. (b) If $\alpha/2 > \pi - \theta/2$ the profile width from behind is $2r \sin(\theta/2) \sin(x_2)$.

- (4) When $x_4 = \pi/2$ the angle of approach is from behind the sensor, but we can once again be detected on the right side of the sensor (Fig. S2.5b). Therefore the width of the profile is $r r\cos(x_4)$.
- (5) Finally, we have the x_2 profile, but from behind.

$$\bar{p}_{\text{NW1}} = \frac{1}{\pi} \left(\int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{0}^{\theta - \frac{\pi}{2}} r - r \cos\left(-x_4 + \theta\right) \, dx_4 \right)$$

$$+ \int_{\theta - \frac{\pi}{2}}^{\frac{\pi}{2}} r \, dx_4 + \int_{\frac{\pi}{2}}^{\theta} r - r \cos(x_4) \, dx_4 + \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2$$
eqn S15
$$\bar{p}_{\text{NW1}} = \frac{r}{\pi} (\theta + 2)$$
eqn S16

- S2.7. **Models NW2–4.** The models NW2–4 have the five potential profiles in NW1 but not all profiles occur in each model, and the angle at which transitions occur are different. Furthermore, there is one extra profile possible.
 - (1) When approaching the sensor from behind, there is a period where the profile is *r* wide as in NW1 profile (3).
 - (2) At some point after profile (1) animals to the right of the sensor can be detected again. If this occurs in the x_4 region, the profile width becomes $r r \cos(x_4)$ as in NW1.
 - (3) However, as α is now less than 2π , animals to the right of the sensor may be undetectable until we are in the second x_2 region. In this case, when we first enter the second x_2 region, the profile has a width of $r\cos(x_2 \theta/2)$. This occurs only if $\alpha \le 3\pi 2\theta$. This inequality is found by noting that animals to the right of the sensor can be detected again at $x_4 = 3\pi/2 \alpha$ but the x_2 region starts at $x_4 = \theta$. The new profile in x_2 will only occur if $\theta < 3\pi/2 \alpha/2$ which is rearranged to find the inequality above. This defines the boundary between NW2 and NW3.
 - (4) As $\alpha \le 2\pi$ it is possible that when the angle of approach is from directly behind the sensor the animal will not be detected at all. This is the case if $\alpha/2 \le \pi \theta/2$ (Fig. S2.6). This inequality (simplified as $\alpha \le 2\pi \theta$) defines the boundary between NW3 and NW4.
- S2.7.1. *Model NW2*. NW2 is bounded by $\alpha \ge 3\pi 2\theta$, $\alpha \le 2\pi$ and $\theta \le \pi$ (Fig. S2.1). NW2 has all five profiles as found in NW1. However, the change from the r profile (third integral) to the $r r\cos(x_4)$ profile (fourth integral) occurs at $x_4 = 3\pi/2 \alpha/2$ instead of at $x_4 = \theta$.

$$\bar{p}_{\text{NW2}} = \frac{1}{\pi} \left(\int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{0}^{\theta - \frac{\pi}{2}} r - r \cos\left(-x_4 + \theta\right) \, dx_4 \right)$$

$$+ \int_{\theta - \frac{\pi}{2}}^{\frac{3\pi}{2} - \frac{\alpha}{2}} r \, dx_4 + \int_{\frac{3\pi}{2} - \frac{\alpha}{2}}^{\theta} r - r \cos(x_4) \, dx_4 + \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2$$

$$\bar{p}_{\text{NW2}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S18

S2.7.2. *Model NW3.* NW3 is bounded by $\alpha \le 3\pi - 2\theta$, $\alpha \ge 2\pi - \theta$ and $\theta \ge \pi/2$ (Fig. S2.1).

NW3 does not have the fourth integral from NW2 as animals are not detectable to the right of the sensor until after the x_4 region has ended and the x_2 region has begun. Therefore the second x_4 integral has an upper limit of θ and the profile after has a width of $r\cos(x_2-\theta/2)$ and is integrated with respect to x_2 . The final integral starts at $x_4 = 3\pi/2 - \alpha/2 - \theta/2$ and has the full width of $2r\sin(x_2)\sin(\theta/2)$.

$$\bar{p}_{\text{NW3}} = \frac{1}{\pi} \left(\int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{0}^{\theta - \frac{\pi}{2}} r - r \cos\left(-x_4 + \theta\right) \, dx_4 \right)$$

$$+ \int_{\theta - \frac{\pi}{2}}^{\theta} r \, dx_4 + \int_{\frac{\theta}{2}}^{\frac{3\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) \, dx_2 + \int_{\frac{3\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2$$

$$\bar{p}_{\text{NW3}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S20

S2.7.3. *Model NW4*. Finally, NW4 is bounded by $\alpha \ge \pi$, $\theta \ge \pi/2$ and $\alpha \le 2\pi - \theta$ (Fig. S2.1). NW4 is the same as NW3 except that the final profile width is zero and this profile is reached at $\alpha/2 + \theta/2 - \pi/2$.

$$\bar{p}_{\text{NW4}} = \frac{1}{\pi} \left(\int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_2\right) dx_2 + \int_{0}^{\theta - \frac{\pi}{2}} r - r \cos\left(-x_4 + \theta\right) dx_4 \right)$$

$$+ \int_{\theta - \frac{\pi}{2}}^{\theta} r dx_4 + \int_{\frac{\theta}{2}}^{\frac{\alpha}{2} + \frac{\theta}{2} - \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) dx_2$$

$$= \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S22

S2.8. **Model REM.** REM is the model from (?). It has $\alpha = 2\pi$ and $\theta \le \pi/2$ (Fig. S2.1). It has three profile widths, two of which are repeated, once as the animal approaches from in front of the sensor and once as the animal approaches from behind the sensor.

- (1) Starting with an approach direction of directly towards the sensor, and examining focal angle x_2 , the profile width is $2r\sin(x_2)\sin(\theta/2)$.
- (2) When the profile is perpendicular to the radius on the right hand of the sector sensor region, we instead examine x_3 where the profile width is $r \sin(x_3)$.
- (3) At $x_3 = \pi/2$ the profile becomes simply r and this continues for θ radians of x_4 .
- (4) The x_3 profile is then repeated with an approach direction from behind the sensor.
- (5) Finally the x_2 profile is repeated, again with an approach direction from behind the sensor.

$$\bar{p}_{\text{REM}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3 + \int_{\theta}^{\theta} r \, dx_4 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 \right)$$

$$\bar{p}_{\text{REM}} = \frac{r}{\pi} (\theta + 2)$$
eqn S24

S2.9. **Models NW5–7.** In the models NW5–7, the sensor has $\theta \le \pi/2$ as in the REM. As $\alpha \ge \pi$ a lot of the profiles are similar to the REM. Specifically, the first three profiles are always the same as the first three profiles of the REM. This is because when an animal is moving towards the sensor, the $\alpha \ge \pi$ signal is no different to a 2π signal. However, when approaching the sensor from behind, things are slightly different. The animal can only be detected by the sensor if the signal width is large enough that it can be detected once it has passed the sensor.

- (1) Starting with an approach direction of directly towards the sensor, and examining focal angle x_2 , the profile width is $2r\sin(x_2)\sin(\theta/2)$.
- (2) When the profile is perpendicular to the radius edge of the sector sensor region, we instead examine x_3 where the profile width is $r \sin(x_3)$.
- (3) At $x_3 = \pi/2$ the profile becomes simply r and this continues for θ radians of x_4 .
- (4) If $\alpha \le 2\pi + 2\theta$, the animal becomes undetectable during this profile when x_3 has decreased in size to $\pi \alpha/2$. This inequality marks the boundary between NW7 and NW6.
- (5) If instead $\alpha \ge 2\pi + 2\theta$ then the animal does not become undetectable during the x_3 focal angle. Instead the profile has width greater than zero for the whole of the x_3 angle. The x_2 profile starts with width $r\cos(x_2 \theta/2)$ as only animals approaching to the left of the sensor are detectable.
- (6) During this second x_2 profile the signal width needed for animals to be detected to the left of the detector is increasing while the angle needed for animals to be detected to the right of the detector is decreasing. Therefore, either the left side becomes undetectable, making both sides undetectable (this occurs if $\alpha \le 2\pi \theta$ as in NW6)
- (7) or the right becomes detectable (if $\alpha \ge 2\pi \theta$ as in NW5), making both sides detectable and giving a profile width of $2r\sin(x_2)\sin(\theta/2)$.

S2.9.1. *Model NW5.* NW5 is bounded by $\alpha \ge 2\pi - \theta$, $\alpha \le 2\pi$ and $\theta \le \pi/2$ (Fig. S2.1).

It is the same as REM except that it includes the extra profile in x_2 (the fifth integral) where only animals approaching to the left of the profile are detected.

$$\bar{p}_{\text{NW5}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\theta}^{\frac{\pi}{2}} r \sin\left(x_{3}\right) dx_{3} + \int_{0}^{\theta} r dx_{4} \right) + \int_{\theta}^{\frac{\pi}{2}} r \sin\left(x_{3}\right) dx_{3} + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{3\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \cos\left(\frac{\theta}{2} - x_{2}\right) dx_{2} + \int_{\frac{3\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2}$$

$$\bar{p}_{\text{NW5}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S26

S2.9.2. *Model NW6.* NW6 is bounded by $\alpha \le 2\pi - \theta$, $\alpha \ge 2\pi + 2\theta$ and $\theta \le \pi/2$ (Fig. S2.1).

NW6 is the same NW5 except that as $\alpha \le 2\pi - \theta$, animals that approach from directly behind the detector are not detected. Therefore at $x_2 = \alpha/2 + \theta/2 - \pi/2$ the profile width goes to zero and therefore the last integral in NW5 is not included.

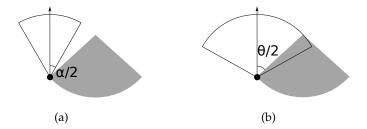


Figure S2.7. The first profile in SW models is limited by either α or β depending on whether $\alpha < \beta$. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. (a) As $\alpha/2 < \theta/2$ the profile width is limited by the signal width rather than the sensor region. The profile width is $2r\sin{(\alpha/2)}$ (b) As $\alpha/2 > \theta/2$ the profile width is limited by the sensor region, not the signal width. The profile width is $2r\sin{(\theta/2)}\sin(x_2)$.

$$\bar{p}_{\text{NW6}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3 \right)$$

$$+ \int_{0}^{\theta} r \, dx_4 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\alpha}{2} + \frac{\theta}{2} - \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) dx_2$$

$$= \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S28

S2.9.3. *Model NW7*. NW7 is bounded by $\alpha \ge 2\pi + 2\theta$, $\alpha \ge \pi$ and $\theta \ge 0$ (Fig. S2.1).

It is similar to NW6 but does not include the last integral as during the x_3 profile, at $x_3 = \pi - \alpha/2$ the signal width is too small for any animals to be detected, so the profile width goes to zero.

$$\bar{p}_{\text{NW7}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) \, dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3 \right)$$

$$+ \int_{0}^{\theta} r \, dx_4 + \int_{\pi - \frac{\alpha}{2}}^{\frac{\pi}{2}} r \sin(x_3) \, dx_3$$

$$\bar{p}_{\text{NW7}} = \frac{r}{\pi} \left(\theta - \cos\left(\frac{\alpha}{2}\right) + 1 \right)$$
eqn S29

S2.10. **Model SW1–3.** The models in SW1–3 are described with the two focal angles used in models NW2–4, x_2 and x_4 . As $\alpha \le \pi$ an animal can never be detected if it is approaching the detector from behind. This makes these models simpler in that they go through the x_2 and x_4 profiles only once each

There are five potential profile sizes.

- (1) At the beginning of x_2 , with an approach direction directly towards the sensor, the parameter that limits the width of the profile can either be the sensor width, in which case the profile width is $2r \sin(\theta/2) \sin(x_2)$.
- (2) Or the signal width can be the limiting parameter, in which case the profile width is instead $2r\sin(\alpha/2)$ (Fig. S2.7)
- (3) The next potential profile in x_2 has a width of $r \sin(\alpha/2) r \cos(x_2 + \theta/2)$ as the right side of the profile is limited by the width of the sensor region while the left side is limited by the signal width. However, the angle at which the profile starts depends on whether the first profile was 1) or 2) above. If the first profile is profile 1) then the profile is limited on both sides by the sensor region and then the left side of the profile becomes limited by the signal width. This happens at $x_2 = \pi/2 \alpha/2 + \theta/2$. If however the first profile was 2) then the first profile is

limited by the signal width. We move into the new profile when the right side of the profile becomes limited by the sensor region. This occurs at $x_2 = \pi/2 + \alpha/2 - \theta/2$.

- (4) In the x_4 region the left side of the profile is always $r \sin(\alpha/2)$ while the right side is either 0, giving a profile of $r \sin(\alpha/2)$.
- (5) Or limited by the sensor giving a profile of size $r \sin(\alpha/2) r \cos(x_4 \theta)$.

S2.10.1. *Model SW1*. SW1 is bounded by $\alpha \ge \theta$, $\alpha \le \pi$ and $\theta \le \pi$ (Fig. S2.1).

As α is large the first profile is limited by the size of the sensor region giving it a width of $2r\sin(\theta/2)\sin(x_2)$. It is the only one of the three SW models to start in this way. Later on, still with x_2 as the focal angle the left side of the profile does become limited by the signal width. So at $x_2 = \pi/2 - \alpha/2 + \theta/2$ the profile width becomes $r\sin(\alpha/2) - r\cos(x_2 + \theta/2)$.

As we enter the x_4 region, the profile remains limited by the signal on the left and by the sensor on the right, giving a profile width of $r \sin(\alpha/2) - r \cos(x_4 - \theta)$. Finally, at $x_4 = \theta - \pi/2$ the right side of the profile becomes zero and the profile is width is $r \sin(\alpha/2)$.

$$\bar{p}_{SW1} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\frac{\theta}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_{2}\right) dx_{2} \right)$$

$$+ \int_{0}^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\theta - x_{4}\right) dx_{4} + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_{4}$$

$$= \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S32

S2.10.2. *Model SW2*. SW2 is bounded by $\theta \ge \pi/2$, $\alpha \le \theta$ and $\alpha \ge 2\theta - \pi$ (Fig. S2.1).

SW2 is largely similar to SW1. However, as $\alpha \le \theta$ the first profile is limited by α and not by the detection region. Therefore the first profile has width $2r\sin(\alpha/2)$. This also means the transition to the second profile occurs at $x_2 = \pi/2 + \alpha/2 - \theta/2$ instead of $x_2 = \pi/2 - \alpha/2 + \theta/2$.

$$\bar{p}_{SW2} = \frac{1}{\pi} \left(\int_{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{\frac{\theta}{2}}^{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right)$$

$$+ \int_{0}^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\theta - x_4\right) dx_4 + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4$$

$$= \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S34

S2.10.3. *Model SW3*. SW3 is bounded by $\alpha \le 2\theta - \pi$ and $\theta \le \pi$ (Fig. S2.1).

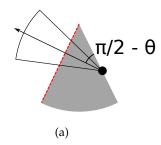
SW3 is similar to SW2 except that the profile does not become limited by sensor at all during the the x_4 regions. Therefore, at $x_4 = 0$ the profile is still of width $2r\sin(\alpha/2)$. Only at $x_4 = \theta - \pi/2 - \alpha/2$ does the profile become limited on the right by the sensor region.

$$\bar{p}_{\text{SW3}} = \frac{1}{\pi} \left(\int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{0}^{-\frac{\pi}{2} + \theta - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_4 \right)$$

$$+ \int_{-\frac{\pi}{2} + \theta - \frac{\alpha}{2}}^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\theta - x_4\right) dx_4 + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4$$

$$= \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S36

S2.11. **Model SW4–9.** As $\alpha < \pi$, animals approaching the sensor from behind can never be detected, so unlike REM, the second x_2 and x_3 profiles are always zero. The six models are split by three inequalities that relate to the models as follows.



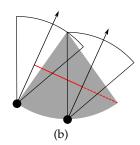


Figure S2.8. Description of two profiles in SW models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile p is shown with a red line. Dashed red lines indicate areas where animals cannot be detected. (a) At $x_4 = 0$, if $\alpha/2 < \pi/2 - \theta$ then $\alpha/2$ is too small for an animal to be detected at all during the x_4 profile (shown with dashed red.) This inequality simplifies to $\alpha < \pi - 2\theta$. (b) The right of the profile is limited by the signal width, not the sensor. On the left, the profile is limited by the sensor and not the signal. Overall the profile width is $r \sin(\alpha/2) - r \cos(x_2 + \theta/2)$.

- (1) Models with $\alpha \le \pi 2\theta$ have no x_4 profile. This is because at $x_4 = 0$, the signal width is already too small to be detected as can be seen in Fig. S2.8a where $\alpha/2 < \pi/2 \theta$ which simplifies to give the previous inequality.
- (2) Models with $\alpha \le \theta$ are limited by α in the first, x_2 region (Fig. S2.7), rather than being limited by θ . Therefore this first profile is of width $2r\sin(\alpha/2)$ rather than $2r\sin(\theta/2)\sin(x_2)$.
- (3) Finally, models with $\alpha \le 2\theta$ have a second profile in x_2 where to one side of the sensor α is the limiting factor of profile width, while on the other side θ is (Fig. S2.8b). This gives a width of $r \sin(\alpha/2) r \cos(x_2 + \theta/2)$. This profile does not occur in models with $\alpha \ge 2\theta$.

S2.11.1. *Model SW4*. SW4 is bounded by $\alpha \le \theta$, $\alpha \ge \pi - 2\theta$ and $\theta \le \pi/2$ (Fig. S2.1). Therefore it does contain a x_4 profile, starts with an α limited profile and does contain the $r \sin(\alpha/2) - r \cos(x_2 + \theta/2)$ profile in x_2 .

$$\bar{p}_{\text{SW4}} = \frac{1}{\pi} \left(\int_{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right)$$

$$+ \int_{\theta}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_3 + \int_{0}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4$$

$$= \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S38

S2.11.2. *Model SW5*. SW5 is the only model with a tetrahedral bounding region. It is bounded by $\alpha \ge \theta$, $\alpha \ge \pi - 2\theta$, $\alpha \le 2\theta$ and $\theta \le \pi/2$ (Fig. S2.1). Therefore it does contain a x_4 profile, but starts with a θ limited profile. It does contain the $r \sin(\alpha/2) - r \cos(x_2 + \theta/2)$ profile in x_2 .

$$\bar{p}_{\text{SW5}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_{2}\right) dx_{2} \right)$$

$$+ \int_{\theta}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_{3} + \int_{0}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_{4}$$

$$= \exp \text{S39}$$

$$\bar{p}_{\text{SW5}} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$

$$= \exp \text{S40}$$

S2.11.3. *Model SW6*. SW6 is bounded by $\alpha \ge \pi - 2\theta$, $\alpha \ge 2\theta$ and $\alpha \le \pi$ (Fig. S2.1). It starts with a θ limited profile and has a x_4 profile. However, it does not contain the $r \sin(\alpha/2) - r \cos(x_2 + \theta/2)$ profile.

$$\bar{p}_{SW6} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\theta}^{\frac{\alpha}{2}} r \sin\left(x_{3}\right) dx_{3} \right)$$

$$+ \int_{\frac{\pi}{2}}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_{3} + \int_{0}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_{4}$$

$$= \exp\left(\frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$

$$= \exp\left(\frac{3\pi}{2}\right)$$

$$= \exp\left(\frac{3\pi}{2}\right) + \exp\left(\frac{3\pi}{2}\right) + \exp\left(\frac{3\pi}{2}\right)$$

$$= \exp\left(\frac{3\pi}{2}\right)$$

$$= \exp\left(\frac{3\pi}{2}\right) + \exp\left(\frac{3\pi}{2}\right)$$

$$= \exp\left(\frac{3\pi}{2}\right$$

S2.11.4. *Model SW7*. SW7 is bounded by $\alpha \le \pi - 2\theta$, $\alpha \le \theta$ and $\alpha < 0$ (Fig. S2.1). Therefore it does not contain a x_4 profile. It starts with an α limited profile and contains the $r \sin(\alpha/2) - r \cos(x_2 + \theta/2)$ profile in x_2 .

$$\bar{p}_{SW7} = \frac{1}{\pi} \left(\int_{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 + \int_{\theta}^{\frac{\alpha}{2} + \theta} r \sin\left(\frac{\alpha}{2}\right) dx_3 \right)$$
eqn S43
$$\bar{p}_{SW7} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right)$$
eqn S44

S2.11.5. *Model SW8*. SW8 is bounded by $\alpha \le \pi - 2\theta$, $\alpha \ge \theta$ and $\alpha \le 2\theta$ (Fig. S2.1). It starts with a θ limited profile. It does contain the $r\sin(\alpha/2) - r\cos(x_2 + \theta/2)$ profile in x_2 but does not have a x_4 profile.

$$\bar{p}_{\text{SW8}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_{2}\right) dx_{2} + \int_{\theta}^{\frac{\alpha}{2} + \theta} r \sin\left(\frac{\alpha}{2}\right) dx_{3} \right) \text{ eqn S45}$$

$$\bar{p}_{\text{SW8}} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1\right) \text{ eqn S46}$$

S2.11.6. *Model SW9*. Finally, SW9, the last model, is bounded by y $\alpha \le \pi - 2\theta$, $\alpha \ge 2\theta$ and $\theta \ge 0$ (Fig. S2.1). Therefore it starts with a θ limited profile. However it does not contain the extra x_2 profile nor a x_4 profile.

$$\bar{p}_{\text{SW9}} = \frac{1}{\pi} \left(\int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin\left(x_{2}\right) dx_{2} + \int_{\theta}^{\frac{\alpha}{2}} r \sin\left(x_{3}\right) dx_{3} + \int_{\frac{\alpha}{2}}^{\frac{\alpha}{2} + \theta} r \sin\left(\frac{\alpha}{2}\right) dx_{3} \right)$$
eqn S47
$$\bar{p}_{\text{SW9}} = \frac{r}{\pi} \left(\theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right)$$
eqn S48

S3. SUPPLEMENTARY SCRIPT: SYMBOLIC ALGEBRA PYTHON SCRIPT

This script uses the SymPy package (?), a computer algebra system to calculate the equations for *p* in the various models and to perform unit checks on the results.

The script can also be found in the plain text file lucas_et_al_S3.py. It will also be hosted and given a DOI on Github at the time of publication (https://github.com/timcdlucas/lucasMoorcroftManuscript/tree/postPeerReview).

```
...
   Supplementary Python script from
   A generalised random encounter model for estimating animal density with remote sensor data
   Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
   Systematic analysis of REM models
   Tim Lucas
   01/10/13
   This script contains:
   1. The integration of all models (lines 24 - 603)
   2. Some tests that the models are correct (606 - 738)
   3. A python function to calculate \bar{p} given any parameters (743 - 763)
   4. Creation of a plot of \bar{p} over all parameter space (767 - 794)
   5. Code to create supplementaryS4.R, an R implementation of the model (797 - 873)
   from sympy import *
   import numpy as np
   import matplotlib.pyplot as pl
   from datetime import datetime
   import os as os
   # Set working directory
   os.chdir('supplementary-material')
36
37
   # Use LaTeX printing
   from sympy import init_printing ;
   init_printing()
     Make LaTeX output white. Because I use a dark theme
   init_printing(forecolor="White")
   # Load symbols used for symbolic maths
   t, a, r, x2, x3, x4, x1 = \frac{\text{symbols}(\text{'theta alpha r x}_2 \text{ x}_3 \text{ x}_4 \text{ x}_1'), positive=True) r1 = {r:1} # useful for lots of checks
48
49
   # Define functions
   # Calculate the final profile averaged over pi.
   def calcModel(model):
          x = pi**-1 * sum( [integrate(m[0], m[1:]) for m in model]).simplify().trigsimp()
   # Do the replacements fit within the area defined by the conditions?
   def confirmReplacements(conds, reps):
           if not all([c.subs(reps) for c in eval(conds)]):
                    print('reps' + conds[4:] + ' incorrect')
59
   # is average profile in range Or-2r?
   print('Total ' + prof + ' not in 0, 2r')
   # Are the individuals integrals >0r
   def intsPositive(model, reps):
67
           m = eval(model)
           for i in range(len(m)):
    if not integrate(m[i][0], m[i][1:]).subs(dict(reps, **r1)) > 0:
        print('Integral ' + str(i+1) + ' in ' + model + ' is negative')
   # Are the individual averaged integrals between 0 and 2r
   def intsRange(model, reps):
           m = eval(model)
```

```
75
             for i in range(len(m)):
                     if not 0 <= (integrate(m[i][0], m[i][1:])/(m[i][3]-m[i][2])).subs(dict(reps, **rl)) <=</pre>
                         2:
                             print('Integral ' + str(i+1) + ' in ' + model + ' has averaged integral outside
 78
     # Are the bounds the correct way around
 80
    def checkBounds(model, reps):
            m = eval(model)
             for i in range(len(m)):
                     84
                                  upper bounds')
 8.5
 86
    # create latex strings with the 1) the integral equation that defines it and 2) the final calculated
         model.
 87
    # There's some if statements to split longer equations on two lines and get +s in the right place.
    def parseLaTeX(prof):
    m = eval('m' + prof[1:])
 88
 89
            f = open('/latexFiles/'+prof+'.tex', 'w')
            f.write('\begin{align}\n \ \bar{p}_{(\text{\tiny}' + prof[1:] + '})} = & \frac{1}{\pi} \ \label{eq:f.write} \ \
 92
                  (\;\;')
 93
            for i in range(len(m)):
        # Roughly try and prevent expressions beginning with minus signs. if latex(m[i][2])[0]=='-':
 94
 95
 96
97
          o1 = 'rev-lex'
        else:
          o1 = 'lex'
 99
        if latex(m[i][3])[0]=='-':
        o2 = 'rev-lex'
else:
          o2 = 'lex'
104
        if latex(m[i][0])[0]=='-':
106
107
          o3 = 'rev-lex'
        else:
108
          o3 = 'lex'
109
         if latex(m[i][1])[0]=='-':
          o4 = 'rev-lex'
        else:
          04 = 'lex'
114
                     f.write('\int\limits_{'+latex(m[i][2], order=o1)+'}^{'+latex(m[i][3], order=o2)+'}'+
                     latex(m[i][0], order=o3)+'\;\mathrm{d}' +latex(m[i][1], order=o4))
if len(m)>3 and i==(len(m)/2)-1:
116
117
                             f.write( '\\right.\\notag\\\\n &\left.' )
                     if i<len(m)-1:</pre>
118
119
                             f.write('+')
            f.write('\\right)\label{' + prof + 'Def}\\\\n ')
f.write('\\bar{p}_{\\text{\\tiny{' + prof[1:] + '}}} =& ' + latex(eval(prof)) + '\label{' + prof[1:] + '}}
                 prof + 'Sln}\n\\end{align}')
             f.close()
122
    # Apply all checks.
126 def allChecks(prof):
            model = 'm' + prof[1:]
reps = eval('rep' + prof[1:])
conds = 'cond' + prof[1:]
128
129
            confirmReplacements(conds, reps)
            profileRange(prof, reps)
131
            intsPositive (model, reps)
             intsRange(model, reps)
134
             checkBounds (model, reps)
136
137
     ### Define and solve all models
     138
140
    # NE1 animal: a = 2*pi. sensor: t > pi, a > 3pi - t
141
    mNE1 = [2*r,
142
                                    x1, pi/2, t/2
              [2*r,

[r + r*cos(x1 - t/2), x1, t/2, pi ],

[r + r*cos(x1 + t/2), x1, pi, 2*pi-t/2],

[2*r, x1, 2*pi-t/2, 3*pi/2]]
143
144
146
147
    # Replacement values in range
148
    repNE1 = {t:3*pi/2, a:2*pi}
149
     # Define conditions for model
    condNE1 = [pi  <= t, a >= 3*pi - t]
    # Calculate model, run checks, write output.
154 pNE1 = calcModel(mNE1)
```

```
155 allChecks('pNE1')
156
     parseLaTeX('pNE1')
159
     \# NE2 animal: a > pi. sensor: t > pi Condition: a < 3pi - t, a > 4pi - 2t \#
160
                [2*r, x1, pi/2, t/2 ],

[r + r*cos(x1 - t/2), x1, t/2, 5*pi/2 - t/2 - a/2 ],

[r + r*cos(x1 + t/2), x1, 5*pi/2 - t/2 - a/2, 2*pi-

[2*r, x1, 2*pi-t/2, 3*pi/2 ] ]
     mNE2 = [ [2*r,
                                                                          2*pi-t/2 ],
165
166 # Replacement values in range
167 repNE2 = {t:5*pi/3, a:4*pi/3-0.1}
168
     # Define conditions for model
     condNE2 = [pi <= t, a >= pi, a <= 3*pi - t, a >= 4*pi - 2*t]
# Calculate model, run
pNE2 = calcModel(mNE2)
     # Calculate model, run checks, write output.
     parseLaTeX('pNE2')
178
     \# NE3 animal: a > pi. sensor: t > pi Condition: a < 4pi - 2t \#
               [2*r, x1, pi/2, t/2],

[r + r*cos(x1 - t/2), x1, t/2, t/2 + pi/2],

[r , x1, t/2 + pi/2, 5*pi/2 - t/2 - a/2],

[r + r*cos(x1 + t/2), x1, 5*pi/2 - t/2 - a/2, 2*pi-t/2],
180
     mNE3 = [2*r]
181
183
                                           x1, 2*pi-t/2, 3*pi/2 ] ]
185
     # Replacement values in range
186
187
     repNE3 = \{t:5*pi/4-0.1, a:3*pi/2\}
     # Define conditions for model
190 condNE3 = [pi <= t, a >= pi, a <= 4*pi - 2*t]
191
192 # Calculate model, run
193 pNE3 = calcModel(mNE3)
     # Calculate model, run checks, write output.
     allChecks('pNE3')
parseLaTeX('pNE3')
194
195
196
197
     # NW1 animal: a = 2*pi. sensor: pi/2 \le t \le pi
199
200 mNW1 = [ [2*r*sin(t/2)*sin(x2), x2, t/2,
                                         x4, 0, t - j
x4, t - pi/2, pi/2
               [r - r*cos(x4 - t),
                                                              t - pi/2 ],
202
               [r,
[r - r*cos(x4),
                                                                    1,
                                             x4, pi/2,
203
              [r - r*\cos(x4), 	 x4, p1/2]

[2*r*\sin(t/2)*\sin(x2), 	 x2, t/2,
                                                              pi/2
204
2.05
206 # Replacement value
207 repNW1 = {t:3*pi/4}
     # Replacement values in range
208
209
     # Define conditions for model
210
     condNW1 = [pi/2 \le t, t \le pi]
211
     # Calculate model, run checks, write output.
213
     pNW1 = calcModel(mNW1)
214
     allChecks('pNW1')
215
216
217
218
     parseLaTeX('pNW1')
219
     # NW2 animal: a > pi. Sensor: pi/2 <= t <= pi. Condition: a > 2pi - t
221
                                                                   t - pi/2
222
223
     mNW2 = [ [2*r*sin(t/2)*sin(x2), x2, t/2,
                 [r - r*\cos(x4 - t), x4, 0, t - pi/2],

[r, x4, t - pi/2, 3*pi/2 - a/2],
224
                 [r - r*\cos(x4),
                                            x4, 3*pi/2 - a/2, t
                 [2*r*sin(t/2)*sin(x2), x2, t/2,
     repNW2 = {t:3*pi/4, a:15*pi/8} # Replacement values in range
229
230
231
     # Define conditions for model
     condNW2 = [a > pi, pi/2 <= t, t <= pi, a >= 3*pi - 2*t]
234 # Calculate moder, run.
235 pNW2 = calcModel(mNW2)
     # Calculate model, run checks, write output.
236
237
     allChecks('pNW2')
parseLaTeX('pNW2')
241 # NW3 animal: a > pi. Sensor: pi/2 <= t <= pi. Cond: 2pi - t < a < 3pi - 2t #
```

```
242
243
    mNW3 = [ [2*r*sin(t/2)*sin(x2), x2, t/2,
                                                             pi/2
                                                                               ],
             [r - r*\cos(x4 - t), x4, 0, [r, x4, t - pi/2,
                                                             t - pi/2
                                                                               ],
245
                                    x2, t/2,
                                                             3*pi/2 - a/2 - t/2],
246
             [r*cos(x2 - t/2),
247
             [2*r*sin(t/2)*sin(x2), x2, 3*pi/2 - a/2 - t/2, pi/2]
248
249
    repNW3 = {t:5*pi/8, a:6*pi/4} # Replacement values in range
    # Define conditions for model
    condNW3 = [a > pi, pi/2 \le t, t \le pi, 2*pi - t \le a, a \le 3*pi - 2*t]
254
255
    # Calculate model, run checks, write output.
256 pNW3 = calcModel(mNW3)
257
    allChecks('pNW3')
258
    parseLaTeX('pNW3')
259
260
261
262
    \# NW4 animal: a > pi. Sensor: pi/2 <= t <= pi. Condition: a <= 2pi - t
263
264
    mNW4 = [ [2*r*sin(t/2)*sin(x2), x2, t/2, pi/2],
             265
266
267
268
    repNW4 = {t:3*pi/4, a:9*pi/8} # Replacement values in range
270
271
272
    # Define conditions for model
    condNW4 = [a > pi, pi/2 \le t, t \le pi, a \le 2*pi - t]
273
    # Calculate model, run checks, write output.
275 pNW4 = calcModel(mNW4)
276
277
278
279
    allChecks('pNW4')
    parseLaTeX('pNW4')
280
    # REM animal: a=2pi. Sensor: t <= pi/2.
281
    mREM = [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
                                              pi/2],
283
             [r*sin(x3),
                                    x3, t,
284
                                     x4, 0*t,
             [r*sin(x3),
2.85
                                    x3, t,
                                                     pi/2],
             [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2]]
286
289
    repREM = {t:3*pi/8, a:2*pi} # Replacement values in range
290
291 # Define conditions for model
292 condREM = [ t <= pi/2 ]
293
    # Calculate model, run checks, write output.
295
    pREM = calcModel(mREM)
296
297
    allChecks('pREM')
    parseLaTeX('pREM')
298
301
    # NW5 animal: a>pi. Sensor: t <= pi/2. Condition: 2*pi - t < a
    mNW5 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
                                                    pi/2],
             [r*sin(x3),
                                    x3, t,
                                    x4, 0,
             ſr,
                                                     tl,
                                                    pi/2],
             [r*sin(x3),
                                     x3, t,
                                 x^2, pi/2 - t/2, 3*pi/2 - t/2 - a/2],
             [r*cos(x2 - t/2),
309
             [2*r*sin(t/2)*sin(x2), x2, 3*pi/2 - t/2 - a/2, pi/2]]
    repNW5 = {t:3*pi/8, a:29*pi/16} # Replacement values in range
    # Define conditions for model
315 condNW5 = [a >= pi, t <= pi/2, 2*pi - t <= a ]
316
317
    # Calculate model, run checks, write output.
318 pNW5 = calcModel(mNW5)
    allChecks('pNW5')
    parseLaTeX('pNW5')
321
    \# NW6 animal: a>pi. Sensor: t <= pi/2. Condition: 2*pi - 2*t <= a <= 2*pi - t \#
    mNW6 = [[2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
                                    x3, t,
             [r*sin(x3),
                                               pi/2],
                                     x4, 0,
                                                     t],
```

```
x3, t, pi/2],
x2, pi/2 - t/2, a/2 + t/2 - pi/2]]
329
              [r*sin(x3),
               [r*cos(x2 - t/2),
     repNW6 = {t:3*pi/8, a:3*pi/2} # Replacement values in range
334
335
    # Define conditions for model
    condNW6 = [a \ge pi, t \le pi/2, 2*pi - 2*t \le a, a \le 2*pi - t]
336
     # Calculate model, run checks, write output.
338 pNW6 = calcModel(mNW6)
339
     allChecks('pNW6')
340 parseLaTeX('pNW6')
341
342
343
344
     # NW7 animal: a>pi. Sensor: t <= pi/2. Condition: a <= 2pi - 2t #
345
346
347
    mNW7 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
348
               [r*sin(x3),
                             x3, t, pi/2],
349
                                        x4, 0,
               ſr,
              [r*sin(x3),
                                       x3, pi - a/2, pi/2]
353
354
     repNW7 = \{t:pi/9, a:10*pi/9\} \# Replacement values in range
    # Define conditions for model
355
    condNW7 = [t \le pi/2, a \ge pi, a \le 2*pi - 2*t]
     # Calculate model, run checks, write output.
    pNW7 = calcModel(mNW7)
360
    allChecks('pNW7')
    parseLaTeX('pNW7')
    # SE1 animal: a <= pi. Sensor: t =2pi.
367
    mSE1 = [ [ 2*r*sin(a/2),x1, pi/2, 3*pi/2]
368
371
372
373
    repSE1 = {a:pi/4} # Replacement values in range
     # Define conditions for model
374 condSE1 = [a <= pi]
376
377
     # Calculate model, run checks, write output.
    pSE1 = calcModel(mSE1)
378
    allChecks('pSE1')
parseLaTeX('pSE1')
380
382
383
     # SE2 animal: a <= pi. Sensor: t > pi. Condition: a > 2pi - t, a > 4pi - 2t #
384
385
                                                                                    t/2 + pi/2 - a/2
386 mSE2 = [2*r*sin(a/2),
                                                          x1, pi/2,
                                                         x1, p1/2,

x1, t/2 + pi/2 - a/2

x1, t/2 + pi/2 - a/2, 5*pi/2 - a/2 - t/2],

x1, 5*pi/2 - a/2 - t/2, 3*pi/2]]
            [r*sin(a/2) + r*cos(x1 - t/2),

[2*r*sin(a/2),
387
388
391
392
     repSE2 = {t:19*pi/10, a:pi/2} # Replacement values in range
    \# Define conditions for model condSE2 = [a <= pi, t >= pi, a >= 4*pi - 2*t]
393
396
397
     # Calculate model, run checks, write output.
    pSE2 = calcModel(mSE2)
398
    allChecks('pSE2')
399
    parseLaTeX('pSE2')
401
402
    # SE3 animal: a <= pi. Sensor: t > pi. Condition: 2pi - t < a < 4pi - 2t #
403
    mSE3 = [ [ 2*r*sin(a/2),
                                                         x1, pi/2, t/2 + pi/2 - a/2],

x1, t/2 + pi/2 - a/2, t/2 + pi/2],

x1, t/2 + pi/2, 5*pi/2 - a/2 - t/2],
404
               [ r*sin(a/2) + r*cos(x1 - t/2),
               [r*sin(a/2),
407
              [2*r*sin(a/2),
                                                         x1, 5*pi/2 - a/2 - t/2, 3*pi/2
408
409
    repSE3 = \{t:3*pi/2 + 0.1, a:pi/2\} # Replacement values in range
410
     # Define conditions for model
    condSE3 = [a \le pi, t \ge pi, a \ge 2*pi - t, a \le 4*pi - 2*t]
     # Calculate model, run checks, write output.
415 pSE3 = calcModel(mSE3)
```

```
416 | allChecks('pSE3')
417
    parseLaTeX('pSE3')
420
     \# SE4 animal: a <= pi. Sensor: t > pi. Condition: a <= 4\starpi - 2\start and a < 2\starpi - t \#
421
422
423
    mSE4 = [ [ 2*r*sin(a/2),
                                                                                t/2 + pi/2 - a/2 ],
                                                         x1, pi/2,
              [r*sin(a/2) + r*cos(x1 - t/2),
                                                         x1, t/2 + pi/2 - a/2, t/2 + pi/2
             [ r*sin(a/2),
                                                         x1, t/2 + pi/2,
                                                                                t/2 + pi/2 + a/2 ] ]
425
426
427
42.8
     repSE4 = \{t:3*pi/2, a:pi/3\} \# Replacement values in range
42.9
430
     # Define conditions for model
432 condSE4 = [a <= pi, t >= pi/2, a <= 4*pi - 2*t, a <= 2*pi - t]
433
# Calculate model, run
435 pSE4 = calcModel(mSE4)
     # Calculate model, run checks, write output.
436
    allChecks('pSE4')
    parseLaTeX('pSE4')
438
439
440
     \# SW1 animal: a <= pi. Sensor: pi/2 <= t <= pi. Condition: a >= t and a/2 >= t - pi/2 \#
441
442 \text{ mSW1} = [2*r*sin(t/2)*sin(x2),
                                                       x2, pi/2 - a/2 + t/2, pi/2
               [r*sin(a/2) - r*cos(x2 + t/2),

[r*sin(a/2) - r*cos(x4 - t),
                                                                   pi/2 - a/2 + t/2],
                                                       x2, t/2,
                                                                                t - pi/2 ],
t - pi/2 + a/2 ]]
                                                       x4, 0,
444
445
                [r*sin(a/2),
                                                        x4, t-pi/2,
446
447
     repSW1 = {t:5*pi/8, a:6*pi/8} # Replacement values in range
448
450
     # Define conditions for model
451
     condSW1 = [a \le pi, pi/2 \le t, t \le pi, a \ge t, a/2 \ge t - pi/2]
452
     # Calculate model, run checks, write output.
453
454 pSW1 = calcModel(mSW1)
    allChecks('pSW1')
parseLaTeX('pSW1'
455
457
458
459
     \# SW2 animal: a <= pi. Sensor: pi/2 <= t <= pi. Condition: a <= t and a/2 >= t- pi/2 \#
460
                                                 x2, pi/2 + a/2 - t/2, pi/2
461
    mSW2 = [2*r*sin(a/2),
                [r*sin(a/2) - r*cos(x2 + t/2), x2, t/2,
[r*sin(a/2) - r*cos(x4 - t), x4, 0*t,
                                                                          pi/2 + a/2 - t/2],
463
                                                                           t - pi/2
                                                                          t - pi/2 + a/2 ] ]
464
                [r*sin(a/2),
                                                  x4, t - pi/2,
465
466
467
     repSW2 = \{t:7*pi/8, a:7*pi/8-0.1\} \# Replacement values in range
     # Define conditions for model
470 \mid \text{condSW2} = [\text{a} \le \text{pi}, \text{pi/2} \le \text{t}, \text{t} \le \text{pi}, \text{a/2} \le \text{t/2}, \text{a/2} >= \text{t} - \text{pi/2}]
471
472 # Calculate model, run
473 pSW2 = calcModel(mSW2)
     # Calculate model, run checks, write output.
474
     allChecks('pSW2')
    parseLaTeX('pSW2')
475
476
477
478
479
     \# SW3 animal: a <= pi. Sensor: pi/2 <= t <= pi. Condition: a <= t and a/2 <= t - pi/2 \#
480
     mSW3 = [2*r*sin(a/2),
                                                        x2, t/2,
                                                                             pi/2
481
                                                        x4, 0,

x4, t - pi/2 - a/2, t - pi/2 ],

x4, t - pi/2, t - pi/2 + a/2]
482
                [2*r*sin(a/2),
483
                [r*sin(a/2) - r*cos(x4 - t),
                                                                             t - pi/2 + a/2 ] ]
484
               [r*sin(a/2)]
485
486
     repSW3 = {t:7*pi/8, a:2*pi/8} # Replacement values in range
488
489
     # Define conditions for model
490 condSW3 = [a <= pi, pi/2 <= t, t <= pi, a/2 <= t/2, a/2 <= t - pi/2]
491
492
     # Calculate model, run checks, write output.
493 pSW3 = calcModel(mSW3)
494
     allChecks('pSW3')
495
     parseLaTeX('pSW3')
496
497
498
     # SW4 animal: a <= pi. Sensor: t <= pi/2. Condition: a > pi − 2t & a <= t
    mSW4 = [ [2*r*sin(a/2),
                                                x2, pi/2 - t/2 + a/2, pi/2
              [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2, pi/2 - t/2 + a/2],
502
               [r*sin(a/2),
                                                 x3, t,
                                                                         pi/2
```

```
503
              [r*sin(a/2),
                                                                    a/2 + t - pi/2   1  1
                                             x4, 0,
504
    repSW4 = \{t:pi/2-0.1, a:pi/4\} \# Replacement values in range
507
    # Define conditions for model
508
    condSW4 = [a \le pi, t \le pi/2, a \ge pi - 2*t, a \le t]
510
    # Calculate model, run checks, write output.
    pSW4 = calcModel(mSW4)
512
    allChecks('pSW4')
513
    parseLaTeX('pSW4')
514
515
516
517
    \# SW5 animal: a <= pi. Sensor: t <= pi/2. Condition: a > pi - 2t & t <= a <= 2t
518
    mSW5 = [ [2*r*sin(t/2)*sin(x2),
                                             x2, pi/2 + t/2 - a/2, pi/2
                                                                    pi/2 + t/2 - a/2],
519
              [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2,
              [r*sin(a/2),
                                             x3, t,
                                                                    pi/2
                                                                    a/2 + t -pi/2
521
              [r*sin(a/2),
                                             x4, 0,
    repSW5 = {t:pi/2-0.1, a:pi/2} # Replacement values in range
526
527
528
    \# define conditions for model
    condSW5 = [a \le pi, t \le pi/2, a \ge pi - 2*t, t \le a, a \le 2*t]
    # Calculate model, run checks, write output.
531 pSW5 = calcModel(mSW5)
532
    allChecks('pSW5')
533
    parseLaTeX( pSW5')
534
535
536
    # SW6 animal: a <= pi. Sensor: t <= pi/2. Condition: a > pi - 2t & a > 2t
537
538
    mSW6 = [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2]
                                                 a/2
                                     x3, t,
539
              [r*sin(x3),
540
              [r*sin(a/2),
                                     x3, a/2,
                                                      pi/2
541
                                                     a/2 + t -pi/2
              [r*sin(a/2).
                                     x4, 0,
542
544
    repSW6 = {t:pi/4, a:3*pi/4} # Replacement values in range
545
546
547
    # Define conditions for model
548 condSW6 = [a <= pi, t <= pi/2, a >= pi - 2*t, a > 2*t]
    # Calculate model, run checks, write output.
551
552
    pSW6 = calcModel(mSW6)
    allChecks('pSW6')
553
    parseLaTeX('pSW6')
556
    \# SW7 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & a <= t
557
558 \text{ mSW7} = [2*r*sin(a/2),
                                             x2, pi/2 - t/2 + a/2, pi/2
              [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2,
                                                               pi/2 - t/2 + a/2],
559
560
                                                                    t + a/2
             [r*sin(a/2),
                                             x3, t,
561
563
    repSW7 = {t:2*pi/8, a:pi/8} # Replacement values in range
564
# Define conditions for model condSW7 = [a <= pi, t <= pi/2, a <= pi - 2*t, a <= t]
567
    # Calculate model, run checks, write output.
569 pSW7 = calcModel(mSW7)
570
571
    allChecks('pSW7')
    parseLaTeX('pSW7')
    \# SW8 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & t <= a <= 2t
575
576
577
                                          x2, pi/2 + t/2 - a/2, pi/2
    mSW8 = [ [2*r*sin(t/2)*sin(x2),
                                                               pi/2 + t/2 - a/2],
              [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2,
                                                                    t + a/2
578
              [r*sin(a/2),
                                             x3, t,
    repSW8 = {t:2*pi/8, a:pi/2-0.1} # Replacement values in range
581
582
    # Define conditions for model
583 condSW8 = [a <= pi, t <= pi/2, a <= pi - 2*t, t <= a, a <= 2*t]
     # Calculate model, run checks, write output.
586 pSW8 = calcModel(mSW8)
    allChecks('pSW8')
588 parseLaTeX('pSW8')
589
```

```
591
      # SW9 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & 2t <= a
      mSW9 = [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2]
                                                   x3, t,
594
                   [r*sin(x3),
595
                   [r*sin(a/2),
                                                   x3, a/2,
                                                                         t + a/2 ] ]
      repSW9 = {t:1*pi/8, a:pi/2} # Replacement values in range
599
600
      # Define conditions for model
601
      condSW9 = [a \le pi, t \le pi/2, a \le pi - 2*t, 2*t \le a]
602
      # Calculate model, run checks, write output.
603
604 pSW9 = calcModel(mSW9)
605
      allChecks('pSW9')
606
      parseLaTeX('pSW9')
607
608
609
      ####################
610
      ## Run tests
      ####################
613
      # create gas model object
614
      qas = 2*r
615
616
      # for each model run through every adjacent model.
      # Contains duplicates but better for avoiding missed comparisons.
619
      # Also contains replacement t->a and a->t just in case.
620
621
62.2
      allComps = [
      ['gas', 'pNE1', {t:2*pi}], ['gas', 'pSE1', {a:pi}],
623
      ['pNE1', 'gas', {t:2*pi}], ['pNE1', 'pNW1', {t:pi}],
['pNE1', 'pNE2', {a:3*pi-t}], ['pNE1', 'pNE2', {t:3*pi-a}],
625
626
62.7
      ['pNE2', 'pNE1', {a:3*pi-t}], ['pNE2', 'pNE1', {t:3*pi-a}],
['pNE2', 'pNE3', {a:4*pi-2*t}], ['pNE2', 'pNE3', {t:2*pi-a/2}],
['pNE2', 'pSE2', {a:pi}],
628
629
631
      ['pNE3', 'pNE2',{a:4*pi-2*t}], ['pNE3', 'pNE2',{t:2*pi-a/2}],
['pNE3', 'pSE3',{a:pi}], ['pNE3', 'pNW2',{t:pi}],
632
633
634
635
      ['pNW1','pNE1', {t:pi}], ['pNW1','pNW2', {a:2*pi}],
      ['pNW2','pNE3',{t:pi}], ['pNW2','pNW3',{a:3*pi-2*t}],
['pNW2','pNW3',{t:3*pi/2-a/2}], ['pNW2','pNW1',{a:2*pi}],
637
638
639
      ['pNW3','pNW5',{t:pi/2}], ['pNW3','pNW4',{a:2*pi-t}],
['pNW3','pNW4',{t:2*pi-a}], ['pNW3','pNW2',{a:3*pi-2*t}],
['pNW3','pNW2',{t:3*pi/2-a/2}],
640
641
643
      ['pNW4','pNW6', {t:pi/2}], ['pNW4','pNW3', {t:2*pi-a}],
['pNW4','pNW3', {a:2*pi-t}], ['pNW4','pSW1', {a:pi}],
644
645
646
647
      ['pREM','pNW1', {t:pi/2}], ['pREM','pNW5', {a:2*pi}],
648
      ['pNW5','pREM',{a:2*pi}], ['pNW5','pNW6',{a:2*pi-t}],
['pNW5','pNW6',{t:2*pi-a}], ['pNW5','pNW3',{t:pi/2}],
649
650
651
      ['pNW6','pNW5',{a:2*pi-t}], ['pNW6','pNW5',{t:2*pi-a}],
['pNW6','pNW7',{t:pi-a/2}], ['pNW6','pNW7',{a:2*pi-2*t}],
['pNW5','pNW4',{t:pi/2}],
652
653
      ['pNW7','pNW6',{t:2*pi-2*a}], ['pNW7','pNW6',{a:2*pi-2*t}],
['pNW7','pSW6',{a:pi}],
656
657
658
659
      ['pSE1'.'pSE2'.{t:2*pi}]. ['pSE1'.'gas'.{a:pi}].
660
      ['pSE2','pSE3',{t:2*pi-a/2}], ['pSE2','pSE3',{a:4*pi-2*t}],
['pSE2','pSE1',{t:2*pi}], ['pSE2','pNE2',{a:pi}],
662
663
      ['pSE3','pSE2',{a:4*pi-2*t}], ['pSE3','pSE2',{t:2*pi-a/2}],
['pSE3','pSE4',{a:2*pi-t}], ['pSE3','pSE4',{t:2*pi-a}],
['pSE3','pNE3',{a:pi}],
664
      ['pSE4','pSE3',{t:2*pi-a}], ['pSE4','pSE3',{a:2*pi-t}],
['pSE4','pSW3',{t:pi}],
668
669
670
      ['psW1','psW5',{t:pi/2}], ['psW1','psW2',{a:t}],
['psW1','psW2',{t:a}], ['psW1','pNW4',{a:pi}],
671
      ['pSW2','pSW1',{a:t}], ['pSW2','pSW1',{t:a}],
675 ['psw2','psw4',{t:pi/2}], ['psw2','psw3',{a:2*t-pi}],
676 ['psw2','psw3',{t:a/2+pi/2}],
```

```
['psw3','psw2',{t:a/2+pi/2}], ['psw3','psw2',{a:2*t-pi}],
['psw3','psE4',{t:pi}],
678
680
681
     ['pSW4','pSW7',{a:pi-2*t}], ['pSW4','pSW7',{t:pi/2-a/2}],
['pSW4','pSW5',{t:a}], ['pSW4','pSW5',{a:t}],
['pSW4','pSW2',{t:pi/2}],
682
683
684
     ['pSW5','pSW4',{t:a}], ['pSW5','pSW4',{a:t}],
['pSW5','pSW8',{t:pi/2-a/2}], ['pSW5','pSW8',{a:pi-2*t}],
['pSW5','pSW6',{a:2*t}], ['pSW5','pSW6',{t:a/2}],
['pSW5','pSW1',{t:pi/2}],
686
687
688
689
690
     ['psW6','psW9',{t:pi/2-a/2}], ['psW6','psW9',{a:pi-2*t}],
['psW6','psW5',{a:2*t}], ['psW6','psW5',{t:a/2}],
['psW6','pNW7',{a:pi}],
691
692
693
694
695
     ['psW7','psW8',{t:a}], ['psW7','psW8',{a:t}],
['psW7','psW4',{t:pi/2-a/2}], ['psW7','psW4',{a:pi-2*t}],
696
697
     ['psw8','psw7',{a:t}], ['psw8','psw7',{t:a}],
['psw8','psw9',{a:2*t}], ['psw8','psw9',{t:a/2}],
['psw8','psw5',{a:pi-2*t}], ['psw8','psw5',{t:pi/2-a/2}],
699
     ['pSW9','pSW8',{a:2*t}], ['pSW9','pSW8',{t:a/2}],
['pSW9','pSW6',{a:pi-2*t}], ['pSW9','pSW6',{t:pi/2-a/2}]
703
     # List of regions that touch a=0. Should equal 0 when a=0.
zeroRegions = ['psW9', 'psW8', 'psW7', 'psW4', 'psW2', 'psW3', 'psE4', 'psE3', 'psE1']
708
709
     # Run through all the comparisons. Need simplify(). Even together() gives some false negatives.
714
715
     checkFile = open('checksFile.tex','w')
716
     checkFile.write('All checks evaluated.\nTim Lucas - ' + str(datetime.now()) + '\n')
     for i in range(len(allComps)):
718
              if (eval(allComps[i][0]).subs(allComps[i][2]) - eval(allComps[i][1]).subs(allComps[i][2])).
                    simplify() == 0:
719
                        \label{eq:checkFile.write} \mbox{(str(i) + ': ' + allComps[i][0]+ ' and ' +allComps[i][1]+': OK\n')}
720
              else:
                        checkFile.write(str(i) + ': ' + allComps[i][0]+ ' and ' +allComps[i][1]+': Incorrect\n')
723
     for i in range(len(zeroRegions)):
724
725
              if eval(zeroRegions[i]).subs({a:0}).simplify() == 0:
                        checkFile.write(zeroRegions[i] + ' at a = 0: OK\n')
726
              else:
                        checkFile.write(zeroRegions[i] + ' at a = 0: Incorrect\n')
729 | # pSE2 is slightly different. Only one corner touches a=0, so need theta value as well. I'm not sure why
           this isn't
730 # A problem for some other regions.
731
732
     else:
734
             checkFile.write('pSE2 at a = 0, t = 2pi: Incorrect\n')
     checkFile.close()
738
     # And print to terminal
     #for i in range(len(allComps)):
740
               if not (eval(allComps[i][0]).subs(allComps[i][2]) - eval(allComps[i][1]).subs(allComps[i][2])).
           simplify() == 0:
741
                       print allComps[i][0] + ' and ' + allComps[i][1]+': Incorrect\n'
742
743
744
     745
     ### Define a a function that calculates p bar answer.
746
     747
748 def calcP(A, T, R):
      assert (A <= 2*pi and A >= 0), "a is out of bounds. Should be in 0<a<2*pi" assert (T <= 2*pi and T >= 0), "s is out of bounds. Should be in 0<s<2*pi"
749
751
752
       if A > pi:
753
         if A < 4*pi - 2* T:
           p = pNW7.subs({a:A, t:T, r:R}).n()
755
          elif A <= 3*pi - T:
756
                                 p = pNE2.subs({a:A, t:T, r:R}).n()
         else:
                                  p = pNE1.subs({a:A, t:T, r:R}).n()
         if A < 4*pi - 2* T:
```

```
761
                             p = pSE3.subs({a:A, t:T, r:R}).n()
762
        else:
763
                             p = pSE2.subs({a:A, t:T, r:R}).n()
764
            return p
765
766
767
     #################################
768
     ## Apply to entire grid ###
771
772
     # How many values for each parameter
    nParas = 100
    \mbox{\#} Make a vector for a and s. Make an empty nParas x nParas array.
     # Calculated profile sizes will go in pArray
    tVec = np.linspace(0, 2*pi, nParas)
    aVec = np.linspace(0, 2*pi, nParas)
778
    pArray = np.zeros((nParas, nParas))
780
    # Calculate profile size for each combination of parameters
    for i in range(nParas):
781
            for j in range(nParas):
783
                     pArray[i][j] = calcP(aVec[i], tVec[j], 1)
785
    \ensuremath{\mathtt{\#}} Turn the array upside down so origin is at bottom left.
786
    pImage = np.flipud(pArray)
787
    # Plot and save.
789
    pl.imshow(pImage, interpolation='none', cmap=pl.get_cmap('Blues'))
790
791
792
     # Show or save image.
    # pl.show()
793
    # pl.savefig('/imgs/profilesCalculated.png')
794
795
797
    ***********
798
    800
     # To reduce mistakes, output R function directly from python.
802
     # However, the if statements, which correspond to the bounds of each model, are not automatic.
803
804
805
    Rfunc = open('supplementaryRscript.R', 'w')
806
    Rfunc.write("""
     # S4
808
    # Supplementary R script from
809
    # A generalised random encounter model for estimating animal density with remote sensor data
810
    # Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
811
812
    # calcDensity is the main function to calculate density.
813 # It takes parameters z, alpha, theta, r, animalSpeed, t
    # z - The number of camera/acoustic counts or captures.
    # alpha - Call width in radians.
# theta - Sensor width in radians.
815
816
817
    # r - Sensor range in metres.
818
    # animalSpeed - Average animal speed in metres per second.
819
     # t - Length of survey in sensor seconds i.e. number of sensors x survey duration.
821
    # calcAbundance calculates abundance rather than density and requires an extra parameter
82.2
    # area - In metres squared. The size of the region being examined.
823
824
825
    # Internal function to calculate profile width as described in the text
     calcProfileWidth <- function(alpha, theta, r) {</pre>
827
            if(alpha > 2*pi | alpha < 0)
828
                 stop('alpha is out of bounds. alpha should be in interval 0<a<2*pi')
             if(theta > 2*pi | theta < 0)
829
                 stop('theta is out of bounds. theta should be in interval 0<a<2*pi')
830
831
     if(alpha > pi){
833
              if(alpha < 4*pi - 2*theta){
834
                 p <- ' + str(pNW7) +
835
                        } else if(alpha <= 3*pi - theta) {'
        p <- ' + str(pNE2) +
} else {'</pre>
    '\n
836 '\n
837 '\n
    ∕\n
839 '\n
                                 p <- ' + str(pNE1) +
    \prime \setminus n
              }'
} else {'
840
    '\n
841
    '∖n
842
843
                if(alpha < 4*pi - 2*theta){'
    '\n
                                p <- ' + str(pSE3) +
844 '\n
845 '\n
            } else {'
                                 p <- ' + str(pSE2) +
846 '\n
847 /\n
              }′
```

Lucas et al. A generalised random encounter model for animals

```
848 '\n
849 '\n}' +
                               return(p)'
851
852
          # Calculate a population density. See above for units etc.
         calcDensity <- function(z, alpha, theta, r, animalSpeed, t){</pre>
                         # Check the parameters are suitable.

if(z <= 0 | !is.numeric(z)) stop('Counts, z, must be a positive number.')

if(animalSpeed <= 0 | !is.numeric(animalSpeed)) stop('animalSpeed must be a positive number.')

if(t <= 0 | !is.numeric(t)) stop('Time, t, must be a positive number.')
853
854
855
857
858
859
                         # Calculate profile width, then density.
p <- calcProfileWidth(alpha, theta, r)
D <- z/{animalSpeed*t*p}</pre>
860
861
862
                         return(D)
863
864
865
866
867
         \ensuremath{\text{\#}} Calculate abundance rather than density.
         calcAbundance <- function(z, alpha, theta, r, animalSpeed, t, area){
   if(area <= 0 | !is.numeric(area)) stop('Area must be a positive number')
   D <- calcDensity(z, alpha, theta, r, animalSpeed, t)
   A <- D*area</pre>
868
869
870
871
872
873
                         return(A)
         )
874 Rfunc.close()
```

lucas_et_al_S3.py

S4. SUPPLEMENTARY SCRIPT: R IMPLEMENTATION OF MODELS

This is an implementation of the models derived in the paper in R (?). Once given the parameters θ and α it automatically selects the correct model to apply.

The script can also be found in the plain text file lucas_et_al_S4.R. It will also be hosted and given a DOI on Github at the time of publication (https://github.com/timcdlucas/lucasMoorcroftManuscript/tree/postPeerReview).

```
# Supplementary R script from
   # A generalised random encounter model for estimating animal density with remote sensor data
   # Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
   # calcDensity is the main function to calculate density.
   # It takes parameters z, alpha, theta, r, animalSpeed, t
   \sharp z - The number of camera/acoustic counts or captures.
   # alpha - Call width in radians.
   # theta - Sensor width in radians.
   # r - Sensor range in metres.
   # animalSpeed - Average animal speed in metres per second.
   # t - Length of survey in sensor seconds i.e. number of sensors x survey duration.
   # calcAbundance calculates abundance rather than density and requires an extra parameter
   # area - In metres squared. The size of the region being examined.
   # Internal function to calculate profile width as described in the text
   calcProfileWidth <- function(alpha, theta, r) {</pre>
            if(alpha > 2*pi | alpha < 0)
       stop('alpha is out of bounds. alpha should be in interval 0<a<2*pi')
   if(theta > 2*pi | theta < 0)</pre>
       stop('theta is out of bounds. theta should be in interval 0<a<2*pi')</pre>
     if(alpha > pi){
              if (alpha < 4*pi - 2*theta) {</pre>
                p <- r*(theta - cos(alpha/2) + 1)/pi
                    p \leftarrow r*(theta + 2*sin(theta/2))/pi
            } else {
              if(alpha < 4*pi - 2*theta){</pre>
                             p \leftarrow r*(theta*sin(alpha/2) - cos(alpha/2) + 1)/pi
       } else {
                             p \leftarrow r*(theta*sin(alpha/2) - cos(alpha/2) + cos(alpha/2 + theta))/pi
            }
42
            return(p)
43
   # Calculate a population density. See above for units etc.
   calcDensity \leftarrow function(z, alpha, theta, r, animalSpeed, t){
              Check the parameters are suitable.
            if(z <= 0 | !is.numeric(z)) stop('Counts, z, must be a positive number.')
if(animalSpeed <= 0 | !is.numeric(animalSpeed)) stop('animalSpeed must be a positive number.')</pre>
48
            if(t <= 0 | !is.numeric(t)) stop('Time, t, must be a positive number.')</pre>
            # Calculate profile width, then density.
           p <- calcProfileWidth(alpha, theta, r)
            D <- z/{animalSpeed*t*p}</pre>
            return(D)
   # Calculate abundance rather than density.
   calcAbundance <- function(z, alpha, theta, r, animalSpeed, t, area){</pre>
           if(area <= 0 | !is.numeric(area)) stop('Area must be a positive number')</pre>
           D \leftarrow calcDensity(z, alpha, theta, r, animalSpeed, t) A \leftarrow D*area
62
            return(A)
```

lucas_et_al_S4.R

S5. Supplementary Information: Simulation model results of the gREM precision

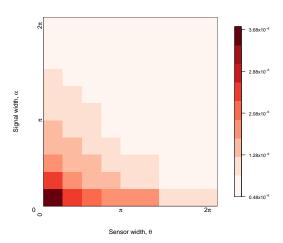


Figure S5.1. Simulation model results of the gREM precision given a range of sensor and signal widths, shown by the standard deviation of the error between the estimated and true densities. Standard deviations are shown from deep red to pink, representing high to low values between 0.483×10^{-6} to 3.74×10^{-6} .

S6. SUPPLEMENTARY INFORMATION: IMPACT OF PARAMETER ERROR

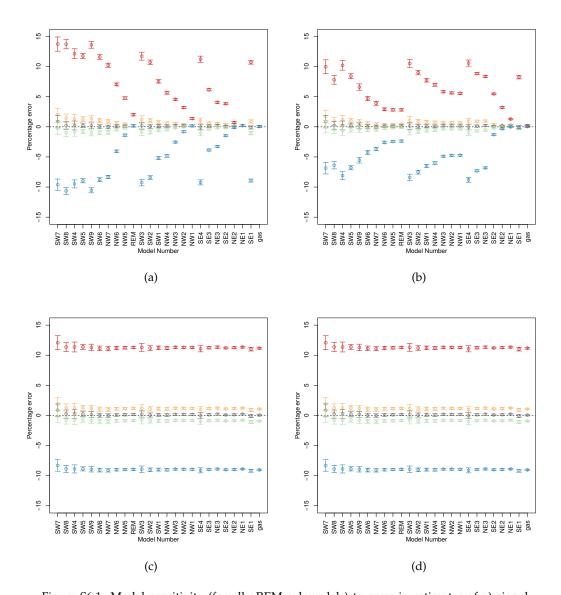


Figure S6.1. Model sensitivity (for all gREM submodels) to error in estimates of a) signal width α , b) sensor width θ , c) detection distance r and d) animal movement speed v. Estimates are -10% (red), -1% (orange), 0% (grey), +1% (green) and +10% (blue) of the true parameter value. The black dashed line indicates zero error in density estimates. The error bars 95% confidence intervals across all simulations.