

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

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S1. TABLE OF SYMBOLS

Symbol	Description	Units
$\theta$	Sensor width	rad
$\alpha$	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	$\text{m s}^{-1}$
$t$	Time	s
$z$	Number of detections	-
$D$	Animal density	$\text{m}^{-2}$
$T$	Step length	s
$N$	Number of steps per simulation	-
$d$	Distance moved in a time step	m
$S$	Probability of remaining stationary	-
$A$	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 Yapp (1956), 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  $\theta$  and  $\alpha$  would be expected to occur (e.g., sensors have different detection widths and animals have different signal widths). For different combinations  $\theta$  and  $\alpha$ , 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  $\theta$  and  $\alpha$  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  $\theta$  and  $\alpha$  for which the derivation of the equations are the same (defined as sub-models in the gREM) (Fig. S2.1). For example, the

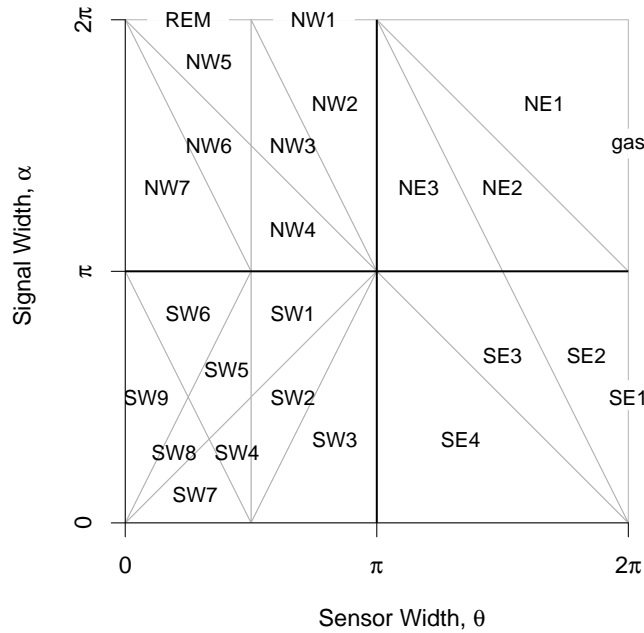


Figure S2.1. Locations where derivation of the average profile  $\bar{p}$  is the same for different combinations of sensor detection and animal signal widths. Symbols within each polygon refer to each gREM submodel named after their compass point, except for Gas and REM which highlight the position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

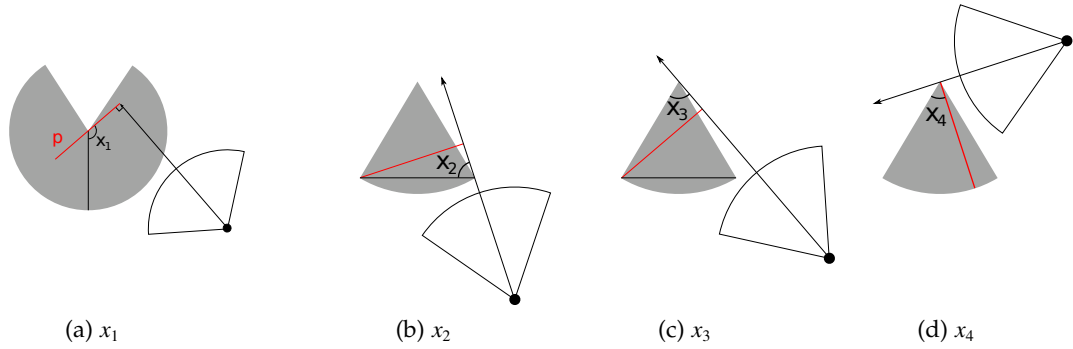


Figure S2.2. The location of the focal angles  $x_{i \in [1,4]}$ .  $x_1$  is used in SE and NE models (including the gas model).  $x_2 - x_4$  are used in NW and SW models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. 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,  $x_3 = \theta$ . (d)  $x_4$  measures the angle between the left side of the detection region and the profile.

gas model becomes the simplest gREM sub-model (upper right in Fig. S2.1) and the REM from (Rowcliffe *et al.*, 2008) 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  $\alpha \leq \pi$  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.

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

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

This profile is integrated over the interval  $[\frac{\pi}{2}, \frac{3\pi}{2}]$  which is  $\pi$  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  $2\pi$  angles of approach and dividing by  $2\pi$ .

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

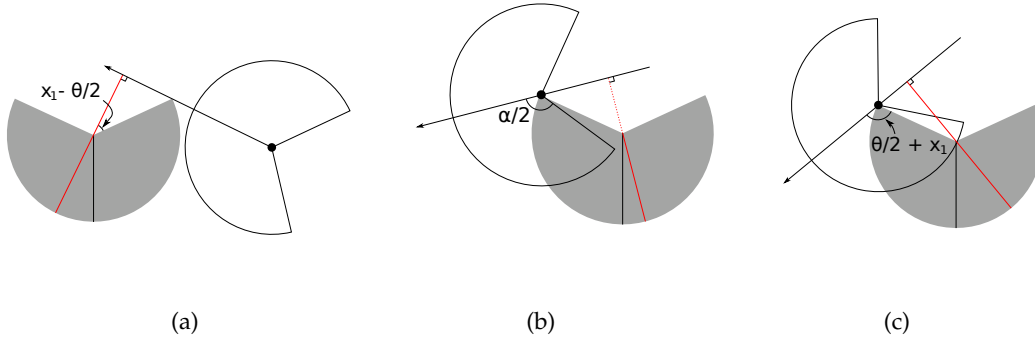


Figure S2.3. Three of the integrals in NE models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile  $p$  is shown with a red line. Dashed red lines 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  $2r$  once again.

**S2.4.1. Model NE1.** Submodel NE1 exists within the area bounded by  $\alpha \leq 2\pi$ ,  $\theta \leq 2\pi$  and  $\alpha \geq 3\pi - \theta$  (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,  $\theta$  is wide enough that the  $r + r \cos(x_1 + \theta/2)$  profile starts at  $\pi$ . This then gives us

$$\bar{p}_{NE1} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, dx_1 + \int_{\frac{\pi}{2}}^{\pi} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, dx_1 + \int_{\pi}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, dx_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, dx_1 \right) \quad \text{eqn S3}$$

$$\bar{p}_{NE1} = \frac{r}{\pi} \left( \theta + 2 \sin\left(\frac{\theta}{2}\right) \right) \quad \text{eqn S4}$$

**S2.4.2. Model NE2.** Model NE2 is bounded by  $\alpha \leq 3\pi - \theta$ ,  $\alpha \geq 4\pi - 2\theta$  and  $\alpha \geq \pi$  (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  $\pi$  which is reflected in the different bounds in the second and third integral.

$$\bar{p}_{NE2} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, dx_1 + \int_{\frac{\theta}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, dx_1 + \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, dx_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, dx_1 \right) \quad \text{eqn S5}$$

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

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

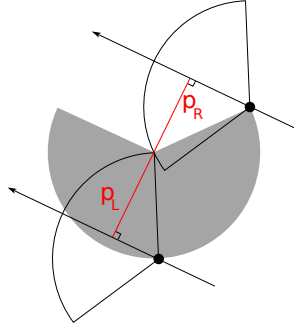


Figure S2.4. The second integral in SE. The right side of the profile ( $p_R$ ) is limited by the size of the sensor region while the left side of the profile ( $p_L$ ) is limited by the size of the signal width. The full profile has width  $p = r \sin(\alpha/2) + r \cos(\theta/2 - x_1)$ . The sector 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.

$$\bar{p}_{NE3} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}}^{\frac{\theta}{2}} 2r \, dx_1 + \int_{\frac{\theta}{2}}^{\frac{\theta}{2} + \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_1\right) + r \, dx_1 + \int_{\frac{\theta}{2} + \frac{\pi}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \, dx_1 + \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{2\pi - \frac{\theta}{2}} r \cos\left(\frac{\theta}{2} + x_1\right) + r \, dx_1 + \int_{2\pi - \frac{\theta}{2}}^{\frac{3\pi}{2}} 2r \, dx_1 \right) \quad \text{eqn S7}$$

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

**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  $\alpha$  is less than  $\pi$  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  $\alpha \leq 2\pi - \theta$ , 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  $\theta/2 + \pi/2 \geq 5\pi/2 - \alpha/2 - \theta/2$  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  $\alpha \leq 4\pi - 2\theta$ .

**S2.5.1. Model SE2.** SE2 is bounded by  $\alpha \geq 4\pi - 2\theta$ ,  $\alpha \leq \pi$  and  $\theta \leq 2\pi$  (Fig. S2.1). As  $\alpha \geq 4\pi - 2\theta$ , there is no  $r \sin(\alpha/2)$  profile. As  $\alpha \leq 4\pi - 2\theta$ , 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.

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

$$\bar{p}_{SE2} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + \cos\left(\frac{\alpha}{2} + \theta\right) \right) \quad \text{eqn S10}$$

**S2.5.2. Model SE3.** SE3 is bounded by  $4\pi - 2\theta \leq \alpha \leq 4\pi - 2\theta$  and  $\alpha \leq \pi$  (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.

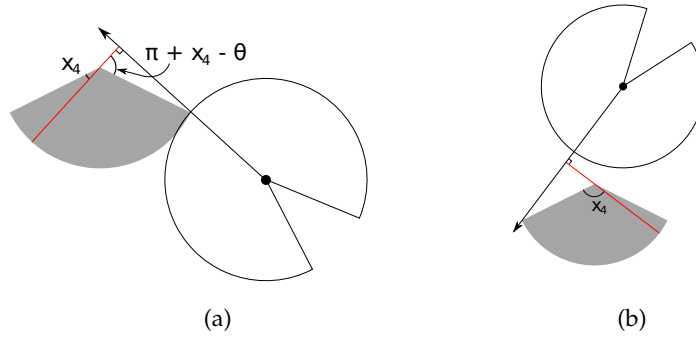


Figure S2.5. The second and fourth profiles of NW1. The left side of both profiles is of width  $r$  while the right side differs. (a) The right side of the profile is  $r \cos(\pi + x_4 - \theta) = -r \cos(\theta - x_4)$  (b) The right side is  $r \cos(\pi - x_4) = -r \cos x_4$  respectively. In both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

$$\bar{p}_{SE3} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 + \int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) + r \cos\left(\frac{\theta}{2} - x_1\right) dx_1 + \int_{\frac{\theta}{2} + \frac{\pi}{2}}^{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_1 + \int_{\frac{5\pi}{2} - \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{3\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_1 \right) \quad \text{eqn S11}$$

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

S2.5.3. *Model SE4.* Finally SE4 is bounded by  $\alpha \leq 4\pi - 2\theta$ ,  $\alpha \leq \pi$  and  $\theta \leq \pi$  (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.

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

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

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  $\gamma_1$  and  $\gamma_2$  in (Rowcliffe *et al.*, 2008) 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  $r$ .

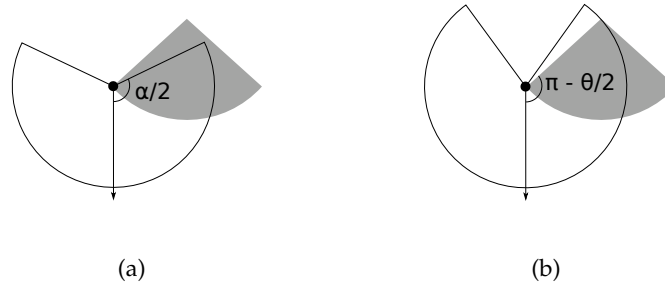


Figure S2.6. Profile sizes when an animal approaches from behind in models NW2–4. If  $\alpha$  is relatively large, animals can be detected when approaching from behind. Otherwise animals cannot be detected. The sector shaped detection region is shown in grey. Animals are filled black circles and the 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.

$$\bar{p}_{NW1} = \frac{1}{\pi} \left( \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_0^{\frac{\theta - \pi}{2}} r - r \cos(-x_4 + \theta) dx_4 \right. \\ \left. + \int_{\frac{\theta - \pi}{2}}^{\frac{\pi}{2}} r dx_4 + \int_{\frac{\pi}{2}}^{\theta} r - r \cos(x_4) dx_4 + \int_{\frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 \right) \quad \text{eqn S15}$$

$$\bar{p}_{NW1} = \frac{r}{\pi} (\theta + 2) \quad \text{eqn S16}$$

**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  $\alpha$  is now less than  $2\pi$ , 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  $\alpha \leq 3\pi - 2\theta$ . 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  $\alpha \leq 2\pi$  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  $\alpha/2 \leq \pi - \theta/2$  (Fig. S2.6). This inequality (simplified as  $\alpha \leq 2\pi - \theta$ ) defines the boundary between NW3 and NW4.

**S2.7.1. Model NW2.** NW2 is bounded by  $\alpha \geq 3\pi - 2\theta$ ,  $\alpha \leq 2\pi$  and  $\theta \leq \pi$  (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  $x_4 = \theta$ .



$$\bar{p}_{\text{NW2}} = \frac{1}{\pi} \left( \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_0^{\theta - \frac{\pi}{2}} r - r \cos(-x_4 + \theta) dx_4 \right. \\ \left. + \int_{\theta - \frac{\pi}{2}}^{\frac{3\pi}{2} - \frac{\alpha}{2}} r dx_4 + \int_{\frac{3\pi}{2} - \frac{\alpha}{2}}^{\theta} r - r \cos(x_4) dx_4 + \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 \right) \quad \text{eqn S17}$$

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

S2.7.2. *Model NW3.* NW3 is bounded by  $\alpha \leq 3\pi - 2\theta$ ,  $\alpha \geq 2\pi - \theta$  and  $\theta \geq \pi/2$  (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  $\theta$  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  $2r \sin(x_2) \sin(\theta/2)$ .

$$\bar{p}_{\text{NW3}} = \frac{1}{\pi} \left( \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_0^{\theta - \frac{\pi}{2}} r - r \cos(-x_4 + \theta) dx_4 \right. \\ \left. + \int_{\theta - \frac{\pi}{2}}^{\theta} r dx_4 + \int_{\frac{3\pi}{2} - \frac{\alpha}{2} - \frac{\theta}{2}}^{\frac{3\pi}{2} - \frac{\alpha}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) dx_2 + \int_{\frac{3\pi}{2} - \frac{\alpha}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 \right) \quad \text{eqn S19}$$

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

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

$$\bar{p}_{\text{NW4}} = \frac{1}{\pi} \left( \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_0^{\theta - \frac{\pi}{2}} r - r \cos(-x_4 + \theta) dx_4 \right. \\ \left. + \int_{\theta - \frac{\pi}{2}}^{\theta} r dx_4 + \int_{\frac{\alpha}{2} + \frac{\theta}{2} - \frac{\pi}{2}}^{\frac{\alpha}{2} + \frac{\theta}{2} - \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) dx_2 \right) \quad \text{eqn S21}$$

$$\bar{p}_{\text{NW4}} = \frac{r}{\pi} \left( \theta - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S22}$$

S2.8. **Model REM.** REM is the model from (Rowcliffe *et al.*, 2008). It has  $\alpha = 2\pi$  and  $\theta \leq \pi/2$  (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  $\theta$  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.

$$\bar{p}_{\text{REM}} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_0^{\theta} r dx_4 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 \right) \quad \text{eqn S23}$$

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

**S2.9. Models NW5–7.** In the models NW5–7, the sensor has  $\theta \leq \pi/2$  as in the REM. As  $\alpha \geq \pi$  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  $\alpha \geq \pi$  signal is no different to a  $2\pi$  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  $\theta$  radians of  $x_4$ .
- (4) If  $\alpha \leq 2\pi + 2\theta$ , 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  $\alpha \geq 2\pi + 2\theta$  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  $\alpha \leq 2\pi - \theta$  as in NW6)
- (7) or the right becomes detectable (if  $\alpha \geq 2\pi - \theta$  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  $\alpha \geq 2\pi - \theta$ ,  $\alpha \leq 2\pi$  and  $\theta \leq \pi/2$  (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.

$$\bar{p}_{\text{NW5}} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_0^{\theta} r dx_4 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 \right) \quad \text{eqn S25}$$

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

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

NW6 is the same NW5 except that as  $\alpha \leq 2\pi - \theta$ , 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.

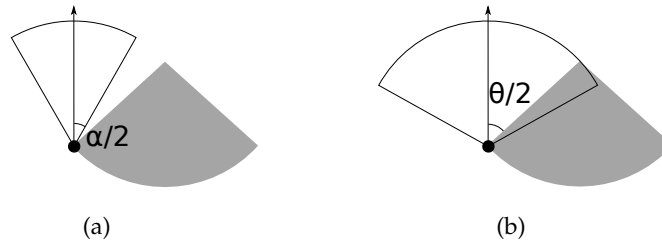


Figure S2.7. The first profile in SW models is limited by either  $\alpha$  or  $\beta$  depending on whether  $\alpha < \beta$ . The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated 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  $2r \sin(\theta/2) \sin(x_2)$ .

$$\bar{p}_{NW6} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_0^{\theta} r dx_4 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\theta}{2} + \frac{\theta}{2} - \frac{\pi}{2}} r \cos\left(\frac{\theta}{2} - x_2\right) dx_2 \right) \quad \text{eqn S27}$$

$$\bar{p}_{NW6} = \frac{r}{\pi} \left( \theta - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S28}$$

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

$$\bar{p}_{NW7} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\pi}{2}} r \sin(x_3) dx_3 + \int_0^{\theta} r dx_4 + \int_{\pi - \frac{\alpha}{2}}^{\frac{\pi}{2}} r \sin(x_3) dx_3 \right) \quad \text{eqn S29}$$

$$\bar{p}_{NW7} = \frac{r}{\pi} \left( \theta - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S30}$$

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  $\alpha \leq \pi$  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  $\alpha \geq \theta$ ,  $\alpha \leq \pi$  and  $\theta \leq \pi$  (Fig. S2.1).

As  $\alpha$  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  $r \sin(\alpha/2)$ .

$$\bar{p}_{SW1} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\frac{\theta}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right. \\ \left. + \int_0^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos(\theta - x_4) dx_4 + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S31}$$

$$\bar{p}_{SW1} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S32}$$

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

SW2 is largely similar to SW1. However, as  $\alpha \leq \theta$  the first profile is limited by  $\alpha$  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  $x_2 = \pi/2 - \alpha/2 + \theta/2$ .

$$\bar{p}_{SW2} = \frac{1}{\pi} \left( \int_{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{\frac{\theta}{2}}^{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right. \\ \left. + \int_0^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos(\theta - x_4) dx_4 + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S33}$$

$$\bar{p}_{SW2} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S34}$$

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

SW3 is similar to SW2 except that the profile does not become limited by sensor at all during 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.

$$\bar{p}_{SW3} = \frac{1}{\pi} \left( \int_{\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_0^{-\frac{\pi}{2} + \theta - \frac{\alpha}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_4 \right. \\ \left. + \int_{-\frac{\pi}{2} + \theta - \frac{\alpha}{2}}^{\theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos(\theta - x_4) dx_4 + \int_{\theta - \frac{\pi}{2}}^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S35}$$

$$\bar{p}_{SW3} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S36}$$

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.

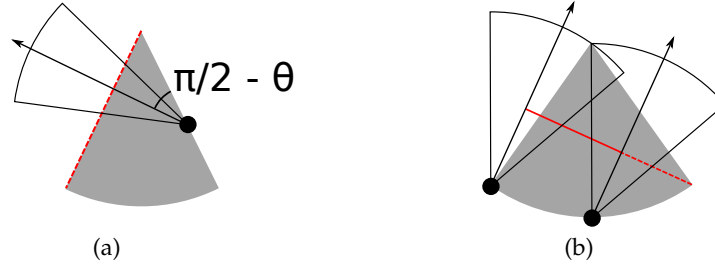


Figure S2.8. Description of two profiles in SW models. The sector shaped detection region is shown in grey. Animals are filled black circles and the animal signal is an unfilled sector. The animals direction of movement is indicated with an arrow. The profile  $p$  is shown with a red line. Dashed red 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  $\alpha \leq \pi - 2\theta$  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  $\alpha \leq \theta$  are limited by  $\alpha$  in the first,  $x_2$  region (Fig. S2.7), rather than being limited by  $\theta$ . Therefore this first profile is of width  $2r \sin(\alpha/2)$  rather than  $2r \sin(\theta/2) \sin(x_2)$ .
- (3) Finally, models with  $\alpha \leq 2\theta$  have a second profile in  $x_2$  where to one side of the sensor  $\alpha$  is the limiting factor of profile width, while on the other side  $\theta$  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  $\alpha \geq 2\theta$ .

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

$$\bar{p}_{SW4} = \frac{1}{\pi} \left( \int_{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\alpha}{2}\right) dx_2 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\alpha}{2} - \frac{\theta}{2} + \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right. \\ \left. + \int_{\theta}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_3 + \int_0^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S37}$$

$$\bar{p}_{SW4} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S38}$$

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

$$\bar{p}_{SW5} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\frac{\pi}{2} - \frac{\theta}{2}}^{\frac{\pi}{2} + \frac{\theta}{2} - \frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 \right. \\ \left. + \int_{\theta}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_3 + \int_0^{\frac{\alpha}{2} + \theta - \frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S39}$$

$$\bar{p}_{SW5} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S40}$$

S2.11.3. *Model SW6.* SW6 is bounded by  $\alpha \geq \pi - 2\theta$ ,  $\alpha \geq 2\theta$  and  $\alpha \leq \pi$  (Fig. S2.1). It starts with a  $\theta$  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.

$$\bar{p}_{SW6} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\alpha}{2}} r \sin(x_3) dx_3 + \int_{\frac{\alpha}{2}}^{\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_3 + \int_0^{\frac{\alpha}{2}+\theta-\frac{\pi}{2}} r \sin\left(\frac{\alpha}{2}\right) dx_4 \right) \quad \text{eqn S41}$$

$$\bar{p}_{SW6} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S42}$$

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

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

$$\bar{p}_{SW7} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S44}$$

S2.11.5. *Model SW8.* SW8 is bounded by  $\alpha \leq \pi - 2\theta$ ,  $\alpha \geq \theta$  and  $\alpha \leq 2\theta$  (Fig. S2.1). It starts with a  $\theta$  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.

$$\bar{p}_{SW8} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}+\frac{\theta}{2}-\frac{\alpha}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}+\frac{\theta}{2}-\frac{\alpha}{2}} r \sin\left(\frac{\alpha}{2}\right) - r \cos\left(\frac{\theta}{2} + x_2\right) dx_2 + \int_{\theta}^{\frac{\alpha}{2}+\theta} r \sin\left(\frac{\alpha}{2}\right) dx_3 \right) \quad \text{eqn S45}$$

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

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

$$\bar{p}_{SW9} = \frac{1}{\pi} \left( \int_{\frac{\pi}{2}-\frac{\theta}{2}}^{\frac{\pi}{2}} 2r \sin\left(\frac{\theta}{2}\right) \sin(x_2) dx_2 + \int_{\theta}^{\frac{\alpha}{2}} r \sin(x_3) dx_3 + \int_{\frac{\alpha}{2}}^{\frac{\alpha}{2}+\theta} r \sin\left(\frac{\alpha}{2}\right) dx_3 \right) \quad \text{eqn S47}$$

$$\bar{p}_{SW9} = \frac{r}{\pi} \left( \theta \sin\left(\frac{\alpha}{2}\right) - \cos\left(\frac{\alpha}{2}\right) + 1 \right) \quad \text{eqn S48}$$

## REFERENCES

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- SymPy Development Team (2014) *SymPy: Python library for symbolic mathematics*. 15
- Yapp, W. (1956) The theory of line transects. *Bird study*, **3**, 93–104. 3

## S3. SUPPLEMENTARY SCRIPT: SYMBOLIC ALGEBRA PYTHON SCRIPT

This script uses the SymPy package (SymPy Development Team, 2014), 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 supplementaryS3Python.py. It will also be hosted and given a DOI on Github at the time of publication.

```

1  """
2  S3
3  Supplementary Python script from
4  A generalised random encounter model for estimating animal density with remote sensor data
5  Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
6  """
7
8  """
9  Systematic analysis of REM models
10 Tim Lucas
11 01/10/13
12 """
13
14
15
16 """
17
18 This script contains:
19 1. The integration of all models (lines 24 - 603)
20 2. Some tests that the models are correct (606 - 738)
21 3. A python function to calculate  $\bar{p}$  given any parameters (743 - 763)
22 4. Creation of a plot of  $\bar{p}$  over all parameter space (767 - 794)
23 5. Code to create supplementaryS4.R, an R implementation of the model (797 - 873)
24
25 """
26
27 from sympy import *
28 import numpy as np
29 import matplotlib.pyplot as plt
30 from datetime import datetime
31 import os as os
32
33 # Set working directory
34 os.chdir('supplementary-material')
35
36 # Use LaTeX printing
37 from sympy import init_printing ;
38 init_printing()
39 # Make LaTeX output white. Because I use a dark theme
40 init_printing(forecolor="White")
41
42
43 # Load symbols used for symbolic maths
44 t, a, r, x2, x3, x4, x1 = symbols('theta alpha r x_2 x_3 x_4 x_1', positive=True)
45 r1 = {r:1} # useful for lots of checks
46
47
48
49 # Define functions
50 # Calculate the final profile averaged over pi.
51 def calcModel(model):
52     x = pi**1 * sum( [integrate(m[0], m[1:]) for m in model] ).simplify().trigsimp()
53     return x
54
55 # Do the replacements fit within the area defined by the conditions?
56 def confirmReplacements(conds, reps):
57     if not all([c.subs(reps) for c in eval(conds)]):
58         print('reps' + conds[4:] + ' incorrect')
59
60 # is average profile in range 0r-2r?
61 def profileRange(prof, reps):
62     if not 0 <= eval(prof).subs(dict(reps, **r1)) <= 2:
63         print('Total ' + prof + ' not in 0, 2r')
64
65 # Are the individuals integrals >0r
66 def intsPositive(model, reps):
67     m = eval(model)
68     for i in range(len(m)):
69         if not integrate(m[i][0], m[i][1:]).subs(dict(reps, **r1)) > 0:
70             print('Integral ' + str(i+1) + ' in ' + model + ' is negative')
71
72 # Are the individual averaged integrals between 0 and 2r
73 def intsRange(model, reps):
74     m = eval(model)
75     for i in range(len(m)):

```

```

76         if not o <= (integrate(m[i][0], m[i][1:])/m[i][3]-m[i][2])).subs(dict(reps, **r1)) <=
77             2:
78                 print('Integral ' + str(i+1) + ' in ' + model + ' has averaged integral outside
79                     0<p<2r')
80 # Are the bounds the correct way around
81 def checkBounds(model, reps):
82     m = eval(model)
83     for i in range(len(m)):
84         if not (m[i][3]-m[i][2]).subs(reps) > 0:
85             print('Bounds ' + str(i+1) + ' in ' + model + ' has lower bounds bigger than
86                 upper bounds')
87 # create latex strings with the 1) the integral equation that defines it and 2) the final calculated
88     model.
89 # There's some if statements to split longer equations on two lines and get +s in the right place.
90 def parseLaTeX(prof):
91     m = eval( 'm' + prof[1:] )
92
93     f = open('/latexFiles/'+prof+'.tex', 'w')
94     f.write('\begin{align}\n    \bar{p}_\text{\tiny{ ' + prof[1:] + ' }} = &\frac{1}{\pi} \left
95         (\; ; \; )
96         for i in range(len(m)):
97             # Roughly try and prevent expressions beginning with minus signs.
98             if latex(m[i][2])[0]=='-':
99                 o1 = 'rev-lex'
100             else:
101                 o1 = 'lex'
102
103             if latex(m[i][3])[0]=='-':
104                 o2 = 'rev-lex'
105             else:
106                 o2 = 'lex'
107
108             if latex(m[i][0])[0]=='-':
109                 o3 = 'rev-lex'
110             else:
111                 o3 = 'lex'
112
113             if latex(m[i][1])[0]=='-':
114                 o4 = 'rev-lex'
115             else:
116                 o4 = 'lex'
117
118             f.write('\int\limits_{'+latex(m[i][2], order=o1)+'}^{' + latex(m[i][3], order=o2)+'}' +
119                 latex(m[i][0], order=o3)+'\; \mathrm{d}' + latex(m[i][1], order=o4))
120             if len(m)>3 and i==(len(m)/2)-1:
121                 f.write( '\right.\notag\\\n &\left.' )
122             if i<len(m)-1:
123                 f.write(' + ')
124             f.write('\right)\label{' + prof + 'Def}\\\\n    ')
125             f.write('\bar{p}_\text{\tiny{ ' + prof[1:] + ' }} = &' + latex(eval(prof)) + '\label{' +
126                 prof + 'Sln}\n\end{align}')
127             f.close()
128
129 # Apply all checks.
130 def allChecks(prof):
131     model = 'm' + prof[1:]
132     reps = eval('rep' + prof[1:])
133     conds = 'cond' + prof[1:]
134     confirmReplacements(conds, reps)
135     profileRange(prof, reps)
136     intsPositive(model, reps)
137     intsRange(model, reps)
138     checkBounds(model, reps)
139
140 #####
141 ### Define and solve all models
142 #####
143 # NE1 animal: a = 2*pi. sensor: t > pi, a > 3pi - t #
144
145 mNE1 = [ [2*r, x1, pi/2, t/2 ],
146          [r + r*cos(x1 - t/2), x1, t/2, pi ],
147          [r + r*cos(x1 + t/2), x1, pi, 2*pi-t/2 ],
148          [2*r, x1, 2*pi-t/2, 3*pi/2 ] ]
149
150 # Replacement values in range
151 repNE1 = {t:3*pi/2, a:2*pi}
152
153 # Define conditions for model
154 condNE1 = [pi <= t, a >= 3*pi - t]
155
156 # Calculate model, run checks, write output.
157 pNE1 = calcModel(mNE1)
158 allChecks('pNE1')

```



```

156 parseLaTeX('pNE1')
157
158
159 # NE2 animal: a > pi. sensor: t > pi Condition: a < 3pi - t, a > 4pi - 2t #
160
161 mNE2 = [ [2*r, x1, pi/2, t/2 ],
162          [r + r*cos(x1 - t/2), x1, t/2, 5*pi/2 - t/2 - a/2 ],
163          [r + r*cos(x1 + t/2), x1, 5*pi/2 - t/2 - a/2, 2*pi-t/2 ],
164          [2*r, x1, 2*pi-t/2, 3*pi/2 ] ]
165
166 # Replacement values in range
167 repNE2 = {t:5*pi/3, a:4*pi/3-0.1}
168
169 # Define conditions for model
170 condNE2 = [pi <= t, a >= pi, a <= 3*pi - t, a >= 4*pi - 2*t]
171
172 # Calculate model, run checks, write output.
173 pNE2 = calcModel(mNE2)
174 allChecks('pNE2')
175 parseLaTeX('pNE2')
176
177
178 # NE3 animal: a > pi. sensor: t > pi Condition: a < 4pi - 2t #
179
180 mNE3 = [ [2*r, x1, pi/2, t/2 ],
181          [r + r*cos(x1 - t/2), x1, t/2, t/2 + pi/2 ],
182          [r, x1, t/2 + pi/2, 5*pi/2 - t/2 - a/2 ],
183          [r + r*cos(x1 + t/2), x1, 5*pi/2 - t/2 - a/2, 2*pi-t/2 ],
184          [2*r, x1, 2*pi-t/2, 3*pi/2 ] ]
185
186 # Replacement values in range
187 repNE3 = {t:5*pi/4-0.1, a:3*pi/2}
188
189 # Define conditions for model
190 condNE3 = [pi <= t, a >= pi, a <= 4*pi - 2*t]
191
192 # Calculate model, run checks, write output.
193 pNE3 = calcModel(mNE3)
194 allChecks('pNE3')
195 parseLaTeX('pNE3')
196
197
198 # NW1 animal: a = 2*pi. sensor: pi/2 <= t <= pi #
199
200 mNW1 = [ [2*r*sin(t/2)*sin(x2), x2, t/2, pi/2 ],
201          [r - r*cos(x4 - t), x4, 0, t - pi/2 ],
202          [r, x4, t - pi/2, pi/2 ],
203          [r - r*cos(x4), x4, pi/2, t ],
204          [2*r*sin(t/2)*sin(x2), x2, t/2, pi/2 ] ]
205
206 # Replacement values in range
207 repNW1 = {t:3*pi/4}
208
209 # Define conditions for model
210 condNW1 = [pi/2 <= t, t <= pi]
211
212 # Calculate model, run checks, write output.
213 pNW1 = calcModel(mNW1)
214 allChecks('pNW1')
215 parseLaTeX('pNW1')
216
217
218
219
220 # NW2 animal: a > pi. Sensor: pi/2 <= t <= pi. Condition: a > 2pi - t #
221
222 mNW2 = [ [2*r*sin(t/2)*sin(x2), x2, t/2, pi/2 ],
223          [r - r*cos(x4 - t), x4, 0, t - pi/2 ],
224          [r, x4, t - pi/2, 3*pi/2 - a/2],
225          [r - r*cos(x4), x4, 3*pi/2 - a/2, t ],
226          [2*r*sin(t/2)*sin(x2), x2, t/2, pi/2 ] ]
227
228
229 repNW2 = {t:3*pi/4, a:15*pi/8} # Replacement values in range
230
231 # Define conditions for model
232 condNW2 = [a > pi, pi/2 <= t, t <= pi, a >= 3*pi - 2*t]
233
234 # Calculate model, run checks, write output.
235 pNW2 = calcModel(mNW2)
236 allChecks('pNW2')
237 parseLaTeX('pNW2')
238
239
240
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 ],
244           [r - r*cos(x4 - t), x4, 0, t - pi/2 ],
245           [r, x4, t - pi/2, t ],
246           [r*cos(x2 - t/2), x2, t/2, 3*pi/2 - a/2 - t/2],
247           [2*r*sin(t/2)*sin(x2), x2, 3*pi/2 - a/2 - t/2, pi/2 ] ]
248
249
250 repNW3 = {t:5*pi/8, a:6*pi/4} # Replacement values in range
251
252 # Define conditions for model
253 condNW3 = [a > pi, pi/2 <= t, t <= pi, 2*pi - t <= a, a <= 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           [r - r*cos(x4 - t), x4, 0, t - pi/2],
266           [r, x4, t - pi/2, t],
267           [r*cos(x2 - t/2), x2, t/2, a/2 + t/2 - pi/2] ]
268
269 repNW4 = {t:3*pi/4, a:9*pi/8} # Replacement values in range
270
271 # Define conditions for model
272 condNW4 = [a > pi, pi/2 <= t, t <= pi, a <= 2*pi - t]
273
274 # Calculate model, run checks, write output.
275 pNW4 = calcModel(mNW4)
276 allChecks('pNW4')
277 parseLaTeX('pNW4')
278
279
280 # REM animal: a=2pi. Sensor: t <= pi/2. #
281
282 mREM = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
283           [r*sin(x3), x3, t, pi/2],
284           [r, x4, 0*t, t],
285           [r*sin(x3), x3, t, pi/2],
286           [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2] ]
287
288
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
294 # Calculate model, run checks, write output.
295 pREM = calcModel(mREM)
296 allChecks('pREM')
297 parseLaTeX('pREM')
298
299
300
301 # NW5 animal: a>pi. Sensor: t <= pi/2. Condition: 2*pi - t < a #
302
303
304 mNW5 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
305           [r*sin(x3), x3, t, pi/2],
306           [r, x4, 0, t],
307           [r*sin(x3), x3, t, pi/2],
308           [r*cos(x2 - t/2), x2, pi/2 - t/2, 3*pi/2 - t/2 - a/2],
309           [2*r*sin(t/2)*sin(x2), x2, 3*pi/2 - t/2 - a/2, pi/2] ]
310
311
312 repNW5 = {t:3*pi/8, a:29*pi/16} # Replacement values in range
313
314 # 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)
319 allChecks('pNW5')
320 parseLaTeX('pNW5')
321
322
323 # NW6 animal: a>pi. Sensor: t <= pi/2. Condition: 2*pi - 2*t <= a <= 2*pi - t #
324
325
326 mNW6 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2],
327           [r*sin(x3), x3, t, pi/2],
328           [r, x4, 0, t],
329           [r*sin(x3), x3, t, pi/2],

```

```

330     [r*cos(x2 - t/2),      x2, pi/2 - t/2, a/2 + t/2 - pi/2] ]
331
332 repNW6 = {t:3*pi/8, a:3*pi/2} # Replacement values in range
333
334 # Define conditions for model
335 condNW6 = [a >= pi, t <= pi/2, 2*pi - 2*t <= a, a <= 2*pi - t]
336
337 # 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          [r,                    x4, 0,             t ],
350          [r*sin(x3),             x3, pi - a/2,      pi/2] ]
351
352
353 repNW7 = {t:pi/9, a:10*pi/9} # Replacement values in range
354
355 # Define conditions for model
356 condNW7 = [t <= pi/2, a >= pi, a <= 2*pi - 2*t]
357
358 # Calculate model, run checks, write output.
359 pNW7 = calcModel(mNW7)
360 allChecks('pNW7')
361 parseLaTeX('pNW7')
362
363
364
365 # SE1 animal: a <= pi.  Sensor: t =2pi.  #
366
367 mSE1 = [ [ 2*r*sin(a/2),x1, pi/2, 3*pi/2      ],
368          ]
369
370
371 repSE1 = {a:pi/4} # Replacement values in range
372
373 # Define conditions for model
374 condSE1 = [a <= pi]
375
376 # Calculate model, run checks, write output.
377 pSE1 = calcModel(mSE1)
378 allChecks('pSE1')
379 parseLaTeX('pSE1')
380
381
382
383
384 # SE2 animal: a <= pi.  Sensor: t > pi. Condition: a > 2pi - t, a > 4pi - 2t  #
385
386 mSE2 = [ [ 2*r*sin(a/2),          x1, pi/2,          t/2 + pi/2 - a/2      ],
387          [ r*sin(a/2) + r*cos(x1 - t/2), x1, t/2 + pi/2 - a/2, 5*pi/2 - a/2 - t/2 ],
388          [ 2*r*sin(a/2),          x1, 5*pi/2 - a/2 - t/2, 3*pi/2] ]
389
390
391 repSE2 = {t:19*pi/10, a:pi/2} # Replacement values in range
392
393 # Define conditions for model
394 condSE2 = [a <= pi, t >= pi, a >= 4*pi - 2*t]
395
396 # Calculate model, run checks, write output.
397 pSE2 = calcModel(mSE2)
398 allChecks('pSE2')
399 parseLaTeX('pSE2')
400
401
402 # SE3 animal: a <= pi.  Sensor: t > pi. Condition: 2pi - t < a < 4pi - 2t #
403
404 mSE3 = [ [ 2*r*sin(a/2),          x1, pi/2,          t/2 + pi/2 - a/2 ],
405          [ r*sin(a/2) + r*cos(x1 - t/2), x1, t/2 + pi/2 - a/2, t/2 + pi/2 ],
406          [ r*sin(a/2),             x1, t/2 + pi/2,      5*pi/2 - a/2 - t/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
411 # Define conditions for model
412 condSE3 = [a <= pi, t >= pi, a >= 2*pi - t, a <= 4*pi - 2*t]
413
414 # Calculate model, run checks, write output.
415 pSE3 = calcModel(mSE3)
416 allChecks('pSE3')

```

```

417 parseLaTeX('pSE3')
418
419
420 # SE4 animal: a <= pi. Sensor: t > pi. Condition: a <= 4*pi - 2*t and a < 2*pi - t #
421
422
423 mSE4 = [ [ 2*r*sin(a/2), x1, pi/2, t/2 + pi/2 - a/2 ],
424 [ r*sin(a/2) + r*cos(x1 - t/2), x1, t/2 + pi/2 - a/2, t/2 + pi/2 ],
425 [ r*sin(a/2), x1, t/2 + pi/2, t/2 + pi/2 + a/2 ] ]
426
427
428 repSE4 = {t:3*pi/2, a:pi/3} # Replacement values in range
429
430
431 # Define conditions for model
432 condSE4 = [a <= pi, t >= pi/2, a <= 4*pi - 2*t, a <= 2*pi - t]
433
434 # Calculate model, run checks, write output.
435 pSE4 = calcModel(mSE4)
436 allChecks('pSE4')
437 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 mSW1 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - a/2 + t/2, pi/2 ],
443 [r*sin(a/2) - r*cos(x2 + t/2), x2, t/2, pi/2 - a/2 + t/2],
444 [r*sin(a/2) - r*cos(x4 - t), x4, 0, t - pi/2 ],
445 [r*sin(a/2), x4, t-pi/2, t - pi/2 + a/2 ] ]
446
447
448 repSW1 = {t:5*pi/8, a:6*pi/8} # Replacement values in range
449
450 # Define conditions for model
451 condSW1 = [a <= pi, pi/2 <= t, t <= pi, a >= t, a/2 >= t - pi/2]
452
453 # Calculate model, run checks, write output.
454 pSW1 = calcModel(mSW1)
455 allChecks('pSW1')
456 parseLaTeX('pSW1')
457
458
459 # SW2 animal: a <= pi. Sensor: pi/2 <= t <= pi. Condition: a <= t and a/2 >= t - pi/2 #
460
461 mSW2 = [ [2*r*sin(a/2), x2, pi/2 + a/2 - t/2, pi/2 ],
462 [r*sin(a/2) - r*cos(x2 + t/2), x2, t/2, pi/2 + a/2 - t/2],
463 [r*sin(a/2) - r*cos(x4 - t), x4, 0*t, t - pi/2 ],
464 [r*sin(a/2), x4, t - pi/2, t - pi/2 + a/2 ] ]
465
466
467 repSW2 = {t:7*pi/8, a:7*pi/8-0.1} # Replacement values in range
468
469 # Define conditions for model
470 condSW2 = [a <= pi, pi/2 <= t, t <= pi, a/2 <= t/2, a/2 >= t - pi/2]
471
472 # Calculate model, run checks, write output.
473 pSW2 = calcModel(mSW2)
474 allChecks('pSW2')
475 parseLaTeX('pSW2')
476
477
478
479 # SW3 animal: a <= pi. Sensor: pi/2 <= t <= pi. Condition: a <= t and a/2 <= t - pi/2 #
480
481 mSW3 = [ [2*r*sin(a/2), x2, t/2, pi/2 ],
482 [2*r*sin(a/2), x4, 0, t - pi/2 - a/2 ],
483 [r*sin(a/2) - r*cos(x4 - t), x4, t - pi/2 - a/2, t - pi/2 ],
484 [r*sin(a/2), x4, t - pi/2, t - pi/2 + a/2 ] ]
485
486
487 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 #
499
500 mSW4 = [ [2*r*sin(a/2), x2, pi/2 - t/2 + a/2, pi/2 ],
501 [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), x4, 0, a/2 + t - pi/2 ] ]

```

```

504
505 repSW4 = {t:pi/2-0.1, a:pi/4} # Replacement values in range
506
507 # Define conditions for model
508 condSW4 = [a <= pi, t <= pi/2, a >= pi - 2*t, a <= t]
509
510 # Calculate model, run checks, write output.
511 pSW4 = calcModel(mSW4)
512 allChecks('pSW4')
513 parseLaTeX('pSW4')
514
515
516 # SW5 animal: a <= pi. Sensor: t <= pi/2. Condition: a > pi - 2t & t <= a <= 2t #
517
518 mSW5 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 + t/2 - a/2, pi/2 ],
519          [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2, pi/2 + t/2 - a/2],
520          [r*sin(a/2), x3, t, pi/2 ],
521          [r*sin(a/2), x4, 0, a/2 + t -pi/2 ] ]
522
523
524 repSW5 = {t:pi/2-0.1, a:pi/2} # Replacement values in range
525
526 # define conditions for model
527 condSW5 = [a <= pi, t <= pi/2, a >= pi - 2*t, t <= a, a <= 2*t]
528
529
530 # 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 ],
539          [r*sin(x3), x3, t, a/2 ],
540          [r*sin(a/2), x3, a/2, pi/2 ],
541          [r*sin(a/2), x4, 0, a/2 + t -pi/2 ] ]
542
543
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]
549
550 # Calculate model, run checks, write output.
551 pSW6 = calcModel(mSW6)
552 allChecks('pSW6')
553 parseLaTeX('pSW6')
554
555
556 # SW7 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & a <= t #
557
558 mSW7 = [ [2*r*sin(a/2), x2, pi/2 - t/2 + a/2, pi/2 ],
559          [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2, pi/2 - t/2 + a/2],
560          [r*sin(a/2), x3, t, t + a/2 ] ]
561
562
563 repSW7 = {t:2*pi/8, a:pi/8} # Replacement values in range
564
565 # Define conditions for model
566 condSW7 = [a <= pi, t <= pi/2, a <= pi - 2*t, a <= t]
567
568 # Calculate model, run checks, write output.
569 pSW7 = calcModel(mSW7)
570 allChecks('pSW7')
571 parseLaTeX('pSW7')
572
573
574 # SW8 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & t <= a <= 2t #
575
576 mSW8 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 + t/2 - a/2, pi/2 ],
577          [r*sin(a/2) - r*cos(x2 + t/2), x2, pi/2 - t/2, pi/2 + t/2 - a/2],
578          [r*sin(a/2), x3, t, t + a/2 ] ]
579
580 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]
584
585 # Calculate model, run checks, write output.
586 pSW8 = calcModel(mSW8)
587 allChecks('pSW8')
588 parseLaTeX('pSW8')
589
590

```

```

591 # SW9 animal: a <= pi. Sensor: t <= pi/2. Condition: a <= pi - 2t & 2t <= a #
592
593 mSW9 = [ [2*r*sin(t/2)*sin(x2), x2, pi/2 - t/2, pi/2 ],
594          [r*sin(x3), x3, t, a/2 ],
595          [r*sin(a/2), x3, a/2, t + a/2 ] ]
596
597
598 repSW9 = {t:l*pi/8, a:pi/2} # Replacement values in range
599
600 # Define conditions for model
601 condSW9 = [a <= pi, t <= pi/2, a <= pi - 2*t, 2*t <= a]
602
603 # Calculate model, run checks, write output.
604 pSW9 = calcModel(mSW9)
605 allChecks('pSW9')
606 parseLaTeX('pSW9')
607
608 #####
609 ## Run tests ###
610 #####
611
612 # create gas model object
613 gas = 2*r
614
615
616
617 # for each model run through every adjacent model.
618 # Contains duplicates but better for avoiding missed comparisons.
619 # Also contains replacement t->a and a->t just in case.
620
621
622 allComps = [
623 ['gas', 'pNE1', {t:2*pi}], ['gas', 'pSE1', {a:pi}],
624
625 ['pNE1', 'gas', {t:2*pi}], ['pNE1', 'pNW1', {t:pi}],
626 ['pNE1', 'pNE2', {a:3*pi-t}], ['pNE1', 'pNE2', {t:3*pi-a}],
627
628 ['pNE2', 'pNE1', {a:3*pi-t}], ['pNE2', 'pNE1', {t:3*pi-a}],
629 ['pNE2', 'pNE3', {a:4*pi-2*t}], ['pNE2', 'pNE3', {t:2*pi-a/2}],
630 ['pNE2', 'pSE2', {a:pi}],
631
632 ['pNE3', 'pNE2', {a:4*pi-2*t}], ['pNE3', 'pNE2', {t:2*pi-a/2}],
633 ['pNE3', 'pSE3', {a:pi}], ['pNE3', 'pNW2', {t:pi}],
634
635 ['pNW1', 'pNE1', {t:pi}], ['pNW1', 'pNW2', {a:2*pi}],
636
637 ['pNW2', 'pNE3', {t:pi}], ['pNW2', 'pNW3', {a:3*pi-2*t}],
638 ['pNW2', 'pNW3', {t:3*pi/2-a/2}], ['pNW2', 'pNW1', {a:2*pi}],
639
640 ['pNW3', 'pNW5', {t:pi/2}], ['pNW3', 'pNW4', {a:2*pi-t}],
641 ['pNW3', 'pNW4', {t:2*pi-a}], ['pNW3', 'pNW2', {a:3*pi-2*t}],
642 ['pNW3', 'pNW2', {t:3*pi/2-a/2}],
643
644 ['pNW4', 'pNW6', {t:pi/2}], ['pNW4', 'pNW3', {t:2*pi-a}],
645 ['pNW4', 'pNW3', {a:2*pi-t}], ['pNW4', 'pSW1', {a:pi}],
646
647 ['pREM', 'pNW1', {t:pi/2}], ['pREM', 'pNW5', {a:2*pi}],
648
649 ['pNW5', 'pREM', {a:2*pi}], ['pNW5', 'pNW6', {a:2*pi-t}],
650 ['pNW5', 'pNW6', {t:2*pi-a}], ['pNW5', 'pNW3', {t:pi/2}],
651
652 ['pNW6', 'pNW5', {a:2*pi-t}], ['pNW6', 'pNW5', {t:2*pi-a}],
653 ['pNW6', 'pNW7', {t:pi-a/2}], ['pNW6', 'pNW7', {a:2*pi-2*t}],
654 ['pNW5', 'pNW4', {t:pi/2}],
655
656 ['pNW7', 'pNW6', {t:2*pi-2*a}], ['pNW7', 'pNW6', {a:2*pi-2*t}],
657 ['pNW7', 'pSW6', {a:pi}],
658
659 ['pSE1', 'pSE2', {t:2*pi}], ['pSE1', 'gas', {a:pi}],
660
661 ['pSE2', 'pSE3', {t:2*pi-a/2}], ['pSE2', 'pSE3', {a:4*pi-2*t}],
662 ['pSE2', 'pSE1', {t:2*pi}], ['pSE2', 'pNE2', {a:pi}],
663
664 ['pSE3', 'pSE2', {a:4*pi-2*t}], ['pSE3', 'pSE2', {t:2*pi-a/2}],
665 ['pSE3', 'pSE4', {a:2*pi-t}], ['pSE3', 'pSE4', {t:2*pi-a}],
666 ['pSE3', 'pNE3', {a:pi}],
667
668 ['pSE4', 'pSE3', {t:2*pi-a}], ['pSE4', 'pSE3', {a:2*pi-t}],
669 ['pSE4', 'pSW3', {t:pi}],
670
671 ['pSW1', 'pSW5', {t:pi/2}], ['pSW1', 'pSW2', {a:t}],
672 ['pSW1', 'pSW2', {t:a}], ['pSW1', 'pNW4', {a:pi}],
673
674 ['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}],
677

```

```

678 ['pSW3', 'pSW2', {t:a/2+pi/2}], ['pSW3', 'pSW2', {a:2*t-pi}],
679 ['pSW3', 'pSE4', {t:pi}],
680
681
682 ['pSW4', 'pSW7', {a:pi-2*t}], ['pSW4', 'pSW7', {t:pi/2-a/2}],
683 ['pSW4', 'pSW5', {t:a}], ['pSW4', 'pSW5', {a:t}],
684 ['pSW4', 'pSW2', {t:pi/2}],
685
686 ['pSW5', 'pSW4', {t:a}], ['pSW5', 'pSW4', {a:t}],
687 ['pSW5', 'pSW8', {t:pi/2-a/2}], ['pSW5', 'pSW8', {a:pi-2*t}],
688 ['pSW5', 'pSW6', {a:2*t}], ['pSW5', 'pSW6', {t:a/2}],
689 ['pSW5', 'pSW1', {t:pi/2}],
690
691 ['pSW6', 'pSW9', {t:pi/2-a/2}], ['pSW6', 'pSW9', {a:pi-2*t}],
692 ['pSW6', 'pSW5', {a:2*t}], ['pSW6', 'pSW5', {t:a/2}],
693 ['pSW6', 'pNW7', {a:pi}],
694
695
696 ['pSW7', 'pSW8', {t:a}], ['pSW7', 'pSW8', {a:t}],
697 ['pSW7', 'pSW4', {t:pi/2-a/2}], ['pSW7', 'pSW4', {a:pi-2*t}],
698
699 ['pSW8', 'pSW7', {a:t}], ['pSW8', 'pSW7', {t:a}],
700 ['pSW8', 'pSW9', {a:2*t}], ['pSW8', 'pSW9', {t:a/2}],
701 ['pSW8', 'pSW5', {a:pi-2*t}], ['pSW8', 'pSW5', {t:pi/2-a/2}],
702
703 ['pSW9', 'pSW8', {a:2*t}], ['pSW9', 'pSW8', {t:a/2}],
704 ['pSW9', 'pSW6', {a:pi-2*t}], ['pSW9', 'pSW6', {t:pi/2-a/2}]
705 ]
706
707
708 # List of regions that touch a=0. Should equal 0 when a=0.
709 zeroRegions = ['pSW9', 'pSW8', 'pSW7', 'pSW4', 'pSW2', 'pSW3', 'pSE4', 'pSE3', 'pSE1']
710
711
712 # Run through all the comparisons. Need simplify(). Even together() gives some false negatives.
713
714 checkFile = open('checksFile.tex', 'w')
715
716 checkFile.write('All checks evaluated.\nTim Lucas - ' + str(datetime.now()) + '\n')
717 for i in range(len(allComps)):
718     if (eval(allComps[i][0]).subs(allComps[i][2]) - eval(allComps[i][1]).subs(allComps[i][2])):
719         simplify() == 0:
720             checkFile.write(str(i) + ': ' + allComps[i][0] + ' and ' + allComps[i][1] + ': OK\n')
721         else:
722             checkFile.write(str(i) + ': ' + allComps[i][0] + ' and ' + allComps[i][1] + ': Incorrect\n')
723
724 for i in range(len(zeroRegions)):
725     if eval(zeroRegions[i]).subs({a:0}).simplify() == 0:
726         checkFile.write(zeroRegions[i] + ' at a = 0: OK\n')
727     else:
728         checkFile.write(zeroRegions[i] + ' at a = 0: Incorrect\n')
729
730 # pSE2 is slightly different. Only one corner touches a=0, so need theta value as well. I'm not sure why
731 # this isn't
732 # A problem for some other regions.
733 if pSE2.subs({a:0, t:2*pi}) == 0:
734     checkFile.write('pSE2 at a = 0, t = 2pi: OK\n')
735 else:
736     checkFile.write('pSE2 at a = 0, t = 2pi: Incorrect\n')
737
738 checkFile.close()
739
740 # And print to terminal
741 #for i in range(len(allComps)):
742 #    if not (eval(allComps[i][0]).subs(allComps[i][2]) - eval(allComps[i][1]).subs(allComps[i][2])):
743 #        simplify() == 0:
744 #            print allComps[i][0] + ' and ' + allComps[i][1] + ': Incorrect\n'
745
746 #####
747 ### Define a a function that calculates p bar answer. #####
748 #####
749
750 def calcP(A, T, R):
751     assert (A <= 2*pi and A >= 0), "a is out of bounds. Should be in 0<a<2*pi"
752     assert (T <= 2*pi and T >= 0), "s is out of bounds. Should be in 0<s<2*pi"
753
754     if A > pi:
755         if A < 4*pi - 2*T:
756             p = pNW7.subs({a:A, t:T, r:R}).n()
757         elif A <= 3*pi - T:
758             p = pNE2.subs({a:A, t:T, r:R}).n()
759         else:
760             p = pNE1.subs({a:A, t:T, r:R}).n()
761     else:
762         if A < 4*pi - 2*T:
763             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 ## Apply to entire grid   ###
768 #####
769
770 # How many values for each parameter
771 nParas = 100
772
773 # Make a vector for a and s. Make an empty nParas x nParas array.
774 # Calculated profile sizes will go in pArray
775 tVec = np.linspace(0, 2*pi, nParas)
776 aVec = np.linspace(0, 2*pi, nParas)
777 pArray = np.zeros((nParas,nParas))
778
779 # Calculate profile size for each combination of parameters
780 for i in range(nParas):
781     for j in range(nParas):
782         pArray[i][j] = calcP(aVec[i], tVec[j], 1)
783
784 # Turn the array upside down so origin is at bottom left.
785 pImage = np.flipud(pArray)
786
787 # Plot and save.
788 pl.imshow(pImage, interpolation='none', cmap=pl.get_cmap('Blues'))
789
790 # Show or save image.
791 # pl.show()
792 # pl.savefig('/imgs/profilesCalculated.png')
793
794 #####
795 ##### Output R function. #####
796 #####
797
798 # To reduce mistakes, output R function directly from python.
799 # However, the if statements, which correspond to the bounds of each model, are not automatic.
800
801 Rfunc = open('supplementaryRscript.R', 'w')
802
803 Rfunc.write("""
804 # S4
805 # Supplementary R script from
806 # A generalised random encounter model for estimating animal density with remote sensor data
807 # Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
808 #
809 # calcDensity is the main function to calculate density.
810 # It takes parameters z, alpha, theta, r, animalSpeed, t
811 # z - The number of camera/acoustic counts or captures.
812 # alpha - Call width in radians.
813 # theta - Sensor width in radians.
814 # r - Sensor range in metres.
815 # animalSpeed - Average animal speed in metres per second.
816 # t - Length of survey in sensor seconds i.e. number of sensors x survey duration.
817 #
818 # calcAbundance calculates abundance rather than density and requires an extra parameter
819 # area - In metres squared. The size of the region being examined.
820
821 # Internal function to calculate profile width as described in the text
822 calcProfileWidth <- function(alpha, theta, r){
823     if(alpha > 2*pi | alpha < 0)
824         stop('alpha is out of bounds. alpha should be in interval 0<a<2*pi')
825     if(theta > 2*pi | theta < 0)
826         stop('theta is out of bounds. theta should be in interval 0<a<2*pi')
827
828     if(alpha > pi){
829         if(alpha < 4*pi - 2*theta){
830             "" +
831             '
832                 p <- ' + str(pNW7) +
833                 '
834                 } else if(alpha <= 3*pi - theta){
835                 p <- ' + str(pNE2) +
836                 '
837                 } else {
838                 p <- ' + str(pNE1) +
839                 '
840                 }
841             } else {
842                 if(alpha < 4*pi - 2*theta){
843                 p <- ' + str(pSE3) +
844                 '
845                 } else {
846                 p <- ' + str(pSE2) +
847                 '
848                 }
849             }
850         }
851     }
852     return(p)
853 """

```



```

849 '\n}' +
850 """
851 # Calculate a population density. See above for units etc.
852 calcDensity <- function(z, alpha, theta, r, animalSpeed, t){
853     # Check the parameters are suitable.
854     if(z <= 0 | !is.numeric(z)) stop('Counts, z, must be a positive number.')
855     if(animalSpeed <= 0 | !is.numeric(animalSpeed)) stop('animalSpeed must be a positive number.')
856     if(t <= 0 | !is.numeric(t)) stop('Time, t, must be a positive number.')
857
858     # Calculate profile width, then density.
859     p <- calcProfileWidth(alpha, theta, r)
860     D <- z/{animalSpeed*t*p}
861     return(D)
862 }
863
864 # Calculate abundance rather than density.
865 calcAbundance <- function(z, alpha, theta, r, animalSpeed, t, area){
866     if(area <= 0 | !is.numeric(area)) stop('Area must be a positive number')
867     D <- calcDensity(z, alpha, theta, r, animalSpeed, t)
868     A <- D*area
869     return(A)
870 }
871 """
872 )
873
874 Rfunc.close()

```

supplementaryS3Python.py

## S4. SUPPLEMENTARY SCRIPT: R IMPLEMENTATION OF MODELS

This is an implementation of the models derived in the paper in R (R Development Core Team, 2010). Once given the parameters  $\theta$  and  $\alpha$  it automatically selects the correct model to apply.

The script can also be found in the plain text file supplementaryS4R.R. It will also be hosted and given a DOI on Github at the time of publication.

```

1
2 # S4
3 # Supplementary R script from
4 # A generalised random encounter model for estimating animal density with remote sensor data
5 # Tim C.D. Lucas, Elizabeth A. Moorcroft, Robin Freeman, Marcus J. Rowcliffe, Kate E. Jones
6 #
7 # calcDensity is the main function to calculate density.
8 # It takes parameters z, alpha, theta, r, animalSpeed, t
9 # z - The number of camera/acoustic counts or captures.
10 # alpha - Call width in radians.
11 # theta - Sensor width in radians.
12 # r - Sensor range in metres.
13 # animalSpeed - Average animal speed in metres per second.
14 # t - Length of survey in sensor seconds i.e. number of sensors x survey duration.
15 #
16 # calcAbundance calculates abundance rather than density and requires an extra parameter
17 # area - In metres squared. The size of the region being examined.
18
19
20 # Internal function to calculate profile width as described in the text
21 calcProfileWidth <- function(alpha, theta, r){
22   if(alpha > 2*pi | alpha < 0)
23     stop('alpha is out of bounds. alpha should be in interval 0<a<2*pi')
24   if(theta > 2*pi | theta < 0)
25     stop('theta is out of bounds. theta should be in interval 0<a<2*pi')
26
27   if(alpha > pi){
28     if(alpha < 4*pi - 2*theta){
29       p <- r*(theta - cos(alpha/2) + 1)/pi
30     } else if(alpha <= 3*pi - theta){
31       p <- r*(theta - cos(alpha/2) + cos(alpha/2 + theta))/pi
32     } else {
33       p <- r*(theta + 2*sin(theta/2))/pi
34     }
35   } else {
36     if(alpha < 4*pi - 2*theta){
37       p <- r*(theta*sin(alpha/2) - cos(alpha/2) + 1)/pi
38     } else {
39       p <- r*(theta*sin(alpha/2) - cos(alpha/2) + cos(alpha/2 + theta))/pi
40     }
41   }
42   return(p)
43 }
44 # Calculate a population density. See above for units etc.
45 calcDensity <- function(z, alpha, theta, r, animalSpeed, t){
46   # Check the parameters are suitable.
47   if(z <= 0 | !is.numeric(z)) stop('Counts, z, must be a positive number.')
48   if(animalSpeed <= 0 | !is.numeric(animalSpeed)) stop('animalSpeed must be a positive number.')
49   if(t <= 0 | !is.numeric(t)) stop('Time, t, must be a positive number.')
50
51   # Calculate profile width, then density.
52   p <- calcProfileWidth(alpha, theta, r)
53   D <- z/(animalSpeed*t*p)
54   return(D)
55 }
56
57 # Calculate abundance rather than density.
58 calcAbundance <- function(z, alpha, theta, r, animalSpeed, t, area){
59   if(area <= 0 | !is.numeric(area)) stop('Area must be a positive number')
60   D <- calcDensity(z, alpha, theta, r, animalSpeed, t)
61   A <- D*area
62   return(A)
63 }

```

supplementaryS4R.R

S5. SUPPLEMENTARY INFORMATION: SIMULATION MODEL RESULTS OF THE gREM PRECISION

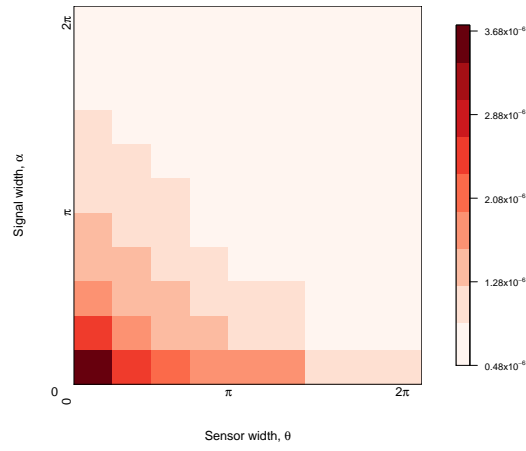


Figure S5.1. Simulation model results of the gREM precision given a range of sensor and signal widths, shown by the standard deviation of the error between the estimated and true densities. Standard deviations are shown from deep red to pink, representing high to low values between  $0.483 \times 10^{-6}$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

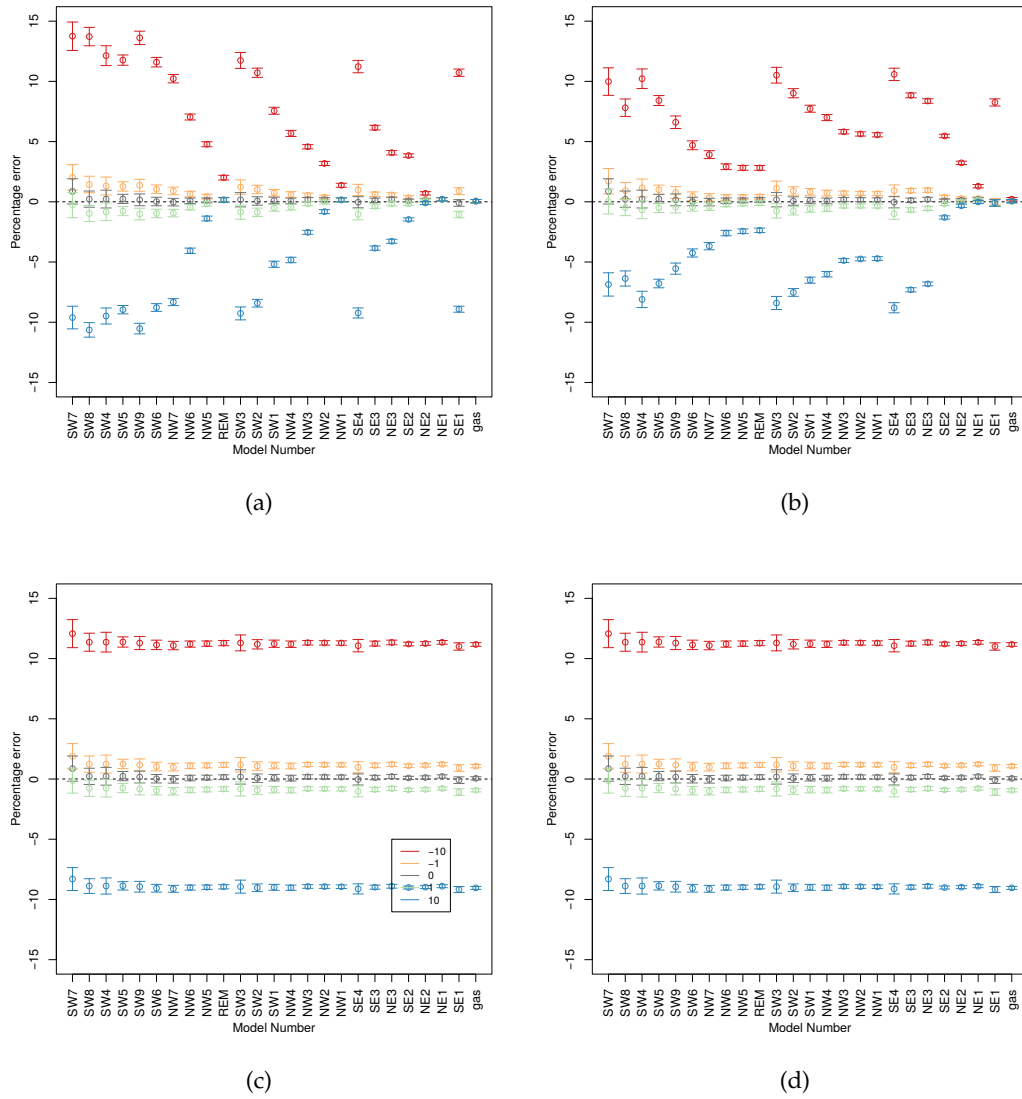


Figure S6.1. Model sensitivity (for all gREM submodels) to error in estimates of a) signal width ( $\alpha$ ) b) sensor width ( $\theta$ ), c) sensor radius ( $r$ ), d) animal movement speed ( $v$ ). Estimates are -10% (red), -1% (orange), 0% (yellow), +1% (green) and +10% (blue) of the true parameter value. The grey dashed line indicates zero error in density estimates. The error bars indicate standard error across all simulations.

## S6. SUPPLEMENTARY INFORMATION: IMPACT OF PARAMETER ERROR