



# Roe Deer show an affinity for woodland and reluctance to cross roads

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## Abstract

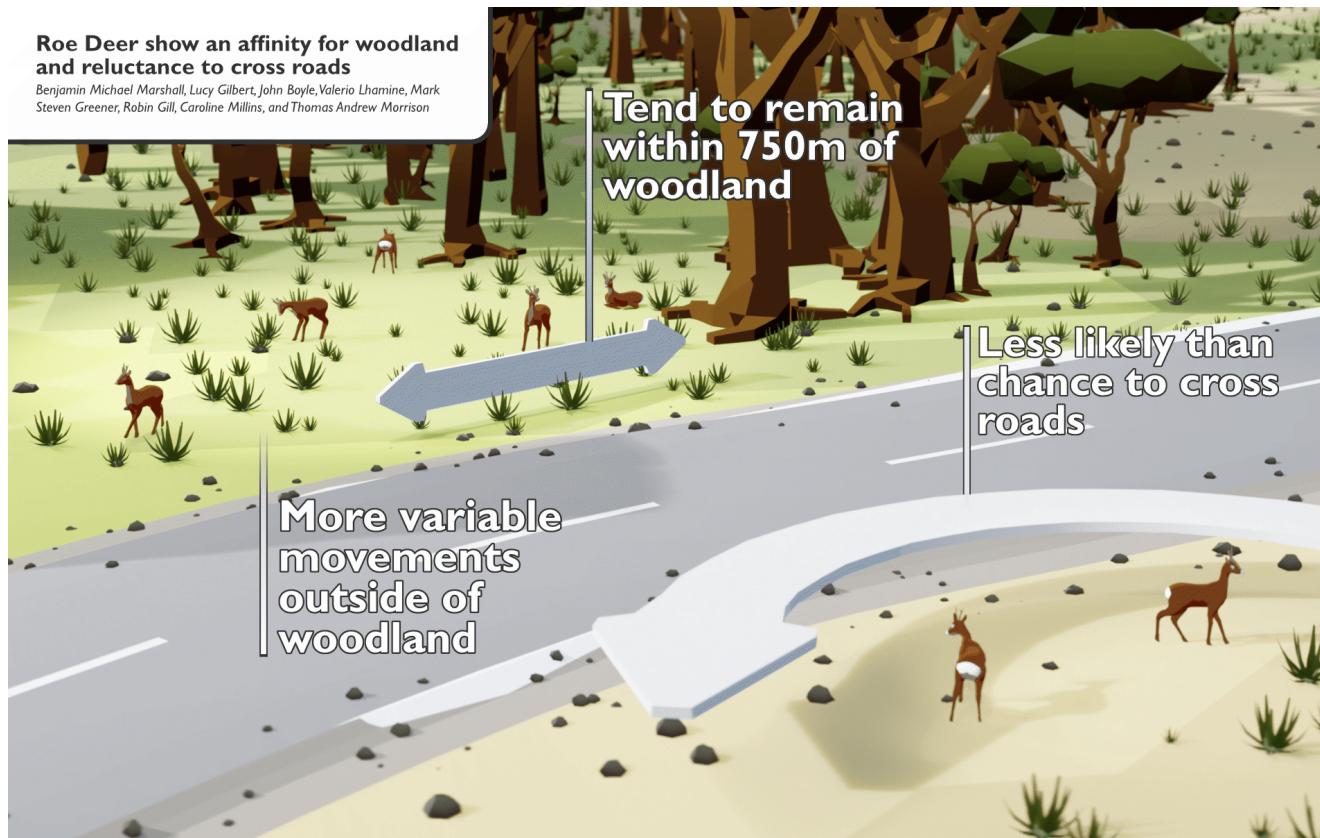
Animals use landscapes unequally and have differential responses to anthropogenic changes such as land cover modification. Predicting such responses can be challenging, requiring knowledge of animal movements. This knowledge is particularly valuable where human-animal interactions have implications for either's well-being. Large herbivores, with relatively high mobility, often come in contact with humans as habitats become more restricted leading to grazing pressures, animal-vehicle collisions, and disease transfer risks. In much of Europe, Roe Deer (*Capreolus capreolus*) are abundant herbivores often finding themselves in proximity to anthropogenic infrastructure and activity. In the United Kingdom Roe Deer are the most abundant deer species, but there have been few efforts to track their movements. By examining Roe Deer movements in two mixed use landscapes in the United Kingdom, we explore how Roe Deer home range size and how they use the available land covers. Through the GPS tracking of 15 deer, we reveal reasonably limited home ranges that are centred on woodland patches. We observed forays outside of the woodland patches into open shrub-, crop- and grass- lands, but only to distances up to 750m. When outside of the woodland, deer movements appeared to be lower but more variable, potentially indicative of a behavioural change when in open areas. We documented a consistent, albeit slight, aversion to crossing roads indicating that roads contribute to deer habitat fragmentation. Overall, our results highlight UK Roe Deer's connection to woodland patches, and suggest their movements and overall landscape connectivity are closely tied to woodland presence.

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## Keywords

Movement ecology, step selection function, habitat preference, habitat selection, animal movement, Roe Deer, *Capreolus capreolus*, road crossing, United Kingdom

## Graphical Abstract



## 1 Introduction

Animals disproportionately use and move through some areas more than others. The study of these processes, i.e., habitat selection and movement, provides insights into the dynamic relationship between an animals' physiology and the set of resources and risks occurring in landscapes (Manly, McDonald & Thomas, 1993; Gallagher et al., 2017). The balance between habitat, energy, risk, and resources determine the spatial distribution of populations, while also fundamentally underpinning demographic performance (Matthiopoulos et al., 2015; Gallagher et al., 2017). Accelerating landscape changes in the form of habitat destruction, creation and transformation increases the urgency for understanding if and how animals adjust movements to accommodate landscape changes.

Animal responses to landscapes can be hard to predict. At the global scale, wildlife movement is significantly impacted by human presence (Tucker et al., 2018). Mammals and reptiles broadly experience reductions in movement, with mobility limited by human-created barriers [e.g., roads or fencing; Tucker et al. (2018); Jerina (2012); Jones et al. (2019)]. Similarly, examinations of 586 mammal species show that home range sizes are roughly five times smaller in areas with high versus low human disturbance (Broekman et al., 2024). All else being equal, habitat fragmentation can also increase travel time for resource acquisition, resulting in higher energy expenditure (Doherty & Driscoll, 2018), shifts in diets (Redpath, 1995), and lower breeding success (Saunders, 1982).

Large herbivores, with relatively high mobility and large home ranges, often experience interactions with humans, domesticated animals, and infrastructure. Due to their economic importance and ecological impact, the space use patterns of large herbivores and their responses to anthropogenic features have been relatively well studied. Infrastructure such as roads and fences tend to degrade and fragment habitat, altering animal movement paths (Sawyer et al., 2013; Schwandner et al., 2025) and behaviour (Xu et al., 2021). However, movement responses to specific landscape features can be complex. For example, GPS-tagged Guanacos in Argentina were attracted to roads for grazing resources and low predation risk, but they strongly avoided crossing roads (Serota et al., 2024). By understanding movement responses, we may be better placed to predict and mitigate harmful interactions between wildlife and humans.

Deer live near humans in many temperate regions of the world and, as such, receive considerable attention from land managers (Cederlund, 1983; Dupke et al., 2017). In many areas, efforts are taken to control deer space use and population density, depending on the local management objectives (Pepper, Barbour & Glass, 2020). Objec-

tives include reducing deer vehicular collisions, reducing damage to agriculture through disease transmission to livestock or herbivory, reducing vectors of disease to humans (e.g., ticks such as *Ixodes ricinus*), improving forestry productivity, enhancing biodiversity and woodland regeneration, and increasing hunting conditions (Putman et al., 2011). Deer population management could be improved by more precise knowledge of the movement behaviour and space use of deer; such knowledge could provide a more complete understanding from which controversial approaches to deer management can be assessed. While there is a history of individual-level tracking of deer in mainland Europe and North America (Morellet et al., 2013; Mysterud et al., 2025), only a handful of similar studies have been carried out on British deer (10 Red Deer *Cervus elaphus* in Glenbranter Forest, South Argyll Catt & Staines (1987); 36 Reeves's Muntjac Deer *Muntiacus reevesi*, 7 Roe Deer *Capreolus capreolus* in The King's Forest, Suffolk Chapman et al. (1993); 23 Muntjac Deer, 19 Fallow Deer *Dama dama* in Rockingham Forest and Monks Wood, Northamptonshire Staines et al. (1998)), with most research attention in the UK focused on other deer species (Staines et al., 1998; Broekman et al., 2023). Comparatively, Roe Deer movements remain understudied in parts of the UK such as Scotland (Mitchell, Staines & Welch, 1977) despite being the most abundant species in the UK, living most closely with humans, and the most frequently involved in deer-vehicle collisions (Lush & Lush, 2026). The need for individual level tracking of Roe Deer is apparent to gain a fuller understanding of their habitat requirements and reactions to human activities.

Roe deer are the smallest native deer species in the United Kingdom and the most ubiquitous, covering almost all the British mainland from the northern highlands of Scotland to the south coast of England (Staines et al., 1998; Burbaić & Csányi, 2009). Roe deer are woodland edge species that use isolated fragments of woodland, open or cultivated habitats near woody cover, and even gardens in and near urban areas (Staines et al., 1998). Their flexibility in terms of diet and space use allow widespread acclimatisation to a variety of ecological contexts and tolerate high levels of human disturbance (Jepsen & Topping, 2004; Ewald et al., 2014; Basak et al., 2020). When this adaptability is paired with a lack of natural predators and infrastructure expansion, Roe Deer can become frequently embroiled in human-wildlife conflict over forest management and deer-vehicle collisions (Putman et al., 2011; Lush & Lush, 2026).

Here we expand the knowledge of Roe Deer movement, targeting two different landscapes in the UK. We aim to document space use of UK Roe Deer, while exploring their movement in relation to various anthropogenic land cover types (cropland and settlements) and anthropogenic features, with a particular focus on roads.

## 2 Methods

### 2.1 Roe Deer Tracking

The study was conducted in two regions: in Northeastern Scotland (Aberdeenshire) and in Southern England in and around the New Forest National Park (Wiltshire and Hampshire, England; Fig. 1). Broadly, both locations represent mixed-use landscapes comprised of patches of woodland surrounded by cultivated agriculture, livestock pasture and buildings. Deer were captured to attach GPS collars, all being captured in woodland patches, during winter months (January - March in 2023) and (January - March 2024).

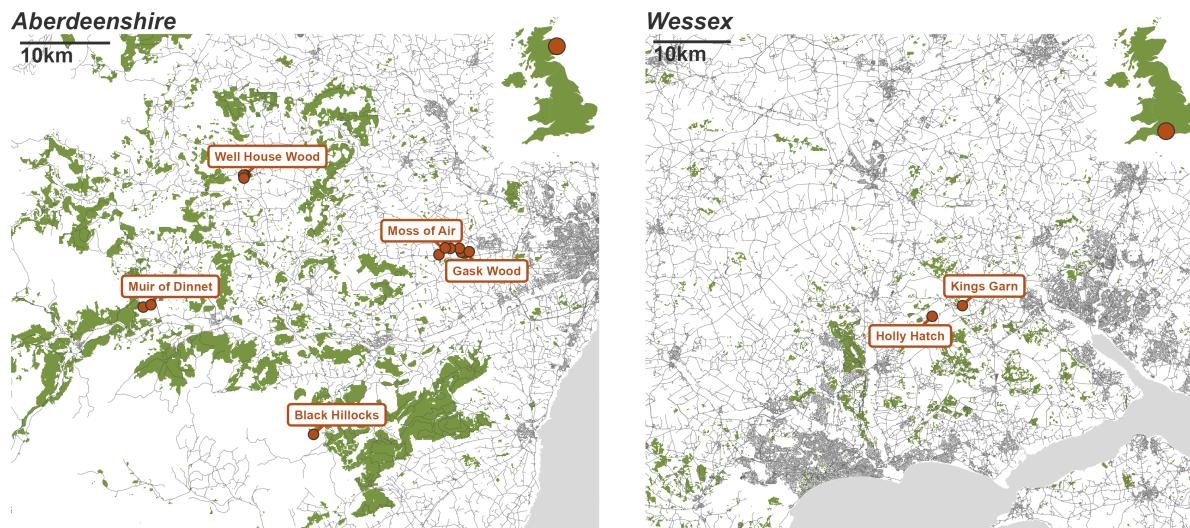
More specifically, we focused deer capture efforts in five Aberdeenshire locations: Muir of Dinnet National Nature Reserve (2 female deer), Black Hillocks (1 male deer), Wellhouse Woods (2 female and 1 male deer), Moss of Air (2 female deer), and Gask Woods (3 female and 1 male deer). In Hampshire, we focused deer capture at Holly Hatch (1 female deer) and Kings Garn (1 male deer). These sites were selected to gain a range of habitats to enable us to address questions about Roe Deer habitat selection in woodlands, and to be representative of the types of forest and landscape typical of Aberdeenshire and Hampshire.

- The Muir of Dinnet is a National Nature Reserve managed by NatureScot, the Scottish Government's statutory conservation agency, in the east of the Cairngorms National Park. The reserve consists of mainly mixed woodland dominated by birch (*Betula spp.*) and Scot's Pine (*Pinus sylvestris*) with extensive areas of naturally regenerating Aspen (*Populus tremula*) and patches of heath and bogs.
- Black Hillocks is managed by the Glendye Estate and includes a patch of coniferous forest dominated by Scot's Pine and European Larch (*Larix decidua*) surrounded by open upland heathland. Within 300m of the site, is the edge of a large tract of commercial, densely planted, mature conifer forest dominated by exotic species such as Sitka Spruce (*Picea sitchensis*).
- Wellhouse Woods is a commercial Sitka Spruce plantation surrounded by farmland pasture.
- Moss of Air, near Garlogie is a mixed woodland including Scot's Pine and birch, surrounded by a mosaic landscape consisting of patches of commercial conifer forest, arable and pasture farmland.
- Gaskwood, also near Garlogie, is a commercial conifer forest including Scot's Pine and Sitka spruce, within a mosaic landscape of forest and farmland patches, located 1 km across open farmland from Moss of Air.
- Holly Hatch and Kings Garn in Hampshire are both within the New Forest National Park. Both sites are a mix of mature deciduous (Oak *Quercus spp.* and Beech *Fagus spp.*) and coniferous trees. Holly Hatch also has dense stands of spruce saplings, surrounded by heathland. Both areas also support other large herbivores, including cattle, ponies, donkeys, Fallow Deer, Red Deer and Muntjac Deer.

We captured adult Roe Deer under Home Office Licence Project licence (PP1285683); a Nature Scot Research Authorisation licence to capture Roe Deer (16759); and a Natural England licence To Take deer Alive for Science, Education and Relocation to capture Roe and Fallow Deer (2022-63493-SPM-WLM). We employed a “long net” capture method (Cockburn, 1976), which has been used extensively and safely to capture Roe Deer in the UK (Gill et al., 1996) and elsewhere (Morellet et al., 2009).

Briefly, we set up 1-2 km of 2-metre-high nylon nets placed along paths, clearings and the edges of woodland patches. Nets were suspended on evenly spaced flexible bamboo poles. These nets then dropped off the poles to the ground and entangled the deer when they ran into them. Nets were placed in the shape of a horseshoe. Capture personnel were spaced every ~50m on the inside part of the netting area. A team of beaters, spaced every 10-15m, moved slowly from the open side of the horseshoe towards the top of the horseshoe. When a deer was caught in the net, the capture team restrained the animal, checked for any injuries, and for general health, estimated the age as either adult or juvenile and determined the sex. Juvenile deer were released. If the animal was judged to be in adequate health, and an adult it was injected intramuscularly with a sedative (Acepromazine (ACP Injection 2mg/ml, Elanco UK AH Limited), at a dose rate of 0.75ml (1.5mg) per deer. The heart rate, respiratory rate and temperature were monitored at capture, and every 5 minutes during subsequent handling. The deer was transferred to a wooden crate designed and built to safely retain the deer for at least 30 minutes for the sedative to take effect, which helped to calm the animal and reduced stress during subsequent handling. After 30 minutes to 1 hour, we removed the deer from the box and attached a GPS collar. Due to the use of sedative, an ear tag with a unique numeric identifier and “do not eat” was attached, along side an ear notch. This tag and notch were used to identify the animal following detachment of the collar and indicate that the deer should not enter the food chain if killed by a stalker. The deer was then released in a safe area away from any hazards.

We fitted captured adult Roe Deer with a GPS collar (30mm reinforced Tellus GP Light Iridium by Followit) that weighed 276g, representing <2% of the average Roe Deer total body mass. Following release, deer were closely monitored for ~10 days with high frequency (30 minute) GPS fixes and daily data transmission to detect any unusual behaviour (e.g., erratic movements, or small daily displacement) indicative of collar or capture trauma. After the initial monitoring period, collars collected GPS fixes every 3 hours in Aberdeenshire, and every 30 minutes in Hampshire. Collars in Hampshire were detached after a short duration (~1 month) due to evidence from one animal that the collar may have caused some friction to the neck. As a result, the Hampshire dataset is more limited in overall duration.



**Figure 1.** Locations of the study sites in Aberdeenshire and Hampshire. Points show the mean location of GPS collared Roe Deer. Green areas in right panel depict woodland, and grey lines show roads. All maps are north orientated.

**Table 1.** Summary of tracking data used in analysis. Deer ID suffix denotes the sex of the individual deer.

Location	Forest Area	Deer ID	Duration (days)	Number of fixes	Fixes per day	Mean lag between fixes (hours)
Aberdeenshire	Gask Wood	Roe01_F	252.50	1799	7.12	3.37 ± 1.49
Aberdeenshire	Gask Wood	Roe02_F	255.50	1805	7.06	3.4 ± 1.61
Aberdeenshire	Gask Wood	Roe04_F	248.50	1787	7.19	3.34 ± 1.45
Aberdeenshire	Muir of Dinnet	Roe05_F	101.88	676	6.64	3.62 ± 1.18
Aberdeenshire	Moss of air	Roe06_F	282.50	1938	6.86	3.5 ± 1.57
Aberdeenshire	Gask Wood	Roe08_M	250.50	1799	7.18	3.34 ± 1.42
Aberdeenshire	Black Hillocks	Roe09_M	254.50	1821	7.16	3.36 ± 1.44
Aberdeenshire	Muir of Dinnet	Roe10_F	270.50	1873	6.92	3.47 ± 1.6
Aberdeenshire	Well House Wood	Roe11_F	67.88	548	8.07	2.98 ± 0.15
Aberdeenshire	Well House Wood	Roe12_F	139.88	1120	8.01	3 ± 0.21
Aberdeenshire	Moss of air	Roe13_F	280.50	1931	6.88	3.49 ± 1.53
Aberdeenshire	Well House Wood	Roe14_M	149.88	1201	8.01	3 ± 0.18
Aberdeenshire	Moss of air	Roe15_F	282.50	1932	6.84	3.51 ± 1.59
Hampshire	Kings Garn	Roe03_M	49.88	305	6.11	3.94 ± 1.81
Hampshire	Holly Hatch	Roe07_F	52.92	368	6.95	3.46 ± 1.67

We retrieved movement data from 15 GPS collars worn by Roe Deer, 13 in Aberdeenshire, 2 in Hampshire. We re-sampled the Roe Deer data to a more consistent rate, aiming for a standard 3 hour time lag between locations (with a 1 hour tolerance). We additionally filtered out the first weeks' worth of data to avoid the impacts of capture/immediate post-release movements that may have been atypical (Morellet et al., 2009).

## 2.2 Home Range Estimation

We estimated Roe Deer home range using Autocorrelated Kernel Density Estimators [AKDE; Fleming & Calabrese (2023); Calabrese, Fleming & Gurarie (2016); Fleming et al. (2015); Fleming & Calabrese (2017)]; a method of home range area estimation that accounts for the particular structures present in movement data such as auto-correlation and data gaps. This process consisted of fitting a number of continuous time movement models to an individual deer's movement data, selecting the best fitting movement model, and extracting a suitable range contour from the resulting utilisation distribution. We fit the following models (following the default process provided by the ctmm package): Ornstein-Uhlenbeck (OU), Ornstein-Uhlenbeck Foraging (OUF), and Independent Identically Distributed (IID), all in both isotropic and anisotropic forms. We elected to use perturbative hybrid residual maximum likelihood (pHREML) (Fleming et al., 2019; Silva et al., 2022) and AICc to determine the best fitting movement model on an individual basis, and used that single best fitting model for all further estimations.

Before committing to the estimations of home range size we examined whether the Roe Deer exhibited stable ranges through the visual inspection of variograms. A stable range should be revealed by a clear asymptote in variogram, where the semi-variance flattens as time lags increase. We paired these visual inspections with a judgement of effective sample size to help gauge our confidence in the home range area estimates [effective sample size approximating the overall tracking duration divided by the mean time taken to cross the home range; Silva et al. (2022)]. All our individuals showed effective sample sizes between 154.4 and 926.7, indicating that the movement data contained a large number of complete home range crossing events, allowing us to be confident in overall home range estimates.

Having determined the data's suitability for home range estimations, we extracted the 95% and 99% contours from the weighted AKDE estimate, alongside 95% confidence interval surrounding that contour. We selected 95% for further estimate mean ranges as it struck a balance between a generous estimate of home range, while also avoiding the undue influence of the most extremely outlying movements. To generate an overall home range estimate for Aberdeenshire Roe Deer, we averaged the home ranges using the weighted mean function provided by the ctmm package (Silva et al., 2022; Fleming & Calabrese, 2023). This way the home range mean is weighted by the confidence (i.e., effective sample size) surrounding each home range.

We retained 99% contour estimates to help quantify the distance from which Roe Deer will range away from woodland patches. We calculated the widest dimension of each 99% home range polygon (or largest polygon if the home range area was non-contiguous), and halved that value to quantify the distance deer would be willing to travel beyond their resident patch. To support this approach and to produce an overall distribution deer distance away from woodland patches, we determined how far every deer location was from the nearest patch and plotted those values against the distance estimated by the home range models.

## 2.3 Habitat Selection

We used the reformulated Poisson model approach described by Muff, Signer & Fieberg (2020) to generate population level estimates of habitat selection as well as gauge deer's movement capabilities in relation to different aspects of the landscape.

The model required data pertaining to the used locations (i.e., GPS locations of the deer) and comparable data on randomly generated available points (i.e., randomly generated locations the deer could have travelled to). For each confirmed deer location, we generated 10 random alternative locations they could have travelled to. The location of these random locations was governed by two distributions. A Gamma distribution from which random step lengths were drawn from, and a Von Mises distribution from which random turn directions were drawn from. Both distribution were calibrated (e.g., shape, size, mu, and kappa) by the underlying movement data.

Once all random locations had been generated, we extracted a suite of environmental variables at all those locations, in order to relate Roe Deer space use with land cover and anthropogenic features. First was the land cover type as described by the 2023 UKCEH land cover maps, which is a 25m resolution classified raster originally based on Sentinel-2 imagery (Morton et al., 2024). Validation of these data suggest 83% accuracy (Morton et al., 2024). The UKCEH land cover data comprises of 21 land cover classes, broadly following the Biodiversity Broad Habitats (Jackson, 2000).

We recategorised these 21 land cover types categories into 10 more general categories (Tab. 2). This reduced instances of limited interaction with the deer movement data thereby aiding habitat selection model convergence and avoided extreme, unstable selection estimates.

**Table 2.** Overview of the reclassification of UKCEH land cover classes for inclusion into the habitat selection models.

UKCEH land cover class	UKCEH land cover identifier	Reclassified category
Deciduous woodland	1	Deciduous Broadleaf Forest
Coniferous woodland	2	Evergreen Needleleaf Forest
Arable	3	Cropland
Improved grassland	4	Tall Grassland
Neutral grassland	5	Short Grassland
Calcareous grassland	6	Short Grassland
Acid grassland	7	Short Grassland
Fen	8	Permanent Wetland
Heather	9	Open Shrubland
Heather grassland	10	Open Shrubland
Bog	11	Permanent Wetland
Inland rock	12	Barren
Saltwater	13	Other
Freshwater	14	Other
Supralittoral rock	15	Barren
Supralittoral sediment	16	Barren
Littoral rock	17	Barren
Littoral sediment	18	Barren
Saltmarsh	19	Permanent Wetland
Urban	20	Human Settlements
Suburban	21	Human Settlements

In addition to land cover classes, we also acquired “woody linear feature” (i.e., hedgerows) data from UKCEH (Scholfield et al., 2016), which maps hedgerows across the UK (e.g., woodland) as polylines, based on Ordnance Survey maps and the 2007 UKCEH Land Cover Map (Morton et al., 2011).

We converted the polygon spatial data into a raster, where 1 == hedgerow, and used that rasterisation to generate a distance to hedgerow raster for the entire study landscape. We conducted the same process to create a distance to woodland raster, where we calculated the distance from any area the UKCEH land cover data classed as deciduous or coniferous woodland. These distance rasters allowed for easy extraction of the distance to the nearest hedgerow and woodland for all locations. This allowed us to investigate the influence of woodland or hedgerow on the use of open habitats by Roe Deer, as well as the extent to which expanses of open habitats might act as barriers to Roe Deer dispersal.

We acquired road data from Ordnance Survey Open Roads dataset (Ordnance Survey, 2024). We created a binary variable describing crossing events for all steps, with all steps that crossed one or more of the roads being classed as 1. This binary variable allowed us to estimate the likelihood Roe Deer cross a road and therefore the extent to which roads may present a barrier (Serota et al., 2024).

To ensure compatibility between all data sources, we projected all data into the British National Grid (BNG) coordinate reference system (OSGB36, epsg: 27700) before undertaking analysis.

The population level model consisted of land cover (a 8-term category variable formed into 7 dummy variables, with deciduous woodland placed as the reference category, barren and other excluded), distance to woodland (continuous in m), distance to hedgerow (continuous in m), road crossing (binary). In addition to these habitat selection focused predictors, we included several movement predictors: step length, log step length, and cos turn angle, as well as the interaction between step length and log of step length with all land cover types.

To account for the structure originating from having multiple individuals in the model, we included fixed Gaussian processes for the time step and the individual in keeping with the approach described by Muff, Signer & Fieberg (2020). This formulation, namely the fixed Gaussian processes, as described by Muff, Signer & Fieberg (2020) allows for the efficient estimation of population level selection using integrated nested Laplace approximation (INLA) (Martins et al., 2013; Lindgren & Rue, 2015). Our final formula was:  $y \sim -1 + \text{Distance to woodland (continuous)} + \text{Distance to hedgerows (continuous)} + \text{Land cover categories (7 binary variables)} + \text{Road crossed (binary)} + \text{Step length (continuous)} + \text{Log step length (continuous)} + \text{Cos turn angle (continuous)} + \text{Step length interactions with Land cover categories} + \text{Log step length interactions with Land cover categories} + \text{Gaussian process for deer ID} + \text{Gaussian process for time step}$ .



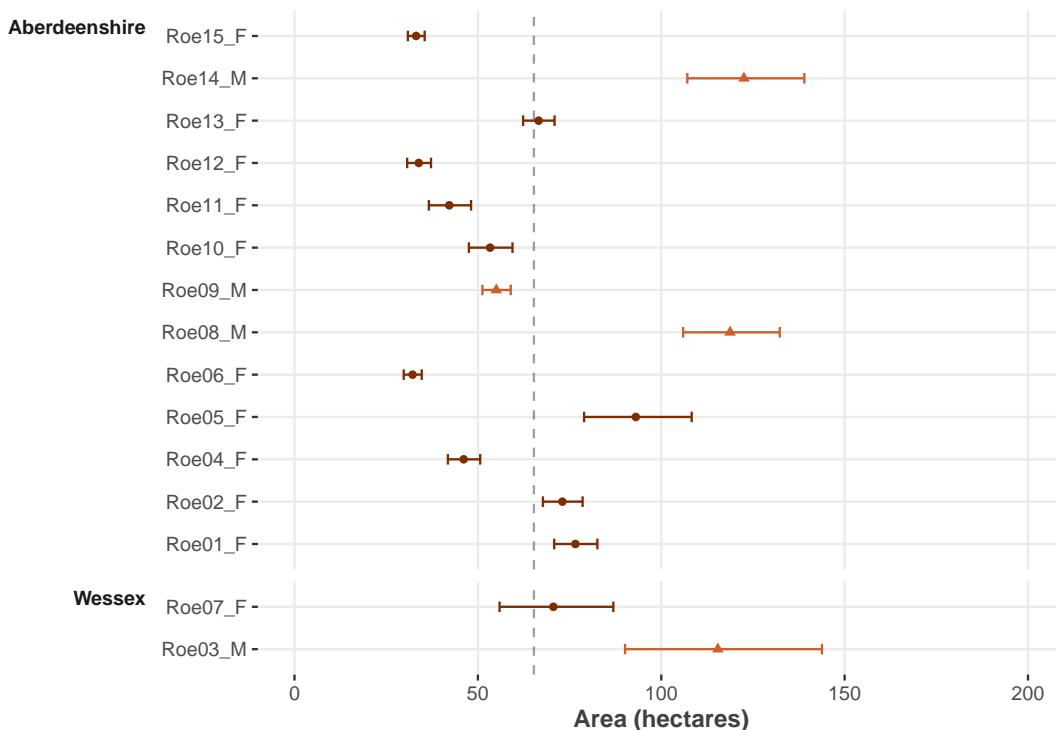
To supplement the population model, we ran individual level step-selection models for each Roe Deer separately. These models used the same data as the population level model, but focused on individual level responses relative to the population mean. Past studies of wild Roe Deer have found substantial levels of individual variability in behavioural responses to risk (Bonnot et al., 2015). For the individual models, we used a formula that included land cover class, distance to woodland, distance to hedgerows, a binary describing whether they crossed a road, step length, log step length, and cos of turn angle. Once we had estimated individual responses to the environmental variables, we explore how the variation in individual coefficients could highlight individual variability in selection. We used the IndRSA package (Bastille-Rousseau, 2025) to explore the variation of resulting coefficients (i.e., selection or attraction towards environmental characteristics), producing population-level estimates of specialisation, heterogeneity, and a weighted population mean (Bastille-Rousseau & Wittemyer, 2022). Specialisation is the absolute magnitude of the coefficients; differences compared to the population mean coefficient could highlight diverging responses to the habitat covariates and a bimodal response to a given environmental characteristic. This can be particularly informative when the diverging responses have resulted in a “neutral” population mean for the coefficient. Heterogeneity is the standard deviation of the coefficients; therefore, larger values indicate greater variation between individual Roe Deer in their response to land cover or landscape features. The weighted population mean provides an alternative measure of population level selection to our Poisson model. To carry forward the uncertainty surrounding the initial habitat coefficients in the weighted population mean, 10,000 replicates of each metric were generated from a normal distribution centred on the original coefficient with a standard deviation equal to the standard error of the coefficient (Bastille-Rousseau & Wittemyer, 2022).

### 3 Results

Overall, the 15 GPS collared Roe Deer resulted in 20903 location fixes, with a mean of  $1394 \text{ SD} \pm 627.9$  per individual, spread across a mean of  $196 \text{ SD} \pm 90.99$  days per individual (Fig. S1). This resulted in an average of  $7.133 \text{ SD} \pm 0.5359$  fixes per day per individual (Fig. S2; Tab. 1).

For our Aberdeenshire Roe Deer home range sizes ranged from 32.2 to 122.5ha (95% contour point estimates), with a weighted mean of 65.3 ha (95% CI 55.5-75.8) (Fig. 2). The deer appeared range resident; however, a couple of individuals may show evidence of a range shift during the tracking period (Roe Deer 8 and Roe Deer 3; Fig. S3). Effective sample sizes were all high (154.4 and 926.7), suggesting we can be confident in the home range estimates. All except for two individuals (Roe Deer 5 and Roe Deer 12) found OU models to fit best, with the remaining two being better described by OUF models (Tab. S1).

All best fitting models were anisotropic, suggesting these Roe Deer are inhabiting non-uniform home ranges (i.e., not being as wide as they are long). The placement of home ranges suggest the importance of woodland, with 95% of all deer movements falling within 756 m of woodland (756 m is half of the longest dimension of the 99% home range area; thereby suggesting ranges tend to centre on woodland; Fig. 3).

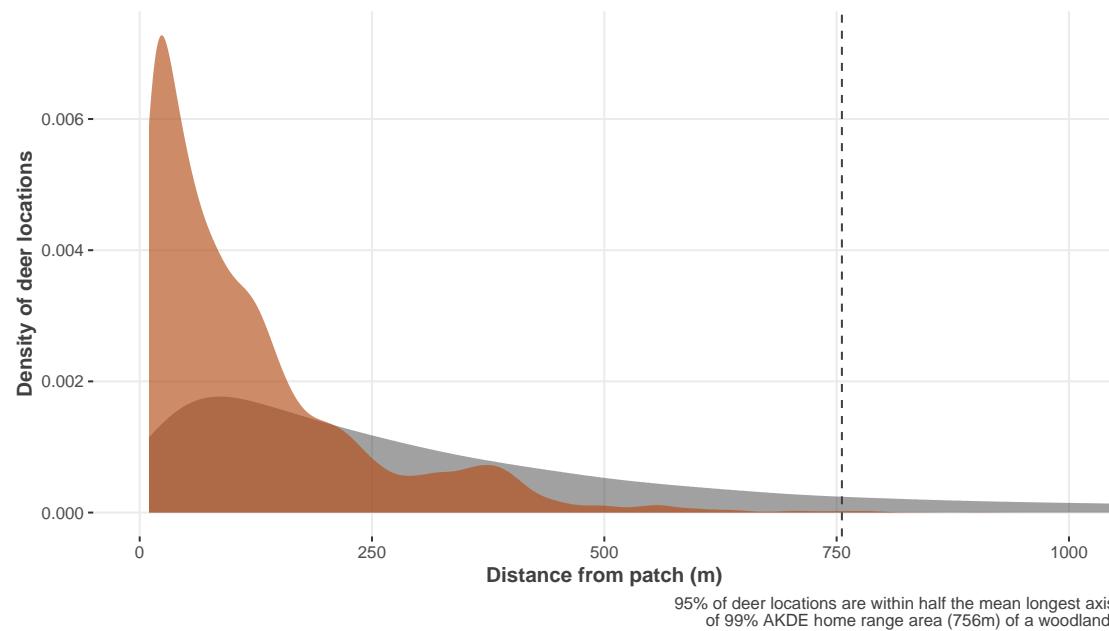


**Figure 2.** Home range size of Roe Deer in two landscapes as estimated via the Autocorrelated Kernel Density Estimators. Depicted are the 95% contour with 95% confidence intervals surrounding estimates for female (red circles) and male (orange triangles). The vertical line shows the weighted mean of Aberdeenshire home range estimates.

#### 3.1 Population level selection

The population level habitat selection model revealed a general tendency for Roe Deer to remain closer to the woodland patches ( $-0.0051$ ; 95% CI  $-0.0082$  to  $-0.0022$ ), with no significant selection for hedgerows ( $-9e-04$ ; 95% CI  $-0.0019$  to  $1e-04$ ; Fig. 4). The model also revealed that roads play a significant role in reducing connectivity across the landscape: observed deer steps were significantly less likely to cross roads than random steps ( $-0.76$ ; 95% CI  $-1.2$  to  $-0.36$ ). The population habitat selection model also revealed significant selection for the land cover classes open shrubland (1.5; 95% CI 0.62 to 2.2) and tall grassland (0.57; 95% CI 0.17 to 0.98); other relationships were less clear (Tab. S2).

The model's inclusion of step length and log step length interactions allowed us to examine whether movement was altered by land cover. Movement was most impacted by short grassland, tall grassland, and cropland, all showing the same pattern. The step lengths in these land covers tended to be lower (i.e., slower movement), as seen in



**Figure 3.** Distribution of Roe Deer locations in relation to their distance from woodland patches (Orange) compared to the distribution of random locations throughout the landscape in relation to their distance from woodland patches (Grey). Locations within woodland are excluded. Vertical dashed line shows the half longest dimension of the 99% home range area.

coefficients for log step length (short grassland -0.76; 95% CI -1.2 to -0.32; tall grassland -0.18; 95% CI -0.24 to -0.12; cropland -0.16; 95% CI -0.24 to -0.078) but with a larger tail to the Gamma distribution (i.e., larger coefficient for step lengths; short grassland 0.0044; 95% CI 0.0018 to 0.007; tall grassland 0.00048; 95% CI 5.1e-05 to 0.00092; cropland 0.0012; 95% CI 0.00064 to 0.0017). Combined this could be indicative of more stop-start movements and behaviours.

Estimates regarding human settlements were paired with very wide confidence intervals (-70; 95% CI -180 to 36), likely a result of minimal overlap between the Roe Deer movement data (and any associated random available points) and human settlements making estimation difficult.

### 3.2 Individual level selection and variation in selection

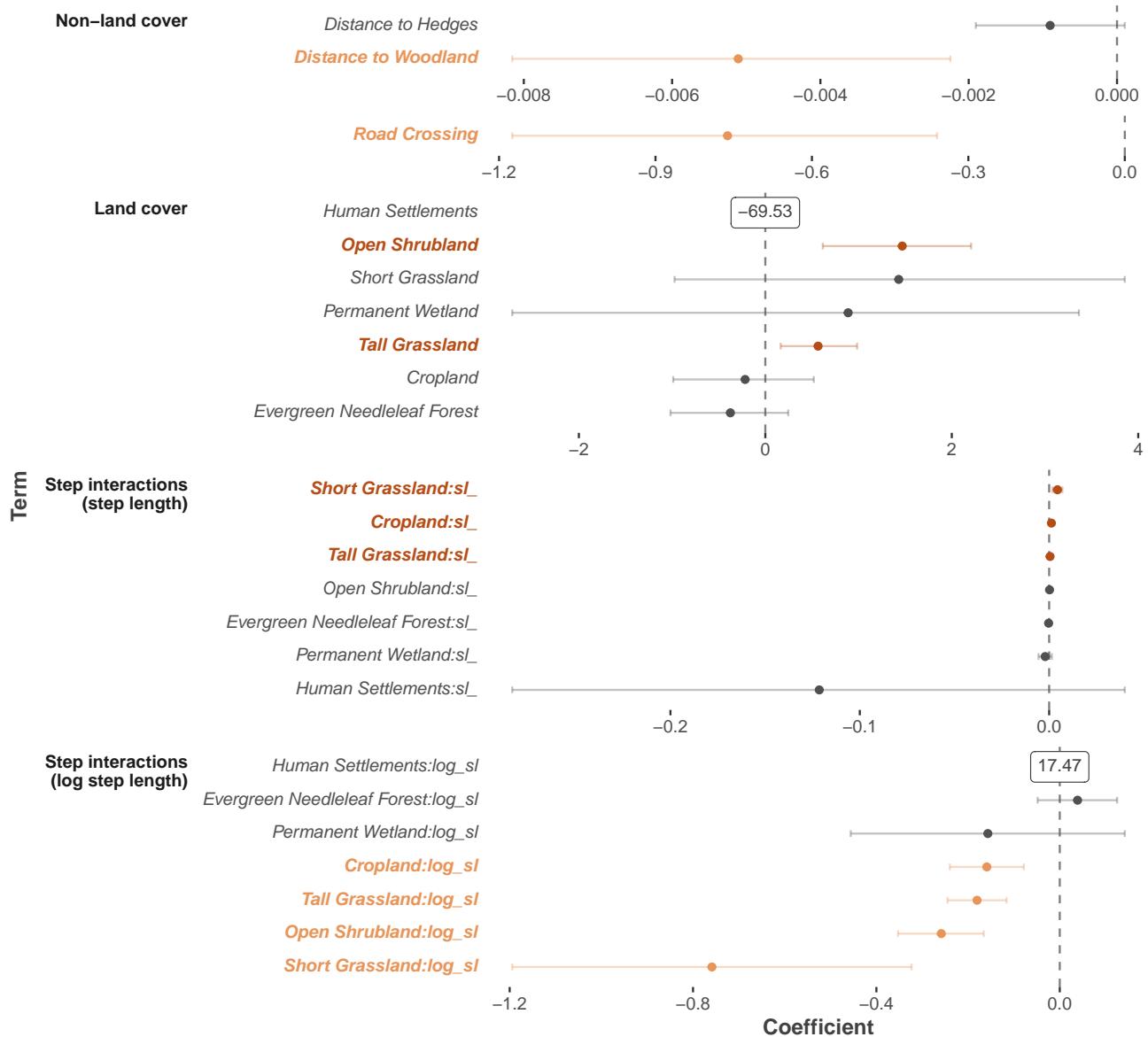
Further exploration of individual habitat selection models highlights whether the uncertainty in the population level model stems from weak responses or diverging responses that average towards a zero effect (Tab. S3; Tab. S4). Distance to woodland shows a marginally higher specialisation (0.01) compared to the weighted population mean (-0.005; 95% CI -0.009 to -0.001; Fig. 5). This difference is likely explained by the deviating Roe 10 who expressed an opposite response to the majority of other deer by expressing an aversion to woodland ( $0.006 \pm 0.001$ ). Apart from Roe 10, Roe Deer are exhibiting the same clear preference for remaining near woodland as seen in the population level model. Roe 10 is also likely the reason the estimated heterogeneity in response to distance to woodland (0.0134) is greater than distance to hedgerows (0.0019; Fig. S4).

Distance to hedgerows does not see the same consistent response, instead with a number of individuals showing a opposing responses. This is reflected in the population mean being close to zero (-0.001; 95% CI -0.002 to 0), while the specialisation is greater (0.002; Fig. S5), the combination of which could indicate two differing responses to hedgerows in the sampled Roe Deer.

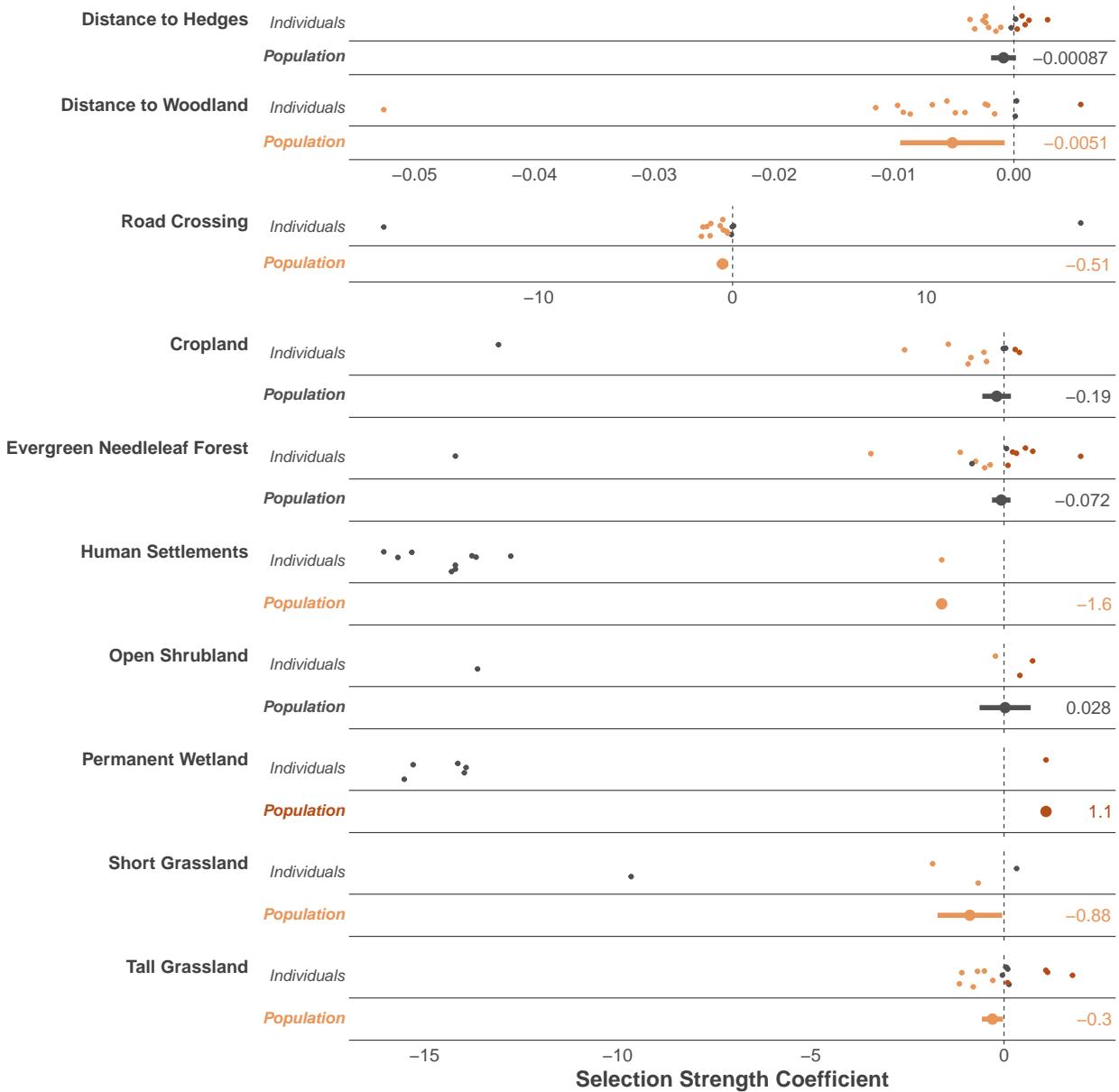
The chance of crossing a road is considerably more consistent, with the vast majority of individuals preferring not to cross roads; this is reflected in a significantly negative population mean (-0.514; 95% CI -0.807 to -0.222). The elevated specialisation (134.97) and heterogeneity (417.65) values are almost entirely driven by two very uncertain estimates from Roe Deer 09 ( $17.957 \pm 888.36$ ) and Roe Deer 11 ( $-17.988 \pm 1620.795$ ). This could have been the result of a lack of exposure; the Roe Deer 09 only crossed roads on two occasions, while Roe Deer 11 never did. Other than those individuals, we can be confident in a consistently negative response to road crossing in Roe Deer.

It was difficult to obtain confident estimates for land cover categories due to the variable levels of availability for

each individual; some individuals only rarely moved close enough to certain land cover classes so estimations are based on a small section of the movement dataset. Cropland and Evergreen Needleleaf Forest showed the high rates of significant estimates, and both weighted population estimates (-0.191; 95% CI -0.561 to 0.179; -0.072; 95% CI -0.311 to 0.168) matched the results from the Poisson population level model (-0.22; 95% CI -0.99 to 0.52; -0.37; 95% CI -1 to 0.25). In both cases a single individual revealed a strong but very uncertain negative response that can explain the heterogeneity (353.18; 252.83) and specialisation values (107.22; 68.37) and additionally explain why the effect overlapped zero in the Population level model. The response to Evergreen Needleleaf Forest appears the least consistent, with multiple individuals expressing significantly negative and positive responses to the land cover. Tall Grassland showed similar levels of diverging estimates, with individuals showing a mix of positive and negative responses. Unlike the population estimates for Cropland and Evergreen Needleleaf Forest, the weighted population mean for Tall Grassland (-0.299; 95% CI -0.572 to -0.026) does not match the population level model (0.57; 95% CI 0.17 to 0.98). This may be indicative that the interaction effects included in the population level model are mediating the responses to Tall Grassland. A similar reason may explain the clear Open Shrubland response in the population level model (1.5; 95% CI 0.62 to 2.2) that is absent in the individual models and the resulting weighted population mean (0.028; 95% CI -0.633 to 0.69); however, this is more likely to be caused by the limited number of individuals exposed to Open Shrubland and a different handling of Roe Deer 03's strongly negative response. The other land covers are all harder to confidently interpret given the frequency of very uncertain extreme estimates. The uncertainty surrounding Human Settlements and Permanent Wetland is well reflected in the population level model. The high levels of specialisation and heterogeneity are driven by these same extreme estimates. The lack of consistent availability of these land covers and potentially inconsistent response leaves a lot of uncertainty concerning Roe Deer response to these land covers.



**Figure 4.** Coefficient estimates, with 95% confidence intervals, from the population level model of habitat selection for all 15 Roe Deer. For distance to \* variables, lower coefficients indicate that Roe Deer are selecting areas with lower distances from the landscape feature. For the road crossing variable, lower coefficients indicate a lower than chance to cross the road. For land cover variables, positive coefficients indicate of Roe Deer preferentially selecting to be in those areas. For step length interactions, positive coefficient interactions with step lengths indicate a longer tail to the overall distribution of step lengths when Roe Deer are in a given land cover class. A positive coefficient with log step lengths indicate a larger step lengths when Roe Deer are in a given land cover class. Central numeric labels report majorly outlying estimates to aid visualisation. Colour highlights and bolding reflect the significantly negative (light orange) and significantly positive (dark orange) coefficients. Note x axes are different per variable type.



**Figure 5.** Coefficient estimates for all individual step-selection models for all 15 Roe Deer, along side weighted population means per covariate. Points indicate the estimated coefficients for each individual. Error bars with the population estimates are the 95% confidence interval. Colour highlights reflect the estimates whose standard errors do not overlap zero, either negatively (light orange) and positively (dark orange). For distance to \* variables, lower coefficients indicate that Roe Deer are selecting areas with lower distances from the landscape feature. For the road crossing variable, lower coefficients indicate a lower than chance to cross the road. For land cover variables, positive coefficients indicate of Roe Deer preferentially selecting to be in those areas.

## 4 Discussion

Our tracking of 15 Roe Deer revealed that they have limited home ranges, that are heavily skewed towards remaining close to deciduous woodland. They are willing to exit woodland, making use of open shrubland and grasslands, but these movements tend to be limited to within ~750m and the vast majority far closer. When entering these more open non-wooded areas they tend to slow down, but also exhibit more variable movements. Exploration of individual level habitat selection models highlight the consistency of preferring to remain close to woodland. However, their limited exposure to certain land cover types (e.g., human settlement) makes it difficult to fully characterise individual-variability. Overall, Roe Deer exhibited a disinclination to cross roads, a pattern found both on a population and individual level. The estimated chances of individuals crossing roads indicate that roads somewhat reduce the permeability of these landscapes for Roe Deer, but do not prohibit movement.

Our findings on Roe Deer home range size are similar to those values reported in the HomeRange database for Roe Deer (approx. 105 Roe Deer; Fig. S6), with the database presenting examples of both ranges larger and smaller than exhibited by our 15 individuals (Broekman et al. (2023); Broekman et al. (2022); see references in data availability section). This similarity in home range sizes is somewhat speculative given it does not account for differences in sampling protocol, duration, or home range estimation method, all of which can modify estimated home range size (Silva et al., 2020). Despite the simple comparison, the UK Roe Deer examined in our study appear to be largely typical in regards to their home range size. Some of the variation in home range has been suggested to be a product of food accessibility and the arrangement of cover from predators (Tufto, Andersen & Linnell, 1996; Saïd & Servany, 2005; Saïd et al., 2009), as well as dynamic in relation to reproduction (Saïd et al., 2005).

Other examinations of Roe Deer have highlighted the importance of risk guiding when Roe Deer leave their core wooded range to make use of more open areas (Venkatesan et al., IN PREP - 2026; Bonnot et al., 2015; Padié et al., 2015). The potential benefits of managing this risk to reap the nutritional benefits afforded by open areas are apparent (Hewison et al., 2009), and may present a direct trade off against the risks posed by humans when in more open areas. Humans are likely the UK Roe Deer's primary concern, as other predators such as Lynx, which require different risk mitigating decisions, are absent in the UK (Lone et al., 2014). The shifts in movements (i.e., step lengths) we report here may be a coarse representation of Roe Deer balancing the risk and benefits in crop- and grasslands. Such trade-offs may be more pronounced in more fragmented or riskier landscapes, and could also explain the lack of consistent selection to remain near hedgerows. Compared to other Roe Deer studies in area such as France, our Roe Deer may have easier access to wooded cover in a less fragmented landscape (Morellet et al., 2011); thereby, being less dependent on hedgerows. To tease apart the UK Roe Deer's response to risk, a more complete quantification of human activity would be required, including presence of walkers, dogs, and hunting pressure.

Roe Deer in our study avoided crossing roads more than expected by chance, suggesting roads reduce landscape connectivity. Thus, in countries with high densities of roads, such as the UK, even limited road crossing avoidance by deer could potentially have impacts on deer population structure, dispersal, and movement of deer-associated parasites (e.g., ticks). Some deer-hosted *Ixodes ricinus* ticks carry pathogens significant for human and livestock health; therefore, this differential landscape permeability may have indirect health implications. Despite this avoidance, it was not complete and deer collisions with vehicles remain a key safety issue (Lush & Lush, 2026). In Scotland, higher rates of deer-vehicle collisions appear to occur in the vicinity of woodland and other semi-natural habitats (Langbein, 2019), with road junctions being key hotspots (Lush & Lush, 2026). Other studies have highlighted the importance of road density in Roe Deer home ranges as a key predictor of road crossing frequency (Kämmerle et al., 2017), further supporting the woodland-road proximity connection. Traffic flow also plays a role in the likelihood of a road section experiencing collisions (Nelli et al., 2018), highlighting that the ultimate impact of roads will be a product of deer and human factors. For example, while urban areas tend to lower deer-vehicle collision risk, the intersection of urban and natural areas (i.e., suburban) may result in risk hotspots where both deer and traffic collide (Nelli et al., 2018). This non-linear relationship between animal densities, their reaction and activity close to roads, and the intensity of traffic makes road-wildlife collision difficult to completely characterise and generalise (Valero et al., 2015; Abraham & Mumma, 2021; Cunningham et al., 2022; Denneboom, Bar-Massada & Shwartz, 2024). The estimation of Roe Deer crossing tendency here, will help refine our understanding of that relationship in the UK. Other Roe Deer investigations have revealed similar preferences for forest and reluctance to cross roads (Passoni et al., 2021), and that the risks fluctuate over the day and year (Cunningham et al., 2022; Märtz, Brieger & Bhardwaj, 2024; Lush & Lush, 2026).

There are several aspects of the study that may limit its generalisability, or that would need to be considered when applying the findings to other contexts. Using the STRANGE framework (Webster & Rutz, 2020), we highlight key limitations (omitting those that are unquantifiable or of limited relevance). **Social background.** We have little information on the individuals not tracked that may be impacting the movements of tracked deer. Roe Deer will maintain territories (Hoem et al., 2007; Pagon et al., 2017), so it is likely that the density of and interactions with

conspecifics could alter the distribution and size of our tracked deer's movements via competitive exclusion or territorial patrolling. **Trappability and self-selection.** While there is not obvious bias to the deer trapping methods we used, there may be an unknown behavioural variation altering the likelihood of a deer being captured. If the trappability of a deer is associated with certain movement or behavioural parameters, our sample may be skewed towards those tendencies (e.g., boldness leads to increased capture likelihood, while also being connected to increased chance of crossing roads). Long-term studies of individually-tagged Roe Deer in France have detected behavioural and body size differences between deer associated with woodlands versus open or cultivated landcover (Hewison et al., 2009). Given deer were exclusively trapped in woodlands in our study, we may have overestimated the strength of woodland preferences for some Roe Deer, and in areas with more continuous woodlands, deer may prefer open habitat more strongly (Hewison et al., 2001). Nonetheless, Roe Deer in general are undoubtedly strongly woodland associated (Gill et al., 1996) and need shrubby or herbaceous cover for hiding and foraging. **Acclimation and habituation.** None of the deer had been previously collared, but there may have been a habituation effect to collars over time. Our removal of the first week of data likely have mitigated the largest impact prior to collar-habituation, but as the Hampshire deer demonstrate there may be ongoing effects we cannot control for. Further explorations for longer tracking periods and with different age groups may elucidate the collar and human habituation effects.

Overall, we documented a tendency for Roe Deer to remain close to woodland in the UK. This pattern apparently limits their ranges to areas within 750m of woodland patches. The resulting home ranges of UK Roe Deer do not appear atypical when compared to other tracked Roe Deer. We see a clear and consistent, albeit slight, aversion to crossing roads that may be limiting the permeability of the landscape for Roe Deer. Further work could benefit from focusing on UK Roe Deer in areas with a greater urban footprint, or comparing movements against more granular quantifications of human activity (Gomez et al., 2025).

## 5 Acknowledgements

We are indebted to the hundreds of volunteers who assisted in the deployment of deer GPS collars. We especially thank Mark Hewison, Jochen Langbein, Tim Dansie, Andy Page, Sandy Shore and Andy Shore for generous advice in the field and training in capture methods. This research forms part of the TickSolve project (<https://ticksolve.ceh.ac.uk/>) and was funded by UK Research and Innovation through the NERC grants NE/W003171/1 and NE/W003244/1. For permission to conduct the work, we thank NatureScot, ForestryEngland and NaturalEngland.

### 5.1 Software availability

For all analysis we used R (v.4.4.2) (R Core Team, 2024), and R Studio (v.2024.12.0+467) (Posit team, 2024). For analysis of animal movement data we used amt (v.0.2.2.0) (Signer, Fieberg & Avgar, 2019), ctmm (v.1.2.0) (Fleming & Calabrese, 2023), and move (v.4.2.6) (Kranstauber, Smolla & Scharf, 2024). For general data manipulation we used glue (v.1.8.0) (Hester & Bryan, 2024), sjmisc (v.2.8.10) (Lüdecke, 2018), tidyverse (v.2.0.0) (Wickham et al., 2019), and units (v.0.8.5) (Pebesma, Mailund & Hiebert, 2016). For project and code management we used here (v.1.0.1) (Müller, 2020), tarchetypes (v.0.11.0) (Landau, 2021a), and targets (v.1.9.0) (Landau, 2021b). For visualisation we used the following as expansions from the tidyverse suite of packages: ggdist (v.3.3.2) (Kay, 2024a,b), ggridges (v.0.5.6) (Wilke, 2024), ggtext (v.0.1.2) (Wilke & Wiernik, 2022), patchwork (v.1.3.0) (Pedersen, 2024), and scales (v.1.3.0) (Wickham, Pedersen & Seidel, 2023). Other packages we used were boot (v.1.3.31) (A. C. Davison & D. V. Hinkley, 1997; Angelo Canty & B. D. Ripley, 2024), circular (v.0.5.1) (Agostinelli & Lund, 2024), doParallel (v.1.0.17) (Corporation & Weston, 2022), foreach (v.1.5.2) (Microsoft & Weston, 2022), knitr (v.1.49) (Xie, 2014, 2015, 2024), and usethis (v.3.0.0) (Wickham et al., 2024). To generate typeset outputs we used bookdown (v.0.42) (Xie, 2016, 2025), and rmarkdown (v.2.29) (Xie, Allaire & Grolemund, 2018; Xie, Dervieux & Riederer, 2020; Allaire et al., 2024). To manipulate and manage spatial data we used gdistance (v.1.6.4) (van Etten, 2017), raster (v.3.6.30) (Hijmans, 2024a), sf (v.1.0.19) (Pebesma, 2018; Pebesma & Bivand, 2023), sp (v.2.1.4) (Pebesma & Bivand, 2005; Bivand, Pebesma & Gomez-Rubio, 2013), terra (v.1.7.83) (Hijmans, 2024b), and tidyterra (v.0.6.1) (Hernangómez, 2023). To run models and explore model outputs we used effects (v.4.2.2) (Fox, 2003; Fox & Hong, 2009; Fox & Weisberg, 2018, 2019), INLA (v.24.6.27) (Martins et al., 2013; Lindgren & Rue, 2015), lme4 (v.1.1.35.5) (Bates et al., 2015), and performance (v.0.12.4) (Lüdecke et al., 2021).

The code used to complete this study can be found at [https://github.com/BenMMarshall/TICKSOLVE\\_DeerMovement](https://github.com/BenMMarshall/TICKSOLVE_DeerMovement) amongst code for the broader examination of deer's role in these landscapes; and is archived at <https://doi.org/10.5281/zenodo.18701055>.

We used Blender 4.5 LTS (Blender Online Community, 2025) and Affinity Publisher 2 (Serif, 2025) to create the graphical abstract visuals.

### 5.2 Data availability

Core data underlying these analyses can be found at: <https://doi.org/10.5281/zenodo.18699746>.

Aberdeenshire Roe Deer movement data can be accessed via Movebank <https://www.movebank.org>; Movebank ID 2890266958.

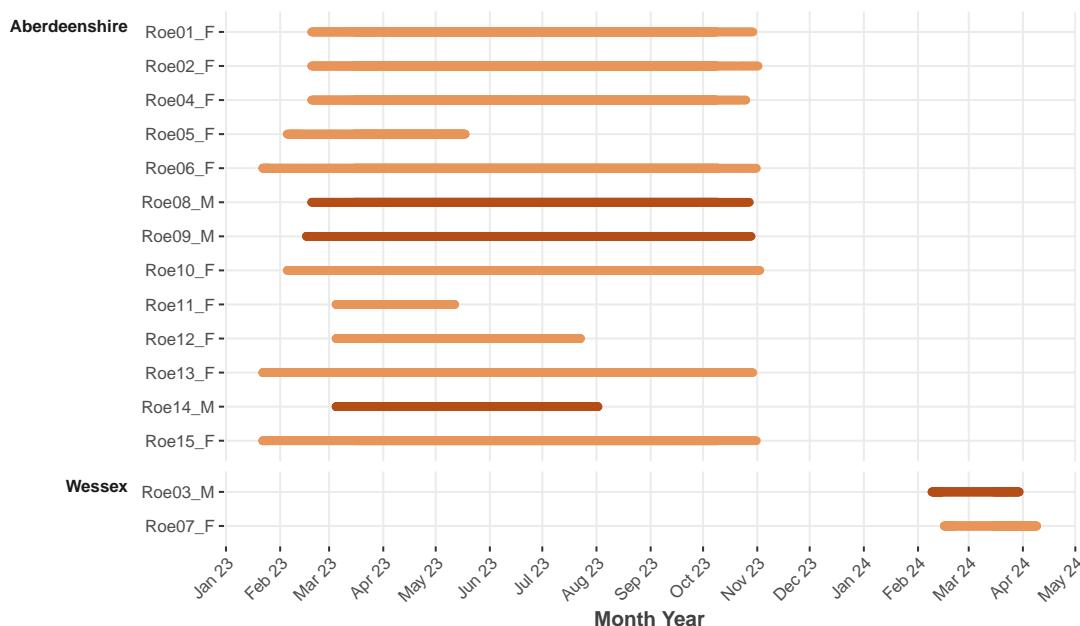
New Forest, Hampshire Roe Deer movement data can be accessed via Movebank <https://www.movebank.org>; Movebank ID 8086497234.

Studies that the HomeRange Database mean was based on: Melis, Cagnacci & Lovari (2005); Biosa et al. (2015); Dupke et al. (2017); Rossi et al. (2003); Focardi et al. (2006); Picardi et al. (2019); Ramanzin, Sturaro & Zanon (2007); Mysterud (1999); Ranc et al. (2020); Richard et al. (2008); Aiello, Lovari & Bocci (2013); Morellet et al. (2013); Vanpé et al. (2009); Pellerin et al. (2016); Kjellander et al. (2004); Van Laere, Boutin & Gaillard (1996); Cederlund (1983); Saïd et al. (2005); Bideau et al. (1993); Cimino & Lovari (2003); Lamberti et al. (2001); Lamberti et al. (2006); Bevanda et al. (2015); Pagon et al. (2017); Debeffe et al. (2012); Chapman et al. (1993); Lamberti, Mauri & Apollonio (2004); Maublanc et al. (2018); Padié et al. (2015); Saïd & Servanty (2005); Carvalho et al. (2008); Malagnino et al. (2021); Linnell & Andersen (1995); Rossi et al. (2001); Jeppesen (1990).

### 5.3 Author Contributions

Conceptualization: Caroline Millins and Thomas A. Morrison. Data curation: Thomas A. Morrison. Formal analysis: Benjamin M. Marshall. Funding acquisition: Lucy Gilbert, Caroline Millins, and Thomas A. Morrison. Investigation: John Boyle, Mark S. Greener, Robin Gill, Caroline Millins, and Thomas A. Morrison. Methodology: Benjamin M. Marshall, John Boyle, Mark S. Greener, Robin Gill, Caroline Millins, and Thomas A. Morrison. Project administration: Lucy Gilbert, Caroline Millins, and Thomas A. Morrison. Resources: Lucy Gilbert. Supervision: Lucy Gilbert and Mark S. Greener. Visualization: Benjamin M. Marshall. Writing - original draft: Benjamin M. Marshall, Valerio V. Lhamine, Caroline Millins, and Thomas A. Morrison. Writing - review & editing: Benjamin M. Marshall, Lucy Gilbert, John Boyle, Valerio V. Lhamine, Mark S. Greener, Robin Gill, Caroline Millins, and Thomas A. Morrison.

### 6 Supplementary Material



**Figure S1.** Dates of data collection and overall duration or deer tracking by individual.

**Table S1.** Overview of the area estimates resulting from the AKDEs. Area units are hectares. ESS = Effective Sample Size

Region	Animal ID	Point Estimate	Lower CI	Upper CI	Contour level	ESS	Movement Model
Aberdeenshire	Roe01_F	58.56	54.15	63.15	0.90	650.33	OU anisotropic
Aberdeenshire	Roe01_F	76.58	70.80	82.57	0.95	650.33	OU anisotropic
Aberdeenshire	Roe01_F	113.75	105.17	122.65	0.99	650.33	OU anisotropic
Aberdeenshire	Roe02_F	54.38	50.43	58.49	0.90	700.24	OU anisotropic
Aberdeenshire	Roe02_F	73.04	67.73	78.54	0.95	700.24	OU anisotropic
Aberdeenshire	Roe02_F	110.31	102.29	118.63	0.99	700.24	OU anisotropic
Wessex	Roe03_M	91.66	71.57	114.20	0.90	70.87	OU anisotropic
Wessex	Roe03_M	115.43	90.13	143.81	0.95	70.87	OU anisotropic
Wessex	Roe03_M	162.18	126.63	202.05	0.99	70.87	OU anisotropic
Aberdeenshire	Roe04_F	36.02	32.66	39.54	0.90	420.40	OU anisotropic
Aberdeenshire	Roe04_F	46.12	41.82	50.63	0.95	420.40	OU anisotropic
Aberdeenshire	Roe04_F	68.67	62.26	75.39	0.99	420.40	OU anisotropic
Aberdeenshire	Roe05_F	74.51	63.22	86.71	0.90	154.42	OUF anisotropic
Aberdeenshire	Roe05_F	93.05	78.95	108.29	0.95	154.42	OUF anisotropic
Aberdeenshire	Roe05_F	130.88	111.05	152.32	0.99	154.42	OUF anisotropic
Aberdeenshire	Roe06_F	21.74	20.11	23.43	0.90	658.77	OU anisotropic
Aberdeenshire	Roe06_F	32.19	29.78	34.69	0.95	658.77	OU anisotropic
Aberdeenshire	Roe06_F	77.81	71.98	83.86	0.99	658.77	OU anisotropic



Wessex	Roe07_F	57.83	45.81	71.22	0.90	79.39	OU anisotropic
Wessex	Roe07_F	70.58	55.91	86.93	0.95	79.39	OU anisotropic
Wessex	Roe07_F	95.50	75.65	117.62	0.99	79.39	OU anisotropic
Aberdeenshire	Roe08_M	90.73	80.94	101.07	0.90	311.78	OU anisotropic
Aberdeenshire	Roe08_M	118.77	105.95	132.31	0.95	311.78	OU anisotropic
Aberdeenshire	Roe08_M	180.55	161.07	201.14	0.99	311.78	OU anisotropic
Aberdeenshire	Roe09_M	40.99	38.16	43.91	0.90	780.51	OU anisotropic
Aberdeenshire	Roe09_M	55.03	51.23	58.95	0.95	780.51	OU anisotropic
Aberdeenshire	Roe09_M	84.50	78.68	90.53	0.99	780.51	OU anisotropic
Aberdeenshire	Roe10_F	40.70	36.28	45.36	0.90	308.73	OU anisotropic
Aberdeenshire	Roe10_F	53.32	47.54	59.43	0.95	308.73	OU anisotropic
Aberdeenshire	Roe10_F	77.06	68.70	85.89	0.99	308.73	OU anisotropic
Aberdeenshire	Roe11_F	35.06	30.44	40.00	0.90	206.39	OU anisotropic
Aberdeenshire	Roe11_F	42.19	36.63	48.13	0.95	206.39	OU anisotropic
Aberdeenshire	Roe11_F	58.53	50.82	66.78	0.99	206.39	OU anisotropic
Aberdeenshire	Roe12_F	26.06	23.62	28.63	0.90	415.67	OUF anisotropic
Aberdeenshire	Roe12_F	33.88	30.70	37.21	0.95	415.67	OUF anisotropic
Aberdeenshire	Roe12_F	51.06	46.27	56.08	0.99	415.67	OUF anisotropic
Aberdeenshire	Roe13_F	49.35	46.22	52.58	0.90	926.67	OU anisotropic
Aberdeenshire	Roe13_F	66.55	62.34	70.91	0.95	926.67	OU anisotropic
Aberdeenshire	Roe13_F	119.95	112.35	127.79	0.99	926.67	OU anisotropic
Aberdeenshire	Roe14_M	83.25	72.76	94.43	0.90	226.68	OU anisotropic
Aberdeenshire	Roe14_M	122.54	107.10	139.00	0.95	226.68	OU anisotropic
Aberdeenshire	Roe14_M	223.61	195.45	253.64	0.99	226.68	OU anisotropic
Aberdeenshire	Roe15_F	25.49	23.76	27.29	0.90	803.99	OU anisotropic
Aberdeenshire	Roe15_F	33.15	30.90	35.48	0.95	803.99	OU anisotropic
Aberdeenshire	Roe15_F	52.71	49.13	56.41	0.99	803.99	OU anisotropic

**Table S2.** All fixed coefficients from the Poisson population-level habitat selection model. Significance base on whether CI overlap zero.

Variable	Mean Estimate	Standard Deviation	Lower CI	Upper CI	Significance
Road Crossing	-0.7618	0.2045	-1.1747	-0.3601	Significant -
cos_ta	-0.3913	0.0834	-0.5567	-0.2256	Significant -
sl_	0.0007	0.0002	0.0003	0.0011	Significant +
log_sl	0.1053	0.0270	0.0531	0.1597	Significant +
Human Settlements	-69.5327	53.9891	-175.3711	36.3680	Not Significant
Evergreen Needleleaf Forest	-0.3746	0.3195	-1.0171	0.2463	Not Significant
Cropland	-0.2166	0.3803	-0.9880	0.5200	Not Significant
Tall Grassland	0.5660	0.2077	0.1653	0.9843	Significant +
Permanent Wetland	0.8883	1.5102	-2.7146	3.3639	Not Significant
Short Grassland	1.4312	1.2195	-0.9736	3.8558	Not Significant
Open Shrubland	1.4684	0.3958	0.6163	2.2074	Significant +
Distance to Woodland	-0.0051	0.0015	-0.0082	-0.0022	Significant -
Distance to Hedges	-0.0009	0.0005	-0.0019	0.0001	Not Significant
Short Grassland:log_sl	-0.7587	0.2221	-1.1943	-0.3234	Significant -
Open Shrubland:log_sl	-0.2584	0.0477	-0.3528	-0.1659	Significant -
Tall Grassland:log_sl	-0.1804	0.0328	-0.2448	-0.1161	Significant -
Cropland:log_sl	-0.1591	0.0412	-0.2398	-0.0784	Significant -
Permanent Wetland:log_sl	-0.1568	0.1524	-0.4560	0.1419	Not Significant
Evergreen Needleleaf Forest:log_sl	0.0388	0.0442	-0.0482	0.1252	Not Significant
Human Settlements:log_sl	17.4673	13.2464	-8.5035	43.4470	Not Significant
Human Settlements:sl_	-0.1214	0.0825	-0.2835	0.0399	Not Significant
Permanent Wetland:sl_	-0.0021	0.0018	-0.0056	0.0014	Not Significant
Evergreen Needleleaf Forest:sl_	-0.0003	0.0003	-0.0009	0.0004	Not Significant
Open Shrubland:sl_	0.0002	0.0004	-0.0005	0.0009	Not Significant
Tall Grassland:sl_	0.0005	0.0002	0.0001	0.0009	Significant +
Cropland:sl_	0.0012	0.0003	0.0006	0.0017	Significant +

**Table S2.** All fixed coefficients from the Poisson population-level habitat selection model. Significance base on whether CI overlap zero. (continued)

Variable	Mean Estimate	Standard Deviation	Lower CI	Upper CI	Significance
Short Grassland:sl_	0.0044	0.0013	0.0018	0.0070	Significant +

**Table S3.** All fixed coefficients from the individual-level habitat selection models. Significance base on whether CI overlap zero. Individual estimates that were NA have been filtered out.

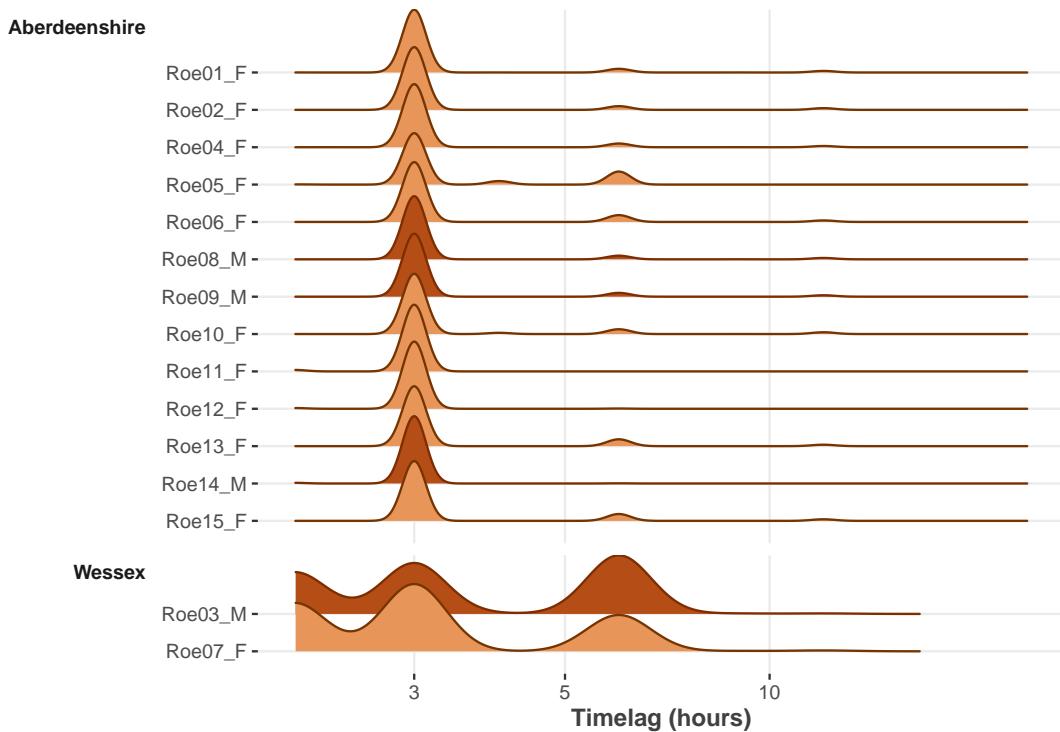
Animal ID	Variable	Mean Estimate	Standard Error	Significance
Roe15_F	Cropland	0.4000	0.1210	Significant +
Roe15_F	Tall Grassland	0.0972	0.0974	Not Significant
Roe15_F	Permanent Wetland	-13.9209	2064.3382	Not Significant
Roe15_F	Other	-13.5494	1041.3412	Not Significant
Roe15_F	Distance to Woodland	-0.0115	0.0015	Significant -
Roe15_F	Distance to Hedges	-0.0023	0.0003	Significant -
Roe15_F	Road Crossing	-1.3295	0.2040	Significant -
Roe15_F	sl_	0.0013	0.0004	Significant +
Roe15_F	log sl_	-0.0273	0.0425	Not Significant
Roe15_F	cos ta_	-0.5156	0.0352	Significant -
Roe14_M	Evergreen Needleleaf Forest	-0.3566	0.1422	Significant -
Roe14_M	Cropland	-2.5724	0.3059	Significant -
Roe14_M	Tall Grassland	-1.1536	0.1202	Significant -
Roe14_M	Short Grassland	-1.8463	0.4051	Significant -
Roe14_M	Human Settlements	-15.6869	947.9684	Not Significant
Roe14_M	Distance to Woodland	-0.0041	0.0006	Significant -
Roe14_M	Distance to Hedges	-0.0021	0.0004	Significant -
Roe14_M	Road Crossing	-0.3079	0.1548	Significant -
Roe14_M	sl_	0.0005	0.0003	Significant +
Roe14_M	log sl_	0.1512	0.0443	Significant +
Roe14_M	cos ta_	-0.2494	0.0444	Significant -
Roe13_F	Evergreen Needleleaf Forest	-0.8290	0.8538	Not Significant
Roe13_F	Cropland	-0.4514	0.1075	Significant -
Roe13_F	Tall Grassland	0.0918	0.0869	Significant +
Roe13_F	Human Settlements	-14.1980	792.3770	Not Significant
Roe13_F	Other	0.4113	0.0990	Significant +
Roe13_F	Distance to Woodland	-0.0049	0.0006	Significant -
Roe13_F	Distance to Hedges	-0.0002	0.0003	Not Significant
Roe13_F	Road Crossing	-1.1608	0.1250	Significant -
Roe13_F	sl_	0.0008	0.0002	Significant +
Roe13_F	log sl_	0.0278	0.0327	Not Significant
Roe13_F	cos ta_	-0.3266	0.0349	Significant -
Roe12_F	Evergreen Needleleaf Forest	0.7455	0.4908	Significant +
Roe12_F	Cropland	-0.0186	0.5504	Not Significant
Roe12_F	Tall Grassland	1.0835	0.4904	Significant +
Roe12_F	Human Settlements	-12.7661	1361.9153	Not Significant
Roe12_F	Distance to Woodland	-0.0097	0.0011	Significant -
Roe12_F	Distance to Hedges	-0.0026	0.0005	Significant -
Roe12_F	Road Crossing	-1.5267	0.2198	Significant -
Roe12_F	sl_	0.0024	0.0006	Significant +
Roe12_F	log sl_	-0.0230	0.0646	Not Significant
Roe12_F	cos ta_	-0.1448	0.0459	Significant -
Roe11_F	Evergreen Needleleaf Forest	1.9865	0.7344	Significant +
Roe11_F	Cropland	-1.4417	1.0209	Significant -
Roe11_F	Tall Grassland	1.7737	0.7350	Significant +
Roe11_F	Short Grassland	0.3272	19511.3285	Not Significant
Roe11_F	Human Settlements	-16.0515	6008.3981	Not Significant
Roe11_F	Distance to Woodland	-0.0056	0.0014	Significant -



Roe11_F	Distance to Hedges	-0.0024	0.0006	Significant -
Roe11_F	Road Crossing	-17.9883	1620.7951	Not Significant
Roe11_F	sl_	0.0027	0.0007	Significant +
Roe11_F	log sl_	-0.0858	0.0798	Significant -
Roe11_F	cos ta_	0.1259	0.0663	Significant +
Roe10_F	Evergreen Needleleaf Forest	-3.4450	1.0068	Significant -
Roe10_F	Tall Grassland	-1.0916	0.1125	Significant -
Roe10_F	Open Shrubland	-0.2218	0.1161	Significant -
Roe10_F	Other	-2.9179	0.5000	Significant -
Roe10_F	Distance to Woodland	0.0056	0.0008	Significant +
Roe10_F	Distance to Hedges	-0.0037	0.0003	Significant -
Roe10_F	Road Crossing	-1.6043	0.2005	Significant -
Roe10_F	sl_	0.0012	0.0004	Significant +
Roe10_F	log sl_	0.0402	0.0442	Not Significant
Roe10_F	cos ta_	-0.7981	0.0366	Significant -
Roe09_M	Evergreen Needleleaf Forest	0.1014	0.0796	Significant +
Roe09_M	Tall Grassland	0.1288	0.1342	Not Significant
Roe09_M	Short Grassland	-0.6686	0.1918	Significant -
Roe09_M	Distance to Woodland	-0.0024	0.0003	Significant -
Roe09_M	Distance to Hedges	0.0001	0.0002	Not Significant
Roe09_M	Road Crossing	17.9570	888.3601	Not Significant
Roe09_M	sl_	-0.0004	0.0003	Significant -
Roe09_M	log sl_	0.0731	0.0447	Significant +
Roe09_M	cos ta_	-0.0604	0.0357	Significant -
Roe08_M	Evergreen Needleleaf Forest	-1.1324	0.5334	Significant -
Roe08_M	Cropland	-0.5180	0.0946	Significant -
Roe08_M	Tall Grassland	-0.6843	0.0997	Significant -
Roe08_M	Permanent Wetland	-14.1362	798.9437	Not Significant
Roe08_M	Human Settlements	-1.6119	0.7258	Significant -
Roe08_M	Distance to Woodland	0.0001	0.0003	Not Significant
Roe08_M	Distance to Hedges	-0.0015	0.0003	Significant -
Roe08_M	Road Crossing	-0.6407	0.0876	Significant -
Roe08_M	sl_	0.0008	0.0003	Significant +
Roe08_M	log sl_	-0.0050	0.0402	Not Significant
Roe08_M	cos ta_	-0.5853	0.0361	Significant -
Roe07_F	Evergreen Needleleaf Forest	0.5556	0.1887	Significant +
Roe07_F	Tall Grassland	0.0465	1.0541	Not Significant
Roe07_F	Open Shrubland	0.7409	0.2894	Significant +
Roe07_F	Distance to Woodland	-0.0068	0.0021	Significant -
Roe07_F	Distance to Hedges	0.0028	0.0007	Significant +
Roe07_F	Road Crossing	-0.2483	0.1817	Significant -
Roe07_F	sl_	0.0019	0.0007	Significant +
Roe07_F	log sl_	-0.0331	0.0668	Not Significant
Roe07_F	cos ta_	-0.1223	0.0799	Significant -
Roe06_F	Evergreen Needleleaf Forest	-14.1975	1168.4078	Not Significant
Roe06_F	Cropland	0.0417	0.0817	Not Significant
Roe06_F	Tall Grassland	-0.0412	0.0776	Not Significant
Roe06_F	Permanent Wetland	1.0868	0.1266	Significant +
Roe06_F	Human Settlements	-13.7704	1151.4231	Not Significant
Roe06_F	Distance to Woodland	-0.0087	0.0007	Significant -
Roe06_F	Distance to Hedges	0.0003	0.0003	Significant +
Roe06_F	Road Crossing	-1.1207	0.0940	Significant -
Roe06_F	sl_	0.0020	0.0004	Significant +
Roe06_F	log sl_	0.0573	0.0464	Significant +
Roe06_F	cos ta_	-0.4989	0.0358	Significant -
Roe05_F	Evergreen Needleleaf Forest	0.2232	0.2018	Significant +
Roe05_F	Cropland	-13.0823	1476.0604	Not Significant
Roe05_F	Tall Grassland	-0.5082	0.1806	Significant -
Roe05_F	Open Shrubland	0.4109	0.1951	Significant +
Roe05_F	Other	-1.5189	0.4414	Significant -



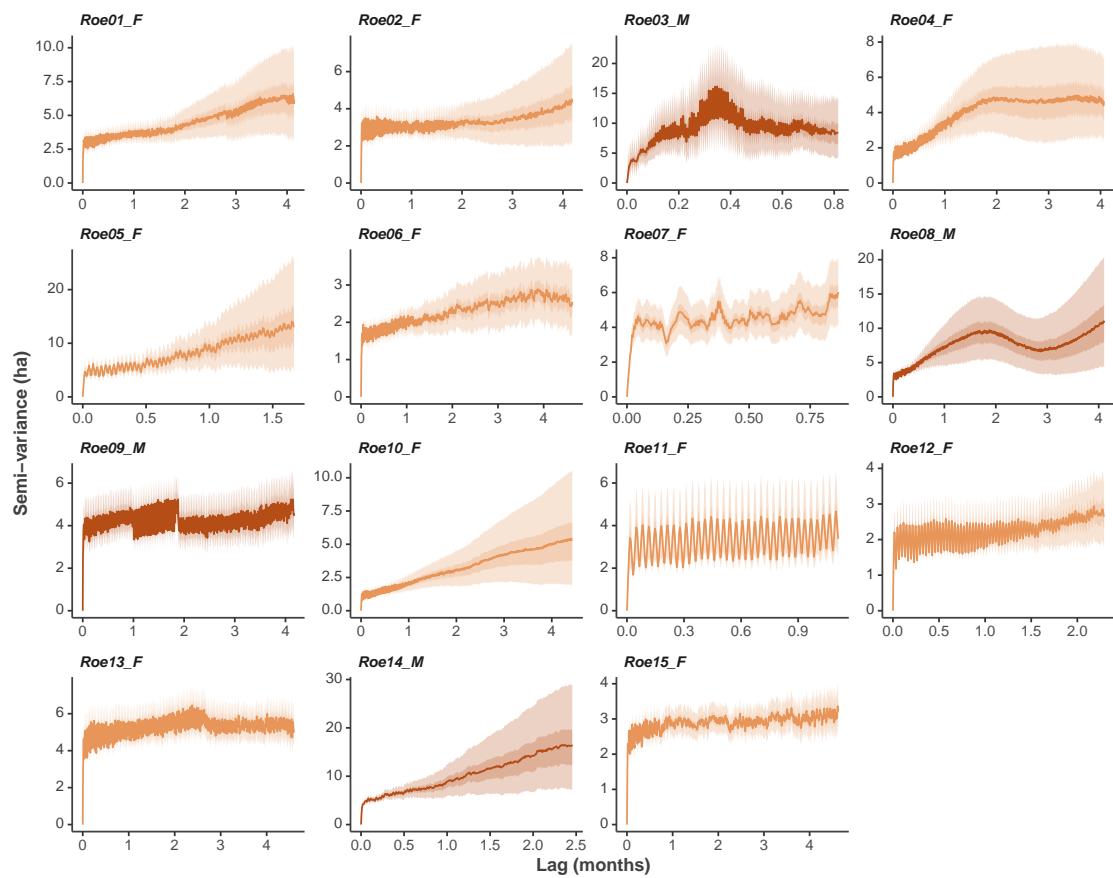
Roe05_F	Distance to Woodland	-0.0092	0.0017	Significant -
Roe05_F	Distance to Hedges	-0.0011	0.0005	Significant -
Roe05_F	Road Crossing	-0.4745	0.2430	Significant -
Roe05_F	sl_	0.0008	0.0004	Significant +
Roe05_F	log sl_	0.0486	0.0579	Not Significant
Roe05_F	cos ta_	0.0288	0.0593	Not Significant
Roe04_F	Evergreen Needleleaf Forest	0.0639	0.0979	Not Significant
Roe04_F	Cropland	0.2897	0.1028	Significant +
Roe04_F	Tall Grassland	0.0710	0.1139	Not Significant
Roe04_F	Permanent Wetland	-15.2921	6251.7909	Not Significant
Roe04_F	Human Settlements	-15.3252	4474.9879	Not Significant
Roe04_F	Other	-14.2898	735.9955	Not Significant
Roe04_F	Distance to Woodland	-0.0016	0.0007	Significant -
Roe04_F	Distance to Hedges	-0.0033	0.0005	Significant -
Roe04_F	Road Crossing	0.0519	0.0708	Not Significant
Roe04_F	sl_	0.0008	0.0004	Significant +
Roe04_F	log sl_	-0.0323	0.0441	Not Significant
Roe04_F	cos ta_	-0.4570	0.0357	Significant -
Roe03_M	Evergreen Needleleaf Forest	0.3222	0.1669	Significant +
Roe03_M	Tall Grassland	1.1273	0.6743	Significant +
Roe03_M	Short Grassland	-9.6499	12303.5977	Not Significant
Roe03_M	Open Shrubland	-13.6255	2092.8520	Not Significant
Roe03_M	Human Settlements	-13.6620	2704.2926	Not Significant
Roe03_M	Distance to Woodland	-0.0526	0.0206	Significant -
Roe03_M	Distance to Hedges	0.0009	0.0007	Significant +
Roe03_M	Road Crossing	-0.0118	0.1714	Not Significant
Roe03_M	sl_	0.0006	0.0006	Significant +
Roe03_M	log sl_	0.0015	0.0994	Not Significant
Roe03_M	cos ta_	-0.8241	0.0904	Significant -
Roe02_F	Evergreen Needleleaf Forest	-0.4986	0.0938	Significant -
Roe02_F	Cropland	-0.9304	0.2062	Significant -
Roe02_F	Tall Grassland	-0.7953	0.1182	Significant -
Roe02_F	Permanent Wetland	-15.5219	2767.4264	Not Significant
Roe02_F	Human Settlements	-14.2981	897.7726	Not Significant
Roe02_F	Other	-14.7043	1090.5249	Not Significant
Roe02_F	Distance to Woodland	0.0002	0.0009	Not Significant
Roe02_F	Distance to Hedges	0.0007	0.0003	Significant +
Roe02_F	Road Crossing	-0.0601	0.0998	Not Significant
Roe02_F	sl_	0.0003	0.0003	Significant +
Roe02_F	log sl_	0.0129	0.0436	Not Significant
Roe02_F	cos ta_	-0.6223	0.0363	Significant -
Roe01_F	Evergreen Needleleaf Forest	-0.7343	0.1979	Significant -
Roe01_F	Cropland	-0.8570	0.1910	Significant -
Roe01_F	Tall Grassland	-0.2884	0.1084	Significant -
Roe01_F	Permanent Wetland	-13.9667	2427.6590	Not Significant
Roe01_F	Human Settlements	-14.1986	799.5390	Not Significant
Roe01_F	Other	-14.3647	1424.3502	Not Significant
Roe01_F	Distance to Woodland	-0.0022	0.0008	Significant -
Roe01_F	Distance to Hedges	0.0013	0.0003	Significant +
Roe01_F	Road Crossing	-0.5058	0.0994	Significant -
Roe01_F	sl_	0.0007	0.0002	Significant +
Roe01_F	log sl_	0.0305	0.0382	Not Significant
Roe01_F	cos ta_	-0.7413	0.0372	Significant -



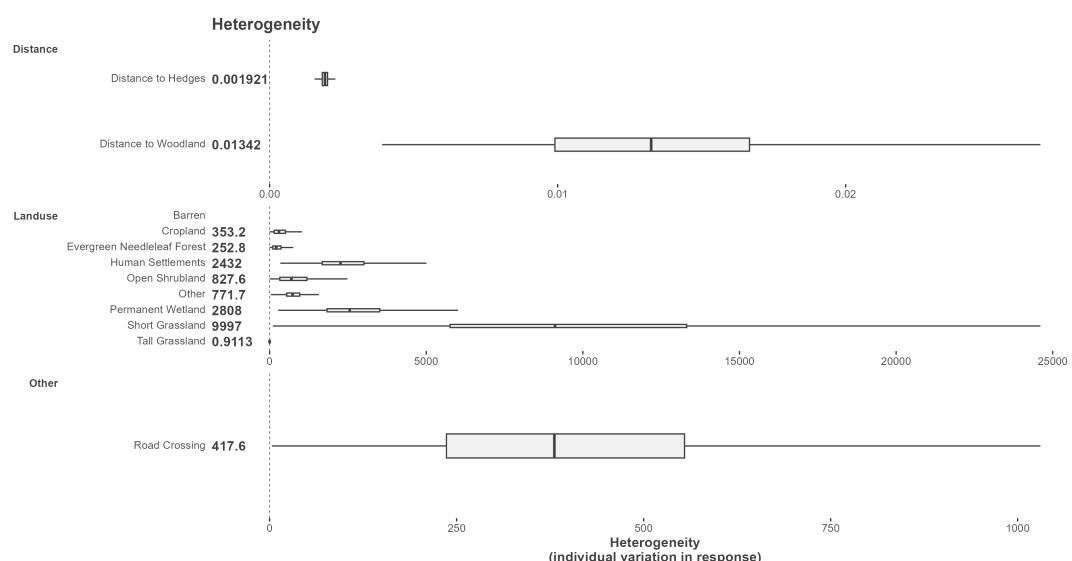
**Figure S2.** Distribution of time lags between Roe Deer location fixes. N.b. x axis is log scaled.

**Table S4.** Resulting coefficients from the population mean selection from all the individual-level habitat selection models. Significance base on whether CI overlap zero. Individual estimates that were NA have been filtered out.

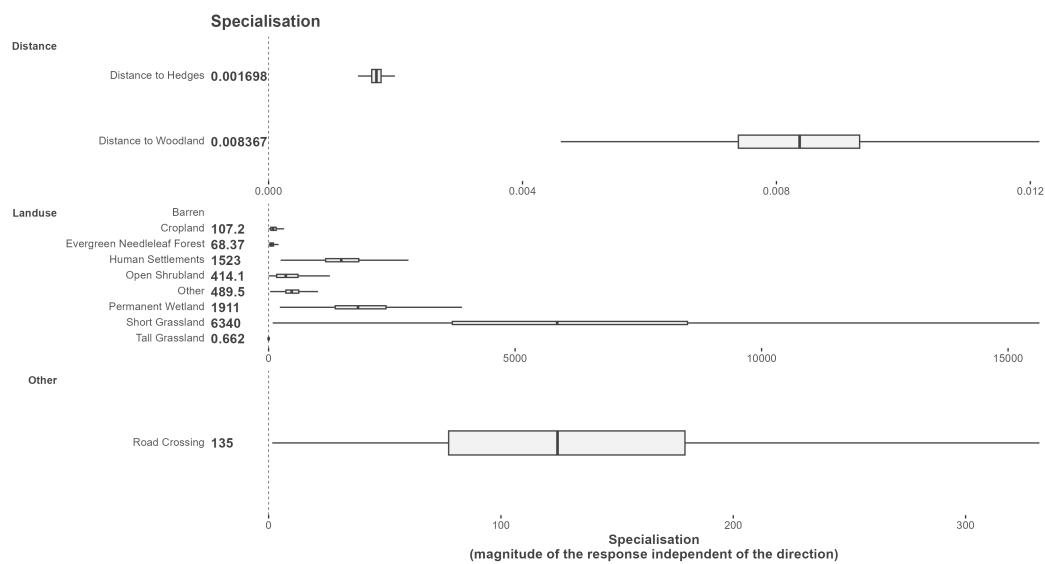
Variable	Mean	Lower CI	Upper CI	Significance
Evergreen Needleleaf Forest	-0.0716	-0.3109	0.1677	Not Significant
Cropland	-0.1911	-0.5608	0.1786	Not Significant
Tall Grassland	-0.2990	-0.5722	-0.0257	Significant -
Short Grassland	-0.8843	-1.7212	-0.0474	Significant -
Open Shrubland	0.0284	-0.6327	0.6895	Not Significant
Permanent Wetland	1.0868	1.0836	1.0901	Significant +
Human Settlements	-1.6119	-1.6303	-1.5935	Significant -
Other	0.2025	-0.5225	0.9275	Not Significant
Distance to Woodland	-0.0051	-0.0095	-0.0008	Significant -
Distance to Hedges	-0.0009	-0.0019	0.0002	Not Significant
Road Crossing	-0.5144	-0.8068	-0.2221	Significant -
sl_	0.0011	0.0006	0.0015	Significant +
log sl_	0.0245	-0.0042	0.0533	Not Significant
cos ta_	-0.4358	-0.5811	-0.2904	Significant -



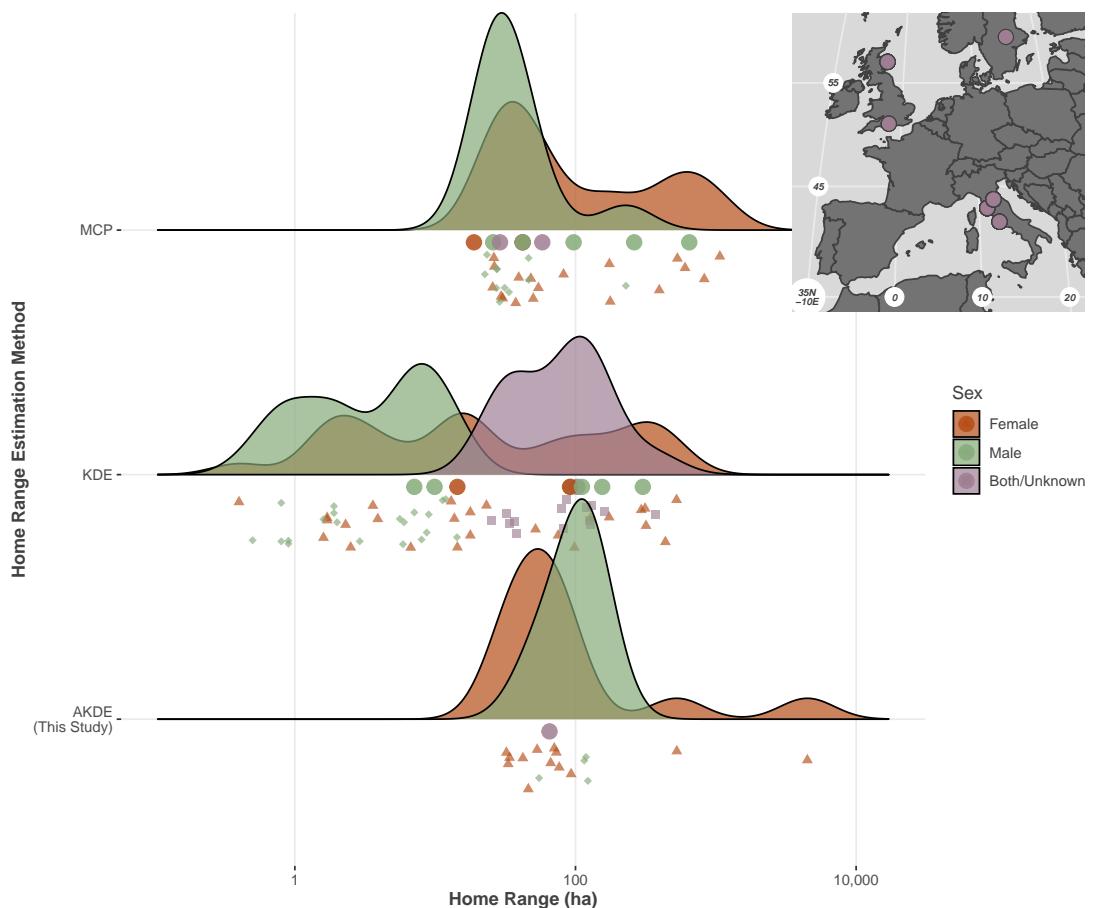
**Figure S3.** Variograms showing the autocorrelative structure of the Roe Deer movement data; The semi-variance (average square displacement) is show over a specific time lag. Examination reveals the level of range residency displayed by each individual (i.e., flattens to an asymptote).



**Figure S4.** The simulated heterogeneity (standard deviation) values from coefficients and standard errors of the individual step-selection models.



**Figure S5.** The simulated specialisation (absolute coefficients) values from individual coefficients and standard errors of the step-selection models.



**Figure S6.** A comparison between reported annual home range sizes present in the HomeRange dataset for Roe Deer, split by estimation method. MCP = Minimum convex polygon, KDE = Kernel Density Estimation, AKDE = Autocorrelated Kernel Density Estimation. Distributions show the spread of individual home range estimates, split by sex. Small points below show the individual estimates (triangles = female; diamonds = male, squares = both/unknown). Note that individuals may appear multiple times if home range estimates were year/season specific. Larger circle points show the estimates provided as population means (total or split between sexes). Inset map shows the study locations, both points within the UK originate from this study. Note x axis is log scaled to accommodate the spread of ranges, particularly the high outliers.

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