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Invasive wild pig carcass availability does not affect coyote abundance in South Carolina, USA

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Short summary

In the United States, control programs often leave wild pig carcasses on the landscape where they are freely available for consumption by scavengers such as coyotes. This study tested if coyote abundance is affected by wild pig carcasses created by control programs, and found coyote abundance did not change in response to wild pig carcass availability. We conclude that wild pig carcass removal will not be an effective coyote management strategy. Photograph by M. Vukovich.

Abstract

Context: Control of invasive wild pigs (*Sus scrofa*) is a growing management concern throughout their invaded range. In the United States, control programs often leave wild pig carcasses on the landscape where they are freely available for consumption by scavengers such as coyotes (*Canis latrans*). Coyotes consume wild pigs, but no studies have investigated the importance of pig carcasses for maintaining coyote abundance.

Aims: We tested the hypothesis that coyote populations may be bolstered by wild pig carcasses created by control programs.

Methods: From July 2014 to July 2017, we surveyed road transects for scat to monitor coyote abundance in response to changing levels of pig carcass availability. From June 2015 to July 2017, wild pigs were removed from the eastern half of the study area and placed at one of four carcass dump sites on the western side. We analysed the scat data using an open population spatially-explicit capture-mark recapture model to estimate seasonal abundance and movement of coyotes in response to the carcass treatment.

Results: Coyote density across the entire study area declined from 95 coyotes/100 km² in July 2014 to 65 coyotes/100 km² by July 2016, but declines were unrelated to the carcass availability treatment. Additional analysis on non-genotyped scat data showed no apparent effect of carcass availability on coyote density. Coyote activity centers moved fewer than 100 meters towards the nearest carcass dump site in response to the carcass treatment.

Conclusions: Wild pig carcass availability is likely not a primary driver of coyote abundance or territory selection at the study location.

Implications: We conclude that the availability of wild pig carcasses had little effect on coyote abundance and thus wild pig carcass removal will not be an effective coyote management strategy.

Keywords: abundance, *Canis latrans*, coyote, invasive wild pig, feral hog, Savannah River Site, supplemental forage, *Sus scrofa*

Introduction

Wild pigs (*Sus scrofa*) are a highly abundant and destructive invasive species throughout their introduced range (Barrios-Garcia and Ballari 2012; Bevins *et al.* 2014; Lewis *et al.* 2019). Originally from Eurasia, wild pigs are now present on all continents except Antarctica as well as several oceanic islands (Mayer *et al.* 2020). Wild pigs cause millions of dollars of damage annually, negatively impacting agricultural crop production, removing native understory cover, and transmitting disease (Barrios-Garcia and Ballari 2012; Lewis *et al.* 2019; Pepin *et al.* 2022). Globally, wild pigs threaten over 670 taxa and have had a major contribution to at least 14 species extinctions (Risch *et al.* 2021). Due to their high fecundity, extensive distribution, and

ability to disperse in response to trapping pressure, wild pig populations present a global management concern (Bevins *et al.* 2014; Risch *et al.* 2021; Garabedian and Kilgo 2024).

Most wild pig control programs typically involve lethal removal of pigs. Common removal methods include trapping, professional sharp-shooting, hunting with dogs, sport hunting, and aerial gunning (Ditchkoff and Bodenchuk 2020, Gaskamp *et al.* 2021, Kilgo *et al.* 2023). Due to food safety concerns and regulations, meat from removed pigs usually cannot be sold or donated (West *et al.* 2009). Wild pigs are known to carry almost 40 parasites that can be transmitted to humans or other wildlife as well as at least 30 viral and bacterial diseases (Miller *et al.* 2017). Regulations on wild pigs vary by country and region, but donation and sale of wild pig meat is banned or discouraged across most of the United States (West *et al.* 2009). As a result, carcasses of removed pigs are typically buried or simply left in the woods.

In the Southeastern United States, where wild pigs are well established and highly abundant (Barrios-Garcia and Ballari 2012, Lewis *et al.* 2019), pigs can be removed in all seasons, potentially providing scavengers with a year-round, low effort food resource. One dominant scavenger that may benefit is the coyote (*Canis latrans*), a highly adaptable and controversial predator in North America. Prior to 1900, coyotes were restricted to the western two-thirds of North America (Hody and Kays 2018), but have since expanded their range to encompass the majority of the continent (Sillero-Zubiri *et al.* 2004). In the absence of red wolves (*Canis rufus*), coyotes are now the dominant carnivore in much of the eastern United States (Berger and Gese 2007), with eastern densities often exceeding those in the western portion of the range (Knowlton *et al.* 1999; Schrecengost *et al.* 2009; Way and Timm 2011). The expansion and increase of coyote populations has led to several management concerns including predation of game and livestock (Knowlton *et al.* 1999; Kilgo *et al.* 2010), interference competition with

other carnivores (Dyck *et al.* 2022), and negative human-wildlife interactions (Poessel *et al.* 2017). Though levels of predation vary throughout the eastern United States, coyotes are the main predator of white-tailed deer (*Odocoileus virginianus*) fawns, and reduced recruitment has been instrumental in localized declines of many deer populations in the Southeast (Kilgo *et al.* 2012; Chitwood *et al.* 2015; Kilgo *et al.* 2019). Consequently, considerable interest in coyote populations exists among hunters and wildlife managers.

Coyotes have a high degree of dietary plasticity, altering their diets in response to seasonal food availability, prey characteristics, and the availability of carrion on the landscape (Knowlton *et al.* 1999; Schrecengost *et al.* 2008; Jensen *et al.* 2022). In addition to ungulates, coyotes have been documented consuming a wide variety of food items, including small mammals, vegetation, birds, and invertebrates (Schrecengost *et al.* 2008; Jensen *et al.* 2022). Coyotes also consume invasive wild pigs, particularly in the early spring when vegetation and ungulate carcasses are less readily available (Turner *et al.* 2017; Watine and Giuliano 2017; Jensen *et al.* 2022). As invasive wild pig populations increase throughout the Southeast (Davis *et al.* 2020; Garabedian and Kilgo 2024), coyote diets may shift to take advantage of this abundant food resource.

In addition to direct predation, scavenging appears to be an important foraging strategy for coyotes, particularly when other food options are scarce or population density is high (Prugh 2005; Schrecengost *et al.* 2008; Jensen *et al.* 2023). Previous studies have noted the importance of carrion in coyote diets (Mastro 2011; Ruprecht *et al.* 2021; Parsons and Young 2024), and wild pig remains were detected in 24.7% of coyote stomach samples in Florida (Watine and Giuliano 2017). However, it remains unclear if wild pig carcass availability directly contributes to higher coyote abundance. Our objectives were to evaluate if wild pig carcasses produced by wild pig control efforts are bolstering local coyote densities.

Methods

Study area

We conducted the study at the United States Department of Energy's Savannah River Site (SRS), a 78,000-ha National Environmental Research Park located in the Southeastern Plain ecoregion of South Carolina (Fig. 1). Upland sites were dominated by planted pine forests comprised of loblolly pine (*Pinus taeda*) and longleaf pine (*P. palustris*). Longleaf pine stands were managed on 120-year rotations, with loblolly stands managed on 50 – 100-year rotations. River drainages and floodplains were occupied by bottomland hardwoods and cypress (*Taxodium distichum*)-tupelo (*Nyssa aquatic* and *N. sylvatica* var. *biflora*) forests. Terrain was generally flat to gently rolling, with elevations ranging from 20 m to 130 m. The climate was humid subtropical, with a mean annual temperature of 18°C and mean annual rainfall of 122.5 cm (Kilgo *et al.* 2017).

Dominant mammalian wildlife in the study area included white-tailed deer, coyotes, bobcats (*Lynx rufus*), raccoon (*Procyon lotor*), Virginia opossum (*Didelphis virginiana*), and wild pig. The wild pig population at the SRS descended from domestic pigs released on the property prior to 1950 and pigs that colonized since that time containing Eurasian wild boar ancestry. Thus, the current population consists of domestic x Eurasian wild boar hybrids (Mayer *et al.* 2020). A targeted control program was implemented in 1985, and since 2010, 1,100 – 1,900 individuals/year have been removed through trapping, shooting, and hunting with dogs. Additionally, an average of 69 individuals/year were taken incidental to recreational deer hunting. Across all age classes, average body mass of removed pigs at the SRS is 39.3 kg, resulting in an average 52,976 kg of carrion annually. At the time of this study, the estimated wild pig density varied across the SRS from 3-20 pigs/km² (Kilgo *et al.* 2021; Kilgo *et al.* 2023).

Coyotes were first reported on site in 1986, with densities reaching an estimated 103 -192 coyotes/100 km² by 2006 (Schrecengost 2007). Prior to 2010, coyotes at the SRS were mostly unexploited, with incidental harvest during white-tailed deer hunting resulting in 20-30 coyotes/year harvested (Kilgo *et al.* 2017). Experimental coyote trapping was conducted from 2010 – 2012, removing 471 coyotes from the property in addition to incidental harvest (Kilgo *et al.* 2017).

Carcass treatments and sampling methodology

We structured our study as a BACI design (before-after controlled impact; Stewart-Oaten *et al.* 1986). Prior to May 2015, wild pig carcasses were left on the landscape at the location where they died and were freely available for scavengers. Total annual culling during 2014–2017 averaged 1,419 pigs. Starting June 1, 2015, all wild pigs killed on the eastern side of the property (hereafter, removal area) were transported to the western side and placed in one of four dump locations (hereafter, carcass area; Fig. 1), while those killed on the western side remained where they were died. Thus, the removal area had essentially no carcasses and the carcass area had approximately double the baseline number of carcasses, both distributed across the landscape and concentrated at dump locations.

From July 2014 to July 2017, we monitored coyote abundance using road transect surveys for scat. We drove transects at approximately 10 km/hr, using the edges of the road as the boundary of our sampling area and collected any scat detected. Sampling was split into nine primary sampling occasions, separated by three-month intervals, to capture potential seasonal changes in coyote abundance. Within each primary occasion, we sampled each transect for four days (secondary occasions). Sampling days were usually three days apart to allow time for scat

to accumulate on the roads. On several occasions, weather or other field conditions impeded sampling, in which case sampling was delayed an additional day. All detected scat samples were collected using tongs and placed in Whirl-pak® bags, which were stored in a cooler with ice packs until the end of the day, whereupon they were placed in a -20°C freezer. For each sample, we recorded the GPS coordinate, date, and transect name.

Laboratory methodology

Due to staffing shortages, we only performed genetic analysis on samples collected from July 2014 to July 2016. We extracted scat samples using Qiagen Mini Stool Kits (Qiagen, Valencia, California, USA) following the manufacturer's protocol. Following the collection of an outer segment of each scat sample to capture epithelial cells, we performed all extractions using two QiaCube automated machines (Qiagen, Valencia, California, USA). We then genotyped each sample across a 9 loci microsatellite panel to identify individuals for subsequent recapture histories. We conducted polymerase chain reaction (PCR) using microsatellite primers FH2001, FH2004, FH2010, FH2054, FH2161 (Francisco et al. 1996); FH2328 (Hellmann et al. 2006); FH2457, FH2062, and FH2289. For every 96-well PCR plate run, we included two negatives of molecular grade water to screen for cross-contamination. The PCR master mix included 1.7 µl of 1x Amplitaq gold PCR Buffer, 1.7 µl 1x Bovine Serum Albumin (BSA), 1.5 µl 2.2µM MgCl₂, 1 µl 0.59 mM Deoxynucleotide (dNTP) Solution Mix (Fisher Scientific #FERR1122), 0.5 µl each of two microsatellite primers, 0.1 µl Tween 20 (Fisher Scientific #BP337-100), 0.06 µl 0.3-unit Amplitaq gold Taq DNA Polymerase (Fisher Scientific #4311814), and 4.94 µl molecular grade water. We used 5 µl of extracted DNA for a total reaction mix of 17 µl. We ran our PCR on a thermocycler for an initial denaturation step of 95°C for 5 minutes and then 20 annealing cycles

at 95°C for 30 seconds, followed by 65°C for 30 seconds with a decrease of 0.5°C every cycle to 55°C and then 72°C for 30 seconds. We ran 20 extension cycles of 95°C for 30 seconds, 55°C for 30 seconds, and 72°C for 30s. We finished PCR at 72°C for 5 minutes.

We genotyped samples on a 3130x capillary machine and used GeneMapper 4.1 (Applied Biosystems) to visualize and score alleles. We required that all homozygous loci be observed in two separate replicates to confirm consensus. Additionally, we selected 500 samples and ran three separate replicates to calculate allelic dropout and false allele rates using RELIOTYPE (Miller et al. 2002). To further test for null alleles and large allele dropout, we selected 304 samples to use MICROCHECKER (Van Oosterhout et al. 2004). Finally, we used DROPOUT (McElvey and Schwartz 2005) to check our genotype dataset for errors caused by low quality DNA. We required that seven loci be confirmed per individual to identify recaptures and built capture histories for downstream analysis from consensus genotypes.

Statistical analysis

We fit the genotyped scat data (scats collected from July 2014 to July 2016) to an open population spatially-explicit capture-mark recapture (SCR) model in a Bayesian framework (Gardner et al. 2009; Gardner et al. 2010). Due to uncertainty associated with the unmarked data, we then fit unmarked spatial capture-recapture models for each primary sampling period to all the scat data collected (July 2014 to July 2017) as an index of population abundance across time. We begin by describing the model for the marked scat data.

Marked spatial capture recapture

We discretized all transects into ‘traps’ spaced 500 m apart and associated each coyote scat sample to the closest point. To reduce bias in SCR analysis, the probability of detecting animals outside the study area needs to be negligible (Borchers and Efford 2008). In previous work on coyotes at the SRS, coyote home ranges averaged 30 – 31 km² (Schrecengost *et al.* 2009); We therefore buffered the boundary of the SRS by 10 km, approximately half the width of a circular coyote home range.

We calculated abundance for each of the 9 primary periods (4 pre-treatment and 5 post-treatment periods). We assumed the population of coyotes was closed during secondary occasions, but open between primary periods. Since we could not separate coyotes by age, we did not distinguish between immigration and reproduction. Instead, we modeled the probability of a coyote entering the population, either by reproduction or immigration, as a single ‘entrance probability’ parameter that varied by occasion. We assumed constant apparent survival throughout the study period.

Individual level data presents a problem when working with Markov chain Monte Carlo because the dimension of the parameter space depends on abundance, N , which changes with each Monte Carlo iteration. To resolve this, we use a data augmentation framework (Royle *et al.* 2007; Royle *et al.* 2013) where we propose a value M , which is some integer value much larger than N . Let $z_{i,t}$ be a binary variable that indicates if an individual is real and present in the population. The probability that individual i is real ($z_{i,t} = 1$) is just $\psi = N/M$. As long as M is much larger than N , the posterior for N is unaffected by the value of M . However, extremely large values of M will greatly increase computation time. Thus, under data augmentation the model for abundance is simply:

$$N_t = \sum_{i=1}^M z_{i,t}$$

$$z_{i,t} \sim \text{Bernoulli}(\psi)$$

Although carcasses were widely distributed throughout the carcass area, to assess the possibility of coyote territories moving in response to carcass availability at carcass dump locations specifically, we fit a biased, correlated random walk model to the activity centers of all coyotes in the study area. Let $\mathbf{X}_{i,t}$ represent the inverse distance weighted average direction of the nearest carcass dump location with respect to each individual's location in primary period $t-1$. If in each time period, individuals were attracted with strength δ towards the carcass dump locations, then the location of individual i 's activity center at primary period t is given by:

$$s_{i,t} \sim \text{Normal}(s_{i,t-1} + \delta \mathbf{X}_{i,t}, \kappa^2)$$

with random error κ^2 . This formulation allows for a strong movement response from coyotes located close to carcass piles, while accounting for the (likely negligible) effect of carcass placement on coyotes with territories hundreds of kilometers away. If carcass dump location had no impact on individual locations ($\delta = 0$), the movement process reverted to a Gaussian random walk.

We modeled detection at each discretized trap location using a half-normal function for detection probability. To allow for seasonal changes in coyote home range size, we modeled σ , the decay of detection probability with increasing distance (d) between the trap and an individual's activity center, as varying by season. We assumed the baseline detection probability, g_0 , was constant across time. The expected number of detections ($\lambda_{i,j,k,t}$) of individual i ($i = 1, \dots, M$) at trap j during survey k in primary period t was:

$$\lambda_{i,j,k,t} = g_0 e^{-d_{ijt}^2 / 2\sigma_t^2}$$

where d_{ijt}^2 is the distance between individual i 's activity center and trap j . Following standard SCR assumptions, home ranges of all individuals were assumed to follow a bivariate normal

distribution, as implied by the capture probability function (Royle *et al.* 2013). The code and data needed to run this analysis can be found in the data archive associated with this manuscript.

We used MCMC to draw posterior samples in NIMBLE 1.0.1 with the 'nimble' package in R 3.6.3 (de Valpine *et al.* 2017; R Core Team 2022; NIMBLE Development Team 2023). We ran three chains for 20,000 iterations each with 15,000 burnin iterations and a thinning rate of two. At each iteration, we calculated abundance in each half of the study area by summing the number of individual activity centers that fell within each region of the study area. We assessed convergence using the Gelman-Rubin statistic (Gelman and Rubin 1992) and visual inspection of the chains.

Unmarked spatial capture recapture

Without genetic analysis, we could not confirm that scats collected after July 2016 were deposited by coyotes. To address this challenge, we analyzed all scats collected (July 2014 to July 2017) using closed-population unmarked spatial capture-recapture models for each primary period to obtain an index of abundance over time. We assumed the proportion of scats incorrectly identified as coyote were constant overtime and did not model movement of coyotes between primary periods. Using the same trap discretization and boundary buffering protocol as in our marked SCR analysis, we calculated abundance for each of the 13 primary periods (4 pre-treatment and 9 post-treatment periods). As demonstrated by (Chandler and Royle 2013), SCR analysis on small datasets of completely unmarked populations may produce biased inference; We therefore only consider the results of these models as an index of population trends, rather than a true estimate of population abundance.

We modeled detection of scats using a binary value to indicate if at least one scat was detected at each discretized trap location j on each secondary occasion k ($k = 1, \dots, K$). Following the binary formulation used in (Margenau *et al.* 2022), the total number of scats collected at each trap in each primary occasion becomes:

$$n_{j,t} \sim \text{Binomial}(K, 1 - e^{-\Lambda_{j,k,t}})$$

where $\Lambda_{j,k,t} = \sum_{i=1}^M \lambda_{i,j,k,t} z_{i,t}$.

We used MCMC to draw posterior samples in NIMBLE 1.0.1 with the 'nimble' package in R 3.6.3 (de Valpine *et al.* 2017; R Core Team 2022; NIMBLE Development Team 2023). We ran three chains for 4,000 iterations each with 3,000 burnin iterations and a thinning rate of one. We assessed convergence using the Gelman-Rubin statistic (Gelman and Rubin 1992) and visual inspection of the chains.

Results

Across 123 km of transect, we collected 1404 animal scats (Fig. S1), 1086 of which were analyzed for species and individual identity. We collected an average of 108 scats per primary period (range: 34 – 216). We successfully obtained individual coyote identification on 509 samples (46.9% genotype success), corresponding to 398 individual coyotes. For our genotyped scats, we detected an average of 56.5 coyote scats per survey (range: 6 – 131) and 1.3 scats per coyote (range: 1 – 6). The majority of coyotes (83%) were only detected in a single primary period.

Median estimates of coyote density from our marked SCR model were consistently higher in the carcass removal area (Fig. 2). The difference in coyote density between carcass treatment areas was more pronounced prior to carcass removal, though there was considerable

overlap in the 95% credible interval estimates between the areas. Across the SRS, coyote density declined from 95 (95% CI: 87 to 105) coyotes/100 km² in July 2014 to 65 (95% CI: 56 to 76) coyotes/100 km² in July 2016.

Median seasonal estimates of σ ranged from 1248.39 in Winter to 457.56 in Fall, suggesting ad-hoc home range sizes (Efford *et al.* 2015) of approximately 4 – 29 km² throughout the annual cycle (Table 1). The estimate of the movement parameter δ was 0.0075 (95% CI: 0.0052 to 0.012), suggesting coyote activity centers shifted on average fewer than 100 meters towards the nearest carcass dump site in response to the carcass treatment. Similarly, κ was estimated at 34.73 (95% CI: 22.77 to 44.25), implying coyote activity centers moved less than 150 meters between primary occasions. Throughout the study period, coyote density was highest in the southwestern region of the study area (Fig. 3).

Density indices from our unmarked spatial capture recapture model suggested that coyote density declined between July 2014 and July 2017, though estimates were much more variable between seasons than the marked SCR estimates. Unmarked estimates were similar between the treatment and control areas, with no obvious differences between the two halves of the study area (Fig. 4). There was also no apparent difference in the average distance between estimated activity centers and pig carcass locations before and after treatment (Fig. S2).

Discussion

Understanding the consequences of leaving wild pig carcasses on the landscape remains an ongoing effort. We used spatially-explicit capture recapture to estimate changes in the abundance and spatial distribution of coyotes in response to the removal of a potential food resource - wild pig carcasses. Though coyote abundance decreased during the study period,

declines began prior to the treatment application and did not substantially change in response to carcass removal. The overall impact on both coyote abundance and spatial distribution was negligible, suggesting the current strategy of leaving wild pig carcasses in the woods has minimal impacts on local coyote populations.

If coyotes were dependent on wild pig carcasses as a primary food source, we would have expected to see the highest coyote densities closest to the carcass dump sites. Ecological theory suggests clumped high quality food resources will result, on average, in smaller territories with increased space use near the food resource (Young *et al.* 2008; Sells and Mitchell 2020). Yet, despite the abundant carrion available at carcass dump sites, we found little evidence to suggest coyotes were attracted to these locations. Conversely, if removal of carrion reduced the carrying capacity of coyotes on the landscape, we would have expected coyote densities to decline in the treatment areas. Resident coyotes can rapidly transition to transient behavior irrespective of age and social status in response to ecological demands (Hinton *et al.* 2015). However, the spatial distribution of coyotes remained stable throughout the study period, even as overall abundance declined. Though previous studies at the SRS have found evidence of pig consumption in coyote scat (Schrecengost *et al.* 2008, Ward *et al.* 2018, Youngmann *et al.* 2023), wild pig does not appear to be a primary driver of population dynamics for this species.

The importance of scavenging for maintaining coyote populations remains an open question. Ward *et al.* (2018) argued that scavenging was not an important foraging strategy for coyotes in the Southeast, noting the infrequency of both roadkill species and wild pig in coyote scat, despite their widespread distribution within the study area. Previous research on western coyotes found that carrion represented the most commonly used prey when primary prey species were in decline, but only comprised a small proportion of the diet when primary prey species

were abundant (Prugh 2005). These findings suggest that scavenging is most important when food resources are limited and may mostly be employed by coyotes experiencing food scarcity. Given the observed decline in coyote abundance within our study area and previous research suggesting eastern coyote populations are often below carrying capacity (Morin *et al.* 2016), food scarcity may be uncommon for coyotes in this region. Thus, while coyotes on the SRS may opportunistically consume wild pigs, the availability of white-tailed deer and other preferred food resources may negate the need for reliance on scavenging.

Few of the individual coyotes detected by our scat surveys were detected in more than one primary period, suggesting a high proportion of transient individuals. Both transience and dispersal are common in coyote populations (Harrison 1992; Windberg *et al.* 1997; Morin and Kelly 2017), and individual coyotes can disperse distances upwards of 390 km (Harrison 1992; Hinton *et al.* 2012). Transient coyotes are less reliant on local food sources and may exhibit different patterns of prey selection compared to resident coyotes (Mills and Knowlton 1991; Gese *et al.* 1996). While resident coyotes may be more directly impacted by removal of pig carrion on the landscape, there is little evidence to suggest wild pigs are a primary food resource for Southeastern coyotes (Ward *et al.* 2018, Jensen *et al.* 2022, but see Watine and Giuliano 2017). Moreover, the high proportion of transient individuals in many eastern coyote populations (Hinton *et al.* 2015; Webster *et al.* 2022) implies that local removal of a non-primary food resource is unlikely to result in meaningful reductions in coyote abundance.

One difficulty with assessing coyote response to environmental change is the large study area required to accurately capture local coyote dynamics. Coyote home range sizes vary seasonally and depend in part on the abundance of food resources within a pack's territory (Schrecengost *et al.* 2009; Ward *et al.* 2018; Youngmann *et al.* 2022). Previous work at the SRS

found home range size ranged from 5.4 – 39.2 km² across the annual cycle with a core area size of 1 – 6.1 km² (Ward *et al.* 2018). These estimates suggest the size of our study area was sufficient to detect changes between treatment blocks, but would not have been large enough to allow for spatial replication of the carcass removal treatment. Thus, while our results suggest current pig management efforts are not subsidizing coyote populations on the SRS, we acknowledge these findings may be limited to this geographic region.

One limitation to our study is the lack of direct diet information from the coyotes in our study area. Previous research has demonstrated that coyote populations can respond rapidly to changes in food availability or habitat conditions, often in the form of prey switching (Andelt and Andelt 1984; Prugh 2005). Without additional information on the diets of the coyote population during our study period, we cannot rule out the possibility that the apparent lack of response to the carcass treatment was simply the product of coyotes shifting their diets to include alternative food sources. However, if coyotes on the SRS can shift their diets to alternative food sources without suffering noticeable declines in abundance, the pig carcasses produced by current pig management efforts are unlikely to represent a meaningful subsidy for the coyote population.

Though we found no effect of wild pig carcass availability on coyote abundance, our findings suggest a decline in the coyote population at the SRS between 2014 and 2016. Schrecengost (2007) estimated the 2006 coyote density at the SRS as 103 -192 coyotes/100 km², compared to our July 2014 estimate of 95 (87 - 105) coyotes/100 km². Recent research by Youngmann (2023) reported densities of 15 – 36 coyotes/100 km² in 2019, much lower than our July 2016 estimate of 65 (56 - 76) coyotes/100 km². Similarly, estimates of the numbers of coyotes shot by deer hunters statewide in South Carolina declined between 2014 and 2021

(South Carolina Department of Natural Resources 2022). Yet to our knowledge, little evidence exists to suggest coyotes are declining elsewhere in the Southeast. On the contrary, most coyote studies report increasing populations, even in areas with substantial hunting or trapping pressure (Conner and Morris 2015; Morin *et al.* 2016). Although we did not investigate the drivers of the decline we observed, as this was beyond the scope of our study, we suggest additional monitoring is warranted to determine whether coyote abundance may be moderating in the state after initially increasing post-colonization.

Invasive wild pigs are numerous and pig management efforts create a large volume of carrion on the landscape. Carrion produced by human activities can increase scavenger presence and alter predator dynamics on the landscape (Newsome *et al.* 2015; Turner *et al.* 2017; Spencer *et al.* 2021). Additional food resources for apex predators can decouple interactions between predators and prey populations (Letnic and Dickman 2010), potentially leading to surges in mesoscavenger populations (O'Bryan *et al.* 2019). Though the consequences of increased pig carcasses for other scavengers remains unclear, our results suggest that pig carcasses produced by current pig management efforts are not subsidizing coyote populations in the southeastern United States. Our work highlights the need to study how invasive animal carcasses are impacting other scavenger populations, both within the southeastern United States and globally.

Data Availability Statement

All data and code are publicly available at: <https://zenodo.org/records/13912786>

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Conflict of interest statement

The authors declare no conflict of interest.

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627

628 Figure captions:

629 Figure 1. Location of 123 km of scat transects at the Savannah River Site, South Carolina, USA.
630 Transects were located on existing roads with surveys repeated over 9 periods from 2015 to 2016
631 to estimate density and spatial distribution of coyotes. Pigs hunted on the eastern half of the
632 property (blue polygon) were placed at carcass dump sites (pig icons) on the western side (beige
633 polygon) starting June 1, 2015.

634

635 Figure 2 Density of coyotes per 100 km² at the Savannah River Site, South Carolina, USA from
636 July 2014 to July 2016 before and after pig carcass removal treatment. Error bars show the
637 median and 95% credible intervals for the eastern (blue) and western (brown) halves of the study
638 area. The vertical dotted line signifies when pig carcasses were first removed from the eastern
639 half of the study area.

640

641 Figure 1. Heat map of coyote activity centers at the Savannah River Site, South Carolina, USA
642 from July 2014 to July 2016. Points represent individual coyote activity centers within each 500
643 meter hex. Areas in white represent hexes with an estimated density lower than 5 coyotes per 100
644 km². Only coyote territory centers located within the study area boundary are shown.

645

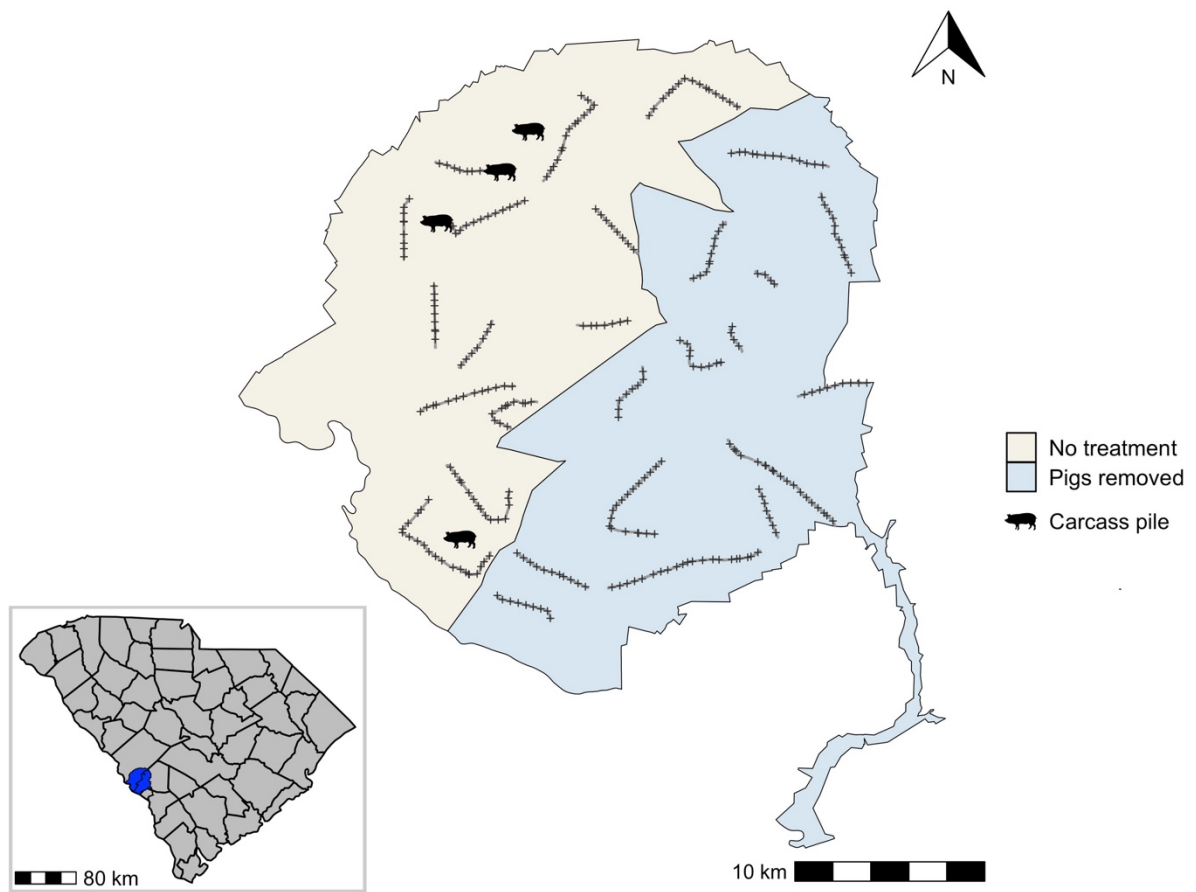
646 Figure 4. Index of coyote density per 100 km² based on unmarked coyote scat data collected at
647 the Savannah River Site, South Carolina, USA from July 2014 to July 2017. Dotted lines show
648 the 95% credible intervals for index estimates. The vertical dotted line signifies when pig
649 carcasses were first removed from the eastern half of the study area.

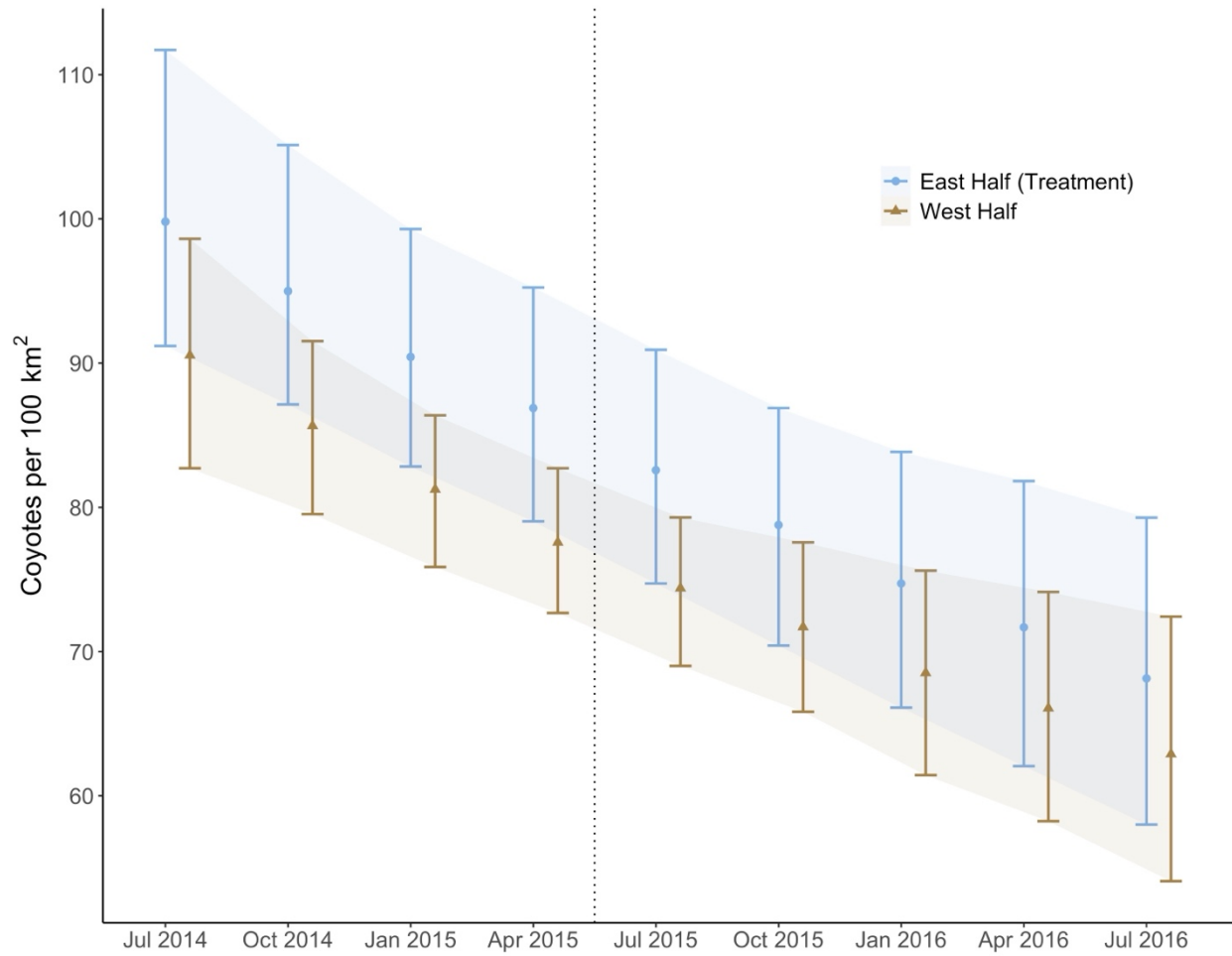
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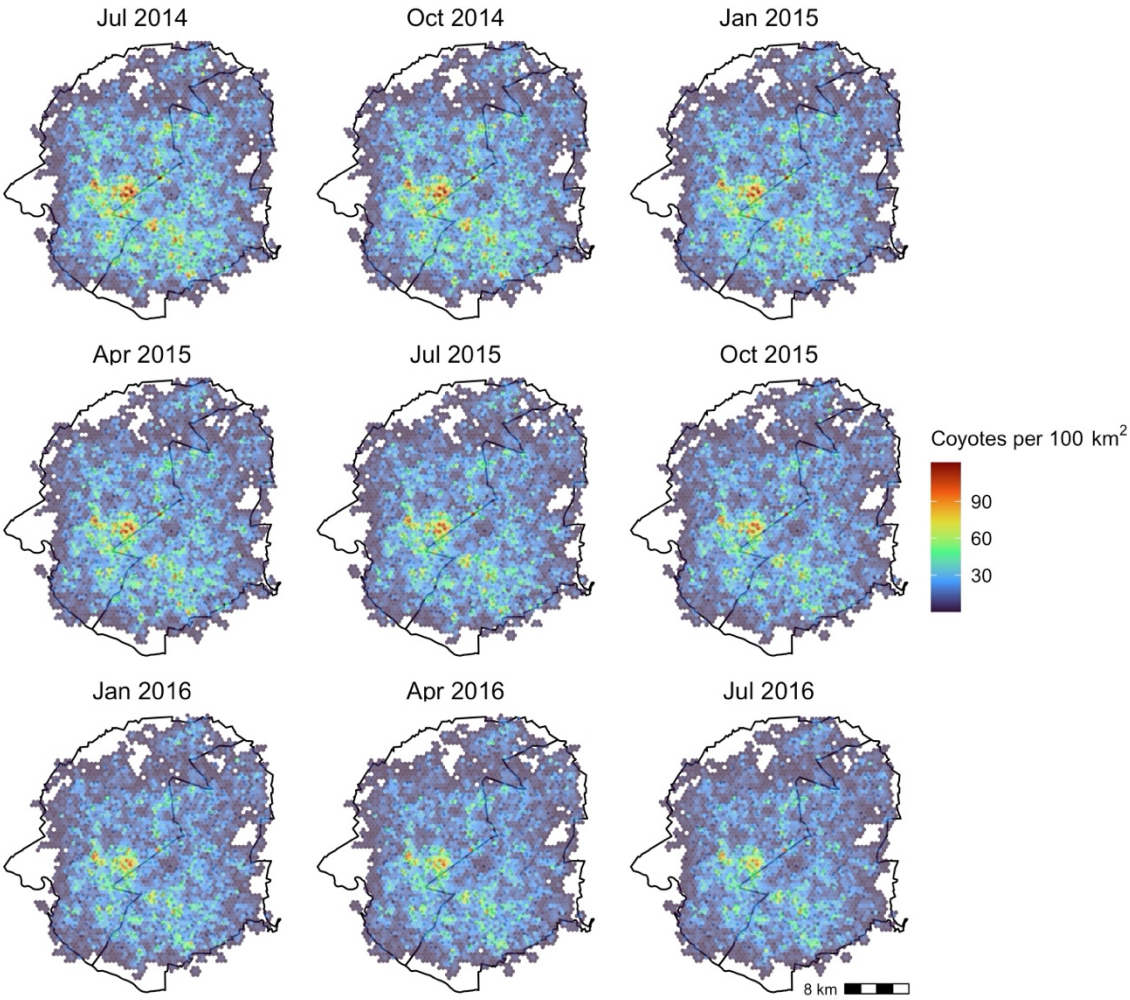
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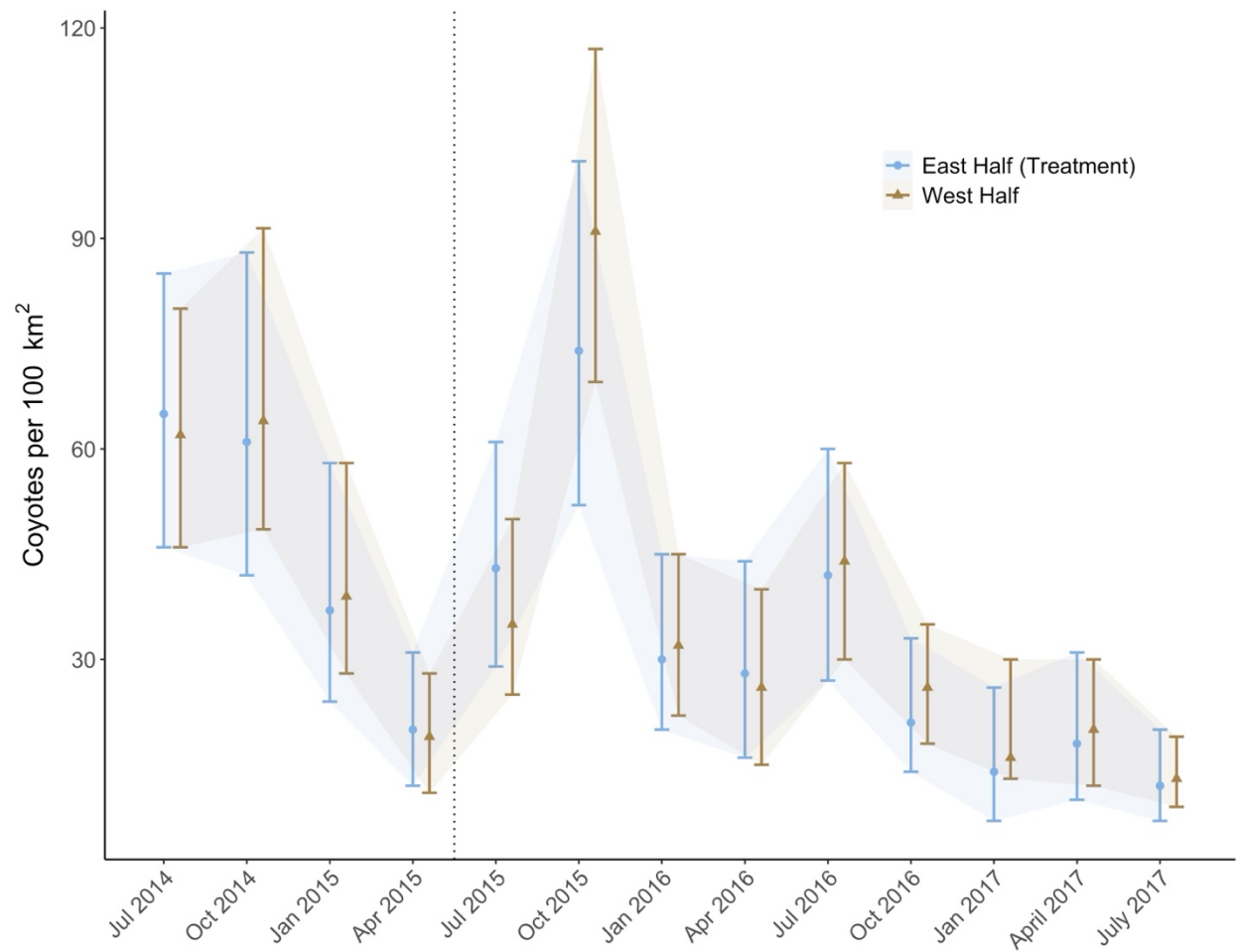
Table 1. Parameter median estimates and 95% credible intervals for an open population spatial capture recapture model of coyote abundance at the Savannah River Site, South Carolina, USA from July 2014 to July 2016.

Parameter	Description	Estimate	95% CI
g_0	Baseline detection	0.01	0.005, 0.02
σ_{Winter}	Detection decay parameter - Winter	1248.39	1116.12, 1310.77
σ_{Spring}	Detection decay parameter - Spring	1197.23	1124.58, 1270.1
σ_{Summer}	Detection decay parameter - Summer	575.56	469.98, 663.74
σ_{Fall}	Detection decay parameter - Fall	457.56	363.37, 575.02
δ	Attraction to carcasses	0.0075	0.0052, 0.012
κ	Movement parameter – sd	34.73	22.77, 44.25









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