# Abstract

# **Introduction**

- 3 Pigs eat everything. But they eat some things more than others. And when those "some things"
- 4 are agricultural crops, it can result in extensive economic damage. [estimate]. However, crops
- 5 are one of many forage resources competing for a feral swine's attention on a landscape. As
- 6 extreme generalists, feral swine consume anything from sapling trees, fungus, and masting seeds
- 7 to small invertebrates and livestock. The availability of these different forage resources varies
- 8 in both space and time, such that pigs will switch their foraging behavior availability of these
- 9 various potential resources

# 10 Methods

### 11 Data

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12 To analyze how forage availability affected pig movement patterns, we used GPS collar data

collected on 500 pigs in the United States of America and Canada (Fig. 1). Of the X pigs, X

were male and Y were female. These data are from X different studies and were collected from

May, 2004 to November, 2017. The median collaring time per pig across all studies was X (95%

quantiles) (see supplementary material for additional details). The average time between GPS

fixes varied by study, with the median fix time being X across all Y studies [95% CI].

For each pig used in the study, we cleaned the movement trajectory using the following

criteria. First, we excluded all 2D GPS fixes from the analysis (Bjorneraas et al. 2010). Second,

we eliminated the first and last 25 fixes for each pig to account for capture effects [citation].

Third, we eliminated all pig movements in which pigs moved faster than 50 km per hour over

any distance or moved greater than 15 km per hour over a distance of 8 km or more. These

movements are unlikely given previously observed patterns of feral swine movement (Mayer &

24 Brisbin 2009). Finally, we also removed large temporal gaps between fix times as these can lead

to a large amount of uncertainty regarding a pig's location in between two temporally distant

fixes. To do this, we split a single pig movement trajectory into multiple sequences when the time

between two adjacent fixes was greater than c. 130 minutes. We then discarded all sequences

that had less than 150 fixes at least 130 minutes apart. The remaining sequences were considered

independent "runs" for a given pig. We chose to only include sequences of 150 fixes as shorter

runs would not always span the length of a day, precluding analysis of diel movement patterns,

and longer runs excluded too many of the pigs from the analysis. After this cleaning, our analysis contained X pigs and a total of X fix times (Fig. 1).

#### Covariates

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The goal of our analysis was to understand how the availability of anthropogenic and natural 34 forage on a landscape affected the movement and resource selection of feral swine. To address 35 these goals, we identified covariates related to natural forage and anthropogenic forage resources 36 that were comparable across the different studies used in this analysis. For the availability of 37 natural forage resources, we considered two proxies: plant productivity as measured by Normal-38 ized Difference Vegetation Index (NDVI) and density of masting trees (Table 1). We included 39 masting tree density as masting events are an important component of pig diets and can have 40 significant implications on population-level pig growth rates (e.g. Bieber & Ruf 2005). NDVI 41 is a general correlate of plant productivity in an area (Pettorelli et al. 2005) and, given that 42 pigs are extreme generalists and plant material often makes up a large percentage of their diet 43 (Mayer & Brisbin 2009), we use NDVI to reflect the availability of herbaceous forage. 44 For anthropogenic forage resources, we used agricultural data available from the National 45 Agricultural Statistics Service (NASS). These data are obtained yearly at a 30m by 30m scale 46 across the contiguous US. Each pixel specifies the dominant type of crop grown in that pixel. 47 While the NASS data enumerates 105 types of crop, we chose not to distinguish between crop 48 types in our initial analyses. Instead, we used a generic grouping of "crop" vs. "no-crop", which 49 allowed us to more easily explore how the effect of agricultural forage on pig movement varied 50 across space and time [though maybe specific croptypes would be better]. We did consider 51 52 specific crop types in our post-hoc analyses, in which we grouped the 105 crop types into 11 groups: cereals, oilseed, tobacco, beverage and spice, leguminous, grasses, sugar, root and tuber, 53 fruit and nuts, vegetables and melons, and other crops [according the criteria outlined in...]. 54 WHAT ABOUT CANADA? 55 While our primary goal of this analysis was to understand the effect of forage availability 56 on pig movement, pig movement and home range size is also driven by a number of other vari-57 ables, including cover, water availability, temperature, pressure, human development, mammal 58 diversity, sex, and age (McClure et al. 2015; Garza et al. 2017; Kay et al. 2017) [MORE]. Of 59 these additional predictors of pig movement, cover and water are critically important as both 60 are necessary for pigs to thermoregulate (Choquenot et al. 1996) and cover, in particular, is 61

necessary for protection from predators [citation] (citations in Mayer2009). We used tree canopy

density data [check wording] from the National Landcover Database (NLCD) as a covariate for cover (Table 1). These data are available at a 30m by 30m scale for the contiguous United States. For a water, we used the National Wetland Inventory (NWI) Database which identifies X types of water bodies larger than X m2 in the continguous United States. For this study, we only considered the water sources that were permanent or semi-permanent as defined by the NWI and computed the distance to the nearest water source at a 30m by 30m resolution (see Table 1 and Sup Mat).

We also included mean monthly temperature, total monthly precipitation, and distance to

We also included mean monthly temperature, total monthly precipitation, and distance to developed land/roads as additional covariates (Table 1). Note that while the inclusion of these covariates is import for capturing pig movement, our questions were focused the foraging resources, such that we considered non-foraging covariates "blocking" covariates and did not exhaustively explore their potential relationships and interactions beyond what had already been shown in the literature.

Finally, we also considered a number of study-level variables to allow us to explore how and why the effect of natural forage resources and anthropogenic forage resources varied across different studies. To this end, we explored study-level covariates such as drought severity, ecoregion, local pig density, and MORE (see Table 1).

# Continuous-time, discrete-space movement model

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To understand role of forage resources on pig movement, we used the modeling framework 81 of Hanks et al. (2015) and Wilson et al. (2018). Generally, this framework leverages auto-82 correlated animal movement data and gridded raster covariates to make inference about the 83 84 resource utilization of an animal (Hanks et al. 2015; Buderman et al. 2018; Wilson et al. 2018). Specifically, this approach can be broken into two distinct steps. Given a trajectory of GPS 85 fixes (not necessarily with equal fix times), the first step of this approach estimates animal 86 movement as a function of continuous time at some particular temporal grain (e.g. 15 minutes). 87 To do this, we used a phenomenological functional movement model (FMM) (Buderman et al. 88 2016; Hooten et al. 2017), which is a non-mechanistic, continuous-time movement model that can 89 capture an animal's movement patterns at some desired-level of detail (Buderman et al. 2016). 90 The phenomenological FMM can be represented as a series of basis functions, which allow for 91 large flexibility in animal movement patterns. In particular, we used a B-spline basis expansion 92 to model the longitude and latitude of an animal as a function of time (see Supplementary 93 Material for additional detail). 94

In the second step of the analysis requires translating our continuous movement model into discrete, rasterized space. To do this we used our fitted FMM to predict a pig's location at 15 minute intervals. We chose 15 minutes as this provided a reasonable trade-off between computational time and computing the exact amount of time spent in each cell, which is equivalent to letting the time between fixes go to zero [NOTE: Would this be hard to do? Double Check interpretation of 15 minute intervals...]. We then converted this predicted pig path into discrete 30m by 30m grid cells such that we were left with a continuous time movement path in discrete space (Fig. X). [We chose 30m by 30m ] To account for the uncertainty in our movement model, we generated 20 continuous-time discrete space movement paths and performed all of the analyses described below on each of the movement paths (Hanks et al. 2015; Buderman et al. 2018).

Given these continuous-time, discrete-space trajectories, we then explored how agricultural and natural forage resources on a landscape affected pig movement using a continuous-time Markov Chain (CTMC) approach Hanks  $et\ al.\ (2015)$ . The CTMC approach considers continuous-time animal movement (i.e. the FMM model described above) through a discrete landscape. From this point-of-view, animal movement can be considered as a series of rates of moving from cell i to an adjacent cell j,  $\lambda_{ij}$ . As for any continuous-time Markov Chain, the process can be decomposed into the waiting time before a state change occurs (i.e. the time an animal spends in a cell) and the new state once a change occurs (i.e. the new cell to which the animal has moved) (Allen 2003). With this interpretation, one can then model the rate of moving between cell i and j  $\lambda_{ij}$  as a function of the environmental covariates in cell i and j. Hanks  $et\ al.\ (2015)$  showed this inference can be re-expressed as a latent-variable, Poisson Generalized Linear Model, where the response variable for adjacent cell j is one if a pig moved to that cell from cell i, and 0 otherwise. Specifically, let  $z_{ij}$  be the zero/one latent variable, then

$$z_{ij} \sim \text{Poisson}(\lambda_{ij})$$
 (1)

$$\log \lambda_{ij} = \log \tau_{ij} + \beta \mathbf{X} \tag{2}$$

where  $\tau_{ij}$  is the waiting time before moving from cell i to cell j,  $\mathbf{X}$  is a vector of landscape covariates, and  $\beta$  is the effect of these covariates on movement.

Considering **X**, we explored two classes of covariates: location-based drivers and directional drivers of movement (Hanks *et al.* 2015). Location-based drivers are a result of the cell that an animal is currently in and affect how long an animal remains in the current cell. For example,

if masting tree density was a negative location-based driver of pig movement then a pig in a cell with high masting tree density would tend to remain in that cell longer than a cell with lower masting tree density. Directional drivers of movement determine the direction that a pig might move once it leaves the cell it is currently occupying. For example, if masting tree density was a positive directional driver of pig movement then, upon leaving the currently occupied cell, a pig tends to move in the direction of increasing masting tree density, relative to its current position [TODO: THIS IS A LOCAL dirIENT EFFECT...SHOULD WE ALSO EXPLORE GLOBAL EFFECTS AT SOME POINT? IN OTHER WORDS, does the presence of masting trees some where in the vicinity affect how a pig uses crop?]. Table 1 shows which of the covariates described in the previous section we considered as location-based drivers, directional drivers, or both.

## 133 Model specification and fitting

- We analyzed three variations of the CTMC model with the goal of understanding the role of foraging resources on the movement of feral swine across populations. Each model is described below
- 137 Model 1: Main effects model

The first, and simplest, model that we analyzed only considered time-invariant main effects of anthropogenic and natural foraging resources as well as three other non-foraging covariates known to affect pig movement: canopy cover, sex, and distance to water. The model is defined as

$$\log(\lambda_{ij}) = \log(\tau_{ij}) + \beta_0 + \beta_1 \text{directional persistence} + \beta_2 \text{male}$$
(3)

$$+ \beta_3 \text{canopy cover}_{\text{loc}} + \beta_4 \text{canopy cover}_{\text{dir}}$$
 (4)

$$+ \beta_5 \text{distance to water}_{\text{loc}} + \beta_6 \text{distance to water}_{\text{dir}}$$
 (5)

$$+ \beta_7 \text{distance to development}_{\text{loc}} + \beta_8 \text{distance to development}_{\text{dir}}$$
 (6)

$$+ \beta_9 \text{distance to } \text{crops}_{\text{loc}} + \beta_{10} \text{distance to } \text{crops}_{\text{dir}}$$
 (7)

$$+ \beta_{11}$$
masting tree density<sub>loc</sub>  $+ \beta_{12}$ masting tree density<sub>dir</sub> (8)

$$+ \beta_{13} \text{NDVI}_{\text{loc}} + \beta_{14} \text{NDVI}_{\text{dir}} \tag{9}$$

where "directional persistence" describes the propensity of pigs to continue to move in the direction they moved in the previous time step. the subscript "loc" refers to a location-based covariate (i.e. how that covariate effects the rate an animal is moving) and the subscript "dir" refers to a directional covariate (i.e. how that covariate effect the direction in which an animal moves). We fit this model in two steps. First, we fit the model using LASSO regularization using the glmnet package (Friedman et al. 2010). LASSO simultaneously performs variable selection and shrinks [non-predictive or highly correlated] covariates to zero James et al. (2013). To select the best-model in terms of predictive performance, we performed 5-fold cross-validation [NOTE: that I probably need to do cross-validation for case-control studies] and chose the model with the maximum penalization parameter for which the Poisson deviance was within one standard error of the minimum out-of-sample deviance (provided in glmnet). This approach increases regularization, with only a small cost to deviance [and allowed us to more easily perform our second set .

While advantageous for model selection, the aforementioned LASSO approach does not provide easy access to parameter uncertainty or incorporation of random effects. We then refit the model model X, excluding the parameters removed by the LASSO, in a Bayesian framework. For each  $\beta_i$  coefficient in the model, we included a random effect of individual. Specifically, given k individual pigs in a population, we allowed  $\beta_{i,k} \sim N(\mu_{\beta_i}, \sigma^2 \phi_i)$ , which indicates that individual-level effects for parameter i are drawn from a population-level distribution (Buderman et al. 2018). When i > 1,  $\log(\phi_i) \sim N(0, 0.04)$ , which weights the variance population-level variance for each i. When i = 1,  $\phi_1 = 1$ . For the population-level means, we specified the hyper-prior  $\mu_{\beta_i} \sim N(0,3)$ . In this model, the random effects of individuals for each covariate account for variation in resource use between individuals that were sampled at different times, in addition to inherent between-individual variability.

Given the size of the "data" for each study (e.g. most studies had > 1 million predicted positions from the FMM model), we used automatic differentiation variational inference (ADVI) to approximate the posterior distributions of this Bayesian model (Kucukelbir *et al.* 2015). [Description of ADVI?]. Combining both ADVI and the preliminary LASSO regularization, allowed us to fit these models in a reasonable amount of time (e.g. X hours).

### Model 2: Time-varying movement model

The second model we considered allowed pig movement and resource use to vary with time.

We allowed overall movement speed to vary on both a daily and monthly time-scale and also allowed selection for forage resource, cover, and water to vary on a monthly time scale. The model is given by

$$\log(\lambda_{ij}(h,m)) = \log(\tau_{ij}) + \beta_0(h) + \beta_0(m) \tag{10}$$

$$+ \beta_1 \text{directional persistence} + \beta_2 \text{male}$$
 (11)

$$+ \beta_3(m)$$
canopy cover<sub>loc</sub>  $+ \beta_4(m)$ canopy cover<sub>dir</sub> (12)

$$+ \beta_5(m)$$
distance to water<sub>loc</sub>  $+ \beta_6(m)$ distance to water<sub>dir</sub> (13)

$$+ \beta_7 \text{distance to development}_{\text{loc}} + \beta_8 \text{distance to development}_{\text{dir}}$$
 (14)

$$+\beta_9(m)$$
distance to  $\operatorname{crops_{loc}} + \beta_{10}(m)$ distance to  $\operatorname{crops_{dir}}$  (15)

+ 
$$\beta_{11}(m)$$
 masting tree density<sub>loc</sub> +  $\beta_{12}(m)$  masting tree density<sub>dir</sub> (16)

$$+ \beta_{13}(m) \text{NDVI}_{\text{loc}} + \beta_{14}(m) \text{NDVI}_{\text{dir}}$$
(17)

where h indicates the hour of the day and m indicates the month of the year.  $\beta_0(h)$  describes how overall movement rate changes as a function of the hour of the day.  $\beta_i(m)$  describes how the effect of a particular covariate varies with the month of the year. We modeled  $\beta_0(h)$  using cyclic splines to ensure movement rate at the start of a 24 hour period was the same as movement rate at the end of the 24 hour period. We modeled  $\beta_i(m)$  using cyclic-cubic regression splines when the data for pigs in a populations spanned 12 months (Wood 2006). This ensured that the effect of a particular covariate was the same at the end December and the beginning of January. When the data for a particular population did not span 12 months [NOTE: rarely, if ever, does a single pig span twelve months! Should we be fitting each pig separately?], we modeled  $\beta_i(m)$  using a non-cyclic cubic spline [check terminology].

Similar to model 1, we performed LASSO regularization followed by an approximate Bayesian analysis using ADVI (see Supplementary Material).

#### Model 3: Environmentally-driven movement model

 The final model we considered still allowed for overall movement speed to vary both daily and monthly, but now sought to describe the seasonal changes in selection for anthropogenic and natural forage in terms of monthly changes in precipitation and temperature. Specifically, we updated the basis function form of model 3 to include interactions between temperature, precipitation and forage resources.

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\log(\lambda_{ij}(h,m)) = \log(\tau_{ij}) + \beta_0(h) + \beta_1' \operatorname{season}_m
+ \beta_2 \operatorname{directional persistence} + \beta_3 \operatorname{male}
+ \operatorname{canopy cover}_{\operatorname{loc}}(\beta_4 + \beta_4' \operatorname{season}_m)
+ \operatorname{canopy cover}_{\operatorname{dir}}(\beta_5 + \beta_5' \operatorname{season}_m)
+ \operatorname{distance to water}_{\operatorname{loc}}(\beta_6 + \beta_6' \operatorname{season}_m)
+ \operatorname{distance to water}_{\operatorname{dir}}(\beta_7 + \beta_7' \operatorname{season}_m)
+ \operatorname{distance to crops}_{\operatorname{loc}}(\beta_8 + \beta_8' \operatorname{season}_m)
+ \operatorname{distance to crops}_{\operatorname{dir}}(\beta_9 + \beta_9' \operatorname{season}_m)
+ \operatorname{masting tree density}_{\operatorname{loc}}(\beta_{10} + \beta_{10'} \operatorname{season}_m)
+ \operatorname{NDVI}_{\operatorname{loc}}(\beta_{12} + \beta_{12'} \operatorname{season}_m)
+ \operatorname{NDVI}_{\operatorname{loc}}(\beta_{13} + \beta_{13'} \operatorname{season}_m)
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where  $\beta'_i$ season<sub>m</sub> is the monthly effect of temperature and precipitation and is specifically given by  $\beta_{i,1}$ temperature<sub>m</sub> +  $\beta_{i,2}$ precipitation<sub>m</sub> +  $\beta_{i,3}$ temperature<sub>m</sub>precipitation<sub>m</sub>. We used the same approach as model 1 when fitting the LASSO model. When incorporating random effects of individuals, we assumed that the effect of season was the same across individuals and that the random effect of individual was manifested in the  $\beta_i$  coefficients (i.e. not the  $\beta'_i$ coefficients). Otherwise, this random effects model was implemented as in model 1.

### 200 Model comparison and population-level analyses

- 201 For LASSO models...just compare CV deviance with standard error.
- For ADVI models...check out LOO techniques. These might work.

## 203 Results

- 204 Effects of non-foraging covariates on movement
- 205 Effects of foraging covariates on movement
- 206 Main effects models
- 207 Time-varying effects
- 208 Seasonal effects model
- 209 Population-level analysis

## 210 Discussion

- Temporal variation in selection for foraging resources...
- 212 With regards to management,

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	Table	Table 1: Description of covariates used in analysis	used in analysis		
Covariate	Description	Data Source	Spatial Resolution	Temporal Resolution	Location or direction?
Normalized Difference Vegetation Index (NDVI)	A proxy for plant productivity and natural forage availability.	MODIS and NASS	$250 \text{ m} \times 250 \text{ m}$	Monthly	Both
Density of hard-masting trees		Tabak et al.	$1~\mathrm{km} \times 1~\mathrm{km}$	Time-invariant	Both
Distance to crops	The distance to the nearest crop field. A measure of anthropogenic forage.	NASS	$30 \text{ m} \times 30 \text{ m}$	Yearly	Both
Distance to water	The distance the nearest permanent or semi-	NWI	$30 \text{ m} \times 30 \text{ m}$	Time-invariant	Both
Canopy density [CHECK]	Canopy density [CHECK] A proxy for habitat cover	NLCD	$30~\mathrm{m} \times 30~\mathrm{m}$	Time-invariant	Both
Distance to developed-land	A measure of human presence	NLCD	$30 \text{ m} \times 30 \text{ m}$	Time-invariant	Both
Temperature	Mean monthly temperature	NOAA	$50~\mathrm{km} \times 50~\mathrm{km}$	Monthly	Location
Precipitation	Total monthly precipitation	NOAA	$50~\mathrm{km} \times 50~\mathrm{km}$	Monthly	Location