

Step by Step: Using Step Selection Analysis to Simulate Dispersal and Assess Landscape Connectivity

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Abstract

This is the abstract.

1 Introduction

An animals movement trajectory can be seen as the result of an interplay between habitat and movement preferences.

In recent decades, least-cost analysis has become the workhorse to study landscape connectivity at large scales. Originally introduced by XX, the concept and relating methods have been refined to more realistically render animal's movement capabilities and to determine valuable movement corridors that require protection. (talk a bit about different methods of LCPs, see methods in gdistance package). More recently, more attention has been assigned to the collection of movement data is better tailored towards assessing landscape connectivity. In particular, it has been proposed and verified that relocation data collected dispersing individuals leads to more accurate and reliable estimates of landscape connectivity. This superiority of dispersal data in comparison to data that stem from residents is mainly owed to the fact that animals behave vastly different during residency. Despite substantial advances in ecologists ability to track animals in space and time, dispersal remains one of the most difficult behavioral modes to observe in wild animals. Especially for wide ranging and long-lived species, dispersal is difficult to predict and observe, such that data remains scarce.

Many connectivity modelling techniques, especially least-cost analysis, implicitly assume that the studied animal has partial or even complete knowledge of the landscape and associated movement costs. While this assumption may be reasonable for migrants that move between a limited number of habitats, yet it is unlikely to hold for dispersers. In contrast to migrants, dispersers move into unknown territory and are therefore confronted with novel landscapes. Consequently, dispersers are more likely to adjust their movement behavior *ad hoc* instead of preplanning an entire trajectory. As a result, methods that assume complete knowledge of the landscape and subsequent optimal movement behavior likely misrepresent true dispersal behavior. As such, methods that allow a more random approach to a dispersers movement behavior may more realistically render the movement corridors of dispersing individuals.

In addition, individual based simulations will allow to explicitly model dispersal between subpopulations in models of population dynamics.

Reliable identification of dispersal corridors will become increasingly important with the uprise of ever-growing and often transboundary conservation areas. One such instance is the KAZA-TFCA, a massive conservation area spanning five countries and over 520'000 km². The KAZA holds the potential of re-establishing dispersal routes for many of its protected species, including the african wild dog *Lycaon pictus*. This species has experienced severe

population declines caused by human induced mortality and deadly diseases. In result, the species currently marks the KAZA's most endangered large carnivore and has been assigned a very high conservation priority. Importantly, due to their inherent mobility and intrinsic need for vast undisturbed landscapes, AWDs have been proposed as surrogate species for landscape connectivity (see recent paper on multispecies connectivity). Nevertheless, the species has received little attention in the connectivity literature, mainly due to the difficulty in observing wild dog dispersal. In a previous paper, we addressed this issue and developed a habitat selection model based on which we predicted landscape connectivity using least-cost corridor analysis. We now expand on this knowledge and develop a more detailed movement model of dispersing wild dogs. We then use this model to simulate thousands of dispersers moving throughout the KAZA. Based on said simulations, we compute heatmaps and identify potential dispersal hotspots and compare them to the dispersal routes identified in (Hofmann). We also apply the data to feed a network-analysis, based on which we determine network-metrics pertinent to landscape connectivity.

2 Methods

2.1 Study Area

The study area (centered at $-17^{\circ}13'9''S$, $23^{\circ}56'4''E$; Figure 1a) stretched over 1.3 Mio km² and encompassed the entire KAZA-TFCA (Figure 1b). The KAZA-TFCA is the world's largest transboundary conservation area and comprises parts of Angola, Botswana, Namibia, Zimbabwe, and Zambia, covering a total area of over 520'000 km². Its landscape varies regionally and ranges from savanna, to grassland, and from dry to moist woodland habitats. A dominant hydrogeographical feature in its center is the Okavango Delta, the earth's largest inland delta and home to a vast diversity of mammal species. The delta and its surroundings are considered a stronghold for African wild dogs which may act as a source for the recolonization of surrounding habitats. The wet season in the region lasts from November to March, albeit the main floodwaters reach the Delta between July and August after descending through the Angolian highlands (McNutt, 1996; Wolski et al., 2017).

2.2 GPS Relocation Data

Between 2011 and 2020, we collected GPS relocation data on free-ranging wild dogs inhabiting the Okavango Delta in northern Botswana. We identified potential dispersers based on criteria reported in Behr et al. (2020), immobilized them according to protocols described

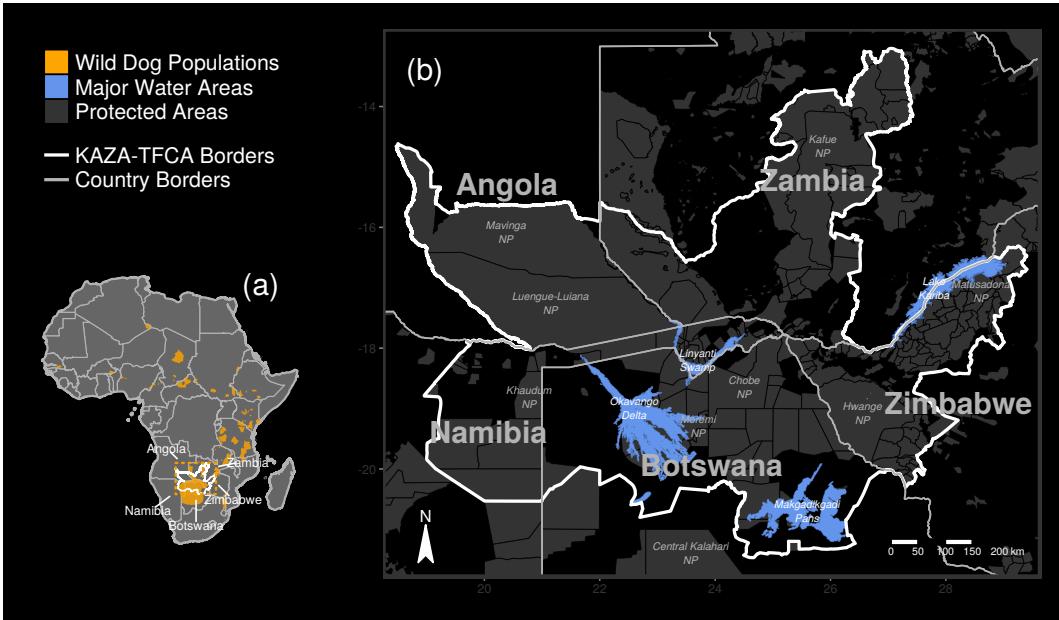


Figure 1: Study area

in Osofsky et al. (1996), and fitted them with GPS/Satellite radio collars (*Vertex Lite; Vectronic Aerospace GmbH, Berlin*). Handling and collaring of all individuals was carried out and supervised by a Botswana-registered wildlife veterinarian. Of all collared individuals, a total of 16 individuals dispersed, each in a same-sex coalition (7 female and 9 male coalitions). When we observed dispersal we remotely increased the GPS fixrate from 24 to 4 hours. Collected relocations were then regularly transmitted via Iridium satellite system to a base station. This allowed remote tracking of dispersers even if they left the main study area or crossed international borders. To effectively distinguish between resident and dispersing individuals, we applied the net-squared displacement metric. This metric measures the Euclidean distance of a collared individual to a reference point (Börger and Fryxell, 2012). In our case, these reference points resembled the center of the dispersing coalition's natal home range. Hence, dispersal was deemed to have started when an individual left its natal home range and ended when the individual became stationary again. Previous research found no differences between males and females during dispersal (Woodroffe et al., 2019; Cozzi et al., 2020), so we did not distinguish sexes in our analyses. GPS relocations were then converted to steps, where a step represented the straight-line distance travelled between two consecutive GPS relocations (Turchin, 1998).

2.3 Covariates

To represent environmental covariates spatially, we prepared a set of raster layers depicting water-cover (dynamically updated), square rooted distance to water (dynamically updated), tree-cover, and shrub/grassland-cover. We also created a proxy for human influence, rendering anthropogenic pressures stemming from human-density, agricultural sites, and roads. We prepared all layers at a resolution of 250m by 250m for the extent of the entire KAZA-TFCA. A more detailed description of the derivation and preparation of each environmental covariate is given in ?. Because wild dogs follow a diurnal activity pattern, we coded a binary variable indicating whether an observed step started during phases of main wild dog activity (07:00 a.m. and 07:00 p.m.) or low wild dog activity (anything else).

2.4 Movement Model

We used integrated step selection functions (iSSF) to parametrize a mechanistic movement model of dispersing wild dogs. In the iSSF framework, observed steps receive a selection score $w(x)$ that depends on the covariates X experienced along a step, as well as the animals preferences towards these covariates β . In contrast to regular step selection analysis, *integrated* step selection analysis allows simultaneous inference on movement and habitat preferences of the studied animal. Moreover, potential interactions between habitat and movement preferences can be modelled. Thus, the method produces more accurate selection estimates and allows to use the resulting models as proper mechanistic movement models from which movement can be generated. To conduct iSSF analysis, we paired each observed step with 24 random steps. Random steps basically resembled potential alternatives that the animal could have realized but decided not to. We generated random steps by sampling random turning angles from a uniform distribution $(-\pi, +\pi)$ and step lengths from a gamma distribution that was fitted to observed steps (scale = 6308, shape = 0.37). While the number of random steps is inversely proportional to the sampling error (Avgar et al., 2016), we found only minor changes in inference when sampling further random steps. Along each step we extracted spatial covariates using the *velox* package. We also calculated step metrics, namely the natural logarithm of the step length $\log(sl_)$ and the cosine of the turning angle $\cos(ta_)$. We scaled all continuous covariates to a mean of zero and a standard deviation of one. We then used the r-package *glmmTMB* to fit mixed effects conditional logistic regression models as proposed by (Muff et al., 2020). Our movement model was built around a previously published habitat selection model of dispersing wild dogs. Because the earlier model was used to predict landscape permeability, interactions between

environmental and spatial covariates were impossible to model. Our current movement model, in contrast, allowed the inclusion of additional interactions that describe differences in movement behavior induced through different environments. Hence, we started with the habitat model and iteratively increased model complexity by including additional interactions between movement metrics and environmental covariates. That is, we proposed all two-way interactions between spatial covariates and movement metrics. For instance, for the covariate *water* we proposed the interactions Water:cos(ta₋) and Water:log(sl₋). Despite interactions between environmental factors and movement metrics, we also proposed the interaction log(sl₋) : *MainActivity* to account for the diurnal activity pattern of wild dogs. A preliminary revealed little differences in turning angles during main and low activity, hence we did not include the interaction cos(ta₋) : *MainActivity*. We then ran stepwise modal forward selection based on Akaike's Information Criterion (AIC, Burnham and Anderson, 2002) values and identified the most parsimonious movement model.

2.5 Dispersal Simulation

We used the most parsimonious movement model to simulate dispersing wild dogs departing from predefined source points. The simulation basically resembled an inverted iSSF function and was set up as follows. First, we determined a source point at which a disperser was initiated. To allow calculation of turning angles, we assumed a random initial orientation of the animal. Second, we proposed 25 random steps originating at the predefined source point. We generated random steps by sampling turning angles from a uniform distribution ($-\pi, +\pi$) and step lengths from a gamma distribution fitted to observed step lengths. To prevent unrealistically large steps, we capped the gamma distribution 35km, the farthest distance travelled in four hours according to our data. Third, along each random step we extracted environmental covariates calculated step metrics. Fourth, we applied the parametrized dispersal model to predict selection scores $w(x)$ and to determine the probability of a step being realized. Fifth, we sampled one of the proposed based on their probabilities and calculated the animal's new position. We then repeated steps two to five until 2000 steps were realized. In case a proposed random step left our study extent, we removed the step from the set of random steps, thereby forcing the disperser to stay within the boundaries of the study area.

2.6 Source Points

Simulations from step selection functions are known to be sensitive to the location of source points, which is why we followed a twofold approach to sample release points across the

KAZA-TFCA. In a first approach, we used the same 68 source points between which we already computed least-cost paths and least-cost corridors (see xx). These source points were regularly spaced 100 km apart, located within protected areas ($> 700 \text{ km}^2$). Because of the regular spacing, not all protected areas received source points. Hence, we placed additional source points at the center of such protected areas. At each of the generated source point we released 1000 dispersers, implying a total of 68'000 simulated dispersers. In a second approach, we used the same 68 source points to define catchment areas inside protected areas. That is, for each of the 68 source points we defined its voronoi polygon within the point's protected area. Within these catchment areas we then randomly sampled 1000 new source points from which we released another 68'000 dispersers. Overall, we simulated 134'000 dispersers for a total of 268 Mio. steps.

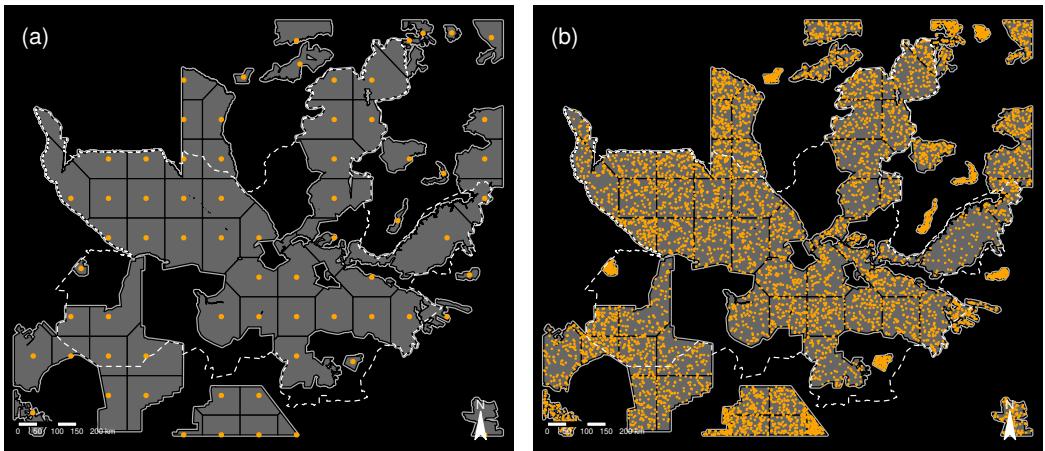


Figure 2: Illustration of source points from which dispersal was simulated. Although in reality we simulated 1'000 per source point, we illustrate an example assuming 10 dispersers from each source point. (a) Static source points similar to the source points reported in ... (b) Source points that were randomized within the catchment areas (dark gray, delineated by solid black lines).

2.7 Heatmaps

Using the simulated trajectories we created heatmaps to examine through which areas most simulated individuals dispersed. We rasterized all trajectories and calculated how often each raster-cell was traversed by simulated dispersers. If the same trajectory crossed a pixel twice, it was only counted once. We achieved high performance rasterization of spatial lines using the recently developed R-package *terra* (Hijmans, 2020). To examine if and how “heat” changes in response to changes in the location of source points and the number of simulated steps, we followed a 2 x 6 design and created heatmaps for both point sampling regimes, as well as for 68, 125, 250, 500, 1000, and 2000 dispersal steps. We quantified the similarity

of the resulting 12 heatmaps to the permeability and least-cost corridor maps presented in (Hofmann ...) we used Bhattacharyya's affinity. Bhattacharyya's affinity ranges from zero (complete separation) to one (perfect match) and has earlier been proposed to compare the overlap of utilisation distributions (Fieberg).

2.8 Network Analysis I: Areas Reached

Based on the simulated trajectories we identified to which other areas each source area is connected. For instance, if a trajectory originated at source point one and intersected with source areas two and three, we assumed that source areas two and three were within reach of source area one. To get a sense of the strength of connections, we also calculated how often each of these connections was realized. This procedure resulted in 6 x 2 edge lists which we further used to generate weighted networks and to calculate network metrics.

2.9 Network Analysis II: Betweenness

We coerced all simulated trajectories into a network consisting of vertices (relocations) and edges (connections between relocations). To do so, we created raster layers at multiple resolutions and identified each trajectory's transition matrix on these rasters see figure xx (Bastille-Rousseau et al., 2018). We then merged the transition matrices of all trajectories and calculated cumulative transitions between all raster-cells. This resulted in an edge-list, containing all observed from-to connections as well as their frequency. Using this edge-list, we generated a weighted graph using the r-package *igraph*. Based on this graph we calculated betweenness scores as well as the degree of each raster cell. Betweenness indicates how often a specific raster-cell lies on a shortest path between two other raster-cells and is a useful metric to detect movement corridors. Degree, on the other hand, indicates how many connections a raster cell has and therefore serves to illustrate the xxx of specific nodes. When calculating betweenness, we used the transition frequency as weighting factor. That is, a higher transition frequency contributed to a higher betweenness score.

2.10 Network Analysis III: Transitions

3 Results

Compared to the base model, the most parsimonious movement model included several additional interactions (Figure 3 and Table S1). The model indicates that dispersers move directional, particularly when distant to water, yet less so in human dominated landscapes.

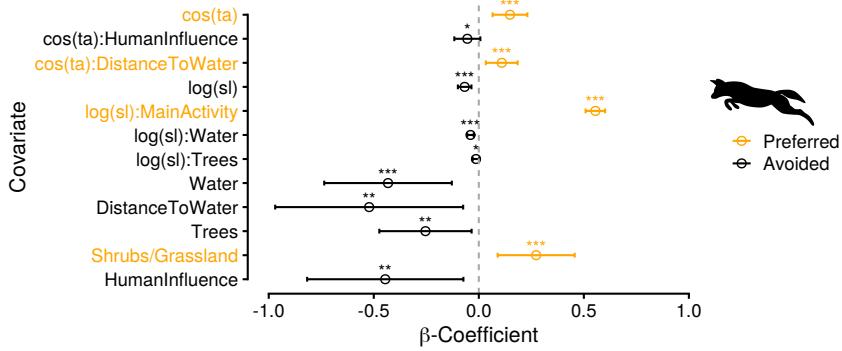


Figure 3: Most parsimonious movement model. Whiskers delineate the 95% Confidence-Intervals. Significance codes: * $p < 0.10$, ** $p < 0.05$, *** $p < 0.01$.

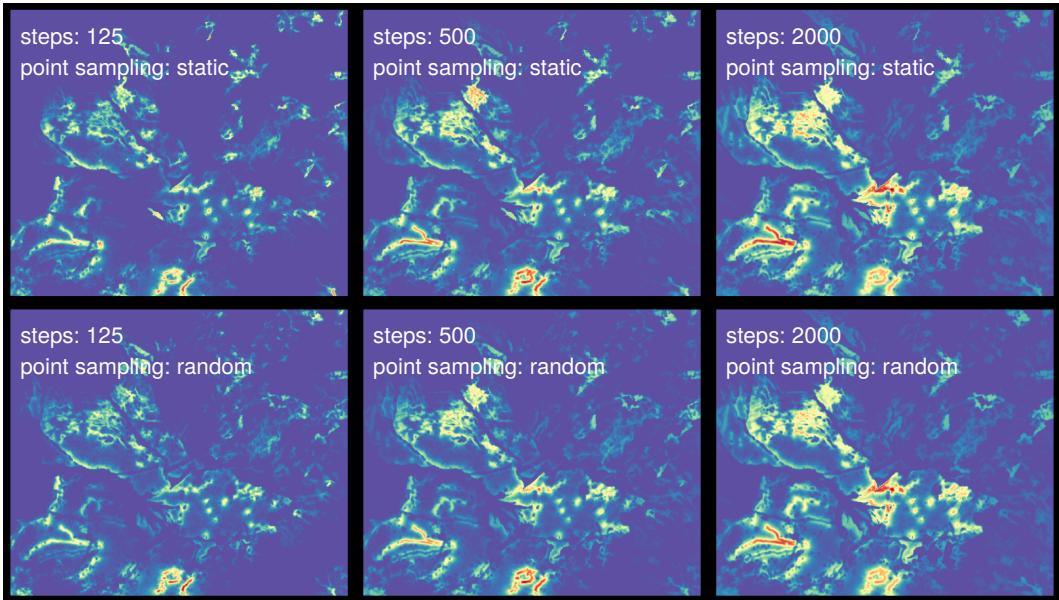


Figure 4

Furthermore, dispersers prefer large steps, especially during main activity. In contrast, step lengths tend to be shorter when water- or tree-cover is high. In general, dispersers avoid water, prefer proximity to water, avoid dense tree-cover, prefer shrubs/grassland, and, finally, avoid human dominated landscapes.

3.1 Heatmaps

Six of the twelve rasterized dispersal trajectories are presented as heatmaps in Figure 4. As can be seen, differences that stem from the method of point sampling disappear as more steps are simulated. This is to be expected as the influence of the origin becomes smaller and smaller as the animal moves through the landscape. However, there are striking differences when simulations are only run for few iterations.

Bhattacharyya’s affinity index supports the notion that heatmaps become more similar to the earlier developed permeability and corridor maps as the number of simulated steps increases. Furthermore, it appears that randomly sampled source points contribute to a higher similarity too, albeit differences due to the sampling regime vanish as the number of simulated steps increases. In fact, both maps are almost identical to each other after 2000 steps. Furthermore, the connectivity networks become increasingly similar to the previously published permeability and corridor maps. Still, some differences remain even after 2000 simulated steps, highlighting that some severe impediments in the landscape exist.

Table 1: Summary statistics of all GPS relocations that have been recorded on dispersing coalitions.

sampling	Metric	Map	68	125	250	500	1000	2000
Static	Affinity	Corr	0.64	0.70	0.76	0.80	0.82	0.84
Static	Affinity	Perm	0.67	0.74	0.80	0.85	0.89	0.91
Static	Correlation	Corr	0.32	0.38	0.45	0.51	0.56	0.60
Static	Correlation	Perm	0.47	0.57	0.66	0.72	0.78	0.82
Random	Affinity	Corr	0.70	0.75	0.79	0.82	0.84	0.85
Random	Affinity	Perm	0.74	0.79	0.83	0.87	0.90	0.92
Random	Correlation	Corr	0.38	0.43	0.48	0.53	0.57	0.61
Random	Correlation	Perm	0.58	0.64	0.69	0.75	0.79	0.83
n.a.	Affinity	Heat	0.76	0.86	0.94	0.97	0.99	0.99
n.a.	Correlation	Heat	0.90	0.94	0.97	0.98	0.99	1.00

4 Discussion

Our connectivity network further suggests that dispersers from the Okavango Delta more likely disperse towards east than west. Indeed, only x out of our y observed dispersers ever reached the western part of the delta. Only when the flood retracts a small pathway between the city of Maun and the floodwaters of the delta emerges and enables dispersers to move towards the delta’s western part.

Our work suggests that the selection of source points significantly impacts resulting connectivity networks. Especially when dispersal durations are short, wrongly placed source points lead to vastly different results. Signer et al. used estimated utilisation distributions by means of simulated movements. They used a rather long burn in period prior to alleviate the problem of selecting meaningful source points. However, this approach only works when individuals move around a point of attraction. This is typically not the case when simulating dispersers, introducing an important trade-off. The researcher can decide to increase the

number of simulated steps, hence reducing the influence of starting locations, yet this also inevitably increases estimated connectivity.

5 Authors' Contributions

D.D.H., D.M.B., A.O. and G.C. conceived the study and designed methodology; D.M.B., G.C., and J.W.M. collected the data; D.D.H. and D.M.B. analysed the data; G.C. and A.O. assisted with modelling; D.D.H., D.M.B., and G.C. wrote the first draft of the manuscript and all authors contributed to the drafts at several stages and gave final approval for publication.

6 Data Availability

GPS movement data of dispersing coalitions will be made available on dryad at the time of publication.

7 Acknowledgements

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