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The search behavior of terrestrial mammals

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Summary

Animals moving through landscapes need to strike a balance between finding sufficient resources to grow and reproduce while minimizing encounters with predators ^{1,2}. Because encounter rates are determined by the average distance over which directed motion persists ^{1,3–5}, this trade-off should be apparent in individuals' movement. Using GPS data from 1,396 individuals across 62 species of terrestrial mammals, we show how predators maintained directed motion ~7 times longer than for similarly-sized prey, revealing how prey species must trade off search efficiency against predator encounter rates. Individual search strategies were also modulated by resource abundance, with prey species forced to risk higher predator encounter rates and resource availability in shaping broad patterns mammalian movement strategies.

Main

As motile organisms move through landscapes in search of food, mates, and cover, they need to strike a balance between finding sufficient resources to grow and reproduce, while also minimizing the rate at which they encounter predators^{1,2}. Because of the fitness consequences of foraging success^{6,7} and predator-prey dynamics^{8,9}, there should be strong selection pressure on movement strategies that maximize resource encounter rates while minimizing encounters with predators. Although it is well recognised that animals will adjust the size of their home-range areas based on resource availability^{10–12}, there remains the question of how to optimally find resources within these ranges. In this context, random search models have proven influential in understanding how individual movement strategies translate to encounter rates^{1,3,4,13}. The consensus from these models is that directed (i.e., ballistic) movement leads to higher encounter rates than more tortuous (i.e., diffusive) movement^{1,3,4}. This occurs because individuals that exhibit tortuous movement will tend to repeatedly search over the same areas, whereas directed motion allows individuals to search over a larger area within the same amount of time. The average distance over which ballistic motion persists, l_{ν} (in m), is thus a key determinant of encounter rates¹ and a potent trait that individuals can optimize.

An individual's ballistic length scale, l_v , is a function of the spatial variance of their movement $(\sigma_p, \text{ in } m^2)$ and their positional and velocity autocorrelation timescales $(\tau_p \text{ and } \tau_v \text{ respectively, in sec})$, given by

Because directed movement is the more efficient search strategy^{1,4,5}, bottom-up pressure exerted by the need to encounter resources should select for more ballistic movement. Importantly, however, increasing l_v will also increase the rate at which individuals encounter predators¹. Topdown predation pressure should thus select for shorter ballistic length scales. Prey species searching for immobile vegetation must therefore optimize their movement against the opposing forces of their energetic requirements selecting for longer l_v , and predation pressure selecting for shorter l_v^{-1} . Predators also benefit from maintaining longer ballistic length scales¹, but without the intense top-down predation pressure experienced by prey species. The combination of bottom-up and top-down regulation is thus expected to select for longer ballistic length scales in predators, versus more diffusive movement in prey species, all else being equal¹ (Fig. 1).

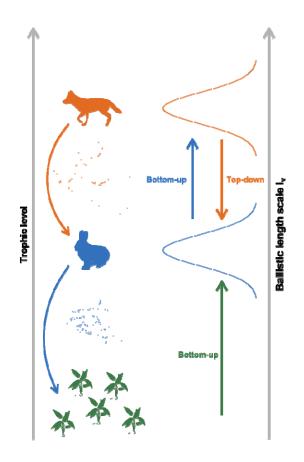


Figure 1 Selection pressures on predator and prey ballistic length scales. Schematic representation of bottom-up energetic requirements selecting for longer ballistic length scales in mammalian movement paths, and top-down

predation pressure selecting for shorter ballistic length scales. The simulated prey movement path (blue) has a ballistic length scale of 10m, whereas the predator's movement (orange) has a ballistic length scale of 100m.

Notably, 'all else' is rarely equal in ecological systems, and the relative importance of bottom-up versus top-down regulation is expected to be context specific. In resource-poor ecosystems, individuals need to spend a substantial amount of time searching for food and moving between patches^{6,14}. The bottom-up driven need to find sufficient resources to survive should outweigh top-down pressure when resources are scarce. In productive environments, in contrast, bottom-up pressure on resource acquisition rates should be relaxed¹², providing individuals with the capacity to respond more directly to top-down pressure and maintain relatively more diffusive movement. These considerations lead to the expectation of a negative relationship between l_v and environmental productivity.

Although the importance of l_v in governing encounter rates is suggested by theoretical models^{1,5}, there has, to date, been no empirical demonstration of systematic differences in l_v between comparably sized predators and prey for any taxonomic group. Here, we leverage the rapid advances in the capacity to collect¹⁵ and work with^{16,17} animal movement data that have enabled l_v to be estimated for a broad range of species. We annotated Global Positioning System (GPS) location data on 1,396 individuals from 62 species of terrestrial mammals (Fig. 2a) with mean adult body size and trophic group (prey n = 41, and predator n = 21). We restricted our analyses to range-resident animals and used continuous-time stochastic models¹⁶ to estimate l_v for each individual (Fig 2b). We also annotated each data point with the mean Normalized Difference Vegetation Index (NDVI), a satellite-derived measure of resource abundance¹⁸, to which each individual was exposed. Finally, as measures of habitat permeability we quantified the mean percent forest cover, terrain roughness, and human footprint index at each sampled location.

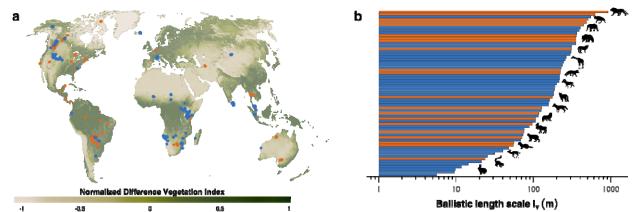


Figure 2 The distribution of mammalian movement data. In a, the GPS locations of 1,396 prey (blue) and predatory (orange) mammals across 62 species are plotted on the global map of Normalized Difference Vegetation Index ranging from low (-1) to high (1) productivity; and b shows the median ballistic length scales, l_{ν} , for each species.

Our analyses revealed allometric scaling in ballistic length scales, with larger mammals tending to have more directed movement, all else being equal ($P < 10^{-7}$; Fig. 3a). The parameters of the body-mass scaling relationships are shown in Table S2. The residuals of the body mass relationship followed theoretical predictions, with predator l_v being 7.1 times longer than that of

comparably sized prey species ($P < 10^{-6}$; Fig. 3b). Least-squares regression also revealed a strong negative correlation between NDVI and the residuals of the allometric relationships in l_{v} for prev species ($P < 10^{-6}$; Fig. 3c). In other words, when resource abundance was ignored, predictions from the simple body-size relationships tended to under-estimate observed l_v in low-productivity ecosystems, and over-estimate l_{ν} in highly productive ecosystems. That bottom-up pressures appear to outweigh top-down pressure for prey species living in resource-poor ecosystems is made all the more poignant when contrasted against known patterns in trophic structure scaling. Resource-poor environments tend to be more trophically top heavy than productive ecosystems¹⁹, effectively increasing the number of predators per individual prey on the landscape. Nonetheless, prey species living in resource-poor ecosystems exhibited significantly longer ballistic length scales than those in resource-rich environments, on average. Because predation rates depend on numerous factors beyond encounter rates, (e.g., capture efficiency, predator hunger levels at the time of the encounter, the presence of suitable cover, prey defenses, etc.²⁰), we would expect prey species to respond more directly to bottom-up factors than predators. In line with this expectation, we found no evidence that predators adjusted their ballistic length scales as a function of NDVI (P = 0.26; Fig. 3C).

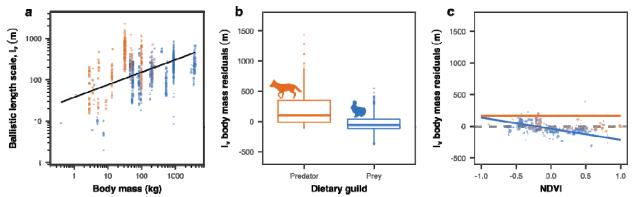


Figure 3 Trends in mammalian ballistic length scales. Mammalian l_v scales with body size, but predators exhibit more ballistic motion than prey, and search behavior is modulated by environmental productivity. The scatterplot in **a** show the allometric scaling of the ballistic length scale, l_v for prey (blue), as well as predatory (orange) mammals. The boxplots in **b** show the residuals of the body-mass scaling of l_v for predators and prey. When body size is accounted for, predatory species have longer ballistic length scales on average ($P < 10^{-5}$). The scatterplot in **c** show the body mass residuals as a function of the normalized difference vegetation index (NDVI). The solid lines in **c** show how prey adjusted their ballistic length scales based on NDVI ($P < 10^{-6}$), whereas predators did not (P = 0.26).

Although predation pressure and resource abundance will influence the relative importance of search times and movement rates^{6,12,14,21}, they are not the only factors that will influence animal movement. For instance, memory²², socially transmitted information²³, and the 'patchiness' of resources^{6,14} have all been shown to influence foraging behavior. Similarly, landscape permeability can impact the capacity for individuals to maintain directed motion, with implications for foraging efficiency²⁴. Indeed, we found complex, non-linear and heteroskedastic relationships between habitat structure and l_{ν} (Fig. S2). Yet, even with all of these other factors at play within the individual datasets we analyzed, our capacity to identify a clear signal for the interplay between encounter rates and resource availability in shaping mammalian movement strategies highlights just how important risk-reward trade-offs are to animals.

One question that arises from these results is 'Why don't predators have longer ballistic length scales?'. Without top-down regulation, there should be little preventing longer l_v in predators, yet we found that predator l_v was only ~7 times longer than that of comparably sized prey on average. There are two main reasons why predator ballistic length scales are not longer than observed. The first is that individuals need to perceive resources within the area they travel through, and while more directed motion might be the more efficient movement strategy, it does not necessarily imply that animals will perceive their targets. Individuals are thus constrained by their perceptual range^{1,13} and the time and effort required to find resources in the landscapes they move through²⁰. Increasing l_v thus only benefits predators up to a certain point, and beyond some optimal value encounter rates may actually decrease^{4,5,14}. The second reason is that while many adult mammalian predators have few predators themselves, their movement can be constrained by intra-guild and conspecific encounters^{25,26}, or parasite avoidance²⁷. Longer l_v in predators might increase prey encounter rates, but at the expense of increasing the rates at which they encounter these negative factors.

Conclusions

The behavioral consequences of foraging within a landscape of fear have been theorized about extensively^{28,29}, yet demonstrated for only a handful of systems, such as elephant seals (*Mirounga angustirostris*), which optimize their diving behavior against predation risk and their energetic state². Here, we demonstrate how the optimization of search strategies and encounter rates underpin broad patterns in the movement of terrestrial mammals spanning orders of magnitude in body size and distributed in multiple different ecosystems around the world. Prey species searching for immobile vegetation under the threat of predation must trade off the efficiency of their search strategies against the risk of encountering predators. However, ballistic length scales were negatively correlated with resource abundance, revealing how prey species are forced to risk higher predator encounter rates when resources are scarce. In contrast, the ballistic length scales of predatory species searching for mobile prey experience less top-down selection pressure, allowing them to maintain more efficient search behavior. These results highlight the interplay between encounter rates and resource availability in shaping mammalian movement strategies.

Methods Empirical analyses

Tracking data analysis

To investigate pattern in the ballistic length scales of terrestrial mammals, we compiled openly available GPS tracking data from the online animal tracking database Movebank³⁰, or from co-authors directly. Individual datasets were selected based on the criterion of range resident behavior, as evidenced by plots of the semi-variance in positions as a function of the time lag separating observations (i.e., variograms) with a clear asymptote at large time lags^{16,31}. All data from migratory, or dispersing periods were excluded as their measured movement strategies would not be representative of the normal foraging dynamics we aimed to describe. The visual verification of range-residency via variogram analysis³¹ was conducted using the R package ctmm (version 1.1.0)¹⁶.

As noted in the main text, the average length scale over which ballistic motion is maintained $(l_{\nu},$ in m) is a function of the spatial variance of an animal's movement ($\sigma_{position}$, in m²) and its positional and velocity autocorrelation timescales ($\tau_{position}$ and $\tau_{velocity}$ respectively, in sec). While σ_p can be well estimated from coarse data³², autocorrelation structures are only revealed when the time scale of measurement is less than or equal to the autocorrelation timescales ³³. In particular, τ_{v} tends to be on the order of minutes to hours for medium to large terrestrial mammals^{32,34}. Estimating l_{ν} for these data thus first required estimating the autocorrelation structure in each of the individual tracking datasets and identifying those for which there was sufficient information to estimate σ_p , τ_p , and τ_v . To do this, we fit a series of range-resident, continuous-time movement models to the data. The fitted models included the Independent and Identically Distributed IID process, which features uncorrelated positions and velocities; the Ornstein-Uhlenbeck (OU) process, which features correlated positions but uncorrelated velocities³⁵; and an OU-Foraging (OUF) process, featuring both correlated positions, and correlated velocities^{31,36}. We then employed AICc based model selection to identify the best model for the data^{36,37}, from which the σ_v , τ_v , and τ_v parameter estimates were extracted. To fit and select the movement models, we used the R package ctmm, applying the workflow described by¹⁶. Because only the OUF process included information on all of the parameters required to estimate l_{ν} , we further restricted our analyses to only those individuals for which the OUF model was selected. This latter threshold was based on the sampling resolution of the GPS data, as only data of a sufficiently fine sampling resolution allow for the estimation of l_{v} . In other words, we used AICc based model selection to identify the individual datasets with data of sufficient resolution to allow for an estimate of l_v , rather than relying on an arbitrary sampling threshold. The final dataset included data from 62 species, comprising a total of 8,613,485 locations for 1,396 individuals. Finally, l_{ν} was calculated from the parameter estimates for each of these individuals as

$$l_{v} = \sqrt{\frac{\tau_{v}}{\tau_{p}}}\sigma_{p}$$

Covariate data

For each of the species in our dataset we compiled covariate data on that species' mean adult mass, in kilograms, and diet taken from the EltonTraits database³⁸. This dataset includes species with body masses covering five orders of magnitude (0.4 - 4000 kg). Dietary class was then used to categorize species as being either a predator or prey species. Predators were species that specialized primarily on mobile animal prey, whereas prey were herbivores and frugivores that specialized primarily on sessile vegetation. A summary of the dataset is shown in Table S1. To assess ecological factors that may have influenced l_{ν} , we also annotated each estimate with four satellite-derived habitat metrics: i) mean Normalized Difference Vegetation Index (NDVI), as a measure of local resource abundance¹⁸; ii) percent forest cover³⁹; iii) terrain roughness⁴⁰; and iv) machine-learning human footprint index (ml-HFI)⁴¹, as measures of habitat permeability. Details on the annotation process for the habitat covariate data were as follows:

- 1. **NDVI**. To annotate the location data with NDVI we first compiled NDVI data from the publicly accessible NASA MODIS archive. We used global NDVI rasters sampled at a 250m resolution at 16-day intervals between 2000-2022 (i.e., a total of 518 raster layers). For each individual GPS location in the dataset, we identified the point in time and space that was closest to the sampled location and extracted the NDVI value. Once each location was annotated with the appropriate NDVI value, we calculated the geometric mean value of NDVI to which each individual was exposed.
- 2. **Percent forest cover**. Consensus land cover data at a 1-km resolution were obtained from the EarthEnv data repository based on the methods presented in³⁹. These data were available as 12 raster layers containing percentages on the prevalence of one of 12 land-cover classes. From these individual raster layers we quantified the percent forest cover as the summation of five layers containing information on the prevalence of evergreen/deciduous needleleaf trees, evergreen broadleaf trees, deciduous broadleaf trees, mixed/other trees, and shrubs. Following a similar process as the NDVI annotation, we identified the point in space that was closest to each sampled GPS location and extracted the percent forest cover value and calculated the geometric mean value to which each individual was exposed.
- 3. **Terrain roughness**. Terrain roughness data at a 1-km resolution were obtained from the EarthEnv data repository based on the methods presented in⁴⁰. Terrain roughness represents a measure of topographic heterogeneity, and was quantified as the largest inter-cell difference in elevation between a focal cell and its 8 surrounding cells. Here again we identified the point in space that was closest to each sampled GPS location and extracted the roughness value and calculated the geometric mean value to which each individual was exposed.
- 4. **ml-HFI**. The machine-learning-based human footprint index (ml-HFI)⁴¹ is an index of human pressure on the landscape that is derived from remotely sensed surface imagery and ranges on a scale between 0 (no human impact), and 1 (high human impact). Briefly, convolutional neural networks, are used to identify patterns of human activity from the Hansen Global Forest Change imagery version 1.7 (GFCv1.7). The raster is available at an approximately 1-km resolution. We identified the point in space that was closest to each sampled GPS location and extracted the ml-HFI value and calculated the geometric mean value to which each individual was exposed.

Table S1 Data summary table. Summary statistics on the GPS tracking data used in the analyses presented in the main text. Values include the species binomial, including subspecies where appropriate, the number of individuals per species (*n*), body mass, trophic group, the median spatial variance of the animals' movement (σ_p), the median positional (τ_p) and velocity (τ_v) autocorrelation timescales, and the median ballistic length scale (l_v).

Binomial	n	Mass (kg)	Trophic Group	σ_p (km ²)	$ au_p$ (hrs)	τ, (min)	/, (m)
Acinonyx jubatus	1	46.7	Predator	14.882	190.21	98.05	357.58
Aepyceros melampus	20	52.5	Prey	0.229	33.43	8.05	31.12

Alces alces	56	541.46	Prey	8.56	706.57	31.62	76.29
Antidorcas marsupialis	10	31.5	Prey	66.451	345.81	42.96	338.58
Antilocapra americana	46	46.08	Prey	20.41	264.17	31.75	215.78
Beatragus hunteri	4	79.13	Prey	4.888	128.43	20.38	101.8
Blastocerus dichotomus	3	86.67	Prey	0.929	145	13.45	37.97
Brachylagus idahoensis	3	0.42	Prey	0.002	13.18	35.18	8.93
Canis latrans	49	13.41	Predator	2.981	17.6	6.18	115.15
Canis lupus	128	32.18	Predator	58.427	88.49	27.43	542.95
Capra ibex	39	85.17	Prey	4.155	353.57	25.02	74.6
Cerdocyon thous	19	5.24	Predator	0.156	7.43	1.63	21.62
Cervus canadensis	14	200	Prey	67.242	1636.76	27.93	167.19
Chlorocebus pygerythrus	10	4.99	Prey	0.024	11.05	3.35	9.94
Connochaetes taurinus	35	180	Prey	41.062	490.7	12.12	118.85
Cuon alpinus	11	14.17	Predator	5.589	51.68	33.05	244.06
Elephas maximus indicus	24	35 00	Prey	50.144	1375.75	48.63	124.34
Elephas maximus maximus	51	3750	Prey	12.971	202.4	64.2	232.89
Elephas maximus sumatranus	9	3000	Prey	26.06	1013.66	51.66	177.53
Equus hemionus hemionus	18	240	Prey	1129.502	4715.34	14.21	234.94
Equus quagga	9	400	Prey	463.718	930.14	34.95	494.23
Eulemur rufifrons	3	2.25	Prey	0.029	7.03	10.83	24.01
Euphractus sexcinctus	4	4.78	Prey	0.024	4.85	1.19	11.23
Felis catus	53	2.88	Predator	2.05	74.06	8.72	80.7

Felis silvestris	3	5.1	Predator	2.634	30.4	23.67	108.93
Giraffa camelopardalis antiquorum	17	899.99	Prey	48.821	676.22	28.25	210.75
Giraffa camelopardalis camelopardalis	51	899.99	Prey	18.352	195.06	33.92	232.42
Giraffa camelopardalis peralta	17	899.99	Prey	66.868	253.61	33.64	346.69
Giraffa giraffa angolensis	72	899.99	Prey	39.839	303.38	36.98	255.91
Giraffa giraffa giraffa	31	899.99	Prey	6.082	118.25	34.9	183.03
Giraffa reticulata	35	899.99	Prey	20.083	191.98	31.17	211.52
Giraffa tippelskirchi	18	899.99	Prey	9.298	157.74	27.83	172.82
Hyaena brunnea	3	42.98	Predator	19.798	44.85	25.77	425.29
Lagostrophus fasciatus	1	1.7	Prey	0.005	6.1	34.35	21.17
Leopardus pardalis	3	11.9	Predator	0.246	7.66	9.85	72.59
Loxodonta africana	22	3940.03	Prey	242.666	1025.23	34.06	309.1
Lycalopex culpaeus	8	8.62	Predator	0.5 05	5.26	5.65	95.17
Lynx rufus	13	8.9	Predator	0.506	22.71	2.19	28.56
Madoqua guentheri	15	7.5	Prey	0.003	2.96	1.69	4.86
Neogale vison	5	1.15	Predator	0.548	127.94	53.65	44.45
Odocoileus hemionus	5	54.21	Prey	26.912	1734.41	23.69	91.76
Odocoileus virginianus	33	55.51	Prey	0.187	12.04	16.09	69.39
Oreamnos americanus	4	72.5	Prey	4.618	48.56	53.35	318.42
Oryx dammah	38	200	Prey	38.379	265.81	20.29	203.16
Ovis canadensis	3	74.64	Prey	10.951	822.88	47.67	114.35
Ovis dalli	66	55.65	Prey	23.112	524.31	61.06	204.6

Panthera leo	3	161.5	Predator	31.722	57.14	65.13	748.76
Panthera onca	105	100	Predator	11.749	126.68	45.69	254.45
Panthera pardus pardus	7	48.75	Predator	29.444	359.11	13.47	101.31
Panthera pardus saxicolor	6	52.04	Predator	15.643	185.1	30.3	221.34
Pekania pennanti	17	4	Predator	0.707	20.18	2.93	35.31
Propithecus verreauxi	7	3.48	Prey	0.006	< 0.01	< 0.01	6.11
Puma concolor	31	51.6	Predator	46.227	428.86	52.92	264.8
Rangifer tarandus	19	86.03	Prey	55.734	1521.59	62.11	182.18
Rangifer tarandus tarandus	8	86.03	Prey	285.467	1749.55	72.86	402.56
Sus scrofa	23	96.12	Prey	0.234	7.53	2.18	43.85
Syncerus caffer	6	580	Prey	23.035	255.58	29.75	240.2
Tapirus terrestris	41	207.5	Prey	0.367	9.11	21.41	117.39
Tremarctos ornatus	1	140	Prey	15.035	258.17	35.41	185.38
Ursus arctos horribilis	18	212.5	Predator	30.3	249.65	40.92	253.13
Vulpes lagopus	18	3.58	Predator	1.289	4.42	0.81	64.09
Vulpes vulpes	20	5.48	Predator	1.39	5.89	13.04	199.24

Assessing trends in l_v

The resulting dataset of ballistic length scales was then analyzed to test for differences in l_{ν} between predators and prey, and for any effects of NDVI or forest cover on l_{ν} . Because l_{ν} was correlated with body size (Fig. 3A), we controlled for mass by regressing l_{ν} against body size on a log₁₀-log₁₀ scale using generalized least-squares fitting with Gaussian distributed errors. Due to phylogenetic autocorrelation⁴², closely related species may exhibit similarities in movement due to common descent. Accordingly, we did not treat species data records as independent, but rather corrected for this inertia by adjusting the variance-covariance matrix in our regression model based on the phylogenetic relationships using the R package nlme⁴³. Phylogenetic relationships

between the mammalian species in our dataset were obtained from the VertLife repository⁴⁴. We sampled 1,000 trees and estimated the consensus tree using the R package phytools, version $1.0-3^{45}$. Several of the species and subspecies included in our analyses were not included in the VertLife repository. These were manually inserted into the phylogeny using the most recent estimates of divergence times from the closest related species found in the VertLife repository. This included the Persian leopard (Panthera pardus saxicolor), with divergence time of 0.297 Ma from the African leopard (P. pardus pardus)⁴⁶, the Sumatran elephant (Elephas maximus sumatranus), with divergence time of 190,000 years from the Indian elephant (E. maximus *indicus*)⁴⁷, the Sri Lankan elephant (*E. maximus maximus*), with divergence time of 43,000 years from the Indian elephant⁴⁷, elk (Cervus canadensis), with divergence time of 20 million years from Rangifer tarandus⁴⁸, and Norwegian reindeer (R. tarandus tarandus) with a divergence time from *R. tarandus* of 115,000 years⁴⁹. Divergence times for all of the species of *Giraffa* were taken from⁵⁰. The species G. camelopardalis was assigned a divergence time from G. giraffa of 0.37 Ma. G. tippelskirchi was assigned a divergence time from G. giraffa of 0.23 Ma, and G. reticulata a divergence time from G. camelopardalis of 0.26 Ma. The subspecies G. giraffa angolensis and G. giraffa giraffa had a divergence time of 0.04 Ma. G. camelopardalis antiquorum was assigned a divergence time from G. camelopardalis peralta of 0.15 Ma and G. camelopardalis camelopardalis from G. camelopardalis peralta of 0.12 Ma. The resulting phylogenetic tree is shown in figure S1.

The final step in our analyses was to determine whether individual deviations from the allometric relationship in l_v could be described by trophic group, ecosystem productivity, or habitat permeability. We regressed the residuals of the phylogenetically controlled allometric model described above against trophic group, NDVI, percent forest cover, terrain roughness, and ml-HFI using generalized least-squares fitting with Gaussian distributed errors. The relative support for these models was then assessed by comparing the AICc values of the fitted models against intercept only models. We chose to work with the residuals as it allowed for like-to-like comparisons across all of the individuals in our dataset and clearer visualisations of the partial effects, while still correcting for differences in body sizes and phylogenetic inertia. The parameters of the body-mass scaling relationships in l_v are shown in Table S2. Results of the trophic group and NDVI regressions are presented in the main text and the model selection results are presented in Table S3. The R scripts used to produce the results presented in this work are openly available on GitHub at https://github.com/NoonanM/BallisticMotion.

	βo	95% CI	a	95% CI
All taxa	4.67	1.34 — 15.72	0.30	0.19 — 0.41
Prey	0.84	0.22 — 3.22	0.42	0.30 — 0.53
Predators	1.26	0.19 — 8.56	0.51	0.32 - 0.71

Table S2. Observed scaling relations of ballistic length scales in terrestrial mammals. All three of the scaling relations had the general form $l_v = \beta_0 \text{ mass}^{\alpha}$.

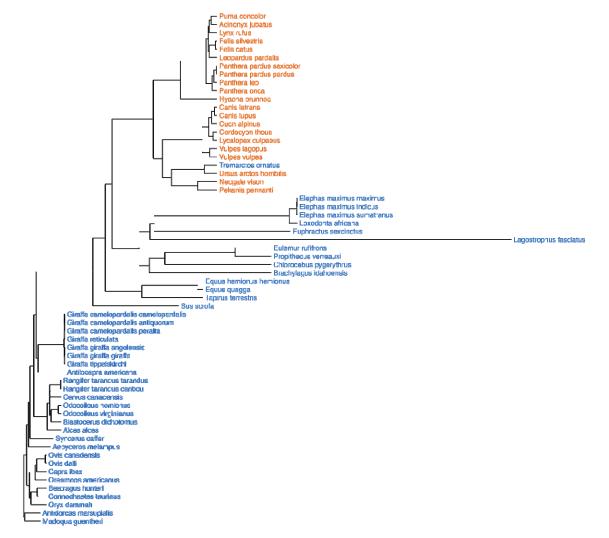


Figure S1 Phylogenetic relationships of the Mammalian species analysed in the main text with species labels coloured by trophic guild (prey in blue, predators in orange).

Table S3. Table showing the model selection results for the effect of NDVI on the residuals of the body mass relationship in ballistic motion length scales, l_v , for predators and prey.

Model	AICc	Δ ΑΙCc
Prey l_v residuals		
-23.0 - 95.7 x NDVI	10552	0
-21.4	10577	25
Predator l_v residuals		
172.9 - 41.8 x NDVI	7145	0

165.3 7153 8

When corrected for body size, mammalian ballistic length scales, l_v , show mixed responses to measures of habitat permeability. We found no relationship between the l_v body mass residuals and percent tree cover within each individual's habitat for prey (P = 0.06) nor predators (P =0.92), but both trophic groups exhibited exponentially variable heteroskedasticity with changing forest cover (Fig. S2a, table S4). We also found no relationship between the l_v body mass residuals and terrain roughness for prey (P = 0.06), but there was a positive relationship for predators (P = 0.001). Here again both trophic groups exhibited exponentially variable heteroskedasticity with changing terrain roughness (Fig. S2b, table S5). Finally, both predators and prey living in human modified landscapes had shorter ballistic length scales than those living in more natural environments, but these relationships were non-linear and heteroskedastic (Fig. S2c, table S6). Although this latter result is somewhat surprising given some empirical studies' findings that the structure present in natural ecosystems can impact the capacity for individuals to maintain directed motion²⁴, it is in line with the reductions in mammalian movement observed in human modified environments generally²¹.

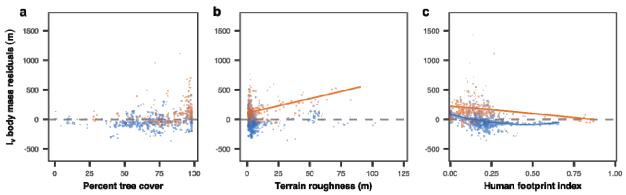


Figure S2 Relationship between mammalian ballistic length scales and habitat permeability. When corrected for body size, mammalian ballistic length scales, l_{ν} , show mixed responses to measures of habitat permeability. The scatterplot in **a** shows the residuals of the allometric scaling of the ballistic length scale, l_{ν} for prey (blue), as well as predatory (orange) mammals as a function of the percent tree cover. In **b** the residuals of the body mass scaling of l_{ν} for predators and prey are shown as a function of terrain roughness. The scatterplot in **c** depicts the body mass residuals as a function of the human footprint index, a satellite derived measure of human disturbance ranging on a scale between 0 and 1. Each point is an individual (n = 1,396) representing 62 species. The solid lines depict the selected regression models and the shaded areas the 95% confidence intervals around the mean trend. Trends are not shown for cases where the intercept-only model was selected.

Table S4. Table showing the model selection results for the effect of percent forest cover on the residuals of the body mass relationship in ballistic motion length scales, l_{ν} , for predators and prey.

	Model	AICc	∆AICc
Prey l_v residuals			
	$-14.5 + \varepsilon \sim N(0, \sigma^2 x e^{-0.004 x \text{ forest cover}})$	10574	0

$$-53.0 + 0.5 \text{ x forest cover} + \varepsilon \sim N(0, \sigma^2 \text{ x e}^{-0.004 \text{ x forest cover}})$$
 10574 0

$$-52.7 + 0.5 \text{ x forest cover} + \varepsilon \sim N(0, \sigma^2)$$
 10577 3

$$-21.4 + \varepsilon \sim N(0, \sigma^2)$$
 10577 3

Predator l_v residuals

$158.7 + \varepsilon \sim N(0, \sigma^2 \ge e^{0.014 \le \text{forest cover}})$	7145	0

163.7 - 0.1 x forest cover + $\varepsilon \sim N(0, \sigma^2 x e^{0.014 x \text{ forest cover}})$	7145	0
156.6 + 0.1 x forest cover + $\varepsilon \sim N(0, \sigma^2)$	7154	9
$165.3 + \varepsilon \sim N(0, \sigma^2)$	7154	9

Table S5. Table showing the model selection results for the effect of terrain roughness on the residuals of the body mass relationship in ballistic motion length scales, l_{ν} , for predators and prey.

Model	AICc	∆AICc
Prey l_v residuals		
$-37.4 + \varepsilon \sim N(0, \sigma^2 \ge e^{-0.02 \ge roughness})$	10504	0
-11.8 - 0.6 x roughness + $\varepsilon \sim N(0, \sigma^2 x e^{-0.02 x roughness})$	10505	1
-5.1 - 1.1 x roughness + $\varepsilon \sim N(0, \sigma^2)$	10563	59
$-21.4 + \varepsilon \sim N(0, \sigma^2)$	10577	73
Predator l _v residuals		
106.4 + 4.9 x roughness + $\varepsilon \sim N(0, \sigma^2 x e^{0.02 x roughness})$	7127	0
$130.9 + \varepsilon \sim N(0, \sigma^2 x e^{0.02 x roughness})$	7138	11
$102.2 + 4.3 \text{ x roughness} + \varepsilon \sim N(0, \sigma^2)$	7141	14
$165.3 + \varepsilon \sim N(0, \sigma^2)$	7154	27

Model	AICc	∆AICc
$Prey l_v$ residuals		
-1095.4 - 1876.4 x HFI + 1180.6 x e^{HFI} + $\varepsilon \sim N(0, \sigma^2 x e^{-2.8 x HFI})$	10487	0
-972.6 - 1630.8 x HFI + 1040.8 e^{HFI} + $\epsilon \sim N(0, \sigma^2)$	10512	25
11.5 - 189 x HFI + $\varepsilon \sim N(0, \sigma^2 x e^{-2.4 x \text{ HFI}})$	10512	25
23 - 219.3 x HFI + $\varepsilon \sim N(0, \sigma^2)$	10531	44
$-66.6 + \varepsilon \sim N(0, \sigma^2 \ge e^{-2.7 \ge \text{HFI}})$	10535	48
$-21.4 + \varepsilon \sim N(0, \sigma^2)$	10577	90
Predator l_v residuals		
262.1 - 198.0 x HFI - 39.1 x e^{HFI} + $\varepsilon \sim N(0, \sigma^2 x e^{-3.5 x HFI})$	7094	0
227.7 -262.8 x HFI + $\varepsilon \sim N(0, \sigma^2 x e^{-3.5 x \text{ HFI}})$	7105	11
778.2 + 684.9 x HFI - 598.2 x e^{HFI} + $\varepsilon \sim N(0, \sigma^2)$	7115	21
97.1 + $\varepsilon \sim N(0, \sigma^2 \ge e^{-3.0 \ge \text{HFI}})$	7123	29
201.5 - 152.3 x HFI + $\varepsilon \sim N(0, \sigma^2)$	7129	35
$165.3 + \varepsilon \sim N(0, \sigma^2)$	7154	27

Table S6. Table showing the model selection results for the effect of the human footprint index on the residuals of the body mass relationship in ballistic motion length scales, l_v , for predators and prey.

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Author contributions: MJN and JMC conceived the manuscript; MJN conducted the analyses. MJN, JMC, RMG, CHF, and WHF wrote the first manuscript draft. Co-authors contributed data and assisted with writing the final version of the manuscript.

Competing interests: Authors declare that they have no competing interests.

Data and materials availability: The data reported in this paper and the R scripts used to carry out this study are openly available on GitHub at https://github.com/NoonanM/BallisticMotion.

Materials & Correspondence: Data and/or methods request should be sent to Michael Noonan.

Supplementary Materials

Figs. S1 to S2 Tables S1 to S6 Appendix S1: Extended List of Acknowledgements and Funders Data S1: Data used to generate the results presented in the main text.

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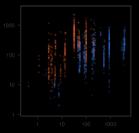
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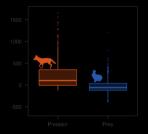
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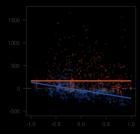
Appendix S1: Extended List of Acknowledgements and Funders

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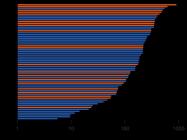
Austrian Science Fund (P18624). Funding was provided by the Utah Division of Wildlife Resources.

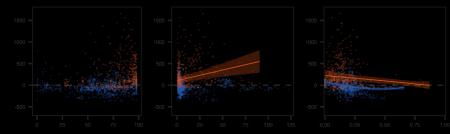


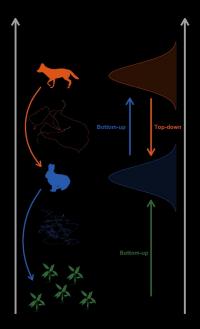












Puma concolor Acinonyx jubatus Lynx rufus Felis silvestris Felis catus Leopardus pardalis Panthera pardus saxicolor Panthera pardus saxicolor Panthera pardus pardus Panthera aloo Panthera onca Hyaena brunnea Canis larunnea Canis lupus Cuon alpinus Cerdocyon thous Lycalopex culpaeus Vulpes lagopus Vulpes vulpes Tremarcos ornatus Ursus arctos horriblis Neogale vison Pekania pennanti

Elephas maximus maximus Elephas maximus indicus Elephas maximus sumatranus Loxodonta africana junbractus sexcinctus

Eulemur rufifrons Propithecus verreauxi Chlorocebus pygerythrus Brachylagus idahoensis hemionus

Equus quagga Tapirus terrestris

Giraffa camelopardalis camelopardalis Giraffa camelopardalis antiquorum Giraffa camelopardalis peralta Giraffa camelopardalis peralta Giraffa giraffa angolensis Giraffa giraffa giraffa Giraffa tippelskirchi Antilocapra americana Rangifer tarandus tarandus Rangifer tarandus caribou Cervus canadensis Odocolleus hemionus Odocolleus hemionus Odocolleus hemionus Blastocerus dichotomus Alces alces Syncerus caffer Aepyceros melampus Ovis canadensis Ovis dalli Capra ibex Oreamnos americanus Beatragus hunteri Connochaetes taurinus Lagostrophus fasciatus