Home range is not constrained by numbers of hippocampal neurons

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Abstract

Numbers of hippocampal neurons vary by over three orders of magnitude across mammalian species. What evolutionary pressures shape this diversity? Given the role of the hippocampus in spatial mapping, the greater spatial navigation demands of larger home ranges may drive selection for more hippocampal neuron. Using data from 342 species, we crossed home range and population density data with cortical and hippocampal neuron counts predicted from clade-specific brain scaling laws to examine whether home range scales universally with estimated hippocampal neuron numbers across mammals. We confirm that home range scales universally with the inverse of population density across species and increases with body mass and metabolic rate. However, home range does not scale universally with hippocampal or cortical neuron numbers. Rather, scaling relationships differ by clade, such that carnivorans and cetartiodactyls traverse home ranges over 1,000-fold larger than primates with equivalent hippocampal neuron numbers. These findings persist across data subsets controlling for study method, duration, and temporal scope. Numbers of hippocampal neurons are thus not limiting to spatial navigation in the wild, calling into question adaptationist explanations for the evolution of more hippocampal neurons based on a supposed need for increased spatial processing capacity. We propose that home range is determined primarily by population density, mediated by field metabolic rate and diet. The diversity in hippocampal neuron numbers across mammals, in turn, arises as a byproduct of clade-specific scaling of numbers of cortical neurons which we suggest is contingent on energetic opportunity, not on navigational or other cognitive demands.

Significance Statement

Why do some animals have so many more brain neurons than others? One explanation is evolution through selection for more neurons as needed for instance to navigate their environments to find food. However, this study finds that carnivorans and whales traverse distances more than 1,000-fold larger than primates with similar numbers of neurons in the hippocampus, the brain structure required for spatial navigation. The authors question the traditional need-based explanation for brain diversity in favor of an opportunity-based account of the evolution of numbers of brain neurons.

Introduction

There is enormous diversity in numbers of cortical neurons across mammalian species. What are the drivers of that diversity, that is, what explains why some species have very few and others have up to a thousand times as many cortical neurons? Ever since Darwin, need-based natural selection is as a rule invoked as the main driver of diversity in evolution. By that logic, and assuming that more neurons increase the information-processing capacity of brain structures, different parts of the brain with different functions should be found to have more neurons as species rely more heavily on those functions.

In what concerns the hippocampus, a cortical structure responsible for the associative learning underlying concept building and spatial navigation[1-6] , those species with increasing needs for spatial navigation should be under selective pressure for increasing numbers of hippocampal neurons. Across mammalian species, the need for spatial navigation capacity should increase with home range, which is the area that an animal traverses over a set period of time in its normal activities of food gathering, mating, and caring for young[7]. Mammals create spatial maps using their hippocampus[1,8,9]It has been suggested that hippocampus size varies with relative selection pressures on cognitive mapping abilities and spatial memory[10-12]. Larger animals have increasingly larger home ranges[13-19], which is considered to be a response to the larger metabolic needs of species with increasing body mass[7,13,14,18,19,21-24]. Increased metabolic needs would also explain the larger home ranges of carnivores compared to herbivores of similar body mass[13,19,24,25]. Accordingly, larger animals with larger home ranges should be under selective pressure for increased hippocampal volume. Indeed, animals who rely more on complex spatial behaviors (e.g., animals with larger home ranges) have larger hippocampi[10,26-30].

Similarly, increasing the spatial complexity of the environment has been shown to increase
hippocampal volume[31-35].

The underlying rationale behind the argument that animals need larger hippocampi to
navigate larger home ranges is that larger hippocampi should consist of proportionately more
neurons; absent direct data on numbers of hippocampal neurons, hippocampal volume has been
expected to serve as a proxy for the number of neurons in the hippocampal network serving
spatial navigation. So far, however, data on actual numbers of neurons in the hippocampus have
not been available. Models shows that increasing numbers of neurons in the hippocampal
network should afford a linear increase in the capacity of this network to represent the area of the
environment that an individual traverses with a given spatial resolution[9,36-38]. If covering the
increased surface areas of larger home ranges requires increased numbers of hippocampal
neurons to the point that it imposes a selective pressure, then one should find a universal
correlation of both hippocampal size and numbers of hippocampal neurons with home range
across a wide variety of mammalian species.

The number of neurons in the mammalian hippocampus was just recently shown to scale
universally across species not with hippocampal mass, but with the total number of neurons in
the cerebral cortex[39]. That finding puts us in position to finally examine directly whether home
range scales universally with the estimated number of hippocampal neurons across species
covering the entire spectrum of mammalian body sizes. Here we use the home range dataset
compiled by Broekman et al.[40] to which we add estimates of population density for those same
species compiled by Santini et al.[41]; and data on body mass and metabolic rate compiled in the
AnAge database by de Magalhães and Costa[42].We then cross these data both with numbers of
neurons in the cerebral cortex predicted from brain mass in each species according to the clade-
specific relationships that have been found to apply separately to primates, glires,
certartiodactyls, afrotherians, marsupials, and carnivorans[43]; and with the numbers of neurons in
the hippocampus of each species predicted from their number of cerebral cortical neurons using
the scaling relationships that apply to their clade[39]. The latter was used as an added check
because the universal scaling of numbers of hippocampal neurons with the square root of total
numbers of cerebral cortical neurons[39] means that any scaling relationships that apply to total
numbers of cerebral cortical neurons also apply to hippocampal neurons with the exponent

divided by 2. We could then test whether the expected universal correlation between home range

and numbers of hippocampal neurons applied across mammalian species.

Results

Across all mammalian species in our dataset, we find that average home range scales as power functions of adult body mass (Fig. 1A) and basal metabolic rate (Fig. 1B) with exponents 1.16 ± 0.04 and 1.44 ± 0.09, respectively. Thus, our dataset reproduces the finding that larger animals, which also have higher metabolic rates, have larger average home ranges, and with scaling exponents above unity, as has been described previously[14,17-19,22,25,44]. With the exception of aquatic carnivores and some bears (asterisks in Fig. 1), this scaling applies across species of all clades, including primates, which have average home ranges well within the 95% prediction interval for their body mass (Fig. 1A) and basal metabolic rate (Fig. 1B). Analysis of the residual variation in $\log_{10}$ average home range after accounting for $\log_{10}$ body mass or $\log_{10}$ basal metabolic rate shows a significant effect of clade (Fig. 1C, D, respectively, both ANOVA $p<0.0001$). However, the apparent effect of clade is due to diet type (Fig. 1E, F, both ANOVA $p<0.0001$); the clear outliers in Fig. 1A are aquatic carnivores and some bears, which stand out in having far greater home ranges than predicted for their body size. Across species of each diet type, average home range scales similarly across clades (all ANOVA $p>0.01$).

Figure 1. Average home range scales universally with body mass and basal metabolic rate across a wide range of mammalian species, with the exception of aquatic carnivorans and some bears. All plotted functions include all species and clades in the dataset. Allometric exponents and r^2 values are shown in the graphs. **A**, average home range scales with adult body mass (p < 0.0001, n = 339) in a manner that leaves aquatic carnivorans and some bear species (asterisks) outside the 95% prediction interval (dotted lines). **B**, average home range scales with basal metabolic rate (p < 0.0001, n = 153) in an apparently universal fashion, but notice that data for aquatic carnivores and some bears are not available in the dataset. **C,E**, residual $\log_{10}$ home range for each species after accounting for $\log_{10}$ adult body mass, grouped by clade (**C**) or by diet type (**E**). **D,F**, residual $\log_{10}$ home range for each species after accounting for $\log_{10}$ basal metabolic rate, grouped by clade (**D**) or by diet type (**F**). Diet types are carnivory (C), herbivory (H) or omnivory (O).

In contrast, we find that the scaling relationships between home range and numbers of neurons in the cerebral cortex (Fig. 2A) and hippocampus (Fig. 2B) do not apply universally across all mammals or clades. Using the functions fitted across all species, primates consistently have smaller home ranges and carnivorans consistently have larger home ranges than expected for their numbers of neurons, as the residuals show (Fig. 2C, D; ANOVA for all clades, p<0.0001; ANOVA for carnivorans and primates, p<0.0001). Importantly, the distinct scaling of home range with numbers of cortical and hippocampal neurons in primates is not due to diet type (Fig. 2E, F), because primate home ranges remain significantly different (smaller) after accounting for diet type (ANOVA for carnivorans, glires and primates, p<0.0001 for each diet type). Aquatic carnivores (all of which are carnivoran species) again are the species with the largest residual home range, now joined by several cetacean species (Fig. 2A, B). The systematic

disparities between carnivorans, cetaceans and primates suggest that the scaling of home range

with numbers of cortical and hippocampal neurons is clade-specific.

Figure 2: Home range does not scale universally with numbers of cortical or hippocampal neurons across
mammalian species. All plotted functions include all species and clades in the dataset. Allometric
exponents and r^2 values are shown in the graphs. A, B, average home range scales with estimated total
numbers of cerebral cortical neurons (A; $p < 0.0001$, $n = 295$) or of hippocampal neurons (B; $p < 0.0001$,
$n = 295$) in a manner that leaves aquatic carnivorans and some bear species (asterisks) outside the 95%
prediction interval (dotted lines), and additionally leaves all primate species below the fitted line,
although mostly still inside the prediction interval. C, residual $\log_{10}$ home range for each species after
accounting for $\log_{10}$ number of cerebral cortical neurons, grouped by clade. D, residual $\log_{10}$ home range
for each species after accounting for $\log_{10}$ number of hippocampal, grouped by clade. Note that primate
species consistently have smaller home ranges and carnivorans consistently have larger home ranges than
predicted for their numbers of neurons.

The clade-specificity of the scaling of home range with numbers of cortical and hippocampal

neurons is evidenced in Figure 3. Glires, eulipotyphlans, marsupials, afrotherians and primates

share scaling exponents below 3.0 and 6.0 (for numbers of cortical and hippocampal neurons,

respectively), whereas carnivorans and artiodactyls (including cetaceans) share exponents above

those values (Fig. 3A, B). Interestingly, the data can be grouped in two different ways that make

either primates or large carnivorans and artiodactyls into outliers. When carnivorans and

cetartiodactyls are excluded from the analysis, yielding a joint power function with an r^2 of

0.639, primates are well aligned with glires, marsupials and eulipotyphlans in the scaling of

home range with numbers of neurons (Fig. 3A, B). In this case, large carnivorans and

cetartiodactyls would be described as having far larger home ranges than predicted for their

numbers of cortical and hippocampal neurons. Alternatively, if only primates are excluded while

carnivorans and cetartiodactyls are included in the analysis together with all other clades (Fig.

3C, D), we find that average home range scales with numbers of all cortical or hippocampal

neurons as power functions of a higher r^2 of 0.790 that however still exclude the aquatic

carnivorans and some bears in the dataset. Using this latter analysis, which provides a better

description of the variation in the dataset, primates are the outlier clade, consistently found to

occupy much smaller home ranges than other mammals of similar numbers of cortical or

hippocampal neurons. Either way, two facts remain: that home range and numbers of cortical

neurons do not vary in a universally correlated manner across mammalian species; and that

aquatic carnivorans and some bears occupy much larger home ranges than can be predicted by

their body size or numbers of neurons.

Figure 3. Home range scales with numbers of cortical or hippocampal neurons in a clade-specific manner. Scaling exponents ± standard error that apply to each clade (not plotted) are indicated in the graphs. Plotted functions include all species and clades in the dataset minus carnivorans and cetartiodactyls (A and B) or minus primates (C and D). Allometric exponents and r^2 values for the plotted functions are shown in black in the top left of each graph. **A and B:** When carnivorans and cetartiodactyls are excluded from analysis, average home range scales with number of cortical neurons and number of hippocampal neurons (p < 0.0001, n = 216) in a manner that includes primates and leaves aquatic carnivorans (asterisks), most cetaceans and some artiodactyls outside the 95% prediction interval (dotted lines). **C and D**: When primates are excluded from analysis, average home range scales as power functions of numbers of cortical and hippocampal neurons (p < 0.0001, n = 216) with a higher r^2 and a 95% prediction interval that excludes primates and still excludes the aquatic carnivorans and some bears in the dataset.

Effect of tracking methods, sample size, and study duration

The average home range estimates used so far in the analysis combine various tracking methods, sample size, and study duration, as has been standard practice in the field[19,40,45,46]. The lack of consistence in the time span (daily, annual, other) and level (individual, group, population) that apply to the values of home range analyzed raises the concern that the finding that primates are outliers in the scaling of home range with numbers of neurons is a measurement artifact. To address this possibility, we determined whether primates are still outliers when the scaling between numbers of cortical or hippocampal neurons and home range is calculated exclusively for subsets of data restricted to the same combination of time span (home range covered in a day or in a year) and level (home range of single individuals or of a population). Figure 4 shows that even in these much more restricted datasets, and despite their inclusion in the fitted data, primates still consistently have smaller home ranges than predicted for their numbers of cortical or hippocampal neurons, well below the 95% prediction interval.

Figure 4. Primates have smaller home ranges than predicted for their numbers of cortical and hippocampal neurons even in data sets restricted to similar time span and population level. Average home range is plotted as a function of numbers of cortical neurons (A-D) or estimated numbers of hippocampal neurons (E-H) using data restricted to daily home range measured in individuals (A, E); annual home range measured in individuals (B, F); daily home range measured in a population (C, G); or annual home range measured in a population (D, H). The plotted functions apply to all species in each data set, including primates, which still fall below the 95% prediction interval (dotted lines) in every case. Allometric exponents (all $p<0.0001$), r^2 values and n are shown in the graphs.

Population density is a universal predictor of home range, and also does not scale universally with numbers of cortical neurons

Our dataset reproduces the previous observations that home range is a negative inverse function of population density[15,18,47-49]. We find that average home range scales with the near inverse of population density, with an exponent indistinguishable from -1 (Fig. 5A), in a manner that applies equally across carnivores, herbivores and omnivores (Fig. 5B, ANOVA $p=0.3789$). While an ANOVA across clades (Fig. 5C) indicates a significant effect of clade ($p<0.0001$), residual home ranges after accounting for population density do not differ across the three clades that share large numbers of hippocampal neurons but differ markedly in average home range (ANOVA for carnivorans, cetartiodactyls, and primates, $p=0.1429$). Interestingly, while the scaling of home range with the inverse of population density includes cetaceans, most aquatic carnivore species remain outliers, with larger home ranges than predicted for their population density (Fig. 5A).

Figure 5. Population density is a better universal predictor of home range than any other variable examined. A, home range scales as a power function of population density (A, p < 0.0001, n = 255). B, residual $\log_{10}$ home range by population density for each species grouped by diet. C residual $\log_{10}$ home range by population density for each species grouped by clade.

In line with the lack of a universal correlation between home range and numbers of hippocampal

neurons, we find that the power function that relates population density to numbers of

hippocampal neurons (and thus also numbers of cortical neurons; not shown) is not universal

across all mammals in the dataset (Fig. 6A, ANOVA $p < 0.0001$), and also does not apply equally

across species of different diet types (Fig. 6B, ANOVA $p < 0.0001$). Primates and glires have

systematically higher and carnivorans have systematically lower population densities than

predicted for their numbers of neurons (Fig. 6C), even when analysis is restricted to herbivorous

and omnivorous diet types (ANOVA, $p < 0.0001$), which argues for clade-specific relationships

between population density and numbers of hippocampal and cortical neurons.

Figure 6. Population density is not a universal predictor of number of hippocampal neurons. Population density scales as a power function of number of hippocampal neurons (A; $p < 0.0001$, $n = 220$) across species in a manner that includes primates but still leaves some aquatic carnivorans (asterisks) and cetaceans outside the 95% prediction interval (dotted lines). B, residual $\log_{10}$ population density by number of hippocampal neurons for each species grouped by diet type. C residual $\log_{10}$ population density by number of hippocampal neurons for each species grouped by clade.

Finally, we sought to determine the source of the larger home ranges and lower population densities of carnivores and carnivorans compared to primates of similar numbers of hippocampal neurons by examining how these three variables scale with the field metabolic rate that indicates how much energy different species must be able to procure per day in order to survive. Figure 7 shows that while field metabolic rate is a good predictor of home range (Fig. 7A), population

density (Fig. 7B) and number of hippocampal neurons (Fig. 7C), it is not a universal predictor of any of these three variables. Carnivoran species (the vast majority of which also have carnivorous diets; Fig. 7D) and also carnivorous species (Fig. 7G) consistently have larger home ranges than predicted for their field metabolic rate. Carnivorous species, which include carnivorans and cetaceans, also have lower than predicted population densities for their field metabolic rates (Figs. 7E, H). Diet, on the other hand, does not differentiate the relationship between numbers of hippocampal neurons and field metabolic rate (Fig. 7I; ANOVA, p=0.0343 versus p<0.001 for all other comparisons); the distinguishing factor here is clade, as primates have systematically more hippocampal neurons than predicted for their field metabolic rates (Fig. 7F), in line with their systematically larger numbers of cortical neurons compared to non-primate mammals of similar body mass[50]. We thus find that while the home range occupied by mammalian species is a universal function of their population density, with both variables affected coordinately by diet, the number

of neurons in the hippocampus of mammalian species is not universally related to the home range,

population density, or field metabolic needs of a species.

Figure 7. Field metabolic rate mediates population density. Field metabolic rate scales as a power
function of average home range, population density, and number of hippocampal neurons across
mammalian species. All plotted functions include all species and clades in the dataset. Allometric
exponents and r^2 values are shown in the graphs. A, average home range scales with field metabolic rate
(A; p < 0.0001, n = 63) population density (B; p < 0.0001, n = 49), and number of hippocampal neurons
(C; p < 0.0001, n = 58) in a manner that realigns aquatic carnivorans and some bear species (asterisks)
and primates inside the 95% prediction interval (dotted lines). D-F, residual log$_{10}$ home range, population
density, and number of hippocampal neurons by field metabolic rate for each species grouped by clade.
G-I, residual log$_{10}$ home range, population density, and number of hippocampal neurons by field
metabolic rate for each species grouped by diet.

Discussion

Obtaining sufficient energy to make it repeatedly through the days, together with

maintaining physical integrity, is the one non-negotiable condition for animal survival. Our

findings are consistent with a scenario in which, given an environmental reality of limited food

availability per surface area, the higher rates of energy use per day of larger animals cause them

to self-organize through food-seeking exploitative behaviors into predictably lower population

densities that in turn necessitate exploratory behaviors and dispersal into inversely larger home

ranges. Diet type affects feeding efficacy and efficiency and thus modulates the relationship

between field metabolic rate and population density (Fig. 7), which, given the necessarily inverse

relationship between population density and home range[20], explains the dependence on diet of

the scaling of home range with body mass[19]. Specifically, carnivory exposes animals to the

relative scarcity of prey compared to vegetable foodstuffs per surface area in the

environment[18,19,49]. As a result, for a given body size and field metabolic rate, carnivorous

animals exist in lower population densities and explore larger home ranges than omnivorous

animals, and herbivorous species exist in larger population densities and explores smaller home

ranges, all the while home range and population density continue to scale universally across all

diet types.

We propose that the relative scarcity of prey combined with higher metabolic costs of

living in cold water and the three-dimensional nature of hunting in an aquatic environment

explain the finding that aquatic carnivores, which includes both carnivorans and cetaceans, are

consistently outliers in our dataset, exploring home ranges that are larger than expected not only

for their body mass and field metabolic rate but also population density. The resting metabolic

costs of aquatic mammals are about twice as high as those of terrestrial mammals, with a scaling

relationship for resting metabolic rate with body mass of $26.9 \ M_{bd}^{0.69}$ in aquatic mammals[51] in

contrast to McNab's allometric prediction for the basal metabolic rate of similarly sized

terrestrial mammals[52] of $10.55 \ M_{bd}^{0.724}$ (after converting for M_{bd} in kg), probably due to heat lost

to cold water. The effect of prey scarcity to drive decreased population density and larger home

range is expected to be especially severe in aquatic environments, where there is an added

dimension to prey movement. However, given the difference between stroke vs. stride energetic

costs, fully submerged mammals are able to avoid hydrodynamic drag and exhibit lower mean

stroke costs compared to surface swimming carnivores[53]. The higher efficiency of fully

submerged carnivores, such as cetaceans, might explain how come their home ranges match the

predicted values for their very low population densities whereas the aquatic carnivorans, as

inefficient surface swimmers, explore much larger home ranges than predicted for their

population densities (Fig. 5A).

To the heart of our central question, we find that the enormous range of coordinated

variation in home range and population density occurs without the expected universal correlation

with numbers of hippocampal neurons typically presumed to be necessary to support spatial

exploration[10,26-35]. To the contrary, we show that carnivoran species are capable of exploring

home ranges of around 1,000 km^2 on land and as much as 100,000 km^2 over water with the same

estimated ca. 30 million hippocampal neurons found in primates who remain in an area of just

about 1 km^2 (Fig. 2B). Conversely, carnivorans and primates that explore home ranges of about

10 km^2 do so with numbers of hippocampal neurons that are 10 times as large in primates than in

carnivorans. If carnivorans and primates alike obtain the energy that they need by occupying the

home range that is predicted for their population density (Fig. 5A), the number of hippocampal

neurons available to represent the surface area covered by a species is clearly not limiting to the

point that it would be under selective pressure to increase together with home range as animals

became larger in evolution. Known mechanisms of multiplexing hippocampal representations of

environments on different scales[38] would suffice to account for how a given number of

hippocampal neurons suffices to allow exploration of home ranges varying by a factor of over

1,000-fold that we uncover here.

Those determined to find adaptationist explanations to the discrepancy in scaling of home

range with numbers of hippocampal neurons between primates and other mammals might turn to

Fig. 3D and argue that non-primate species with larger home ranges do have the correspondingly

larger numbers of hippocampal neurons predicted by a supposed need for more hippocampal

spatial representational power. In that case, however, a new argument would have to be invoked

to explain how come primates evolved much larger numbers of hippocampal neurons than

"necessary" to represent their comparatively small environments. In our view, a much more

parsimonious explanation lies in acknowledging that the scaling relationship between numbers of

neurons and body size is not universal[50,54]. If there is any correlation between home range and

numbers of hippocampal (and cerebral cortical) neurons, we argue that it is simply because

larger animals, which have universally larger home ranges as predicted by their population

densities (according to the combination of field metabolic rate and diet), coincidentally tend to

have more cortical neurons – but following scaling rules that differ between primate and non-

primate clades[50]. In this case, the clade-specific correlation between home range and numbers of

hippocampal neurons is a mathematical contingency, not a biological necessity. This is not to say

that larger numbers of hippocampal neurons do not bring a cognitive advantage. According to

theory, the more the neurons in the hippocampal network, the bigger should be the capacity of

the network to represent an area with a certain spatial resolution[38]. Given the combination of the

predictably small home ranges of primate species and their large numbers of hippocampal

neurons (which are also predictable for their cerebral cortices[39]), we propose that primates

benefit from increased spatial resolution in their hippocampal maps of the environment, as the

radius of their place cells is expected to be reduced[38]. This prediction can be tested with

systematic comparisons of place cell radius across primate and non-primate species of either

similar home ranges or numbers of hippocampal neurons. Evidently, the comparatively lower

spatial resolution of non-primates is perfectly sufficient for their successful navigation in

exploring and exploiting whatever size home range is required for their survival. Thus, the

increased spatial resolution we predict for primates cannot be considered a necessity, and

therefore also not the result of selective pressures – unless one resorts to a convoluted argument

that only primates were subject to such pressures for increased spatial resolution.

Rather, we propose that it is more parsimonious that in exploring their environment,

mammals go as far as they have to go in order to meet their metabolic demands (given how their

diet impacts population density), but their numbers of hippocampal neurons are just not limiting

to how far they can go. The diversity of numbers of hippocampal neurons, which cannot be said

to have been driven by selective pressure for increased spatial navigation capacity, is tied to the

square root of the variation in total numbers of cortical neurons[39]. Thus, what needs to be

explained is what leads to the enormous variation in total numbers of cortical neurons in

mammalian evolution. We favor the simpler alternative explanation that numbers of cortical (and

so also hippocampal) neurons increase in evolution as consistently increased energetic

opportunities present themselves to a population across generations[55].

Materials and Methods

The home range and population densities of 342 mammalian species were compiled using

two databases: "HomeRange: A global database of mammalian home ranges"[40] and

"TetraDENSITY 2.0 – A database of population density estimates in Tetrapods"[41]. Home range

data collection and statistical techniques for each species differed across studies, with often

differing temporal measurements and differing observation techniques employed for any given

species. To combine home range and population density data into a single database, it was

necessary to calculate average home range values for each species.The combined database was

then integrated with data on body mass, basal metabolic rate and diet type compiled by de

Magalhães and Costa[42] in the AnAge database and with data on numbers of cortical neurons

predicted from brain mass[43]. From the numbers of cortical neurons, the number of hippocampal

neurons was estimated for each species according to clade-specific formulas: $e^{7.673} N_{cx}^{0.815}$ for

cetartiodactyls, $e^{3.181} N_{cx}^{0.658}$ for carnivorans, $e^{4.322} N_{cx}^{0.640}$ for marsupials, $e^{9.841} N_{cx}^{0.346}$ for

primates, $e^{3.384} N_{cx}^{0.693}$ for glires, and $e^{7.767} N_{cx}^{0.450}$ for afrotherians. Note that these clade-specific

calculations provide an alternate set of estimates for N_{hp}, which can also be more simply

obtained as roughly $N_{cx}^{0.5}$ [39].

From these data, average home range (measured in km^2) for each species was plotted in

log-transformed bivariate-fit graphs comparing home range against adult body mass (measured

in grams), metabolic rate (W), number of cortical neurons (N_{cx}), estimated number of

hippocampal neurons (N_{hp}), and population density (individuals/km^2) in JMP Pro 18.

Mammalian data points were separated by clade for comparison, as indicated in the graphs. We

purposefully do not use any phylogenetic correction methods because clade effects are addressed

directly in our analysis.

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