Supplementary Note 1: Sensitivity to additional death terms

It should be noted that our set of dynamics (Eqs. 2 and 4, main text) could include a constant death term of the form $-d_F F$ and $-d_H H$ to represent death not directly linked to starvation. Adding terms of this form to our model would simply adjust the effective value of λ and μ , and we could rewrite Eq. 4 with $\lambda' = \lambda - d$ and $\mu' = \mu - d$. These substitutions would not alter the functional form of our model nor the steady-states and qualitative results, however the quantitative values could shift based on the size of d relative to λ and μ .

Survivorship has a well-known functional form which changes systematically with size (e.g. ¹⁷). Typically survivorship is defined using the Gompertz curve

$$F = F_0 e^{(c_0/c_1)\left(1 - e^{c_1 t}\right)} \tag{1}$$

where the parameters have the following allometric dependencies on adult mass $c_0 = a_0 M^{b_0}$ and $c_1 = a_1 M^{b_1}$, with $a_0 = 1.88 \times 10^{-8}$ (s g^{-b₀}), $b_0 = -0.56$, $a_1 = 1.45 \times 10^{-7}$ (s g^{-b₁}), and $b_1 = -0.27$ (see¹⁷ for a review).

We are interested in the specific death rate of the form $\dot{F} = -dF$, and using the derivative of Eq. 1 we find that $d = c_0 e^{c_1 t}$. Our model considers the average rates over a population and lifecycle and the average death rate is given by

$$\bar{d} = \frac{1}{t_{\text{exp}}} \int_0^{t_{\text{exp}}} c_0 e^{c_1 t} dt \tag{2}$$

$$= \frac{c_0 \left(e^{c_1 t_{\rm exp}} - 1\right)}{c_1 t_{\rm exp}} \tag{3}$$

where $t_{\rm exp}$ is the expected lifespan following the allometry of $t_{\rm exp}=a_2M^{b_2}$ with $a_2=4.04\times 10^6$ (s g^{-b₂}) and $b_2=0.30$ ^{17,18}. Given the allometries above we have that

$$\bar{d} = \frac{a_0 \left(e^{a_1 a_2 M^{b_1 + b_2}} - 1\right) M^{b_0 - b_1 - b_2}}{a_1 a_2} \tag{4}$$

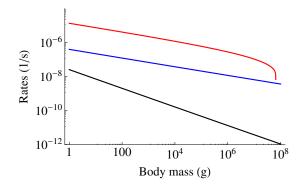
which scales roughly like M^{b_0} because b_1 and b_2 are close in value but opposite in sign. In Supplementary Figure 1 we compare the value of \bar{d} to the reproductive, λ , and starvation-based mortality, μ , rates. The values of \bar{d} are orders of magnitude smaller than these rates for all mammalian masses, and thus, adding this non-starvation based death rate to our model does not shift our results within numerical confidence.

Supplementary Note 2: NSM and the energy equivalence hypothesis

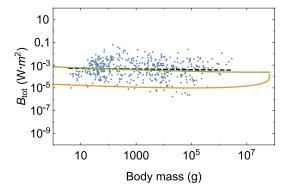
The energy equivalence hypothesis is based on the observation that if one assumes that the total metabolism of an ecosystem B_{tot} is equally partitioned between all species (B_i , the total metabolism of one species, is a constant), then the abundances should follow $N(M)B(M) = B_i$ implying that $N(M) \propto M^{-\eta}$, where η is the metabolic scaling exponent^{19,20}. As $\eta \approx 3/4$ this hypothesis is consistent with Damuth's law¹⁹. However, the actual equivalence of energy usage of diverse species has not been measured at the population level for a variety of whole populations. Supplementary Figure 2 recasts the results of the NSM in terms of this hypothesis and shows that F^*B is nearly constant over the same range of mammalian sizes up to the asymptotic behavior for the largest terrestrial mammals.

Supplementary Note 3: Application of NSM limits to aquatic mammals A theoretical upper bound on mammalian body size is given by $\epsilon_{\sigma} = 0$, where mammals are entirely composed of metabolic reserves, and this occurs at $M = 8.3 \times 10^8$ (g), or 120 times the mass of a male African elephant. We note this particular limit as it may have future relevance to considerations of the ultimate constraints on aquatic mammals.

Supplementary Figures



Supplementary Figure 1: The rates of reproduction λ (blue), starvation-based mortality μ (red), and survivorship-based death \bar{d} (black) as a function of adult mass.



Supplementary Figure 2: Total energetic use $B_{\rm tot}$ of consumer populations at the steady state as a function of body mass (F^* is shown in green and H^* in orange). The data are from Damuth²¹ and have been converted to total population metabolism using the allometric relationships for metabolic rate (e.g. Refs.³⁻⁵).

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