SUPPLEMENTARY INFORMATION

Physiology can influence animal activity, exploration, and dispersal

Nicholas C. Wu, & Frank Seebacher

School of Life and Environmental Sciences, The University of Sydney, NSW 2006, Australia

Supplementary Methods

Data exclusion criteria

- Non-English literature, books without links to primary source, and non-peer-reviewed articles.
- Passive dispersal such as human transportation of wildlife from native to new environments, parasite dispersal, and abiotic driven dispersal. In addition, dispersal metrics related to offspring success such as the production of dispersing gametes/eggs or the colonisation success of offspring from water/air currents.
- Categorised dispersal polymorphism. i.e., winged and non-winged insects (Zera and Denno, 1997).
- Migration as defined by Nathan et al. (2008).
- Studies that did not define specific metabolic rate measurement.
- Response variables measured at test temperatures different from acclimation temperatures (e.g. animals acclimated at warm treatment but response measured at an acute cold treatment).
- Interspecific correlation studies of physiology and movement.
- Experimental manipulations that confound comparisons of movement, such as food restriction, exposure to pollutants.
- Molecular responses not directly related to physiological traits such as mRNA expression or transcriptomics.
- Principal component variables where body condition was grouped with other morphological variables.
- Artificial selection lines for more active, more explorative individuals.
- Within-individual correlation studies which reflect repeatability of measurements.

Conversion statistics

The correlation coefficient (r) was calculated from inferential statistics with the following equations from Lipsey and Wilson (2001), Nakagawa and Cuthill (2007), and Noble et al. (2017):

$$r = \sqrt{[t^2/(t^2 + d.f.)]} = \sqrt{F/(F + d.f.)} = \sqrt{(\chi^2/n)}$$

where t = t-statistics, F = F-statistics, d.f. = model denominator degrees of freedom, $\chi^2 = \text{chi-squared statistics}$, and n = sample size. Effect sizes from inferential statistics were only retained when directions could be determined (e.g., F and χ^2 alone do not contain directional information).

Supplementary Tables

Table S1 Trait categories associated with individual movement and the specific responses within these. Number of effect sizes, studies and species are shown. Responses with asterisks were corrected for direction. MR = metabolic rate.

Categorised trait	Specific response	Effect size (k)	Studies (n)	Species (n)
Cardiovascular	Haematocrit	4	2	2
Condition	Body condition	40	25	28
Condition	Glycogen content	2	1	1
Hormone	Glucocorticoids	7	6	5
Hormone	Testosterone	2	2	2
Immunity	White blood cell (WBC) count	2	1	1
Locomotor capacity	Cost of transport (CoT)*	8	2	1
Locomotor capacity	Endurance	14	7	7
Locomotor capacity	Max force	1	1	1
Locomotor capacity	Max jump	3	1	3
Locomotor capacity	Sprint speed	17	9	9
Locomotor capacity	Critical sustained swimming speed (Ucrit)	31	4	2
Metabolism	Aerobic scope	8	4	4
Metabolism	Basal MR	9	4	4
Metabolism	Field MR	4	3	4
Metabolism	Liver citrate synthase	1	1	1
Metabolism	Liver cytochrome c oxidase	1	1	1
Metabolism	Max MR	17	7	6
Metabolism	Muscle citrate synthase	13	2	2
Metabolism	Muscle cytochrome c oxidase	1	1	1
Metabolism	Muscle lactate dehydrogenase	12	1	1
Metabolism	Mitochondrial P/O ratio	2	1	1
Metabolism	Resting MR	32	22	20
Metabolism	Routine MR	25	11	10
Metabolism	Standard MR	11	8	8
Metabolism	Mitochondria State III respiration (State III)	2	1	1
Metabolism	Mitochondria State IV respiration (State IV)	2	1	1
Musculoskeletal	Muscle biomechanics	1	1	1

Table S2 Trait categories associated with population range expansion and the specific responses within these. Number of effect sizes, studies and species are shown. Responses with asterisks were corrected for direction. MR = metabolic rate. MR = metabolic rate.

Category	Specific response	Effect size (k)	Studies (n)	Species (n)
Activity	Activity	64	11	6
Cardiovascular	Haematocrit	3	2	2
Cardiovascular	Haemoglobin	2	1	1
Cardiovascular	Heart mass	2	1	1
Cardiovascular	Red blood cell (RBC) count	1	1	1
Condition	Body condition	28	6	4
Condition	Fat score	1	1	1
Heat shock protein	Heat shock protein (HSP70)	3	1	1
Hormone	Corticosterone	20	7	4
Immunity	Bacterial killing assay (BKA)	6	3	2
Immunity	Neutrophil/lymphocyte ratio (NLR)	3	2	2
Immunity	White blood cell (WBC) count	1	1	1
Locomotor capacity	Endurance	38	9	7
Locomotor capacity	Sprint speed	30	3	1
Metabolic enzyme	Muscle citrate synthase	2	1	1
Metabolic enzyme	Muscle lactate dehydrogenase	2	1	1
Metabolism	Max MR	2	1	1
Metabolism	Resting MR	6	3	3
Metabolism	Routine MR	4	1	1
Metabolism	Standard MR	4	2	2
Musculoskeletal	Muscle profile	1	1	1
Musculoskeletal	Muscle ratio	3	1	1
Oxidative status	Reactive oxygen species	2	1	1
Oxidative status	Total antioxidant capacity	2	1	1
Thermal tolerance	Critical thermal maximum (CTmax)	10	2	2
Thermal tolerance	Critical thermal minimum (CTmin)*	12	6	4

Table S3 Mean parameter estimates, estimate error, and 95% Bayesian credible intervals for the individual movement model, which includes three movement types (activity, exploration, dispersal), thermal strategy (ectotherm and endotherm), sex (female, male, mixed), age (juvenile and adult), origin (wild caught or lab reared) as fixed effects, and sampling error and publication year as covariates to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), species identity, and phylogenetic relatedness.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Activity	0.29	0.15	-0.003	0.58
Exploration	0.26	0.16	-0.05	0.57
Dispersal	0.35	0.18	-0.01	0.70
Thermal strategy	-0.08	0.11	-0.29	0.15
Male	-0.04	0.08	-0.19	0.12
Mixed	-0.01	0.10	-0.20	0.18
Age	0.09	0.10	-0.11	0.29
Origin	-0.10	0.09	-0.28	0.07
Sampling error	-0.51	0.35	-1.18	0.18
Publication year	-0.01	0.01	-0.02	0.002
Random effects				
σΔeffect size ID	0.4	0.02	0.36	0.45
σ∆study ID	0.15	0.06	0.03	0.25
$\sigma\Delta$ species	0.08	0.06	0	0.21
$\sigma\Delta$ phylogeny	0.08	0.07	0	0.25

Table S4 Heterogeneity of the individual movement analysis and population range expansion analysis. k = number of estimates, and $I^2 =$ heterogeneity. Estimates shown correspond to modes and 95% highest posterior density intervals.

	k (n)	I² [effect size]	<i>I</i> ² [study ID] (%)	I ² [species] (%)	I ² [phylogeny] (%)	I ² [total] (%)
Individual overall model	272	58.6	21.7	1.3	3.5	97.3
		[40.7–72.8]	[4.3–33.6]	[0-23.3]	[0-26.0]	[96.6–98.0]
Population overall model	252	30.3	46.1	6.1	6.3	99.9
		[19.2–38.7]	[32.6–60.7]	[0-23.3]	[0-33.0]	[99.9–100]

Table S5 Mean parameter estimates, estimate error, and 95% credible intervals for the individual movement analysis with separate models for activity, exploration, and dispersal. Models include fixed-effect parameters for all extracted traits with >5 effect sizes, thermal strategy (ectotherm and endotherm), and sampling error with the publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), species identity, and phylogenetic relatedness.

Parameters - Activity only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Condition	0.31	0.22	-0.12	0.74
Locomotor capacity	0.38	0.19	-0.01	0.74
Metabolism	0.23	0.18	-0.15	0.58
Thermal strategy	-0.21	0.23	-0.67	0.23
Sampling error	-0.57	0.47	-1.49	0.37
Publication year	-0.01	0.01	-0.03	0.01
Random effects				
σΔeffect size ID	0.39	0.03	0.33	0.45
σ∆study ID	0.28	0.07	0.15	0.42
σΔspecies	0.11	0.08	0	0.29
σΔphylogeny	0.15	0.14	0.01	0.53
Parameters - Exploration only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Locomotor capacity	0.19	0.23	-0.24	0.66
Metabolism	0.12	0.21	-0.29	0.55
Thermal strategy	-0.03	0.22	-0.44	0.43
Sampling error	-0.59	0.86	-2.33	1.09
Publication year	0.00	0.03	-0.06	0.05
Random effects				
σΔeffect size ID	0.34	0.04	0.27	0.41
σ∆study ID	0.11	0.08	0	0.31
σΔspecies	0.13	0.09	0.01	0.34
σΔphylogeny	0.16	0.15	0.01	0.55
Parameters - Dispersal only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Condition	0.01	0.49	-0.99	0.97
Thermal strategy	0.28	0.45	-0.58	1.23
Sampling error	0.43	2.25	-4.14	4.93
Publication year	-0.04	0.03	-0.09	0.02
Random effects				
σ∆effect size ID	0.38	0.1	0.24	0.62
σ∆study ID	0.21	0.16	0.01	0.61
σΔspecies	0.22	0.16	0.01	0.61
σΔphylogeny	0.37	0.33	0.01	1.2

Table S6 Mean parameter estimates, estimate error, and 95% credible intervals for the individual movement analysis, grouped by metabolism, and locomotor performance traits, which includes fixed-effect parameters for all extracted responses with >5 effect sizes, age (juvenile, adult), thermal strategy (ectotherm and endotherm), and sampling error and publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), species identity, and phylogenetic relatedness.

Parameters - Metabolism	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Aerobic scope	0.22	0.21	-0.2	0.63
Inactive MR	0.04	0.15	-0.25	0.34
Active MR	0.51	0.17	0.18	0.84
Metabolic enzyme	0.15	0.26	-0.37	0.66
Maximum MR	0.21	0.19	-0.15	0.57
Age	-0.07	0.13	-0.32	0.17
Sampling error	-0.79	0.52	-1.84	0.22
Publication year	-0.01	0.01	-0.02	0.01
Random effects				
σΔeffect size ID	0.38	0.03	0.33	0.44
σΔstudy ID	0.13	0.07	0.01	0.26
σΔspecies	0.09	0.07	0	0.25
$\sigma\Delta phylogeny$	0.11	0.1	0	0.37
Parameters - Locomotor cap.	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects			-	-
CoT	-0.12	0.71	-1.55	1.3
Endurance	0.58	0.45	-0.3	1.51
Sprint speed	0.5	0.45	-0.33	1.46
$U_{ m crit}$	0.02	0.71	-1.4	1.47
Thermal strategy	-0.45	0.57	-1.61	0.68
Sampling error	0.02	1.39	-2.76	2.72
Publication year	0.01	0.02	-0.04	0.05
Random effects				
σΔeffect size ID	0.37	0.04	0.3	0.47
σ∆study ID	0.24	0.16	0.01	0.61
σΔspecies	0.27	0.19	0.01	0.69
$\sigma\Delta$ phylogeny	0.44	0.33	0.02	1.27

Table S7 Mean parameter estimates, estimate error, and 95% credible intervals for the individual movement analysis, with separate models for activity, exploration, dispersal. Models include fixed-effect parameters for all extracted taxa with >5 effect size, and sampling error with the publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), and species identity.

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Parameters - Activity only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects		0.50	0.10	0
Amphibia	0.21	0.20	-0.18	0.61
Aves	0.08	0.26	-0.45	0.60
Invertebrate	0.37	0.17	0.04	0.70
Mammalia	0.11	0.19	-0.27	0.48
Osteichthyes	0.28	0.16	-0.03	0.60
Reptilia	0.24	0.30	-0.36	0.83
Sampling error	-0.59	0.51	-1.57	0.44
Publication year	-0.01	0.01	-0.03	0.01
Random effects				
σΔeffect size ID	0.4	0.03	0.35	0.46
σΔstudy ID	0.28	0.07	0.14	0.43
$\sigma\Delta$ species	0.11	0.08	0	0.29
Parameters - Exploration only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Amphibia	0.08	0.24	-0.38	0.55
Aves	-0.03	0.20	-0.42	0.37
Invertebrate	0.09	0.20	-0.30	0.51
Mammalia	0.09	0.20	-0.30	0.48
Osteichthyes	0.17	0.24	-0.31	0.65
Reptilia	0.20	0.28	-0.35	0.76
Sampling error	-0.68	0.88	-2.45	1.02
Publication year	0.01	0.02	-0.04	0.06
Random effects				
σΔeffect size ID	0.33	0.03	0.27	0.4
σΔstudy ID	0.1	0.08	0	0.29
σΔspecies	0.13	0.09	0.01	0.35
Parameters - Dispersal only	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Amphibia	0.19	0.29	-0.39	0.78
Aves	0.35	0.33	-0.28	1.01
Invertebrate	-0.25	0.48	-1.19	0.72
Mammalia	0.19	0.27	-0.35	0.72
Sampling error	0.05	1.64	-3.24	3.27
Publication year	-0.04	0.01	-0.07	-0.01

Random effects

σΔeffect size ID	0.37	0.07	0.25	0.54
σ∆study ID	0.16	0.12	0.01	0.44
σΔspecies	0.18	0.13	0.01	0.48

Table S8 Mean parameter estimates, estimate error and 95% credible intervals for the interaction between dispersal mode (aerial, aquatic, terrestrial) and temperature difference between range core and edge.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Intercept, β_0	3.51	0.09	3.33	3.69
Temperature difference	0.11	0.03	0.06	0.17
Diserpsal mode - Aquatic	-2.07	0.16	-2.39	-1.75
Diserpsal mode - Terrestrial	-0.54	0.14	-0.81	-0.27
Temperature difference x Diserpsal mode - Aquatic	0.81	0.21	0.40	1.23
Temperature difference x Diserpsal mode - Terrestrial	0.01	0.04	-0.07	0.09

Table S9 Mean parameter estimates, estimate error and 95% credible intervals for the interaction between dispersal mode (aerial, aquatic, terrestrial) and precipitation (rainfall) difference between range core and edge.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Intercept, β_0	33.57	2.91	27.83	39.27
Rainfall difference	0.02	0.02	-0.01	0.05
Diserpsal mode - Aquatic	-13.37	4.18	-21.57	-5.12
Diserpsal mode - Terrestrial	-7.87	3.53	-14.80	-0.88
Rainfall difference x Diserpsal mode - Aquatic	-0.04	0.09	-0.22	0.13
Rainfall difference x Diserpsal mode - Terrestrial	-0.02	0.02	-0.05	0.01

Table S10 Mean parameter estimates, estimate error, and 95% credible intervals for the interaction between dispersal mode (aerial, aquatic, terrestrial) and time since divergence between the core range and dispersal front. Random effects include the standard deviations (σ) for study-level observations (study ID), and species identity.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Intercept	-2.92	0.57	-4.04	-1.8
σ Intercept	-0.15	0.14	-0.41	0.14
Time	0.3	0.18	-0.06	0.65
Diserpsal mode - Aquatic	0.24	0.65	-1.05	1.5
Diserpsal mode - Terrestrial	-0.31	0.71	-1.71	1.1
Time x Diserpsal mode - Aquatic	0.01	0.28	-0.54	0.55
Time x Diserpsal mode - Terrestrial	0.08	0.2	-0.31	0.48
σ Time	0.07	0.04	-0.02	0.14
Random effects				
σΔstudy ID	0.86	0.16	0.58	1.21
$\sigma\Delta$ species	0.37	0.27	0.02	1

Table S11 Mean parameter estimates, estimate error ,and 95% credible intervals for the population range expansion analysis, which includes the intercept, and sampling error with the publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), species identity, and phylogenetic relatedness.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Intercept	0.25	0.15	-0.02	0.52
Sampling error	-0.27	0.17	-0.61	0.05
Publication year	0.01	0.02	-0.03	0.04
Random effects				
$\sigma\Delta$ effect size ID	0.24	0.01	0.22	0.27
σ∆study ID	0.39	0.06	0.29	0.52
$\sigma\Delta$ species	0.09	0.07	0	0.27
$\sigma\Delta$ phylogeny	0.13	0.12	0.01	0.42

Table S12 Mean parameter estimates, estimate error, and 95% credible intervals for the population range expansion analysis, which includes fixed-effect parameters for all extracted traits with >5 effect sizes, sex, and sampling error with the publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), species identity, and phylogenetic relatedness.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Activity	0.24	0.19	-0.14	0.65
Cardiovascular	0.10	0.21	-0.31	0.53
Condition	-0.09	0.24	-0.59	0.36
Hormone	0.60	0.23	0.16	1.05
Immunity	0.57	0.23	0.11	1.04
Locomotor capacity	0.25	0.19	-0.13	0.64
Metabolism	0.55	0.20	0.15	0.95
Thermal tolerance	0.33	0.20	-0.065	0.74
Male	-0.04	0.05	-0.15	0.06
Mixed	-0.19	0.08	-0.35	-0.03
Sampling error	-0.002	0.20	-0.39	0.37
Publication year	0.0009	0.02	-0.04	0.04
Random effects				
σΔeffect size ID	0.23	0.01	0.21	0.26
σ∆study ID	0.45	0.07	0.33	0.6
σΔspecies	0.1	0.08	0	0.31
$\sigma\Delta$ phylogeny	0.16	0.14	0.01	0.51

Table S13 Mean parameter estimates, estimate error, and 95% credible intervals for the population range expansion analysis, which includes fixed-effect parameters for all extracted taxa with >5 effect sizes, with sampling error and the publication year to account for sampling and publication bias. Random effects include the standard deviations (σ) for individual-level observations (effect size ID), study-level observations (study ID), and species identity.

Parameters	Estimate	Est.Error	Q2.5	Q97.5
Fixed effects				
Amphibia	0.29	0.14	0.002	0.57
Aves	0.53	0.21	0.10	0.94
Invertebrate	0.22	0.15	-0.07	0.51
Osteichthyes	0.23	0.23	-0.21	0.69
Male	-0.04	0.06	-0.15	0.07
Mixed	-0.15	0.08	-0.31	0.01
Sampling error	-0.13	0.20	-0.52	0.26
Publication year	0.01	0.02	-0.03	0.05
Random effects				
$\sigma\Delta$ effect size ID	0.24	0.01	0.21	0.27
σΔstudy ID	0.39	0.06	0.29	0.52
$\sigma\Delta$ species	0.1	0.08	0	0.31

Supplementary Figures

Publications extracted until 2021 (n = number of papers or species, k = number of effect size)

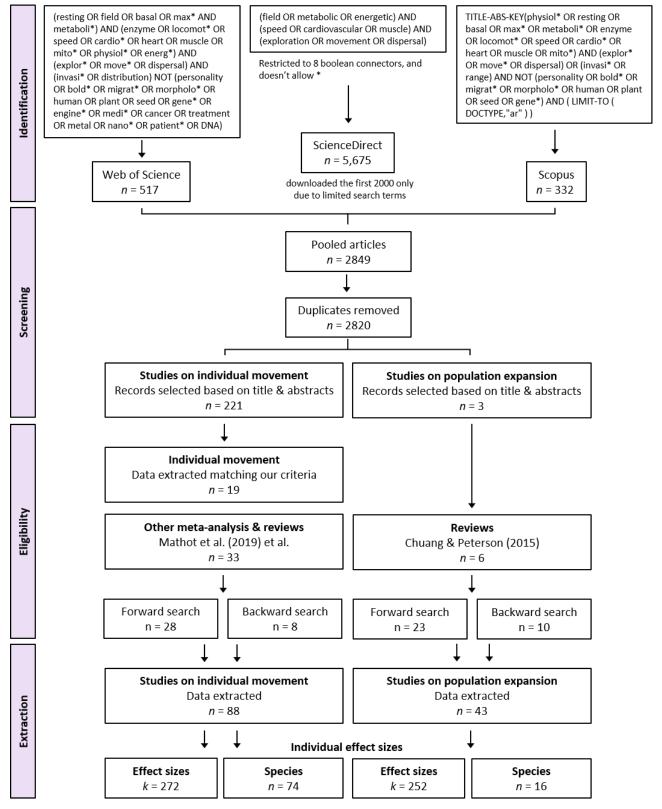


Figure S1 PRISMA flow chart for the data-collection process. Search terms for each online database are shown. n = number of papers remaining after each stage of selection; k = number of effect sizes. Studies on individual movement include among-individual correlation of a physiological trait with activity, exploration, or dispersal. Studies on population expansion include comparison of physiology between population range core and edge.

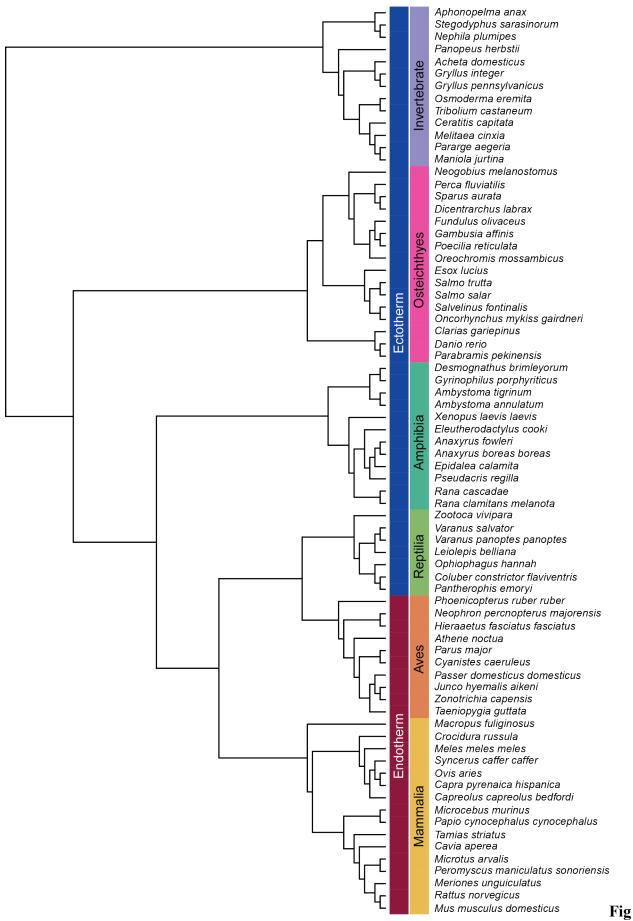


Figure S2 Phylogenetic reconstruction for individual movement studies from the Open Tree of Life (https://tree.opentreeoflife.org/). The phylogeny was converted to a correlation matrix for the phylogenetic multilevel model.

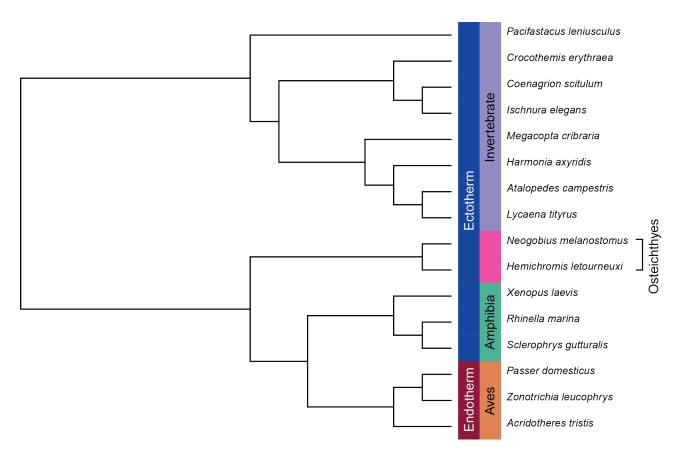


Figure S3 Phylogenetic reconstruction for population range expansion studies from the Open Tree of Life (https://tree.opentreeoflife.org/). The phylogeny was converted to a correlation matrix for the phylogenetic multilevel model.

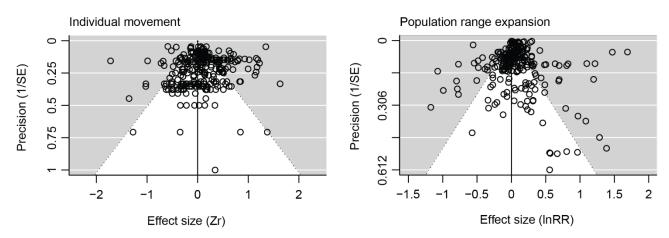


Figure S4 Funnel plots for the individual movement data (left) and population range expansion data (right). The white area bordered by dashed lines represents the region of 95% pseudo confidence intervals where 95% of studies are expected to fall in the absence of bias and heterogeneity.

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