

Supporting Information

Figure S1. An overview of the sample sizes used for each analysis; the area of each circle is proportional to the number of species in that category. (A) The full list of extant mammals included in the database (5420, grey), the subset that were categorized as definitively migratory or non-migratory (1062, black), and the subset of those that were migratory (235, red). (B) Of those 235 migrants, 180 were classified as breeding (B, purple), refuge (R, orange), tracking (T, green), or a combination thereof; 25 species were classified as two types and included in analyses for each type. (C) The subset of species (965) with both definitive movement and phylogenetic data; of those, 158 were migrants. (D) The subset of species (605) with both definitive movement and life history data; of those, 113 were migrants.

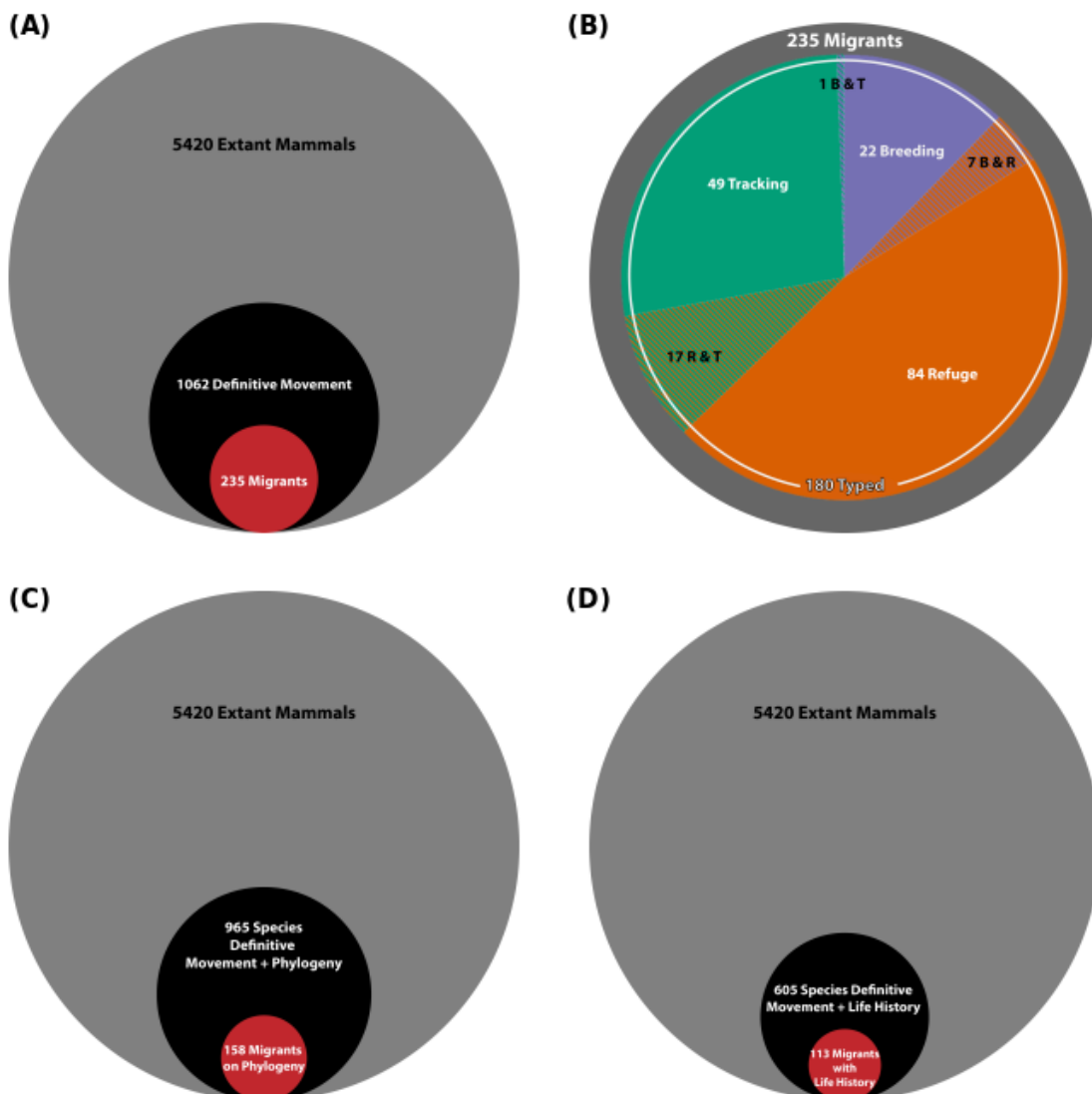


Figure S2. A circular phylogeny of all mammals. Tree branches colors correspond to knowledge of movement patterns: definitively migratory or non-migratory (black; shown in Fig. 1), unclear movement (grey), and data deficient (red). Markers around the outside of the phylogeny indicate breeding (purple), refuge (orange), and tracking (green) migrants, whereas black markers indicate species definitely migrating whose categorization is unclear due to a lack of data. The outer-most ring indicates broad taxonomic groups within the mammals.

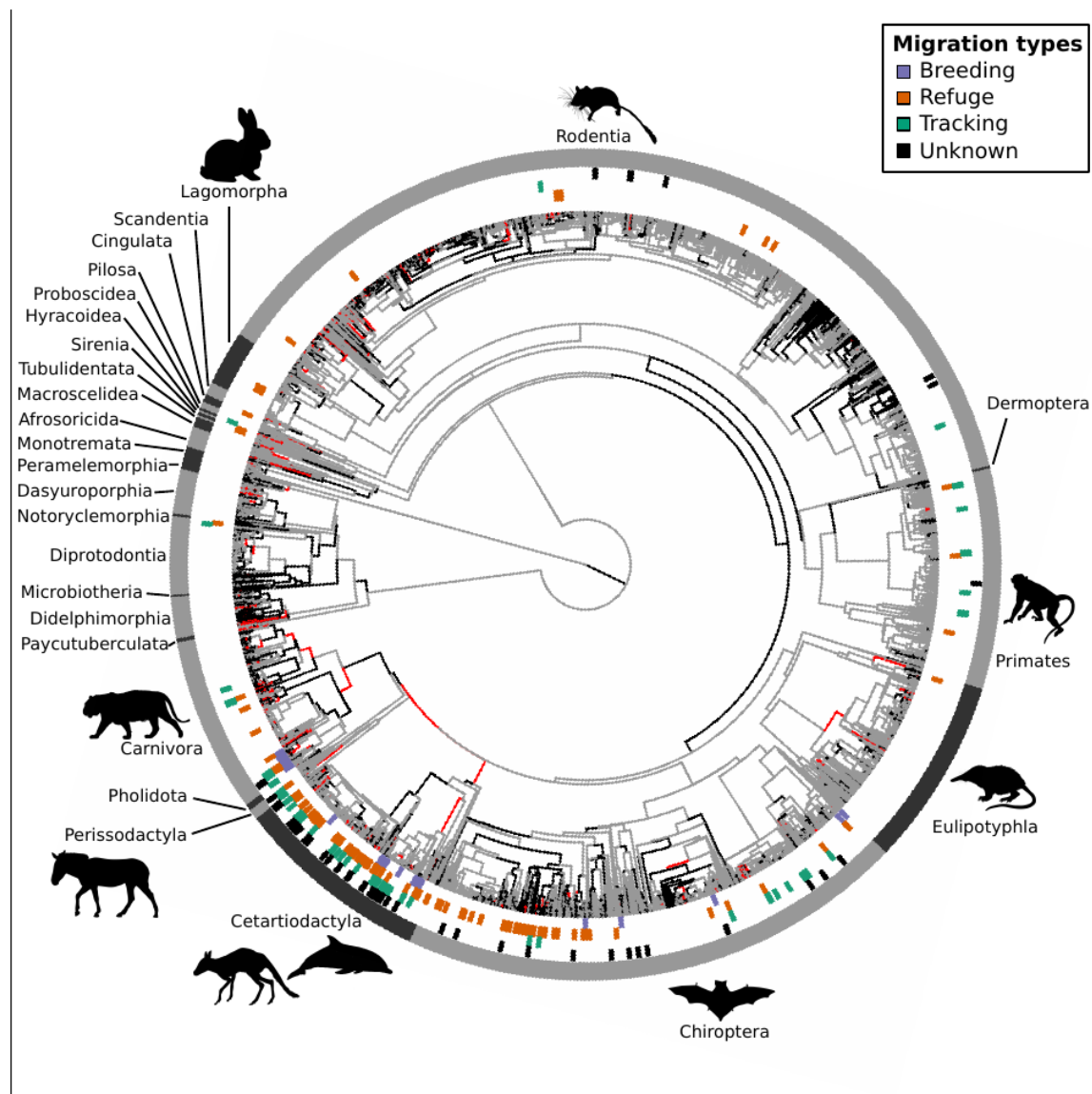


Table S1. The order, family, genus and species of all (235) migratory species considered in our analysis, along with their migration type (B=breeding, R=refuge, T=tracking, R=refuge, U=unknown type), and the ID numbers of the references used to make these classifications (references listed at end). See Dryad repository (DOI: XXX) for data on all mammals species.

Order	Family	Genus	Species	Type	Ref ID
CARNIVORA	CANIDAE	<i>Canis</i>	<i>lupus</i>	T	537; 3642
	CANIDAE	<i>Cerdocyon</i>	<i>thous</i>	R	3497
	FELIDAE	<i>Acinonyx</i>	<i>jubatus</i>	T	3463; 3464
	FELIDAE	<i>Panthera</i>	<i>uncia</i>	RT	1230; 3485
	FELIDAE	<i>Puma</i>	<i>concolor</i>	T	100; 529
	MUSTELIDAE	<i>Pteronura</i>	<i>brasiliensis</i>	R	3461
	ODOBENIDAE	<i>Odobenus</i>	<i>rosmarus</i>	BR	40; 60; 72
	OTARIIDAE	<i>Callorhinus</i>	<i>ursinus</i>	B	38
	OTARIIDAE	<i>Eumetopias</i>	<i>jubatus</i>	B	39
	OTARIIDAE	<i>Neophoca</i>	<i>cinerea</i>	B	41; 42
	OTARIIDAE	<i>Zalophus</i>	<i>californianus</i>	B	45
	PHOCIDAE	<i>Cystophora</i>	<i>cristata</i>	BR	49
	PHOCIDAE	<i>Histriophoca</i>	<i>fasciata</i>	U	58
	PHOCIDAE	<i>Lobodon</i>	<i>carcinophaga</i>	RT	56; 3645
	PHOCIDAE	<i>Mirounga</i>	<i>angustirostris</i>	BR	57
	PHOCIDAE	<i>Mirounga</i>	<i>leonina</i>	T	65
	PHOCIDAE	<i>Ommatophoca</i>	<i>rossii</i>	BR	67
	PHOCIDAE	<i>Pagophilus</i>	<i>groenlandicus</i>	BT	54; 55
	PHOCIDAE	<i>Phoca</i>	<i>largha</i>	U	68; 69
	PHOCIDAE	<i>Pusa</i>	<i>casgica</i>	B	40; 50
	PHOCIDAE	<i>Pusa</i>	<i>sibirica</i>	U	40; 53
URSIDAE	<i>Ursus</i>	<i>arctos</i>	T	40; 100	
URSIDAE	<i>Ursus</i>	<i>maritimus</i>	T	100; 511; 512	
URSIDAE	<i>Ursus</i>	<i>thibetanus</i>	T	1231	
CETARTIODA CTYLA	ANTILOCAPRIDAE	<i>Antilocapra</i>	<i>americana</i>	R	1208
	BALAENIDAE	<i>Balaena</i>	<i>mysticetus</i>	RT	6; 14; 16
	BALAENIDAE	<i>Eubalaena</i>	<i>australis</i>	B	7; 8; 13
	BALAENIDAE	<i>Eubalaena</i>	<i>glacialis</i>	B	13
	BALAENIDAE	<i>Eubalaena</i>	<i>japonica</i>	B	13
	BALAENOPTERIDAE	<i>Balaenoptera</i>	<i>acutorostrata</i>	B	6; 37
	BALAENOPTERIDAE	<i>Balaenoptera</i>	<i>bonaerensis</i>	B	6; 11
	BALAENOPTERIDAE	<i>Balaenoptera</i>	<i>borealis</i>	B	5; 6

CETARTIODA CTYLA	BALAENOPTERIDAE	<i>Balaenoptera</i>	<i>musculus</i>	B	1; 2; 356; 358
	BALAENOPTERIDAE	<i>Balaenoptera</i>	<i>physalus</i>	B	1; 6
	BALAENOPTERIDAE	<i>Megaptera</i>	<i>novaeangliae</i>	B	3; 4; 6; 9; 10; 16
	BOVIDAE	<i>Addax</i>	<i>nasomaculatus</i>	T	1206
	BOVIDAE	<i>Alcelaphus</i>	<i>buselaphus</i>	U	186; 3259
	BOVIDAE	<i>Ammotragus</i>	<i>lervia</i>	U	40; 1221
	BOVIDAE	<i>Bison</i>	<i>bison</i>	RT	1209
	BOVIDAE	<i>Bos</i>	<i>javanicus</i>	T	40
	BOVIDAE	<i>Bos</i>	<i>mutus</i>	U	1235
	BOVIDAE	<i>Budorcas</i>	<i>taxicolor</i>	T	3304
	BOVIDAE	<i>Capra</i>	<i>cylindricornis</i>	R	3307
	BOVIDAE	<i>Capra</i>	<i>falconeri</i>	R	40; 3654
	BOVIDAE	<i>Capra</i>	<i>sibirica</i>	R	1210
	BOVIDAE	<i>Connochaetes</i>	<i>gnou</i>	T	186; 1205
	BOVIDAE	<i>Connochaetes</i>	<i>taurinus</i>	T	186; 1203
	BOVIDAE	<i>Damaliscus</i>	<i>lunatus</i>	RT	186
	BOVIDAE	<i>Eudorcas</i>	<i>thomsonii</i>	RT	40
	BOVIDAE	<i>Gazella</i>	<i>cuvieri</i>	U	40; 1222
	BOVIDAE	<i>Gazella</i>	<i>dorcas</i>	T	1211
	BOVIDAE	<i>Gazella</i>	<i>gazella</i>	T	40; 3321
	BOVIDAE	<i>Gazella</i>	<i>subgutturosa</i>	R	1212
	BOVIDAE	<i>Hemitragus</i>	<i>jemlahicus</i>	U	40; 1223
	BOVIDAE	<i>Kobus</i>	<i>kob</i>	R	3327; 3328
	BOVIDAE	<i>Kobus</i>	<i>leche</i>	R	186
	BOVIDAE	<i>Naemorhedus</i>	<i>goral</i>	U	40; 3370; 3643
	BOVIDAE	<i>Nanger</i>	<i>granti</i>	U	1226
	BOVIDAE	<i>Oreamnos</i>	<i>americanus</i>	RT	1215
	BOVIDAE	<i>Oryx</i>	<i>gazella</i>	U	186; 3259
	BOVIDAE	<i>Ovibos</i>	<i>moschatus</i>	U	3380; 3646
	BOVIDAE	<i>Ovis</i>	<i>ammon</i>	R	1216
	BOVIDAE	<i>Ovis</i>	<i>canadensis</i>	R	1217
	BOVIDAE	<i>Ovis</i>	<i>dalli</i>	R	1218
	BOVIDAE	<i>Pantholops</i>	<i>hodgsonii</i>	B	1219
	BOVIDAE	<i>Procapra</i>	<i>gutturosa</i>	T	1201
	BOVIDAE	<i>Pseudois</i>	<i>nayaur</i>	R	3391; 3392
	BOVIDAE	<i>Rupicapra</i>	<i>pyrenaica</i>	R	3273

CETARTIODA CTYLA	BOVIDAE	<i>Rupicapra</i>	<i>rupicapra</i>	U	3259; 3274
	BOVIDAE	<i>Saiga</i>	<i>tatarica</i>	R	1200
	BOVIDAE	<i>Syncerus</i>	<i>caffer</i>	U	186
	BOVIDAE	<i>Tragelaphus</i>	<i>oryx</i>	T	186
	BOVIDAE	<i>Tragelaphus</i>	<i>scriptus</i>	R	186
	BOVIDAE	<i>Tragelaphus</i>	<i>spekii</i>	R	186
	CAMELIDAE	<i>Camelus</i>	<i>ferus</i>	R	3305; 3653
	CAMELIDAE	<i>Lama</i>	<i>guanicoe</i>	R	3330
	CERVIDAE	<i>Alces</i>	<i>alces</i>	R	1207
	CERVIDAE	<i>Alces</i>	<i>americanus</i>	T	3260
	CERVIDAE	<i>Axis</i>	<i>porcinus</i>	R	3267
	CERVIDAE	<i>Blastocerus</i>	<i>dichotomus</i>	R	3297
	CERVIDAE	<i>Capreolus</i>	<i>capreolus</i>	R	3272
	CERVIDAE	<i>Capreolus</i>	<i>pygargus</i>	R	1202
	CERVIDAE	<i>Cervus</i>	<i>elaphus</i>	U	40
	CERVIDAE	<i>Cervus</i>	<i>nippon</i>	R	3312
	CERVIDAE	<i>Dama</i>	<i>dama</i>	R	186; 3314
	CERVIDAE	<i>Hippocamelus</i>	<i>antisensis</i>	U	40
	CERVIDAE	<i>Hippocamelus</i>	<i>bisulcus</i>	U	1224
	CERVIDAE	<i>Odocoileus</i>	<i>hemionus</i>	RT	1213
	CERVIDAE	<i>Odocoileus</i>	<i>virginianus</i>	R	1214
	CERVIDAE	<i>Przewalskium</i>	<i>albirostris</i>	U	3390
	CERVIDAE	<i>Rangifer</i>	<i>tarandus</i>	R	40; 1220
	CERVIDAE	<i>Rucervus</i>	<i>eldii</i>	T	1228
	DELPHINIDAE	<i>Cephalorhynchus</i>	<i>commersonii</i>	U	26; 33
	DELPHINIDAE	<i>Delphinus</i>	<i>delphis</i>	T	21; 33
	DELPHINIDAE	<i>Globicephala</i>	<i>macrorhynchus</i>	T	19; 33
	DELPHINIDAE	<i>Globicephala</i>	<i>melas</i>	T	19; 33
	DELPHINIDAE	<i>Grampus</i>	<i>griseus</i>	B	33
	DELPHINIDAE	<i>Lagenorhynchus</i>	<i>acutus</i>	T	33
	DELPHINIDAE	<i>Lagenorhynchus</i>	<i>albirostris</i>	U	33
	DELPHINIDAE	<i>Lagenorhynchus</i>	<i>australis</i>	B	33
	DELPHINIDAE	<i>Lagenorhynchus</i>	<i>obliquidens</i>	T	33
	DELPHINIDAE	<i>Lagenorhynchus</i>	<i>obscurus</i>	T	33
	DELPHINIDAE	<i>Lissodelphis</i>	<i>borealis</i>	U	33
	DELPHINIDAE	<i>Orcaella</i>	<i>brevirostris</i>	RT	33
	DELPHINIDAE	<i>Orcinus</i>	<i>orca</i>	T	22; 23; 24; 25; 33

CETARTIODA CTYLA	DELPHINIDAE	<i>Sotalia</i>	<i>fluviatilis</i>	R	33; 3649
	DELPHINIDAE	<i>Sousa</i>	<i>chinensis</i>	T	33; 3651
	DELPHINIDAE	<i>Sousa</i>	<i>teuszii</i>	U	33
	DELPHINIDAE	<i>Stenella</i>	<i>attenuata</i>	U	33
	DELPHINIDAE	<i>Stenella</i>	<i>coeruleoalba</i>	U	33
	DELPHINIDAE	<i>Stenella</i>	<i>longirostris</i>	U	33
	DELPHINIDAE	<i>Tursiops</i>	<i>truncatus</i>	U	27; 33
	ESCHRICHTIIDAE	<i>Eschrichtius</i>	<i>robustus</i>	B	6; 16
	HIPPOPOTAMIDAE	<i>Hippopotamus</i>	<i>amphibius</i>	R	186
	INIIDAE	<i>Inia</i>	<i>geoffrensis</i>	T	28
	MONODONTIDAE	<i>Delphinapterus</i>	<i>leucas</i>	R	6; 17; 533
	MONODONTIDAE	<i>Monodon</i>	<i>monoceros</i>	BR	16; 20; 33
	PHOCOENIDAE	<i>Neophocaena</i>	<i>phocaenoides</i>	U	33
	PHOCOENIDAE	<i>Phocoena</i>	<i>phocoena</i>	R	16; 29; 33
	PHOCOENIDAE	<i>Phocoena</i>	<i>spinipinnis</i>	U	33
	PHOCOENIDAE	<i>Phocoenoides</i>	<i>dalli</i>	B	30; 33
	PHYSETERIDAE	<i>Physeter</i>	<i>macrocephalus</i>	T	6; 15; 16
	PLATANISTIDAE	<i>Platanista</i>	<i>gangetica</i>	T	6; 40
	SUIDAE	<i>Sus</i>	<i>barbatus</i>	R	40; 1233
	ZIPHIIDAE	<i>Berardius</i>	<i>arnuxii</i>	R	32; 33
ZIPHIIDAE	<i>Berardius</i>	<i>bairdii</i>	U	33	
ZIPHIIDAE	<i>Hyperoodon</i>	<i>ampullatus</i>	U	6; 18; 33	
CHIROPTERA	EMBALLONURIDAE	<i>Diclidurus</i>	<i>albus</i>	U	189; 191; 199; 319; 3644
	EMBALLONURIDAE	<i>Taphozous</i>	<i>melanopogon</i>	B	145; 175
	EMBALLONURIDAE	<i>Taphozous</i>	<i>nudiventris</i>	U	132; 145
	HIPPOSIDERIDAE	<i>Asellia</i>	<i>tridens</i>	U	132; 145
	HIPPOSIDERIDAE	<i>Hipposideros</i>	<i>armiger</i>	U	145; 200; 208; 216; 222
	HIPPOSIDERIDAE	<i>Hipposideros</i>	<i>commersoni</i>	RT	145; 167; 235; 3648
	HIPPOSIDERIDAE	<i>Hipposideros</i>	<i>lankadiva</i>	U	145; 216; 239; 251
	MOLOSSIDAE	<i>Nyctinomops</i>	<i>macrotis</i>	R	100; 189; 191
	MOLOSSIDAE	<i>Tadarida</i>	<i>australis</i>	RT	134; 257
	MOLOSSIDAE	<i>Tadarida</i>	<i>brasiliensis</i>	U	100; 106; 120

CHIROPTERA	NATALIDAE	<i>Natalus</i>	<i>stramineus</i>	R	314; 356
	NYCTERIDAE	<i>Nycteris</i>	<i>thebaica</i>	U	102; 186; 206
	PHYLLOSTOMIDAE	<i>Choeronycteris</i>	<i>mexicana</i>	RT	100
	PHYLLOSTOMIDAE	<i>Leptonycteris</i>	<i>curasoae</i>	B	361; 362
	PHYLLOSTOMIDAE	<i>Leptonycteris</i>	<i>nivalis</i>	U	100; 120; 363
	PHYLLOSTOMIDAE	<i>Leptonycteris</i>	<i>yerbabuena</i>	U	100; 101; 364; 387
	PHYLLOSTOMIDAE	<i>Sturnira</i>	<i>lilium</i>	R	189; 191; 356; 404
	PTEROPODIDAE	<i>Eidolon</i>	<i>helvum</i>	T	102; 103; 107; 108; 145; 216
	PTEROPODIDAE	<i>Myonycteris</i>	<i>torquata</i>	T	102; 103; 112; 113; 186
	PTEROPODIDAE	<i>Nanonycteris</i>	<i>veldkampii</i>	T	145; 153
	PTEROPODIDAE	<i>Pteropus</i>	<i>conspicillatus</i>	T	103; 134; 136; 186
	PTEROPODIDAE	<i>Pteropus</i>	<i>poliocephalus</i>	T	103; 134; 137; 138; 145
	PTEROPODIDAE	<i>Pteropus</i>	<i>scapulatus</i>	U	134; 139; 145
	PTEROPODIDAE	<i>Rousettus</i>	<i>aegyptiacus</i>	RT	103; 109; 145
	RHINOLOPHIDAE	<i>Rhinolophus</i>	<i>euryale</i>	BR	145; 217; 250
	RHINOLOPHIDAE	<i>Rhinolophus</i>	<i>mehelyi</i>	R	145; 217
	RHINOPOMATIDAE	<i>Rhinopoma</i>	<i>microphyllum</i>	B	132; 170; 175; 182; 185; 207
	VESPERTILIONIDAE	<i>Antrozous</i>	<i>pallidus</i>	R	100
	VESPERTILIONIDAE	<i>Chalinolobus</i>	<i>morio</i>	R	134
	VESPERTILIONIDAE	<i>Eptesicus</i>	<i>fuscus</i>	R	100; 104; 189; 191; 356
VESPERTILIONIDAE	<i>Eptesicus</i>	<i>nasutus</i>	U	132; 420	
VESPERTILIONIDAE	<i>Lasionycteris</i>	<i>noctivagans</i>	R	100; 120	
VESPERTILIONIDAE	<i>Lasiurus</i>	<i>blossevillii</i>	R	105; 120; 356; 434	

CHIROPTERA	VESPERTILIONIDAE	<i>Lasiurus</i>	<i>borealis</i>	R	100; 435
	VESPERTILIONIDAE	<i>Lasiurus</i>	<i>cinereus</i>	R	22; 33; 100; 189; 190; 191
	VESPERTILIONIDAE	<i>Lasiurus</i>	<i>intermedius</i>	U	100
	VESPERTILIONIDAE	<i>Miniopterus</i>	<i>fuliginosus</i>	U	200; 208; 420; 455
	VESPERTILIONIDAE	<i>Miniopterus</i>	<i>natalensis</i>	R	102
	VESPERTILIONIDAE	<i>Miniopterus</i>	<i>oceanensis</i>	BR	134; 161; 243; 267
	VESPERTILIONIDAE	<i>Miniopterus</i>	<i>schreibersii</i>	R	250; 288; 454
	VESPERTILIONIDAE	<i>Myotis</i>	<i>auriculus</i>	R	100; 356
	VESPERTILIONIDAE	<i>Myotis</i>	<i>austroriparius</i>	R	100; 3655
	VESPERTILIONIDAE	<i>Myotis</i>	<i>blythii</i>	U	208; 288; 420
	VESPERTILIONIDAE	<i>Myotis</i>	<i>brandtii</i>	R	208; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>capaccinii</i>	R	288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>dasycneme</i>	R	208; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>daubentonii</i>	R	208; 250; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>emarginatus</i>	RT	250; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>grisescens</i>	R	100
	VESPERTILIONIDAE	<i>Myotis</i>	<i>lucifugus</i>	R	100; 475
	VESPERTILIONIDAE	<i>Myotis</i>	<i>myotis</i>	RT	250; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>mystacinus</i>	R	250; 288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>nattereri</i>	R	288
	VESPERTILIONIDAE	<i>Myotis</i>	<i>septentrionalis</i>	R	100; 3657
	VESPERTILIONIDAE	<i>Myotis</i>	<i>sodalis</i>	R	100; 418; 510
	VESPERTILIONIDAE	<i>Myotis</i>	<i>velifer</i>	R	100; 356
	VESPERTILIONIDAE	<i>Nyctalus</i>	<i>lasiopterus</i>	U	288
	VESPERTILIONIDAE	<i>Nyctalus</i>	<i>leisleri</i>	R	213; 214; 250; 288
	VESPERTILIONIDAE	<i>Nyctalus</i>	<i>noctula</i>	R	200; 208; 288; 420
	VESPERTILIONIDAE	<i>Nycticeius</i>	<i>humeralis</i>	R	100; 145
	VESPERTILIONIDAE	<i>Pipistrellus</i>	<i>hesperus</i>	R	100; 120
	VESPERTILIONIDAE	<i>Pipistrellus</i>	<i>nathusii</i>	R	288; 509
	VESPERTILIONIDAE	<i>Pipistrellus</i>	<i>subflavus</i>	R	100; 120;

					356
CHIROPTERA	VESPERTILIONIDAE	<i>Vespertilio</i>	<i>murinus</i>	R	208; 250; 288; 420
CINGULATA	DASYPODIDAE	<i>Dasyopus</i>	<i>novemcinctus</i>	R	1949
DIPROTODONTIA	BURRAMYIDAE	<i>Burramys</i>	<i>parvus</i>	R	2048
	BURRAMYIDAE	<i>Cercartetus</i>	<i>lepidus</i>	T	2051
EULIPOTYPHILA	TALPIDAE	<i>Scapanus</i>	<i>townsendii</i>	R	2846
LAGOMORPHA	LEPORIDAE	<i>Lepus</i>	<i>californicus</i>	R	2254; 3641
	LEPORIDAE	<i>Lepus</i>	<i>timidus</i>	R	2249; 3640
PERISSODACTYLA	EQUIDAE	<i>Equus</i>	<i>grevyi</i>	T	542; 543
	EQUIDAE	<i>Equus</i>	<i>quagga</i>	T	40; 97; 98; 99; 538; 539; 543
	EQUIDAE	<i>Equus</i>	<i>zebra</i>	T	40; 538; 543
	RHINOCEROTIDAE	<i>Ceratotherium</i>	<i>simum</i>	RT	40; 92; 3647
	RHINOCEROTIDAE	<i>Dicerorhinus</i>	<i>sumatrensis</i>	U	40
PRIMATES	ATELIDAE	<i>Lagothrix</i>	<i>cana</i>	T	3074
	CEBIDAE	<i>Cebus</i>	<i>albifrons</i>	U	3046; 3074
	CERCOPITHECIDAE	<i>Macaca</i>	<i>fuscata</i>	T	3162; 3209
	CERCOPITHECIDAE	<i>Rhinopithecus</i>	<i>bieti</i>	RT	3210
	CERCOPITHECIDAE	<i>Rhinopithecus</i>	<i>roxellana</i>	T	3208
	HOMINIDAE	<i>Gorilla</i>	<i>beringei</i>	T	1232; 2878
	HOMINIDAE	<i>Homo</i>	<i>sapiens</i>	R	3658
	HOMINIDAE	<i>Pongo</i>	<i>pygmaeus</i>	T	3254
	LEMURIDAE	<i>Eulemur</i>	<i>fulvus</i>	R	2929
	LEMURIDAE	<i>Eulemur</i>	<i>rufifrons</i>	T	2933
	PITHECIIDAE	<i>Cacajao</i>	<i>calvus</i>	T	3075; 3076
PITHECIIDAE	<i>Cacajao</i>	<i>melanocephalus</i>	T	3075	
PROBOSCIDEA	ELEPHANTIDAE	<i>Loxodonta</i>	<i>africana</i>	T	2185; 2186
RODENTIA	CRICETIDAE	<i>Eolagurus</i>	<i>luteus</i>	T	208
	CRICETIDAE	<i>Lemmus</i>	<i>lemmus</i>	U	217
	CRICETIDAE	<i>Lemmus</i>	<i>sibiricus</i>	U	1408
	CRICETIDAE	<i>Microtus</i>	<i>fortis</i>	R	208; 1446
	CRICETIDAE	<i>Microtus</i>	<i>montanus</i>	R	1027; 1429
	CRICETIDAE	<i>Peromyscus</i>	<i>maniculatus</i>	R	668; 1027

RODENTIA	HETEROMYIDAE	<i>Dipodomys</i>	<i>ordii</i>	R	668; 1027
	MURIDAE	<i>Apodemus</i>	<i>flavicollis</i>	U	217; 779
	MURIDAE	<i>Apodemus</i>	<i>sylvaticus</i>	U	217; 778; 779
	MURIDAE	<i>Meriones</i>	<i>libycus</i>	U	208
	MURIDAE	<i>Micromys</i>	<i>minutus</i>	R	200; 208; 1251
	MURIDAE	<i>Otomys</i>	<i>angoniensis</i>	U	186; 1005
	MURIDAE	<i>Rattus</i>	<i>colletti</i>	R	134; 1253
	MURIDAE	<i>Rattus</i>	<i>rattus</i>	R	190; 1254
	SCIURIDAE	<i>Sciurus</i>	<i>arizonensis</i>	U	1741
	SCIURIDAE	<i>Sciurus</i>	<i>niger</i>	U	668; 1748
	SCIURIDAE	<i>Tamias</i>	<i>dorsalis</i>	T	1114
SIRENIA	DUGONGIDAE	<i>Dugong</i>	<i>dugon</i>	R	75; 78; 79
	TRICHECHIDAE	<i>Trichechus</i>	<i>inunguis</i>	R	75; 76
	TRICHECHIDAE	<i>Trichechus</i>	<i>manatus</i>	R	75

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Table S2. *D* (Fritz & Purvis 2010) values for migration and each of the migration types (breeding, refuge, tracking) across only migrating species. The mean and standard deviation values of *D* across all bootstrap phylogenies are given. The last four rows indicate the percentage of times that a significant *p*-value was detected (at the level indicated) in all phylogenies. E.g. 100% for $P(D=0) \leq 0.01$ means in all bootstraps, the *D* values for those migration types were significantly not zero at $\alpha=0.01$ (two-tailed; bootstrapped). The interpretation of *D* values is described in the Methods section.

	Migration (overall)	Breeding (among migrants)	Refuge (among migrants)	Tracking (among migrants)
<i>D</i> - mean	0.42	0.02	0.40	0.46
<i>D</i> - SD	0.01	0.04	0.02	0.02
# models successful	666	666	666	666
% $P(D=0) \leq 0.05$	100.00	0.00	99.70	100.00
% $P(D=0) \leq 0.01$	100.00	0.00	7.96	62.93
% $P(D=1) \leq 0.05$	100.00	100.00	100.00	100.00
% $P(D=1) \leq 0.01$	100.00	100.00	100.00	100.00

Table S3. Number of migratory species by each locomotion and migration type, see also Figure S1.

	Walking	Swimming	Flying	Total
Breeding	1	24	5	30
Refuge	51	16	41	108
Tracking	40	16	11	67
Total	92	56	57	205

Table S4. The fraction of shared branch length (Shared) and the corresponding standard effect size (SES) for each migration type by locomotion type. The contents are described in the Methods section.

	Walking		Swimming		Flying	
	Shared	SES	Shared	SES	Shared	SES
Breeding	0.045	-3.560	0.483	3.118	0.290	0.399
Refuge	0.248	0.576	0.275	-2.098	0.397	1.338
Tracking	0.201	0.802	0.323	-0.290	0.231	-2.953

Table S5. Phylogenetic logistic regressions (Ho & Ané 2014) of migration across all mammals for which we had sufficient data (158 migrants, 965 mammal species in total). All categorical variable estimates, bar the intercept, are statistical contrasts: they are the difference in overall mean for that group from the overall mean of a species with one habitat-type, that is herbivorous, that has one dietary category, and walks. The regression was run two ways: once with IUCN Red List as numeric, and once as binary (threatened vs not). Estimates are reported as means and standard deviations across the 984 bootstrap phylogenies for the first regression and 989 for the second; models in 16 (and 11, respectively) phylogenies had numerical fitting problems (as reported by *phylolm*; Ho & Ané 2014) and their estimates are not reported. The mean and standard deviation of the α parameter, which reports the estimated error due to phylogenetic co-variation, was 0.0124 ± 0.00263 for the first regression and 0.01 ± 0.0027 for the second. Asterisks (*) indicate factors significant at the 0.05 level; for the presentation of coefficients and their associated standard deviations we follow the advice of Cole (2015), while for *p*-values we present at three decimal places following the defaults in the R (R Core Team 2014) statistical software. Coefficients with large standard errors and/or non-significant average *p*-values suggest explanatory variables for which there is little evidence of an association with migration.

	Red List category as numeric			Red List category as binary factor		
	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)
Intercept (1 habitat, herbivore, 1 dietary category,	-5.98 \pm 0.9265	1.07 \pm 0.0961	0 \pm 0			

<i>walking</i>) *						
Intercept (1 habitat, herbivore, 1 dietary category, walking, not Red Listed as threatened) *				-5.89 ± 0.9272	1.06 ± 0.0966	<0.0001 ± 0.0002
log10(body mass) *	1.19 ± 0.1559	0.18 ± 0.0191	0 ± 0	1.16 ± 0.157	0.17 ± 0.0191	<0.0001 ± <0.0001
2 habitat	0.14 ± 0.1137	0.25 ± 0.0226	0.528 ± 0.1702	0.12 ± 0.1078	0.25 ± 0.0217	0.5717 ± 0.1758
3 habitats	-14.46 ± 0.3017	741.01 ± 152.1249	0.983 ± 0.007	-14.5 ± 0.2942	719.68 ± 148.2678	0.9826 ± 0.0066
4 habitats	-13.89 ± 0.3017	3546.95 ± 797.7353	0.997 ± 0.0017	-13.83 ± 0.2942	3375.24 ± 766.3315	0.9964 ± 0.0016
Omnivore	0.6 ± 0.0965	0.35 ± 0.0338	0.098 ± 0.0502	0.61 ± 0.0963	0.34 ± 0.0335	0.0864 ± 0.0469
Carnivore	0.41 ± 0.2105	0.4 ± 0.0254	0.313 ± 0.1943	0.36 ± 0.1927	0.4 ± 0.0248	0.3586 ± 0.1847
2 dietary categories	0.51 ± 0.1048	0.3 ± 0.0193	0.113 ± 0.1019	0.46 ± 0.1105	0.3 ± 0.0192	0.1494 ± 0.1258
3 dietary categories	0.32 ± 0.1121	0.39 ± 0.03	0.427 ± 0.1621	0.27 ± 0.1081	0.38 ± 0.0316	0.4955 ± 0.1626
4 dietary categories	0.21 ± 0.1019	0.41 ± 0.031	0.606 ± 0.1331	0.19 ± 0.1015	0.41 ± 0.0331	0.6204 ± 0.129

5 dietary categories	0.6 ± 0.1591	0.48 ± 0.0403	0.232 ± 0.1213	0.53 ± 0.1655	0.47 ± 0.0423	0.2869 ± 0.1489
6 dietary categories	-0.4 ± 0.1285	0.51 ± 0.04	0.444 ± 0.1443	-0.42 ± 0.1361	0.5 ± 0.0407	0.4154 ± 0.1526
7 dietary categories	0.65 ± 0.1136	0.89 ± 0.0717	0.466 ± 0.0801	0.65 ± 0.1307	0.88 ± 0.0731	0.462 ± 0.0961
8 dietary categories	-13 ± 0.3017	1749.22 ± 443.0844	0.993 ± 0.0042	-12.96 ± 0.2942	1657.13 ± 424.7377	0.9928 ± 0.0041
Red List (slope)	-0.04 ± 0.0221	0.06 ± 0.0051	0.551 ± 0.1905			
Red List (Threatened)				0.06 ± 0.0712	0.19 ± 0.0167	0.7103 ± 0.2119
Flying *	3.29 ± 0.5125	0.79 ± 0.0282	0.002 ± 0.0079	3.23 ± 0.5136	0.79 ± 0.0278	0.0015 ± 0.0059
Swimming	-0.25 ± 0.2175	0.53 ± 0.0682	0.556 ± 0.1505	-0.28 ± 0.202	0.52 ± 0.0639	0.532 ± 0.1395

Table S6. Phylogenetic logistic regressions (Ho & Ané 2014) of whether a species was a breeding migrant ($n = 18$) compared with all mammals we were certain were migrants of some kind ($n = 113$). The regression was run two ways: once with IUCN Red List as numeric, and once as binary (threatened vs not). Estimates are reported as means and standard deviations across the 939 bootstrap phylogenies for the first regression and 999 for the second; models in 61 (and 1, respectively) phylogenies had numerical fitting problems (as reported by *phylolm*; Ho & Ané 2014) and their estimates are not reported. The mean and standard deviation of the α parameter, which reports the estimated error due to phylogenetic co-variation, was 0.0026 ± 0.0035 for the first regression and 0.0052 ± 0.000413 for the second. See Table S4 for presentation details.

	Red List category as numeric			Red List category as binary factor		
	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)
Intercept (1 habitat, herbivore, 1 dietary category, walking)	-42.37 \pm 2.7545	2221.77 \pm 1839.4338	0.961 \pm 0.0457			
Intercept (1 habitat, herbivore, 1 dietary category, walking, not Red Listed as Critical Endangered)				-2.01 \pm 0.1407	2 \pm 0.132	0.3155 \pm 0.0167

log10(body mass) *	1.77 ± 0.4244	0.82 ± 0.1589	0.035 ± 0.0166	0.00 ± 0.0205	0.22 ± 0.0152	0.9858 ± 0.0772
2 habitats	3.37 ± 1.4069	1.99 ± 0.4239	0.135 ± 0.0947	0.02 ± 0.0673	0.43 ± 0.0518	0.9702 ± 0.1227
Omnivore	19.24 ± 1.4055	2004.44 ± 1556.0773	0.980 ± 0.0314	0.02 ± 0.0686	0.43 ± 0.098	0.9787 ± 0.0889
Carnivore	16.66 ± 1.411	2092.79 ± 1761.708	0.984 ± 0.0191	0.01 ± 0.0526	0.63 ± 0.0668	0.9829 ± 0.0684
2 dietary categories	1.68 ± 0.9591	1.14 ± 0.2897	0.206 ± 0.1611	0.02 ± 0.0696	0.26 ± 0.0379	0.9644 ± 0.1439
3 dietary categories	1.07 ± 2.0398	1825.88 ± 2060.9277	0.991 ± 0.0265	0.01 ± 0.0706	0.43 ± 0.0411	0.9731 ± 0.1123
4 dietary categories	6.81 ± 1.9039	3592.57 ± 4556.7071	0.637 ± 0.3547	0.00 ± 0.0294	0.30 ± 0.0913	0.9913 ± 0.0488
5 dietary categories	3.34 ± 1.814	3237.24 ± 4133.4357	0.533 ± 0.4265	0.00 ± 0.0141	0.70 ± 0.0565	0.9971 ± 0.0169
6 dietary categories	-1.4 ± 2.772	1826.18 ± 2060.6628	0.975 ± 0.0446	0.00 ± 0.0159	0.51 ± 0.1052	0.9968 ± 0.0198
7 dietary categories	4.38 ± 1.9008	10828.64 ± 13843.1614	0.862 ± 0.1836	0.01 ± 0.0417	1.57 ± 0.1447	0.9948 ± 0.0296
Red List (slope)	-0.12 ± 0.1736	0.14 ± 0.0559	0.515 ± 0.3521			

Red List (threatened)				0.02 ± 0.0688	0.20 ± 0.0256	0.9583 ± 0.1715
Flying	21.46 ± 2.0387	1528.44 ± 1690.6905	0.884 ± 0.1499	0.03 ± 0.1111	1.58 ± 0.1376	0.9801 ± 0.0857
Swimming	12.87 ± 4.25	1528.38 ± 1690.7476	0.941 ± 0.0856	0.00 ± 0.03	1.11 ± 0.0925	0.9929 ± 0.0322

Table S7. Phylogenetic logistic regressions (Ho & Ané 2014) of whether a species was a refuge migrant ($n = 64$) compared with all mammals we were certain were migrants of some kind ($n = 113$). The regression was run two ways: once with IUCN Red List as numeric, and once as binary (threatened vs not). Estimates are reported as means and standard deviations across the 995 bootstrap phylogenies; models in 5 phylogenies had numerical fitting problems (as reported by *phylolm*; Ho & Ané 2014) and their estimates are not reported. The mean and standard deviation of the α parameter, which reports the estimated error due to phylogenetic co-variation, was 0.018 ± 0.0078 for the first regression and 0.019 ± 0.00295 for the second. See Table S4 for presentation details.

	Red List category as numeric			Red List category as binary factor		
	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)
Intercept (1 habitat, herbivore, 1 dietary category, walking)	2.76 \pm 0.3456	1.55 \pm 0.0771	0.082 \pm 0.0521			
Intercept (1 habitat, herbivore, 1 dietary category, walking, not Red Listed as Critical Endangered)				0.16 \pm 0.1709	1.35 \pm 0.0279	0.8923 \pm 0.0848
log10(body mass)	-0.38 \pm 0.0909	0.26 \pm 0.0102	0.163 \pm	-0.19 \pm 0.0434	0.24 \pm 0.0058	0.4319 \pm

			0.0684			0.097
2 habitats	0.68 ± 0.2042	0.74 ± 0.0598	0.377 ± 0.1321	0.38 ± 0.0914	0.63 ± 0.0225	0.5462 ± 0.1013
Omnivore	-1.23 ± 0.2415	0.94 ± 0.0294	0.204 ± 0.119	0.44 ± 0.1185	0.85 ± 0.0391	0.6069 ± 0.0865
Carnivore	-1.05 ± 0.3415	0.94 ± 0.0352	0.293 ± 0.1946	0.42 ± 0.1496	0.93 ± 0.0273	0.6508 ± 0.1124
2 dietary categories	-0.11 ± 0.2235	0.54 ± 0.0286	0.783 ± 0.2387	0.00 ± 0.1137	0.48 ± 0.0362	0.8568 ± 0.1122
3 dietary categories	-0.19 ± 0.2412	0.82 ± 0.0371	0.815 ± 0.1944	-0.11 ± 0.1166	0.82 ± 0.0393	0.8768 ± 0.0904
4 dietary categories	0.17 ± 0.126	1.05 ± 0.095	0.864 ± 0.0721	0.22 ± 0.1055	0.82 ± 0.0680	0.7912 ± 0.0876
5 dietary categories	-0.46 ± 0.1179	1.02 ± 0.0746	0.652 ± 0.0776	-0.17 ± 0.1131	1.05 ± 0.0368	0.8711 ± 0.0792
6 dietary categories	0.2 ± 0.2204	1.26 ± 0.046	0.849 ± 0.1033	-0.51 ± 0.1248	1.20 ± 0.0498	0.6745 ± 0.0741
7 dietary categories	15.99 ± 0.2657	3134.53 ± 766.8089	0.996 ± 0.0013	0.70 ± 0.2041	2.12 ± 0.1299	0.7457 ± 0.0594
Red List (slope)	-0.05 ± 0.0482	0.11 ± 0.0085	0.589 ± 0.1695			
Red List (threatened)				-0.14 ± 0.1187	0.34 ± 0.0246	$0.6773 \pm$

						0.2178
Flying	-0.36 ± 0.3285	1.31 ± 0.0702	0.760 ± 0.1584	0.41 ± 0.1049	1.17 ± 0.0117	0.7231 ± 0.0616
Swimming	-0.12 ± 0.3324	1.02 ± 0.0445	0.766 ± 0.116	0.33 ± 0.1117	1.06 ± 0.0235	0.752 ± 0.0762

Table S8. Phylogenetic logistic regressions (Ho & Ané 2014) of whether a species was a tracking migrant ($n = 64$) compared with all mammals we were certain were migrants of some kind ($n = 113$). The regression was run two ways: once with IUCN Red List as numeric, and once as binary (threatened vs not). Estimates are reported as means and standard deviations across the 942 bootstrap phylogenies; models in 58 phylogenies had numerical fitting problems (as reported by *phylolm*; Ho & Ané 2014) and their estimates are not reported. The mean and standard deviation of the α parameter, which reports the estimated error due to phylogenetic co-variation, was 0.039 ± 0.016 for the first regression and 0.0066 ± 0.00484 for the second. See Table S4 for presentation details.

	Red List category as numeric			Red List category as binary factor		
	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)	Estimate (mean \pm SD)	Std Err (mean \pm SD)	<i>p</i> -value (mean \pm SD)
Intercept (1 habitat, herbivore, 1 dietary category, walking)	-1.25 \pm 0.1369	1.59 \pm 0.0224	0.431 \pm 0.0462			
Intercept (1 habitat, herbivore, 1 dietary category, walking, not Red Listed as Critical Endangered)				-0.13 \pm 0.2916	1.44 \pm 0.0397	0.8569 \pm 0.1004
log10(body mass)	-0.12 \pm 0.0465	0.27 \pm 0.0061	0.654 \pm	0.10 \pm 0.0748	0.23 \pm 0.0129	0.6395 \pm

			0.1143			0.1833
2 habitats	-1.65 ± 0.1839	0.84 ± 0.0334	0.054 ± 0.0266	-0.91 ± 0.4951	0.53 ± 0.0716	0.1925 ± 0.2159
Omnivore	1.14 ± 0.1193	1.02 ± 0.0192	0.269 ± 0.0533	-0.29 ± 0.1799	0.76 ± 0.0628	0.7051 ± 0.1652
Carnivore *	3.03 ± 0.189	1.36 ± 0.0449	0.026 ± 0.0201	0.98 ± 0.5215	0.93 ± 0.0654	0.3605 ± 0.2405
2 dietary categories	0.91 ± 0.1135	0.63 ± 0.0421	0.159 ± 0.0687	0.14 ± 0.1626	0.44 ± 0.0371	0.7175 ± 0.192
3 dietary categories	1.41 ± 0.0965	1 ± 0.0314	0.161 ± 0.0329	0.07 ± 0.2071	0.76 ± 0.0674	0.8271 ± 0.1423
4 dietary categories	0.87 ± 0.1034	0.78 ± 0.1262	0.263 ± 0.0964	-0.21 ± 0.2181	0.70 ± 0.1353	0.7059 ± 0.1265
5 dietary categories	1.4 ± 0.115	1.22 ± 0.0263	0.255 ± 0.0483	0.61 ± 0.3063	0.99 ± 0.0919	0.5618 ± 0.1754
6 dietary categories	0.29 ± 0.0644	1.54 ± 0.0269	0.851 ± 0.0288	0.28 ± 0.3255	1.11 ± 0.1206	0.7772 ± 0.1989
7 dietary categories	-14.82 ± 0.2362	1867.14 ± 269.8648	0.993 ± 0.003	-0.04 ± 0.2165	1.79 ± 0.2141	0.9432 ± 0.0607
Red List (slope) *	0.36 ± 0.0436	0.15 ± 0.0074	0.017 ± 0.0238			
Red List (threatened)				0.73 ± 0.2301	0.36 ± 0.0286	0.0793 ±

						0.0909
Flying *	-4.33 ± 0.1403	1.58 ± 0.0664	0.006 ± 0.0054	-1.57 ± 0.7962	1.23 ± 0.0609	0.2846 ± 0.233
Swimming *	-2.94 ± 0.1062	1.35 ± 0.0584	0.031 ± 0.0081	-1.19 ± 0.3294	0.95 ± 0.0591	0.2343 ± 0.1413

Table S9. The median and interquartile range (IQR) values of significant life history traits and the IUCN Red List rank for each migration type (breeding, refuge, tracking).

Factor		Breeding	Refuge	Tracking
Mass (g)	Median	137194.86	3185.74	53954.05
	IQR	184679.00	75885.83	195198.60
Habitat breadth	Median	2	1	1
	IQR	0	0	0
IUCN Red List rank	Median	2	1	2
	IQR	2	1	2