

## Supplementary Information

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### Fossorial adaptations in African mole-rats (Bathyergidae) and the unique appendicular phenotype of naked mole-rats

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#### **Abbreviations**

Deltoid tuberosity	<b>DT</b>
Deltopectoral crest	<b>DC</b>
Distal fusion of tibia-fibula	<b>DFTFi</b>
Epicondyles (Humerus)	<b>EH</b>
Fibula	<b>Fi</b>
Femoral head	<b>FH</b>
Humerus	<b>H</b>
Humeral head	<b>HH</b>
Index of fossorial ability	<b>IFA</b>
Grater trochanter	<b>GT</b>
Grater tubercle	<b>GTu</b>
Lesser trochanter	<b>LT</b>
Olecranon process	<b>OP</b>
Relative position of DT	<b>RDT</b>
Third trochanter	<b>TT</b>
Tibia	<b>T</b>
Tibio-fibular junction index	<b>TJI</b>
Trochlear notch	<b>TN</b>

## SUPPLEMENTARY TABLES

**Supplementary Table 1.** Total number of bathyergid individuals analyzed in this study ( $n = 382$ ). Data includes ID code, sex, institution of origin (IO), ontogenetic stage (OS), limb bones and bone superstructures. Outgroup specimens ( $n = 19$ ) from Petromuridae and Hystricidae were also included. The bones analyzed are humerus (H,  $n = 380$ ), ulna (U,  $n = 376$ ), femur (F,  $n = 323$ ), tibia (T,  $n = 312$ ), and fibula (Fi,  $n = 312$ ). Presence/absence of bone superstructures including a projected deltoid tuberosity (DT), olecranon process (OP), third trochanter (TT), and distal tibio-fibular fusion (DFTFi). Some specimens are from captivity (ca). Scanned bones are indicated with an asterisk (\*). Sample size for each species: *Bathyergus suillus* ( $n = 78$ ); *B. janetta* ( $n = 6$ ); *Heliophobius argenteocinereus* ( $n = 38$ ); *Georychus capensis* ( $n = 51$ ); *Cryptomys hottentotus* ( $n = 53$ ); *Fukomys mechowii* ( $n = 32$ ); *Fukomys damarensis* ( $n = 48$ ); *Heterocephalus glaber* ( $n = 76$ ); *Hystrix africae australis* ( $n = 18$ ); *Petromus typicus* ( $n = 1$ ). OS: newborn (0), pup (1), juvenile (2), and adult (3). Sex: males (M) and females (F). Institutions: Kalahari Mole-Rat Project, Kalahari Research Centre (KRC); Department of Biological Sciences, University of Cape Town (UCT); Mammal Research Institute, Department of Zoology and Entomology, University of Pretoria (UP); Department of Zoology, Faculty of Science, University of South Bohemia (USB); Paläontologisches Institut und Museum, Universität Zürich (UZ).

Species	ID Code	Sex	Age	OS	IO	H	U	F	T-Fi	DT	OP	TT	DFTFi
<i>Bathyergus suillus</i>	219	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	307	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	314	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	333	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	365	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	366	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	377	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	717	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	721	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	911	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	913	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	938	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	982	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1085	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1138	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1144	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1153	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1155	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1163	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Bathyergus suillus</i>	1169*	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1171	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1332	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1336	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1373*	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	217	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	220	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	223	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	300	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	313	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	376	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	713	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	765	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	861	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	910	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	964	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	965	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1039	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1050	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1139	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1154	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1338	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1339	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	332	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	337	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	367	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	866	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	1245	F		2	UCT			F	T-Fi			Yes	Yes

<i>Bathyergus suillus</i>	283*	M		2?	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	F10	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	F2	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	F5	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	F7	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S10*	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S11	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S12	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S13	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S14	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S16	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S17	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S2	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S20	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S3	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S6	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S7*	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	S8	F		3	UP	H	U			Yes	Yes		
<i>Bathyergus suillus</i>	F1	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	F4	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S1*	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S15*	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S18	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S19	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S4	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S5	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	S9	M		3	UP	H	U	F		Yes	Yes	Yes	
<i>Bathyergus suillus</i>	GM282*	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Bathyergus suillus</i>	GM283*	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	GM284*	M		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus suillus</i>	GM285*	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus janetta</i>	GM005	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus janetta</i>	GM007	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus janetta</i>	GM013	F		3	UCT	H		F		Yes		Yes	
<i>Bathyergus janetta</i>	GM014	M		3	UCT	H		F		Yes		Yes	
<i>Bathyergus janetta</i>	GM239	-	2 days	1	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Bathyergus janetta</i>	GM503	-		2?	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	GM245	M	a few days		UZ	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Heliophobius argenteocinereus</i>	GM246	F	1 day		UZ	H	U			Yes	Yes		
<i>Heliophobius argenteocinereus</i>	GM247	M	~4 days		UZ	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	GM248	F	~14 days		UZ	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	103	M	2 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Heliophobius argenteocinereus</i>	77	M	2 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	20	M	3.5 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	350*	F	1 month		USB	H	U	F	T-Fi	Yes	Yes	No	Yes
<i>Heliophobius argenteocinereus</i>	526	M	2 years, 4 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	26*	F	1.5 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	11	F	2.5 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	387*	F	at least 5 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	27	F	2 years, 10 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Heliophobius argenteocinereus</i>	386	M	at least 6 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	8	M	4 years, 9 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	6*	F	at least 1 year, 3 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	516	M	at least 8 years, 2 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	361*	F	at least 2 years, 1 month	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	517	M	at least 2 years, 7 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	585	M	at least 5 years, 7 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	377	M	at least 2 years, 2 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	451	F	at least 2 years, 1 month	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	5*	F	at least 3 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	18	M	at least 6 years, 8 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	393	M	at least 1 year, 11 months		USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	473	F	at least 8 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	476	M	at least 2 years, 10 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	508	F	at least 3 years, 8 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	525	F	2 years, 5 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	492	F	at least 14 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	299*	F	2-3 days old		USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Heliophobius argenteocinereus</i>	479*	F	6 years, 10 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Heliophobius argenteocinereus</i>	244	M	at least 2 years, 1 month	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	590*	F	at least 6 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	NB10*	M		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	NB21*	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	NB05*	M		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heliophobius argenteocinereus</i>	NB15	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM528		~2 months	2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM530		3 days	1	UCT	H	U			Yes	Yes		
<i>Georychus capensis</i>	JO475*	F	NB21	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	JO402*	M	GM530	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM295*	-	JO402	2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM296*	F	GM295	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM297*	F	GM296	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	GM298*	F	GM297	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	Z4	F	GM298	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z6	F	Z4	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z9	M	Z9	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z10	F	Z10	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z11	F	Z11	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z12	F	Z12	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z22	M	Z22	1	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z23	F	Z23	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z25	F	Z25	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z27	F	Z27	1	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z28	M	Z28	1	UP	H	U			Yes	Yes		

<i>Georychus capensis</i>	Z31	F	Z31	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z32	F	Z32	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z33	M	Z33	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z35	F	Z35	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z36	F	Z36	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z37	F	Z37	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z38	F	Z38	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z210	M	Z210	1	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z212	M	Z212	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z214	F	Z214	3	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z215	M	Z215	2	UP	H				Yes			
<i>Georychus capensis</i>	Z216	F	Z216	1	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z218	F	Z225	2	UP		U				Yes		
<i>Georychus capensis</i>	Z219	M	A01	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	Z225	-	A03	2	UP	H	U			Yes	Yes		
<i>Georychus capensis</i>	A01	F	A06	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A03	F	A19	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A06	M	A35	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A19	M	A40	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A35	F	A43	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A40	M	A51	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A43	F	A95	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A51	F	A97	2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A95	F	A99	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A97	F	A100	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A99	F	A115	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A100	F	A118	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A115	M	A128	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes



<i>Georychus capensis</i>	A118	M	A130	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A128	F	A131	3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A130	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Georychus capensis</i>	A131	F		2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM102	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM103	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM104	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM105	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM106*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM107	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM108	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM109	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM110	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM111	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM112	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM113	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM114	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM115*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM116	-		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM117	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM118	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM119*	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM120	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM121*	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM122	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM123*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM124	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM125	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Cryptomys hottentotus</i>	GM126	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM127	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM128	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM129*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM130	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM131	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM132*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM133	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM134	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM135	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM136	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM137	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM138	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM139	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM140	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM141	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM142	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM143	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM144	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM145*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM146	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM147	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM148	M		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM149	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM150*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM151	F		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM152*	M		3	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Cryptomys hottentotus</i>	GM153	-		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Cryptomys hottentotus</i>	GM154	F		2	UCT	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	227	F	1 day	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	228*	M	1 day	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	146	F	2 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	148	M	1 day	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	145	F	5 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	147	F	3 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	59	M	2 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	154	F	2 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	55	F	4 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	58	M	4 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	125	F	1 month	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	167	M	21 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	36*	M	5 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	45	F	9 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	153*	M	21 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	78*	M	1 month, 11 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	123	F	4-5 months?	2	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	54	F	17 days	1	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	116	F		1	USB	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys mechowii</i>	14*	M	4 months	2	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	23	F	15 months	2	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	179*	M	13 months	2	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	3	F	1 year, 10 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	1	F	4 years, 4 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	177*	M	2 years, 3 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	497	F	13 years, 6 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	478*	M	3 years, 2 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes

<i>Fukomys mechowii</i>	494	F	7 years, 1 month	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	589	F	10 years, 7 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	242*	M	3 years	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	2*	M	1 year, 8 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys mechowii</i>	531	-	1 year, 5 months	3	USB	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB227	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB228	F		3	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB229	F		2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB230*	M		1-2?	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB231*	M		2	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB232*	M		1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB233*	M		1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB238	-		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	NB422	M	1 day	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	NB423	-	1 day	1	UP	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys damarensis</i>	GAP315	F		3	UCT	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D11	M		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D12	M		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D13	F		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D14	M		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D15	F		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D16	M		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D17	M		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D18	F		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	D19	F		3	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	HD2	-		2-3?	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	FD5	-		2	UP	H	U			Yes	Yes		
<i>Fukomys damarensis</i>	GM039	-		1	UP	H				Yes			

<i>Fukomys damarensis</i> (ca)	G3F035	F	1 Day	1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	No
<i>Fukomys damarensis</i> (ca)	G3M036*	M	1 Day	1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4F030	F	1 Month, 29 Days	1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4F042	F	13 Months, 10 Days	3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M018	M	30 Months, 28 Days	3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M029	M	1 Month, 10 Days	1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M034*	M	20 Months, 27 Days	3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M036	M	1 Month, 29 Days	1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M039*	M	18 Months, 21 Days	3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M041*	M	10 Months, 14 Days	2	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i> (ca)	G4M047*	M	7 Months, 0 Days	2	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	F2M012*	M		1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	F2F013	F		1	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G7F013	F		2	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	F6F002	F		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G4M008*	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G5F003	F		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	L29F002	F		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G4F012	F		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G4F002	F		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G4M009*	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G3M007	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G4M001	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	G3M001*	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Fukomys damarensis</i>	Z3M006*	M		3	KRC	H	U	F	T-Fi	Yes	Yes	Yes	Yes
<i>Heterocephalus glaber</i> (ca)	JJ-402	-	3 days	1	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-405	-	Perinatal	0	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-416	-	2 Months, 2 days	1-2?	UCT	H	U	F	T-Fi	No	Yes	Yes	No

<i>Heterocephalus glaber</i> (ca)	JJ-417	-	5 days	1	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-557	-	Juvenile	1?	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-559	-	Perinatal	0	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-560	-	Perinatal	0	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-001	F	~1 year, 6 months	2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-002	-		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-008	F	~10 years	3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-009	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-010	-	~4 months	1	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-045*	M		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	JJ-046*	F		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-047	-		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-048	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-049	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-050	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-051	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-052*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-053	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-054	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-055	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-056	-		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-057	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-058	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-059	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-060*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-061*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-063	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-064	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No

<i>Heterocephalus glaber</i> (ca)	1200-065	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-066	-		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-067*	F		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-068	F		2	UCT	H		F	T-Fi	No		Yes	No
<i>Heterocephalus glaber</i> (ca)	1200-069	-		3	UCT	H	U	F	T-Fi	No	Yes	Yes	Yes
<i>Heterocephalus glaber</i> (ca)	1000-070	F		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-071	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-073	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-074	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-075	-		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-076	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-077	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-078	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-079	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	Yes
<i>Heterocephalus glaber</i> (ca)	1000-080	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-081*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	Yes
<i>Heterocephalus glaber</i> (ca)	1000-082*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-083	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-084	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-085	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	1000-086	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-087	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-088*	M		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-091	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-092*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-093	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-094	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-095	M		2	UCT	H	U	F	T-Fi	No	Yes	Yes	No

<i>Heterocephalus glaber</i> (ca)	5000-096	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-097	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-098	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-099	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-100	F		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	5000-101*	M		3	UCT	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM498	F		3	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM499	M		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM500	?		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM501	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM505	M		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM506	F		3	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM507	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM508	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM509	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM510	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Heterocephalus glaber</i> (ca)	GM511	F		3?	UP	H	U	F	T-Fi	No	Yes	Yes	No
<i>Hystrix africae australis</i>	38634				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	36199				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	38502				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	38536				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	36738				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	35988				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	39666				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	40427				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	37705				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	40403				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	35137				IZIK O	H	U	F		Yes	Yes	No	



<i>Hystrix africae australis</i>	40310				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	36139				IZIK O	H	U			Yes	Yes		
<i>Hystrix africae australis</i>	38525				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	40429				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	40744				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	36060				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Hystrix africae australis</i>	38256				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No
<i>Petromus typicus</i>	40790				IZIK O	H	U	F	T-Fi	Yes	Yes	No	No

**Supplementary Table 2.** Comparison of the out-of-sample predictive score by means of the Widely Applicable Information Criterion (WAIC) of models with and without body mass (BM). SE gives an approximate standard error of each score,  $\Delta_i$  WAIC gives the difference between the  $i$ th model and the model with the lowest WAIC,  $\Delta_i$  SE gives the approximate standard error of the difference between the  $i$ th model and the model with the lowest WAIC, and  $w_i$  is the weight of the  $i$ th model and indicates a relative support for that model. Model with a subscript  $I$  indicates a model without BM (i.e. intercept-only), and model with a subscript BM is a model with BM. Relative position of the deltoid tuberosity (RDT); Index of fossorial ability (IFA); Tibio-fibular junction index (TJI).

Index	Model	WAIC (SE)	$\Delta_i$ WAIC	$\Delta_i$ SE	$w_i$
RDT					
	RDT <sub>I</sub>	-572.71 (16.62)	0	NA	0.61
	RDT <sub>BM</sub>	-571.78 (16.86)	0.93	2.38	0.39
TJI					
	TJI <sub>I</sub>	-467.85 (20.04)	0	NA	0.62
	TJI <sub>BM</sub>	-466.85 (21.03)	1.01	3.02	0.38
IFA					
	IFA <sub>BM</sub>	-627.95 (24.41)	0	NA	0.96
	IFA <sub>I</sub>	-621.64 (24.64)	6.31	4.43	0.04

**Supplementary Table 3.** Summary of the phylogenetic varying effects regression with estimates of population-level parameters for three morpho-functional indices. Model for each index includes estimate of intercept mean ( $\mu_\alpha$ ) and slope mean ( $\mu_\beta$ ), as well as estimates of phylogenetic ( $\sigma_{phy}$ ) and species-specific ( $\sigma_{spp}$ ) components, with 89% compatibility intervals (CI) in square brackets. Relative position of the deltoid tuberosity (RDT); Index of fossorial ability (IFA); Tibio-fibular junction index (TJI).

Index	$\mu_\alpha$ [89% CI]	$\mu_\beta$ [89% CI]	$\sigma_{phy\alpha}$ [89% CI]	$\sigma_{phy\beta}$ [89% CI]	$\sigma_{spp}$ [89% CI]
RDT	-0.58 [-0.70, -0.45]	0.01 [-0.01, 0.03]	0.02 [0.00, 0.09]	0.00 [0.00, 0.00]	0.07 [0.00, 0.19]
TJI	-0.68 [-0.82, -0.51]	0.01 [-0.01, 0.04]	0.03 [0.00, 0.13]	0.00 [0.00, 0.00]	0.08 [0.01, 0.24]
IFA	-1.31 [-1.49, -1.05]	0.02 [-0.01, 0.06]	0.06 [0.00, 0.23]	0.00 [0.00, 0.01]	0.11 [0.01, 0.33]

**Supplementary Table 4.** Summary of the phylogenetic varying effects regression for three morpho-functional indices. Each species  $j$  has its own estimate of intercept ( $\alpha_j$ ) and slope ( $\beta_j$ ), with 89% compatibility intervals (CI) in square brackets. Relative position of the deltoid tuberosity (RDT); Index of fossorial ability (IFA); Tibio-fibular junction index (TJI).

Index	Species	$\alpha_j$ [89% CI]	$\beta_j$ [89% CI]
RDT			
	<i>B. suillus</i>	-0.58 [-0.71, -0.45]	0.01 [-0.01, 0.03]
	<i>C. hottentotus</i>	-0.60 [-0.74, -0.47]	0.02 [0.00, 0.05]
	<i>F. damarensis</i>	-0.60 [-0.73, -0.46]	0.00 [-0.02, 0.02]
	<i>F. mechowii</i>	-0.63 [-0.80, -0.49]	0.01 [-0.01, 0.04]
	<i>G. capensis</i>	-0.58 [-0.70, -0.45]	0.01 [-0.01, 0.03]
	<i>H. argenteocinereus</i>	-0.53 [-0.66, -0.37]	0.00 [-0.02, 0.03]
TJI			
	<i>B. suillus</i>	-0.63 [-0.80, -0.44]	0.01 [-0.01, 0.04]
	<i>C. hottentotus</i>	-0.71 [-0.88, -0.55]	0.02 [-0.01, 0.05]
	<i>F. damarensis</i>	-0.66 [-0.82, -0.47]	0.00 [-0.03, 0.03]
	<i>F. mechowii</i>	-0.69 [-0.86, -0.51]	0.01 [-0.02, 0.04]
	<i>G. capensis</i>	-0.72 [-0.89, -0.56]	0.02 [-0.01, 0.04]
	<i>H. argenteocinereus</i>	-0.73 [-0.92, -0.56]	0.02 [0.00, 0.06]
IFA			
	<i>B. suillus</i>	-1.25 [-1.49, -0.94]	-0.03 [-0.07, 0.01]
	<i>C. hottentotus</i>	-1.41 [-1.69, -1.17]	0.04 [0.00, 0.10]
	<i>F. damarensis</i>	-1.41 [-1.67, -1.17]	0.04 [0.00, 0.09]
	<i>F. mechowii</i>	-1.29 [-1.53, -0.99]	0.02 [-0.03, 0.06]
	<i>G. capensis</i>	-1.40 [-1.62, -1.17]	0.02 [-0.01, 0.06]
	<i>H. argenteocinereus</i>	-1.31 [-1.53, -1.05]	0.02 [-0.02, 0.07]
	<i>H. glaber</i>	-1.36 [-1.54, -1.16]	0.02 [-0.01, 0.06]

**Supplementary Table 5.** Bone superstructures analyzed in this study and their fossorial functional significance. Data obtained from several sources (see Supplementary References 1-23).

Bone	Character	Morpho-Function	References
Humerus (Stylopod)	Deltoid tuberosity (DT)	An enlarged, protuberant and distally located DT increases the area for the insertion of <i>mm. deltoidei</i> and <i>mm. pectorales</i> , and increases the in-lever arm distance from muscle attachment to joint. This produces a powerful stroke (flexion) of the humerus on the scapula and subsequent retraction of the arm during parasagittal scratch-digging.	1,6,7,9,10,11,13,14,20,21
Ulna (Zeugopod)	Olecranon process (OP)	An enlarged OP increases the area for the insertion of the <i>mm. triceps brachii</i> , hence increase the in-lever arm of forearm extensors and enables greater out-forces during scratch-digging.	4,8,17,22,23
Femur (Stylopod)	Third trochanter (TT)	A mediolaterally enlarged and distally positioned TT increases the insertion area for the <i>m. gluteus superficialis</i> for powerful leg extension and some degree of abduction (to brace against a tunnel or pushing back soil while digging), as well as increases medioalteral cortical thicknss of the diaphysis, which reduces bending strains on the femoral shaft due to rearward motion.	12,16,18
Tibia-Fibula (Zeugopods)	Distal fusion of tibia-fibula (DFTF)	Distal fusion provides a powerful bony base for the attachment of the muscles acting on the paws, as well as increases stabilization of the foot to better withstand load-bearing and bending strains during heavy-impact activities against a resistant medium such as earth.	2,3,5,6,15,18,19,20

**Supplementary Table 6.** Ecological and morpho-functional characteristics of fossorial rodents including 17 genera and 35 species, and the non-fossorial closest relatives of Bathyergidae. All fossorial taxa present a well-developed and projected deltoid tuberosity (DT), except *Heterocephalus glaber*, where such trait is highly reduced. Ecological and morphological data obtained from several sources (see Supplementary References 22,24,28,30,38,40-75).

Family	Genus	Species	Locomotor mode	Digging mode	Social organization	DT
Ctenohystrica						
Bathyergidae	<i>Heterocephalus</i>	<i>glaber</i>	Fossorial	Chisel-tooth	Highly social	No
Bathyergidae	<i>Heliophobius</i>	<i>argenteocinereus</i>	Fossorial	Chisel-tooth	Solitary	Yes
Bathyergidae	<i>Bathyergus</i>	<i>suillus</i>	Fossorial	Scratch	Solitary	Yes
Bathyergidae	<i>Bathyergus</i>	<i>janetta</i>	Fossorial	Scratch	Solitary	Yes
Bathyergidae	<i>Georchus</i>	<i>capensis</i>	Fossorial	Chisel-tooth	Solitary	Yes
Bathyergidae	<i>Cryptomys</i>	<i>hottentotus pretoriae</i>	Fossorial	Chisel-tooth	Social	Yes
Bathyergidae	<i>Cryptomys</i>	<i>hottentotus natalensis</i>	Fossorial	Chisel-tooth	Social	Yes
Bathyergidae	<i>Cryptomys</i>	<i>hottentotus mahali</i>	Fossorial	Chisel-tooth	Social	Yes
Bathyergidae	<i>Fukomys</i>	<i>damarensis</i>	Fossorial	Chisel-tooth	Highly social	Yes
Bathyergidae	<i>Fukomys</i>	<i>mechowii</i>	Fossorial	Chisel-tooth	Highly social	Yes
Hystricidae	<i>Atherurus</i>	<i>macrourus</i>	Semifossorial	Scratch	Gregarious/Solitary	Yes
Hystricidae	<i>Atherurus</i>	<i>africanus</i>	Ambulatory		Solitary	Yes
Hystricidae	<i>Hystrix</i>	<i>africae australis</i>	Semifossorial	Scratch	Gregarious/Social	Yes
Hystricidae	<i>Hystrix</i>	<i>cristata</i>	Semifossorial	Scratch	Gregarious/Solitary	Yes
Hystricidae	<i>Hystrix</i>	<i>indica</i>	Semifossorial	Scratch	Gregarious/Solitary	Yes
Petromuridae	<i>Petromus</i>	<i>typicus</i>	Rock climber		Gregarious/Solitary	Yes
Thryonomyidae	<i>Thryonomys</i>	<i>swinderianus</i>	Semiaquatic/Semifossorial	Scratch	Gregarious/Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>flamarioni</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>lewisi</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>talarum</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>rionegrensis</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>leucodon</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>minutus</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes

Ctenomyidae	<i>Ctenomys</i>	<i>lami</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>torquatus</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Ctenomyidae	<i>Ctenomys</i>	<i>fulvus</i>	Fossorial	Scratch/Chisel-tooth	Solitary	Yes
Octodontidae	<i>Spalacopus</i>	<i>cyaneus</i>	Fossorial	Chisel-tooth	Social	Yes
Geomyoidea/Myomorpha						
Geomyidae	<i>Geomys</i>	<i>bursarius</i>	Fossorial	Scratch	Solitary	Yes
Geomyidae	<i>Thomomys</i>	<i>bottae</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Geomyidae	<i>Thomomys</i>	<i>talpoides</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Geomyidae	<i>Thomomys</i>	<i>mazama</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Geomyidae	<i>Thomomys</i>	<i>towsendii</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Spalacidae	<i>Myospalax</i>	<i>myospalax</i>	Fossorial	Scratch/Head-Lift	Solitary	Yes
Spalacidae	<i>Rhizomys</i>	<i>pruinus</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Spalacidae	<i>Rhizomys</i>	<i>sinensis</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Spalacidae	<i>Rhizomys</i>	<i>sumatrensis</i>	Fossorial	Chisel-Tooth/Scratch	Solitary	Yes
Spalacidae	<i>Spalax</i>	<i>ehrenbergi</i>	Fossorial	Head-Lift/Chisel-tooth	Solitary	Yes
Spalacidae	<i>Spalax</i>	<i>microphthalmus</i>	Fossorial	Head-Lift/Chisel-tooth	Solitary/Gregarious	Yes
Spalacidae	<i>Nannospalax</i>	<i>nehringi</i>	Fossorial	Head-Lift/Chisel-tooth	Solitary	Yes
Cricetidae	<i>Arvicola</i>	<i>(terrestri) scherman</i>	Fossorial	Chisel-tooth	Solitary	Yes
Cricetidae	<i>Ellobius</i>	<i>lutescens</i>	Fossorial	Chisel-tooth	Solitary/Gregarious	Yes
Sciuroidea						
Aplodontidae	<i>Aplodontia</i>	<i>rufa</i>	Fossorial	Scratch	Solitary/Gregarious	Yes

**Supplementary Table 7.** Morpho-functional indices and body masses (BM) of individuals analyzed in this study. Body mass in grams (g). Relative position of the deltoid tuberosity (RDT); Index of fossorial ability (IFA); Tibio-fibular junction index (TJI).

Species	ID	BM (g)	IFA	RDP	TJI
<i>B. suillus</i>	314	998	0.246	0.596	0.595
<i>B. suillus</i>	365	804	0.266	0.594	0.584
<i>B. suillus</i>	366	748	0.237	0.620	0.584
<i>B. suillus</i>	377	918	0.223	0.583	0.583
<i>B. suillus</i>	717	1110	0.233	0.569	0.553
<i>B. suillus</i>	721	640	0.220	0.585	0.612
<i>B. suillus</i>	911	1152	0.231	0.581	0.598
<i>B. suillus</i>	913	366	0.264	0.572	0.580
<i>B. suillus</i>	938	468	0.259	0.570	0.603
<i>B. suillus</i>	982	422	0.263	0.607	0.602
<i>B. suillus</i>	1085	778	0.253	0.612	0.616
<i>B. suillus</i>	1138	1138	0.248	0.603	0.601
<i>B. suillus</i>	1144	914	0.236	0.584	0.606
<i>B. suillus</i>	1153	882	0.265	0.625	0.590
<i>B. suillus</i>	1155	1066	0.240	0.572	0.621
<i>B. suillus</i>	1163	760	0.261	0.632	0.594
<i>B. suillus</i>	1169	910	0.226	0.608	0.631
<i>B. suillus</i>	1171	1346	0.233	0.616	0.576
<i>B. suillus</i>	1332	958	0.232	0.593	0.595
<i>B. suillus</i>	1336	508	0.258	0.578	0.603
<i>B. suillus</i>	217	1278	0.253	0.586	0.616
<i>B. suillus</i>	220	1726	0.232	0.586	0.602
<i>B. suillus</i>	313	1072	0.243	0.585	0.557
<i>B. suillus</i>	713	1614	0.228	0.606	0.630
<i>B. suillus</i>	765	830	0.257	0.607	0.628
<i>B. suillus</i>	861	1636	0.224	0.593	0.593
<i>B. suillus</i>	910	662	0.246	0.567	0.597
<i>B. suillus</i>	964	584	0.280	0.589	0.608
<i>B. suillus</i>	965	1228	0.251	0.582	0.589
<i>B. suillus</i>	1039	1336	0.215	0.604	0.596
<i>B. suillus</i>	1050	1332	0.243	0.629	0.578
<i>B. suillus</i>	1139	752	0.264	0.557	0.613
<i>B. suillus</i>	1154	780	0.271	0.601	0.628
<i>B. suillus</i>	1338	1450	0.246	0.574	0.614
<i>B. suillus</i>	1339	788	0.252	0.597	0.620
<i>B. suillus</i>	GM282	603	0.250	0.572	0.597
<i>B. suillus</i>	GM283	227	0.234	0.596	0.576
<i>B. suillus</i>	GM284	1201	0.232	0.619	0.588
<i>B. suillus</i>	GM285	658	0.253	0.614	0.593



<i>G.capensis</i>	NB21	141	0.280	0.581	0.554
<i>G.capensis</i>	GM530	121	0.277	0.616	0.510
<i>G.capensis</i>	JO402	180	0.268	0.573	
<i>G.capensis</i>	GM295	260	0.255	0.610	
<i>G.capensis</i>	GM296	105	0.212	0.566	
<i>G.capensis</i>	GM297	185	0.271	0.573	
<i>G.capensis</i>	GM298	210	0.265	0.593	
<i>G.capensis</i>	Z4	225	0.261	0.568	
<i>G.capensis</i>	Z9	223.1	0.292	0.573	
<i>G.capensis</i>	Z10	210.2	0.286	0.594	
<i>G.capensis</i>	Z11	105.8	0.251	0.566	
<i>G.capensis</i>	Z12	169.8	0.247	0.605	
<i>G.capensis</i>	Z22	149	0.290	0.594	
<i>G.capensis</i>	Z23	167	0.249	0.562	
<i>G.capensis</i>	Z25	190	0.227	0.574	
<i>G.capensis</i>	Z27	300.8	0.305	0.581	
<i>G.capensis</i>	Z28	171.1	0.262	0.588	
<i>G.capensis</i>	Z31	235.1	0.295	0.595	
<i>G.capensis</i>	Z32	197.4	0.292	0.614	
<i>G.capensis</i>	Z33	126.6	0.267	0.593	
<i>G.capensis</i>	Z35	170.2	0.273	0.573	
<i>G.capensis</i>	Z36	229.5	0.318	0.587	
<i>G.capensis</i>	Z37	135.5		0.577	
<i>G.capensis</i>	Z38	116.6	0.238	0.552	
<i>G.capensis</i>	Z210	141	0.274	0.562	
<i>G.capensis</i>	Z212	156.7	0.269		
<i>G.capensis</i>	Z214	150.8	0.279	0.557	
<i>G.capensis</i>	Z215	298	0.245	0.536	
<i>G.capensis</i>	Z216	175	0.244	0.563	
<i>G.capensis</i>	Z225	75.49	0.277	0.613	0.514
<i>G.capensis</i>	A01	91.85	0.255	0.567	0.537
<i>G.capensis</i>	A03	89.21	0.269	0.566	0.492
<i>G.capensis</i>	A06	95.99	0.270	0.599	0.490
<i>G.capensis</i>	A19	78.05	0.281	0.584	0.515
<i>G.capensis</i>	A35	164.72	0.270	0.572	0.485
<i>G.capensis</i>	A40	166.18	0.283	0.543	0.536
<i>G.capensis</i>	A43	176.27	0.265	0.632	0.508
<i>G.capensis</i>	A51	163.39	0.292	0.581	0.507
<i>G.capensis</i>	A95	175.86	0.273	0.578	0.500
<i>G.capensis</i>	A97	277.75	0.276	0.634	0.496
<i>G.capensis</i>	A99	284.02	0.281	0.587	0.522
<i>G.capensis</i>	A100	274.39	0.288	0.569	0.544
<i>G.capensis</i>	A115	283.3	0.258	0.613	0.533
<i>G.capensis</i>	A118	272.63	0.263	0.595	0.537
<i>G.capensis</i>	A128	177.39	0.260	0.618	0.484

<i>G.capensis</i>	A130	168.25	0.273	0.578	0.532
<i>G.capensis</i>	A131	100.42	0.281	0.606	0.481
<i>H.argenteocinereus</i>	526	101.4	0.311	0.631	0.530
<i>H.argenteocinereus</i>	26	122	0.293	0.621	0.526
<i>H.argenteocinereus</i>	11	129	0.318	0.631	0.537
<i>H.argenteocinereus</i>	387	130	0.307	0.599	0.487
<i>H.argenteocinereus</i>	27	135	0.289	0.590	0.505
<i>H.argenteocinereus</i>	386	145	0.290	0.622	0.501
<i>H.argenteocinereus</i>	8	149	0.283	0.639	0.508
<i>H.argenteocinereus</i>	6	160	0.291	0.620	0.542
<i>H.argenteocinereus</i>	516	166.5	0.330	0.612	0.549
<i>H.argenteocinereus</i>	361	181	0.291	0.573	0.573
<i>H.argenteocinereus</i>	517	190	0.301	0.627	0.583
<i>H.argenteocinereus</i>	585	206	0.305	0.626	0.548
<i>H.argenteocinereus</i>	377	210	0.345	0.607	0.543
<i>H.argenteocinereus</i>	451	235.8	0.285	0.581	0.559
<i>H.argenteocinereus</i>	5	249	0.303	0.601	0.455
<i>H.argenteocinereus</i>	18	298	0.290	0.598	0.592
<i>H.argenteocinereus</i>	473	231	0.325	0.616	0.589
<i>H.argenteocinereus</i>	476	137.6	0.298	0.637	0.579
<i>H.argenteocinereus</i>	508	225.4	0.352	0.612	0.549
<i>H.argenteocinereus</i>	525	112	0.297	0.598	0.539
<i>H.argenteocinereus</i>	492	138.4	0.318	0.625	0.548
<i>H.argenteocinereus</i>	479	176.2	0.293	0.648	0.550
<i>H.argenteocinereus</i>	244	207.5	0.303	0.600	0.528
<i>H.argenteocinereus</i>	590	251.5	0.326	0.607	0.517
<i>H.argenteocinereus</i>	241	100.2	0.298	0.597	0.497
<i>H.argenteocinereus</i>	242	107.3	0.317	0.611	0.480
<i>H.argenteocinereus</i>	243	218.24	0.281	0.630	0.509
<i>H.argenteocinereus</i>	244	187	0.321	0.584	0.454
<i>C.hottentotus</i>	GM102	51	0.255	0.613	0.483
<i>C.hottentotus</i>	GM103	65	0.247	0.582	0.546
<i>C.hottentotus</i>	GM104	78	0.253	0.611	0.511
<i>C.hottentotus</i>	GM105	78	0.277	0.602	0.515
<i>C.hottentotus</i>	GM106	98	0.291	0.602	0.524
<i>C.hottentotus</i>	GM108	62	0.278	0.569	0.510
<i>C.hottentotus</i>	GM109	110	0.283	0.587	0.539
<i>C.hottentotus</i>	GM110	54	0.259	0.605	0.540
<i>C.hottentotus</i>	GM111	68	0.267	0.612	0.478
<i>C.hottentotus</i>	GM112	61	0.274	0.649	0.519
<i>C.hottentotus</i>	GM114	73	0.291	0.638	0.545
<i>C.hottentotus</i>	GM115	52	0.265	0.564	0.509
<i>C.hottentotus</i>	GM116	89	0.290	0.640	0.554
<i>C.hottentotus</i>	GM117	72	0.283	0.572	0.543
<i>C.hottentotus</i>	GM122	77	0.301	0.567	0.480

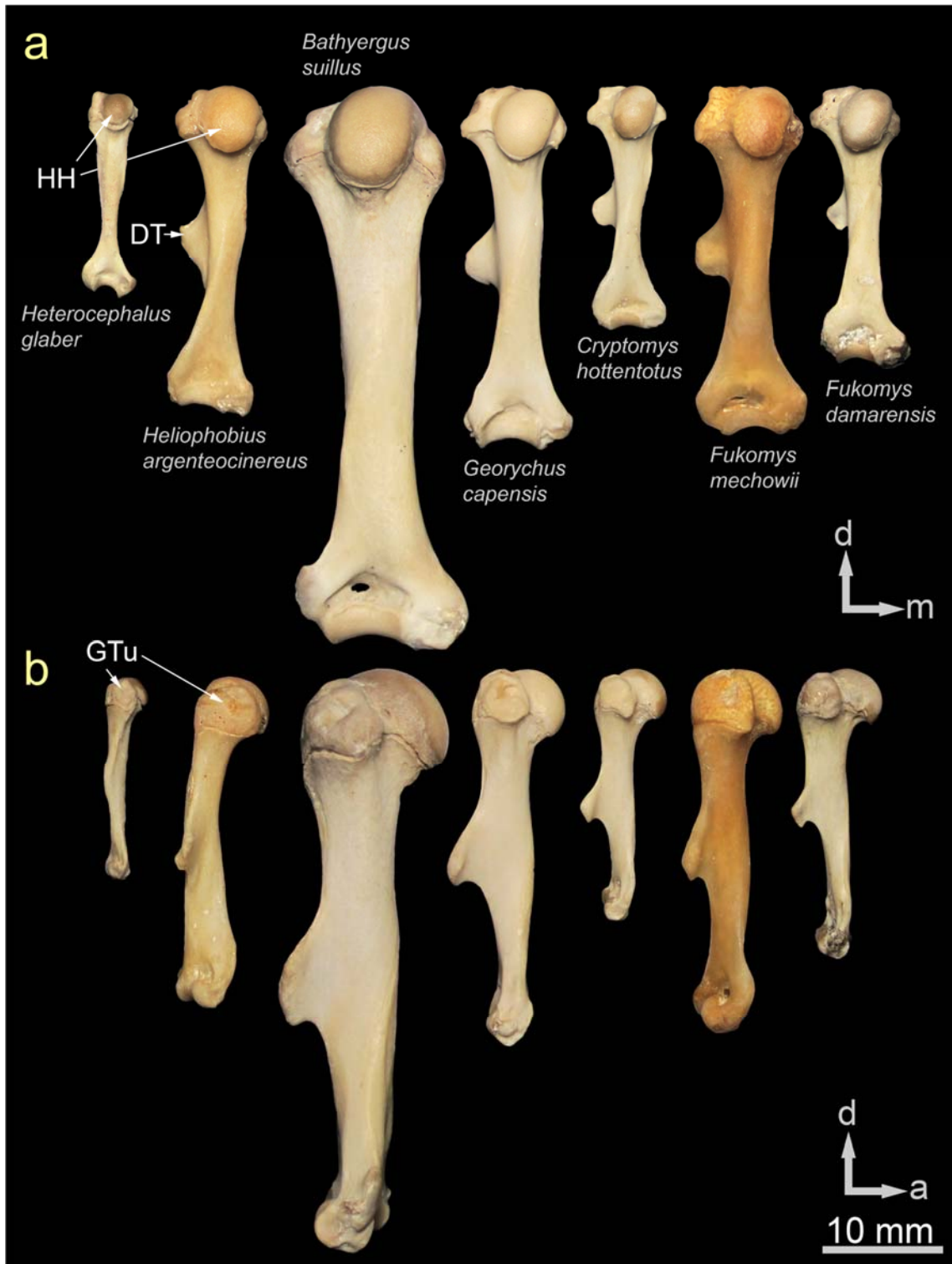
<i>C.hottentotus</i>	GM123	102	0.278	0.595	0.522
<i>C.hottentotus</i>	GM126	67	0.294	0.565	0.563
<i>C.hottentotus</i>	GM127	61	0.290	0.584	0.587
<i>C.hottentotus</i>	GM128	80	0.303	0.613	0.513
<i>C.hottentotus</i>	GM129	130	0.300	0.618	0.569
<i>C.hottentotus</i>	GM136	60.53	0.293	0.581	0.474
<i>C.hottentotus</i>	GM137	76	0.275	0.570	0.509
<i>C.hottentotus</i>	GM138	84	0.296	0.595	0.524
<i>C.hottentotus</i>	GM139	53	0.300	0.588	0.539
<i>C.hottentotus</i>	GM140	54	0.270	0.564	0.527
<i>C.hottentotus</i>	GM142	47	0.276	0.561	0.555
<i>C.hottentotus</i>	GM143	57	0.256	0.571	0.532
<i>C.hottentotus</i>	GM144	71	0.305	0.594	0.528
<i>C.hottentotus</i>	GM145	92	0.260	0.552	0.561
<i>C.hottentotus</i>	GM146	50	0.250	0.607	0.510
<i>C.hottentotus</i>	GM147	68.15	0.263	0.578	0.513
<i>C.hottentotus</i>	GM150	69	0.255	0.571	0.555
<i>C.hottentotus</i>	GM151	70	0.287	0.540	0.533
<i>C.hottentotus</i>	GM152	68	0.256	0.601	0.510
<i>C.hottentotus</i>	GM153	80	0.286	0.611	0.516
<i>F.damarensis</i>	G4F042	87	0.274	0.540	0.506
<i>F.damarensis</i>	G4M018	148	0.260	0.533	0.493
<i>F.damarensis</i>	G4M034	155	0.299	0.563	0.528
<i>F.damarensis</i>	G4M039	141	0.286	0.559	0.497
<i>F.damarensis</i>	G4M041	71	0.281	0.594	0.533
<i>F.damarensis</i>	G4M008	56	0.251	0.530	0.516
<i>F.damarensis</i>	G5F003	71	0.295	0.530	0.561
<i>F.damarensis</i>	L29F002	87	0.299	0.536	0.514
<i>F.damarensis</i>	G4F012	128	0.265	0.529	0.564
<i>F.damarensis</i>	G4F002	133	0.299	0.535	0.553
<i>F.damarensis</i>	G4M009	141	0.318	0.552	0.528
<i>F.damarensis</i>	G3M007	142	0.278	0.533	0.532
<i>F.damarensis</i>	G4M001	147	0.308	0.529	0.536
<i>F.damarensis</i>	G3M001	173	0.298	0.546	0.517
<i>F.damarensis</i>	Z3M006	226	0.305	0.545	0.534
<i>F.damarensis</i>	NB227	67.46	0.278	0.546	0.553
<i>F.damarensis</i>	NB228	63.48	0.256	0.559	0.574
<i>F.damarensis</i>	GAP315	113	0.303	0.567	
<i>F.mechowii</i>	23	133	0.302	0.547	0.535
<i>F.mechowii</i>	179	139	0.325	0.545	0.529
<i>F.mechowii</i>	3	144	0.314	0.538	0.548
<i>F.mechowii</i>	1	158	0.304	0.559	0.536
<i>F.mechowii</i>	177	232	0.278	0.570	0.578
<i>F.mechowii</i>	497	268	0.318	0.592	0.474
<i>F.mechowii</i>	478	380.7	0.303	0.566	0.556

<i>F.mechowii</i>	494	208.8	0.300	0.569	0.501
<i>F.mechowii</i>	589	277.8	0.311	0.557	0.524
<i>F.mechowii</i>	242	405.5	0.295	0.567	0.564
<i>F.mechowii</i>	2	193.4	0.260	0.532	0.590
<i>F.mechowii</i>	531	262.2	0.321	0.570	0.554
<i>H.glaber</i>	JJ-001	32.29	0.245		
<i>H.glaber</i>	JJ-002	14.96	0.246		
<i>H.glaber</i>	JJ-045	56.38	0.256		
<i>H.glaber</i>	JJ-046	16.41	0.265		
<i>H.glaber</i>	1200-047	14.9	0.246		
<i>H.glaber</i>	1200-048	29.84	0.251		
<i>H.glaber</i>	1200-049	19.82	0.288		
<i>H.glaber</i>	1200-050	20.82	0.253		
<i>H.glaber</i>	1200-051	22.41	0.296		
<i>H.glaber</i>	1200-052	16.37	0.267		
<i>H.glaber</i>	1200-053	32.14	0.281		
<i>H.glaber</i>	1200-054	31.5	0.267		
<i>H.glaber</i>	1200-055	26.82	0.260		
<i>H.glaber</i>	1200-056	24.027	0.222		
<i>H.glaber</i>	1200-057	32.47	0.269		
<i>H.glaber</i>	1200-058	30.24	0.279		
<i>H.glaber</i>	1200-059	17.72	0.279		
<i>H.glaber</i>	1200-060	35.81	0.297		
<i>H.glaber</i>	1200-061	39.91	0.247		
<i>H.glaber</i>	1200-063	22.85	0.280		
<i>H.glaber</i>	1200-064	31.29	0.278		
<i>H.glaber</i>	1200-065	32.71	0.287		
<i>H.glaber</i>	1200-066	19.08	0.261		
<i>H.glaber</i>	1200-067	19.72	0.282		
<i>H.glaber</i>	1200-068	23.45			
<i>H.glaber</i>	1200-069	15.29	0.300		
<i>H.glaber</i>	1000-070	31.94	0.275		
<i>H.glaber</i>	1000-071	34.58	0.273		
<i>H.glaber</i>	1000-072	26.77	0.268		
<i>H.glaber</i>	1000-073	28.56	0.281		
<i>H.glaber</i>	1000-074	26.44	0.291		
<i>H.glaber</i>	1000-075	23.95	0.262		
<i>H.glaber</i>	1000-076	29.54	0.258		
<i>H.glaber</i>	1000-077	44.58	0.320		
<i>H.glaber</i>	1000-078	36.08	0.255		
<i>H.glaber</i>	1000-079	23.88	0.271		
<i>H.glaber</i>	1000-080	41.44	0.294		
<i>H.glaber</i>	1000-081	34.45	0.281		
<i>H.glaber</i>	1000-082	43.97	0.302		
<i>H.glaber</i>	1000-083	23.43	0.289		

<i>H.glaber</i>	1000-084	34.49	0.263		
<i>H.glaber</i>	1000-085	27.19	0.254		
<i>H.glaber</i>	1000-086	28.72	0.275		
<i>H.glaber</i>	5000-087	31.06	0.266		
<i>H.glaber</i>	5000-088	22.57	0.264		
<i>H.glaber</i>	5000-089	22.3	0.277		
<i>H.glaber</i>	5000-090	35.17	0.297		
<i>H.glaber</i>	5000-091	28.51	0.259		
<i>H.glaber</i>	5000-092	34.4	0.293		
<i>H.glaber</i>	5000-093	30.4	0.258		
<i>H.glaber</i>	5000-094	33.83	0.275		
<i>H.glaber</i>	5000-095	28.86	0.261		
<i>H.glaber</i>	5000-096	24.63	0.215		
<i>H.glaber</i>	5000-097	31.13	0.292		
<i>H.glaber</i>	5000-098	29.36	0.276		
<i>H.glaber</i>	5000-099	29.82	0.273		
<i>H.glaber</i>	5000-100	21.64	0.297		
<i>H.glaber</i>	5000-101	19.21	0.261		
<i>H.glaber</i>	GM420	22.92	0.264		
<i>H.glaber</i>	GM498	47	0.247		
<i>H.glaber</i>	GM499	66	0.293		
<i>H.glaber</i>	GM500	17.93	0.259		
<i>H.glaber</i>	GM501	63	0.265		
<i>H.glaber</i>	GM505	19.58	0.282		
<i>H.glaber</i>	GM506	16.04	0.263		
<i>H.glaber</i>	GM507	41	0.270		
<i>H.glaber</i>	GM508	35	0.245		
<i>H.glaber</i>	GM509	39	0.272		
<i>H.glaber</i>	GM510	30	0.244		
<i>H.glaber</i>	GM511	33	0.257		

**SUPPLEMENTARY FIGURES**

**Supplementary Figure 1.** Humeral phenotype of African mole-rats analyzed here. a) Posterior view. b) Lateral view. Bones ordered same as in “a”. Abbreviations: anterior (a); deltoid tuberosity (DT); dorsal (d), greater tubercle (GTu); humeral head (HH); medial (m).



## SUPPLEMENTARY METHODS

### Ontogenetic stages

Individuals were classified as newborns, pups, and adults based on chronological age, patterns of molar eruption and tooth wear, and degree of skeletal development (Supplementary Table 1). The ontogeny of *H. argenteocinereus* and *F. mechowii* was based on chronological age. [Gomes Rodrigues et al. \(76\)](#) described that the third molar of *H. argenteocinereus* erupts after the second month of life and the full cheek teeth (four molars) are already developed at two years old, so that individuals older than two years in this study were considered adults and assumed to have developed the fourth molar. Similar classification was used for *F. mechowii*. Newborns are one day old and pups are a few days old. For the rest of the species, individuals having full eruption of all upper or lower molars were considered adults. Almost all genera have four molars (77-80), although *Heterocephalus* has three (sometimes two) (11,81) and some populations of *Heliophobius argenteocinereus* exhibit 4-7 molars and continuous dental replacement (76,82,83). Determination of molar eruption in *Bathyergus* was based on [Hart et al. \(79\)](#) and [Montoya-Sanhueza et al. \(84\)](#): age-classes 5-9 are considered as adults. The classification of *F. damarensis* was based on patterns of molar eruption described for *Fukomys mechowii*: age-classes 5-9 were considered as adults (80). In *Cryptomys hottentotus*, individuals of age-class 3-4 are considered as adults (78). Individuals of *H. glaber* with full molar eruption were considered adults (85). For younger ontogenetic stages (pups), other anatomical features indicating skeletal immaturity were used, including poor development of secondary centers of ossification, presence of chondroepiphyses and unfused distal epiphysis in the humerus (84). Perinatal individuals of unknown age were determined as newborns/pups based on their very small size and incipient skeletal development.

### Bone superstructures

Presence/absence of bone superstructures are presented in Supplementary Table 1. Among the bone superstructures analyzed, the projected and distally located DT is one of the most extensively studied adaptations among fossorial animals, which is principally adapted to accommodate enlarged pectoral and deltoid muscles to increase the power-stroke (retraction) of the arm (Supplementary Table 5). This structure is often referred to in multiple ways, such as tubercle, crest, ridge or process, although proper definition of such terms and the differences among them are rather ambiguous. We define the deltoid tuberosity (= tubercle/process) as a localized and conspicuous protrusion of bone projecting from the diaphysis, with a variable location and size in the diaphysis among mammals. The DT differs from a deltoid crest (DC) (= ridge), which represents an extended bony surface along the diaphysis associated with a wide muscular attachment, but usually not forming a localized protuberance. The DC is observed in some taxa such as tubulidentates, hystricids, tenrecids and solenodontids (34,86). Such structures are not mutually exclusive and a DC can appear at the proximal region of the diaphysis (below the greater tubercle) and extend towards the middle part of the diaphysis fusing with the DT around the midshaft (24), as occurs in hystricids. Two conditions for the DT were recorded in this study: i) the humerus has a smooth diaphyseal surface lacking a projected DT (but not precluding of showing a reduced DT, a small scar for the attachment of the *mm. deltoidei* or a DC); and ii) the humerus has a diaphysis with a conspicuously localized and projected DT.

Regarding the DFTFi, this character has typically been associated with fossoriality, although it is not exclusive to fossorial taxa. [Barnett and Napier \(19\)](#) analyzed the type of tibio-fibular articulation of a wide variety of mammals and found that the fibula of fossorial and aquatic mammals is immobile and united to the tibia at its upper and lower ends by bone (or fibrous) tissue, thus conferring increased robustness and rigidity (Supplementary Table 5). Two conditions were measured: i) tibia and fibula are not distally fused (non-ossified); and ii) tibia and fibula are distally fused and ossified.

Regarding the OP and TT, these features are easily recognizable bony projections, the first located at the proximal ulna, beginning at the base of the anconeal process, and the second located in the proximal femur between the greater trochanter and the midshaft region. Two conditions for these features were recorded, presence or absence, regardless of their cartilaginous or ossified tissue condition.

### Morpho-functional indices

Seven linear measurements were used to calculate morpho-functional indices: total lengths of humerus (HL), ulna (UL) and tibia-fibula (TL); length of the deltoid tuberosity (DLH); length of the olecranon process (OL), functional length of

the ulna (FUL) and length of the distal tibio-fibular junction (DTFJ). Total bone lengths are the maximum distance from the proximal articular surface to the distal articular surface. OL is the length from the tip of the olecranon to the center of the trochlear notch. FUL is the difference between UL and OL. DLH and DTFJ were measured from the proximal articular surface of the bone to the distal origin of the deltoid tuberosity and to the tibio-fibular junction, respectively. All measurements were recorded to the nearest 0.01 mm using a digital caliper. The relative position of the deltoid tuberosity (RDT = DLH/HL) and the index of fossorial ability (IFA = OL/FUL) were calculated following a previous study and references therein (84) (Supplementary Table 7). The tibio-fibular junction index (TJI = DTFJ/TL) is a novel ecomorphological index modified from Montoya-Sanhueza et al. (84) and implemented for the first time in this study to reflect the extension of the distal tibio-fibular fusion, so that lower values (i.e. a more proximal and larger fusion of the tibia and fibula) would suggest a longer bony base and more robust bone diaphysis to increase bone resistance to muscles acting on the paws.

### Phylogenetic varying effects regression

The amount of change in a given index, for example RDT, in response to a one unit increase in BM for individual  $i$  belonging to species  $j$ , was modelled as:

$$\begin{aligned} \log RDT_i &\sim \text{Normal}(\mu_i, \sigma) \\ \mu_i &= \alpha_{j[i]} + \beta_{j[i]} \log BM_i + \varepsilon_{j[i]} \\ \alpha_j &\sim \text{MVNormal}(\mu_\alpha, S_\alpha) \\ S_\alpha &= \sigma_{phy_\alpha} R \\ \beta_j &\sim \text{MVNormal}(\mu_\beta, S_\beta) \\ S_\beta &= \sigma_{phy_\beta} R \\ \varepsilon_j &\sim \text{Normal}(0, \sigma_{spp}) \end{aligned}$$

with intercepts  $\alpha_j$ , i.e. species mean phenotypes, and slopes  $\beta_j$ , i.e. species scaling effects, drawn from multivariate Gaussian distributions, where  $\mu_\alpha$  and  $\mu_\beta$  are intercept mean and slope mean, and  $S_\alpha$  and  $S_\beta$  are covariance matrices for species' intercepts and slopes, respectively. Covariance in both intercepts and slopes was defined as the product of a phylogenetic correlation matrix  $R$ , with expected correlations among species determined by an Ornstein-Uhlenbeck model of trait evolution (87,88), and variance parameter  $\sigma_{phy}$ , giving the magnitude of phylogenetic effect (89). Species-specific effects were captured by a vector of residuals, where each species  $j$  had its own residual  $\varepsilon_j$ , modeled as a normally distributed random variable with mean zero and standard deviation  $\sigma_{spp}$ , giving an overall magnitude of species-specific effects, or in other words, the variation unaccounted for by the phylogenetic component (90,91).

We assigned regularizing priors to all sampled parameters, which reduce the risk of overfitting the data and give more accurate predictions (92). Priors for the sampled parameters were drawn from the following distributions:

$$\begin{aligned} \mu_\alpha &\sim \text{Normal}(0, 0.5) \\ \mu_\beta &\sim \text{Normal}(0, 0.5) \\ \sigma &\sim \text{Exponential}(2) \\ \sigma_{phy_\alpha} &\sim \text{Exponential}(2) \\ \sigma_{phy_\beta} &\sim \text{Exponential}(2) \\ \sigma_{spp} &\sim \text{Exponential}(2) \end{aligned}$$

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