Reproduction triggers adaptive increases in body size in female mole-rats

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JT conceived the study with input from TCB and MZ. JT performed the statistical analyses. JT, NK, KG and MZ organised and/or carried out data collection. JT wrote the paper, with input from the other authors at later stages. All authors gave final approval for publication.

Abstract

In social mole-rats, breeding females are larger and more elongated than nonbreeding female helpers. The status-related morphological divergence is thought to arise from modifications of skeletal growth following the death or removal of the previous breeder and the transition of their successors from a nonbreeding to a breeding role. However, it is not clear what changes in growth are involved, whether they are stimulated by the relaxation of reproductive suppression or by changes in breeding status, or whether they are associated with fecundity increases. Here, we show that, in captive Damaraland mole-rats (Fukomys damarensis) where breeding was experimentally controlled in age-matched siblings, individuals changed in size and shape through a lengthening of the lumbar vertebrae when they began breeding. This skeletal remodelling results from changes in breeding status since i) females removed from a group setting and placed solitarily showed no increases in growth, and ii) females dispersing from natural groups that have not yet bred do not differ in size and shape from helpers in established groups. Growth patterns consequently resemble other social vertebrates where contrasts in size and shape follow acquisition of the breeding role. Our results also suggest that the increases in female body size provide fecundity benefits. Similar forms of socially responsive growth might be more prevalent in vertebrates than is currently recognised, but the extent to which this is the case, and the implications for the structuring of mammalian dominance hierarchies, is as yet poorly understood.

Keywords: Bathyergidae, Growth Plasticity, Morphological Skew, Strategic Growth, Reproductive suppression

Introduction

In several social vertebrates where a single dominant female monopolizes reproduction in each group, subordinates that acquire a dominant breeding position display an increase in their growth rate (fish [1–3], meerkats[4], naked mole-rats [5]). For example, in the social mole-rats (including the naked mole-rat *Heterocephalus glaber* and the Damaraland mole-rat *Fukomys damarensis*), dominant breeding females are both larger and more elongated than nonbreeding subordinates [6,7] and longitudinal studies of individuals show that subordinate females removed from established breeding groups and paired with novel partners increase in size and weight [5]. As body size and weight typically confer competitive advantages, increases in growth in newly dominant individuals may help to consolidate their position and to increase their fecundity, with parallels being drawn between the elongated phenotype of female breeders in mole-rats and the physogastry (enlargement of the abdomen through increasing numbers of ovarioles) observed in queens of some eusocial insect societies [6–8].

While the presence of status-related changes in growth in social mole-rats is well established, their immediate causes are still uncertain, with studies implicating different growth patterns in naked mole-rats and Damaraland mole-rats. In naked mole-rats, longitudinal X-ray sampling of captive newly created breeders revealed upregulated growth of the lumbar vertebrae [5,6] and episodic bursts of vertebral growth in successive periods of pregnancy [9]. This causes breeding females to exhibit a longer body length relative to their skull width. The same status-related morphological difference is seen in Damaraland mole-rats, but repeated measurements of wild females suggested that breeder elongation is not driven by upregulated vertebral growth [7] like in naked mole-rats. Instead it appeared to originate from a decrease in the relative growth of the skull (zygomatic arch width) compared to growth towards total body length [7]. This result prompted the idea that female

Damaraland mole-rats reallocate resources from growth towards reproduction as they become dominant; during this reallocation, skull growth is reduced but growth towards body length is maintained because of the inherent fitness benefits of increased body length [7]. The focus on growth reductions in skull size is particularly pertinent in this argument as the skull of molerats is home to their prominent buccal incisors that are so important for soil excavation (involved in foraging and burrow maintenance), and as a result, reproductive investment in the form of elongation can be argued to trade off directly against investment into work.

As sociality arose independently in naked and Damaraland mole-rats [10], it is plausible that the attainment of elongation (which is assumed to be directly related to their sociality) occurs through alternate developmental routes. However, as the prior analysis of Damaraland mole-rats was based on human measurements of morphological traits taken with callipers and a tape measure, a formal characterisation of morphological divergence at a skeletal level is currently missing. Moreover, although Young and Bennett [7] go to great lengths to remove the possibility that status-related age differences underpin skeletal divergence - because changes in morphology only occurred in females after transitioning to dominance, and because the opportunity to acquire dominance is somewhat stochastic- a contribution of age to the shape and size of breeders has not been definitively ruled out as individuals in the wild were of unknown age.

It is also unclear whether the changes in growth in female Damaraland mole-rats that acquire the breeding position in their group are stimulated by the relaxation of reproductive suppression by the previous dominant female or by the onset of reproduction itself. The degree of reproductive suppression in the social mole-rats is extreme, manifesting itself in a complete blocking of ovulation in nonbreeding Damaraland mole-rats [11]. Even so, the introduction of an unrelated male results in a recrudescence of ovarian activity in nonbreeders [12,13] and stimulates high levels of aggression between females that sometimes leads to the

usurpation of the incumbent breeder [14] (see [15,16] for comparable results in naked molerats). Nonbreeding females also start ovulating in the absence of breeding females [11,17], and in the wild, the reproductive readiness of nonbreeders - measured by the downstream production of luteinising hormone following injection of pituitary gonadotrophin-releasing hormone - is elevated during periods of high rainfall when the likelihood of meeting dispersing males or of dispersing oneself is higher [18]. These physiological changes in nonbreeders in anticipation of reproductive opportunities share similarities with the onset of puberty in other mammals, where sex steroid secretion is a major driver of skeletal longitudinal and radial growth [19], and as such, the removal from reproductive suppression and resultant hormonal changes may stimulate the morphological divergence of nonbreeding Damaraland mole-rats towards a more elongated phenotype. On the other hand, studies of classical rodent lab models highlight that many of the most pronounced changes in skeletal remodelling occur during pregnancy [20,21], and in naked mole-rats, housing females in isolation did not cause them to lengthen their vertebrae, whereas pairing them with a receptive male (with resultant pregnancy) did [6]. Based on these studies, reproduction itself may provide the necessary cue for the skeletal remodelling of Damaraland mole-rat females.

In this study, we used information from X-rays to characterise the morphological divergence of breeders and nonbreeders in Damaraland mole-rats according to three principal aims: 1) to identify the skeletal changes that lead to increases in female size, 2) to identify the precise circumstances that stimulate growth adjustment in females, and 3) to investigate whether growth adjustments are associated with increases in fecundity. To identify the skeletal differences between breeders and nonbreeders, we first carried out cross-sectional comparisons of morphology in captive and wild Damaraland mole-rats. In addition, we experimentally manipulated the life history trajectories of female siblings in captivity to track skeletal development longitudinally within age-matched individuals. By keeping some

females as nonbreeders within their natal group, isolating others by placing them in their own tunnel system, and pairing others with an unrelated male to initiate reproduction, we were able to determine whether removal from reproductive suppression is involved in changes in morphology, whilst controlling for the influence of age on development. If females placed in isolation display growth patterns analogous to newly reproductive females, this would provide strong evidence that the elongation of subordinate females in breeding groups is hindered by reproductive suppression. A similar argument extends to wild females that have dispersed and settled solitarily but have yet to reproduce, so we also compared the morphology of solitary females to breeders and in-group nonbreeders from a wild population of mole-rats. Lastly, we investigated the fecundity implications of increasing body size in breeders using correlative data from litters born in captivity.

Methods:

Study species and X-ray methodology

Our study used information from X-rays taken on both captive and wild Damaraland mole-rats. The wild study population of mole-rats is located around the Kuruman River Reserve in the Northern Cape of South Africa (S26.98706° E21.81229°), with individuals from this population founding a captive study system based at the same location in February 2013. The details of the captive and wild populations are presented in full in the supplementary information (see also [22]). The X-ray data from captive animals were collected between November 2015 and July 2017, and from wild animals between February 2015 and June 2017.

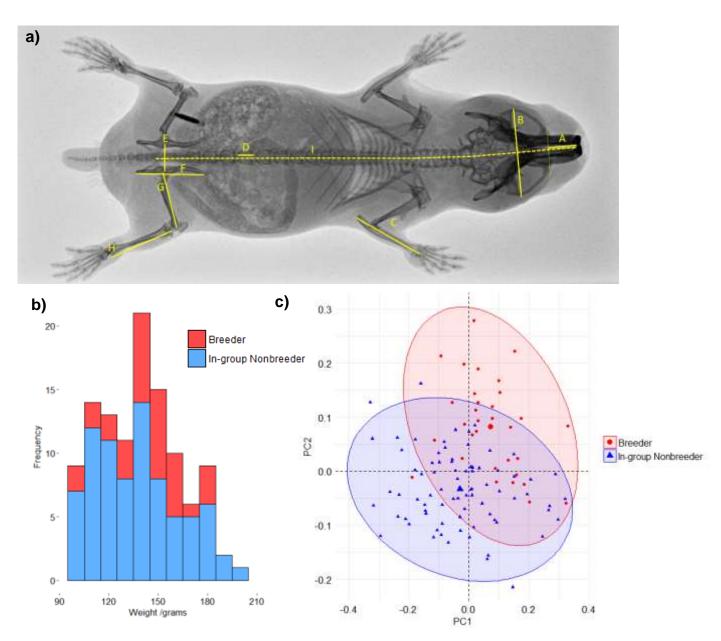


Figure 1a) Dorsoventral X-ray annotated with the 9 morphological traits used in multivariate analysis. A: rostrum length, B: skull width, C: ulna, D: L5 vertebra length, E: pelvic girdle width, F: pelvis length, G: femur length, H: tibia length, I: skeletal body length. b) Weight distribution of captive breeding and nonbreeding in-group females in multivariate analysis. c) PC1 and PC2 separates captive females according to their social status.

X-rays were taken using the Gierth TR 90/20 battery-operated generator unit with portable Leonardo DR Mini plate (OR Technology, Rostock, Germany) under protocols approved by the University of Pretoria ethics committee. For each X-ray, mole-rats were immobilised under isoflurane anaesthesia and gently positioned in a dorsoventral position

with straightened spine and splayed limbs. 9 skeletal traits were measured from each X-ray (Figure 1) using ImageJ software. As in studies that have examined morphology of naked mole-rats, the length of a lumbar vertebra (L5 in our case) served as an index of lumbar length [5,9,23]. Similarly, the log ratio of the L5 vertebra length to zygomatic arch width served as a metric of elongation ('elongation factor' below). Hereafter we refer to zygomatic arch width as skull width. All animals were weighed after anaesthesia. All statistical tests were performed in R version 3.2.3 [24], and standard model assumptions (normal errors and homogeneity of residual variance) were checked throughout.

Morphological divergence

To quantify the morphological divergence of breeders and nonbreeders we analysed a cross-sectional dataset of X-rays taken in captivity. The cross-sectional analysis from captive females was restricted to individuals larger than 100g, as the smallest reproductive female in the captive population was 100g. This resulted in data from 32 breeders and 79 in-group nonbreeders spanning 40 different groups, where group refers either to the group of original capture or group of birth. All 9 skeletal traits were included in a standard principal component analysis (PCA), and a multivariate analysis of variance (MANOVA) was performed on the resulting principal components to test for broad morphological differences between the two classes of female. Follow-up univariate ANOVAs on each of the first five principal components were used to determine on which of the principal components reproductive status was exerting its influence.

To examine whether the bivariate scaling relationships of morphological traits differed with breeding status, we fitted a series of linear models of the form $\log_e(\text{trait}_1) \sim \log_e(\text{trait}_2)*\text{Status}$, where trait_2 represents either the skull width or skeletal body length, two metrics of size. In these models, a significant interaction term denotes significantly different

slope between breeders and in-group nonbreeders. When the slopes did not differ, we tested for a difference in intercept by removing the interaction term from each model: $\log_e(\text{trait}_1) \sim \log_e(\text{trait}_2) + \text{Status}$.

Skeletal changes

To further identify the skeletal changes that lead to increases in female size and to determine the circumstances leading to growth adjustment, we manipulated the life history trajectory of 30 natal sisters born in captivity by altering their social status, and tracked their development longitudinally (originally 32 females, but 2 died shortly after pairing and were excluded throughout). These females came from fourteen litters and were of known age, which also removed any possible confounding effect of age that could have been present in the cross-sectional analysis of captive mole-rats. Females were randomly allocated to one of three treatments: remaining in their natal group as a nonbreeder, placed in a new artificial tunnel system as a solitary nonbreeder, or paired with an unfamiliar male in a new tunnel system to become a breeder. X-rays were taken on these females at 2-month intervals from the point of treatment initiation for 12 months (except in a few cases where we deemed that females were too heavily pregnant to be anaesthetised and X-rayed). Of the females that survived to eight months (n = 28), the mean age at manipulation was 526.8 ± 34.18 , $536.45 \pm$ 30.91 and 526.9 \pm 36.78 days for in-group nonbreeders, solitary nonbreeders, and breeders respectively. Of the 11 females that were paired with a male, only two had failed to produce a litter by 6 months of age, and only one individual by 10 months. The median time to first parturition after pairing was 101.5 days. All females in the "breeder" treatment were included in all analyses.

General linear models were used to investigate skeletal growth trajectories across the three social treatments. The within-individual change in L5 lumbar vertebra length, skull

width, and the elongation factor were each fit as a response, with treatment and the initial trait value specified as fixed covariates. Models were fitted for each response variable at every 2-month sampling interval to determine the point at which skeletal morphological divergence occurred. Pairwise comparisons of significant treatment effects were assessed with Tukey's multiple comparisons (*multcomp* package[25]), and similarities/differences between breeders and nonbreeders were used to assess whether growth adjustment is a result of breeding itself (in which case breeder morphology would differ from both classes of nonbreeding female) or rather due to the relaxation of reproductive suppression (in which case breeder morphology would only differ from females remaining in their natal group).

To control for a possible artefact of captivity on morphological patterns, we also examined X-rays from wild mole-rats. X-rays from wild animals only included females heavier than 101g, the weight of the smallest breeder that was captured for X-ray sampling. This produced a dataset of 56 females captured between February 2016 and June 2017, including 21 breeders, 12 in-group nonbreeders, and 23 solitary nonbreeders. The presence of an unperforated vagina confirmed that these solitary females had not previously engaged in sexual activities, and previous trapping records indicated that many of these solitary nonbreeding females had been solitary for at least two years (and in a few cases four years). The relative elongation (log L5 vertebra / log skull width) of the three classes of female was fitted in a linear mixed effects model with normal errors, with group identity set as a random effect to control for possible morphological similarities within groups. Pairwise comparisons were treated as above.

Body length and fitness

To investigate whether increases in body length in breeders are associated with increases in fecundity we used an extensive dataset of birth events in captivity. The total body length of female breeders is measured during routine sampling in the lab, which made it possible to investigate whether body length was related to three measures of fecundity among a large cohort of breeders: litter size at birth, the total mass of pups at birth (total neonate mass), and individual pup mass at birth. Information was only included from females whose total body length was measured within 90 days of the birth of a litter to ensure that it reflects size around parturition. Further, to remove any uncertainty around birth date, litters were only used if checks of the nest box indicated that the litter must have been born on the day of the check or the day preceding it. The total dataset included 186 litters born to 58 mothers that produced 587 pups. Pups were measured to the nearest gram on an electronic balance. The three traits were fitted to linear mixed effects models, where total neonate mass and individual pup mass were fitted to a normal error structure, and litter size was fitted to a Poisson error structure. For total neonate mass and litter size, a single model was fitted in each case that included breeder body length and whether it was the females first reproductive episode (primiparity) as fixed terms, and maternal identity as a random term. For individual pup mass, the model contained breeder body length, primiparity and litter size as fixed terms, and maternal identity and litter identity as random terms; including litter size allows us to test whether longer mothers produced relatively larger pups; partial residuals of body length were extracted using the *remef* package [26] before being plotted. The significance of fixed terms was assessed by likelihood ratio tests.

To check that any morphological pattern in the results could not be driven by a bias in the X-ray annotation by the lead author, a random subset of 150 X-ray images that formed part of the study were annotated blind by a second person unrelated to the study. The correlation

between the measurements across skeletal traits was consistently high (r > 0.93, except for pelvis length, where r = 0.497), and so it was highly improbable that measurer bias affected the results.

Results

Morphological divergence

In captive Damaraland mole-rats, breeding females are both larger and more elongated than subordinate nonbreeders residing in groups, after controlling for the weight of individuals (Figure 1b; breeder mean = 143.47g \pm 3.56 vs in-group nonbreeder mean 137.91g \pm 2.78; Welch's t-test, $t_{70.1}$ = -1.23, p = 0.22): principal component analysis revealed a morphometric separation of breeders and in-group nonbreeders according to both size and shape (Figure 1c; trait loadings for the first 5 PCs, which together explain 91.8% of the variance, are in Table S1). There was considerable overlap in the morphological space occupied by either class. PC1 revealed positive loading for all traits and is indicative of general size, breeders being generally larger than in-group females. PC2 separated breeders and in-group nonbreeders by shape, and reflects the relatively longer lumbar vertebra, longer skeletal body length, wider pelvis, and shorter femurs of breeders. The MANOVA performed on PCs 1 – 5 revealed a significant effect of social status ($F_{5,104}$ = 26.48, p < 0.001), with follow-up univariate ANOVAs on each PC 1 – 5 suggesting the status effect is most prominent in the first three components (PC1: F_1 = 11.04, p < 0.001; PC2: F_1 = 57.1, p < 0.001; PC3: F_1 = 7.43, p = 0.007; PC4: F_1 = 2.80, p = 0.097; PC5: F_1 = 2.87, p = 0.093).

Breeders in captivity were more elongated than subordinate nonbreeders, as shown by their relatively longer lumbar vertebrae for a given size (skull width, Figure 2a, or skeletal body length, Table S2), as well as being longer overall (total body length: breeder mean = 18.51 ± 0.14 cm, in-group nonbreeder mean 17.72 ± 0.11 cm, Welch's t-test, $t_{64.3} = -4.36$, p <

0.001). Breeders and in-group nonbreeders also showed different bivariate scaling relationships in several other skeletal traits (all bivariate relationships in Table S2), most notable being the wider pelvic girdle of breeders.

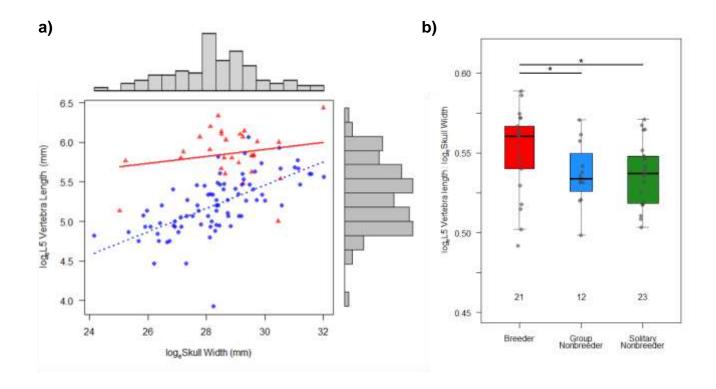


Figure 2 a) The bivariate relationship between the lumbar vertebra length and skull width in captive female mole-rats indicates that breeders (triangles) are relatively more elongated than in-group nonbreeders (circles). b) Breeding females are also more elongated than both in-group females and solitary females in the wild, the latter having dispersed and settled in isolation. Numbers within the plot refer to sample size.

Skeletal changes in breeders

Experimental manipulation of the social status of natal sisters of known age in captive animals showed that the elongation of breeding females was caused by an increased lengthening of the skeletal vertebrae (Figure 3). These increases in growth are a consequence of breeding rather than of the relaxation of reproductive suppression, since females removed from a group setting and placed solitarily displayed a similar growth pattern to nonbreeding

females residing in groups. Females paired with unfamiliar males to become breeders were already more elongated than the two classes of nonbreeding female four months into the experiment (Figure 3a; F_2 = 11.31, both pairwise contrasts p < 0.001). This coincided with the point at which the L5 vertebra of breeders were significantly longer (Figure 3b displays the contrasts in L5 vertebra length at 12 months; LMM contrast = 0.251 mm \pm 0.05 relative to ingroup breeders, p < 0.001, contrast = 0.014mm \pm 0.06, p = 0.025), an effect that persisted thereafter (Figure S1a displays contrasts at each sampling interval). The lengthening effect in breeders was also shown by other lumbar vertebrae (see Figure S2 for growth in L4 and L6 vertebrae). In contrast, there was no evidence that changes in skull width contributed significantly to the relative elongation of breeders (Figure 3c and Figure S1b; p > 0.10 for all pairwise contrasts).

Evidence from wild mole-rats agreed with the role of reproduction in breeder elongation. Here too, breeders displayed a more elongated phenotype than female nonbreeders residing in groups (Figure 2b; elongation factor contrast = 0.013 ± 0.005 , p = 0.027) and solitary nonbreeders (contrast = 0.014 ± 0.006 , p = 0.036), whereas the two classes of nonbreeder did not differ from one another (contrast = -0.0001 ± 0.006 , p = 0.89). Breeders (mean = 18.81 ± 0.17 cm) also exhibited a longer total body length than both solitary nonbreeders (mean 17.86cm ± 0.36 cm; contrast p value < 0.001) and in-group nonbreeders (mean 17.81cm ± 0.27 cm; contrast p value = 0.052).

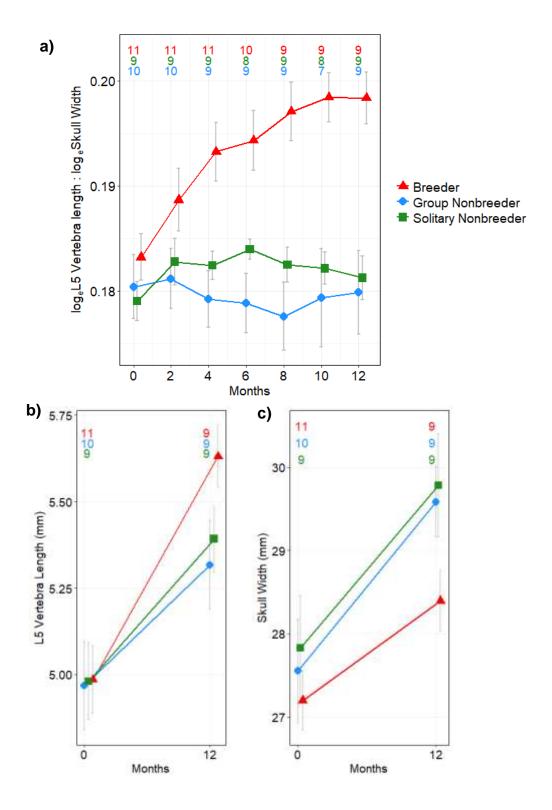


Figure 3: a) Change in the relative elongation b) L5 lumbar vertebra length and c) skull width of captive females experimentally manipulated to follow different social trajectories. Numbers within the plot refer to sample sizes at each timepoint of the experiment.

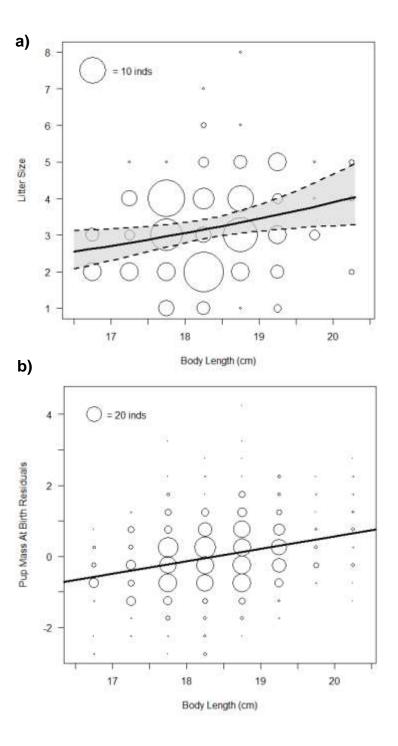


Figure 4: Increases in body length are associated with increases in a) litter size and b) individual pup mass across 186 litters born in captivity. a) Points represent raw data scaled to according to sample size, and lines show predicted litter size \pm 95 confidence intervals from linear mixed effects models. b) Partial residuals of individual pup mass at birth, corrected for litter size and primiparity. Points represent partial residuals estimated from linear mixed effects model. Line displays the partial effect of body length on individual pup mass, but confidence intervals are not provided as the generation of a partial effect removes variation from the fitted model. The partial residuals were standardised as they do not provide meaningful values.

Body Length and fitness

Across breeding females increased body length is associated with higher fecundity (Table S3) as longer females have larger litters (Figure 4a; GLMM: $\beta = 0.12 \pm 0.05$, $\chi^2_1 = 5.81$, p = 0.016). Since litter size is highly correlated with total neonate mass (r = 0.92), longer females also produced litters with larger total mass (LMM: $\beta = 3.83 \pm 1.23$, $\chi^2_1 = 8.89$, p = 0.003). However, longer females do not trade off increasing offspring quantity for reduced quality offspring. On the contrary, controlling for litter size, longer females invest proportionally more per pup (Figure 4b, LMM: $\beta = 0.36 \pm 0.17$, $\chi^2_1 = 4.12$, p = 0.040), which suggests that increases in body length may facilitate increases in both offspring quality and quantity.

Discussion

The morphological divergence of reproductive and nonreproductive individuals has only been documented in two vertebrate societies, those of the naked mole-rat and the Damaraland mole-rat [6,7]. In this study, by altering the life history trajectories of agematched females in captivity, we show that the lengthening of breeding females in Damaraland mole-rats is caused by the upregulated growth of the skeletal vertebrae in breeders, causing breeders to be more elongated than nonbreeders. We also show that it is the onset of breeding, rather than the removal from reproductive suppression, that acts as the driver of this skeletal remodelling. Two lines of evidence support this view. First, we found that reproductively naive females in the wild that dispersed from their natal group and settled solitarily were morphologically equivalent to female nonbreeders still resident in their natal group, despite many of the former being isolated for multiple years. Secondly, we found that individuals in captivity only changed shape and size when they were paired with an unrelated male, whereas no such change took place in females that were housed solitarily. The increase in elongation demonstrated by breeding females presumably serves to enhance fecundity by

allowing the female to be larger without gaining extra girth. It is probably also highly advantageous for a species that occupies a system of narrow subterranean tunnels where increases in girth must impose strong constraints on mobility, as unlike in eusocial insects, reproductive female mole-rats are not bound to the nest (e.g. Ansell's mole-rat *Fukomys anselli* [27]). The fecundity benefits of body lengthening were confirmed by analyses of birth events in captivity which showed that longer breeders produced larger litters and invested more prenatally in each pup after the effect of litter size was statistically controlled for.

In finding that upregulated vertebral growth is central to the elongation of breeding females, the results of this study deviate from a previous result in wild Damaraland mole-rats which suggested that breeder elongation is achieved through a relative reduction in growth of the skull compared to growth towards total body length [7]. The ad libitum feeding of individuals in captivity might also have generated different patterns of growth in captivity compared to the wild. This possibility can only be addressed with longitudinal X-rays of wild animals. Without this information, our study implicates vertebral growth as the key process involved in elongation. As such, similar developmental routes appear to underpin the statusrelated morphological divergence of naked and Damaraland mole-rats, albeit the extent of divergence is greater in naked mole-rats [6]. This may be an indirect result of the much larger group sizes of naked mole-rats (up to 295 individuals [28]); with increasing group size, breeder fecundity will be stronger (larger workforces can rear more offspring; mean and maximum litter sizes in the field = 11.3 ± 6.2 and 28 individuals respectively [29]) and the reproductive potential of helpers declines (because the likelihood of inheriting the dominance position is reduced), which together would favour increased elongation in breeders and developmental arrest in nonbreeders. Although naked and Damaraland mole-rats display the largest group sizes and most extreme forms of sociality in the mole-rats, other members of the Bathyergidae family exhibit comparable social features (group-living, a reproductive

division of labour and delayed dispersal [30]), and one could speculate that if the social environment is important in morphological divergence then morphological divergence may also be present in other mole-rats. Alternatively, if skeletal elongation were principally a consequence of subterranean living and associated constraints on abdominal width, then even the solitary species of mole-rat may undergo vertebral lengthening across reproductive episodes (e.g. the Cape dune mole-rat *Bathyergus suillus*, the Cape mole-rat *Georychus capensis*, the Silvery mole-rat *Heliophobius argenteocinerus*, amongst others); though group-living and sociality could still increase the magnitude of this effect trait. A broader examination of rank-related growth would reveal whether skeletal lengthening is a unique adaptation associated with the highly social mole-rats or a more general feature of the Bathyergidae family and their subterranean habits.

Our study indicates that the vertebral lengthening of formerly subordinate helpers is stimulated by the onset of reproductive activities. The alternative possibility that relaxation of reproductive suppression is sufficient to induce divergence towards an elongated phenotype can be ruled out, since nonbreeding females removed from their natal group and housed solitarily for a year exhibited a growth trajectory equivalent to nonbreeders retained in complete groups (see [6] for analogous result in naked mole-rats). Likewise, solitary females in the wild were morphologically indistinguishable from in-group nonbreeders. Various endocrinological changes that take place throughout pregnancy could be implicated in this parity-driven bone growth. Sex steroids such as estradiol and progesterone increase during pregnancy and are heavily involved in bone formation[25, see 12 for progesterone in DMRs], often by mediating levels of growth hormone and insulin-like growth factor 1 [32]. Testosterone also promotes bone growth [33], and although commonly thought of as a male hormone, the heightened levels of testosterone measured in breeding female Damaraland mole-rats [34] – a pattern also seen in other cooperative breeders [35] – raises the possibility

that testosterone is also involved in vertebral lengthening. The action of such hormones sets up a skeletal mineral reserve that is subsequently resorbed and used in offspring development and milk production [20,36,37]. Yet, despite skeletal demineralisation during lactation, reproduction has been shown to lead to a net increase in total skeletal size. For example, in mice it has been shown that reproduction generates permanent increases in total body length [38], a result bearing obvious relevance to mole-rats, whilst in humans, high parity has been associated with increases in bone size [39,40]. Similar physiological mechanisms might therefore operate to drive the elongation and increased size of female mole-rats, whose unusual physiology could yet offer important biomedical insights concerning skeletal development [41].

The metabolic challenges of reproduction [42,43] can be expected to be particularly large in cooperative breeding mammals like Damaraland mole-rats because dominant females within cooperative societies often breed multiple times per year, and it is not uncommon for a female to conceive during the period that she is lactating for her current litter [8,44,45]. In this context, continued skeletal growth in mole-rats is particularly remarkable because short-inter-birth intervals must necessarily reduce the opportunity to recover calcium lost during lactation. In some cooperative breeders such as the mongooses and the canids, some of the maternal burden of lactation is offset by helpers lactating for non-descendent pups (e.g. African wild dog *Lycaon pictus* [46]; dwarf mongoose *Helogale parvula* [47]; meerkat *Suricatta suricata* [48]), but allo-lactation is absent in mole-rats. In the absence of allo-lactation, the atypically long gestation period of Damaraland mole-rats could be important by allowing more time for pregnant females to accrue mineral reserves prior to lactation (78-92 days [30]). In meeting their calcium requirement, breeding females may also rely on the exceptionally high calcium content of their primary food source - the subterranean tubers of the gemsbok cucumber *Acanthosicyos naudininianus* - and a highly efficient mode of calcium

uptake to drive their skeletal growth [49,50]. In fact, the calcium content of gemsbok cucumber is four to five times higher per unit mass than the sweet potato and cucumber diet given to the animals in the captive population, and this lowered calcium diet may even have led to an underestimation of the degree of vertebral lengthening documented in this study.

Overall, this study confirms the morphological divergence of breeding female Damaraland mole-rats at the skeletal level. By examining the growth trajectories of agematched individuals, it provides the first definitive evidence that the growth patterns underlying skeletal dimorphism cannot be explained by early-life developmental divergence or by status-specific age differences as in some eusocial ants and honeybees [51–53]. Instead, growth patterns resemble those of other cooperative breeding vertebrates (e.g. Meerkats [4], naked mole-rats [5], Haplochromis burtoni cichlid [3]) and other social insects (e.g. Ponerine ants [54], termites [55]) where contrasts in size and shape between breeders and nonbreeders are the result of changes that occur on or around the acquisition of a dominant breeding position. Viewed more broadly, the secondary growth of mole-rat breeders provides a clear example of socially responsive growth adjustment, or what might be termed 'strategic growth' [1,56]. Similar forms of adaptive, socially responsive growth might be more prevalent in mammals than is currently recognised, but the extent to which this is the case, and the implications for the structuring of mammalian dominance hierarchies, is as yet poorly understood. Of the few cases documented in mammals, emphasis has been placed on the status-related upregulation of growth [4,5], but an equally interesting perspective will be to understand the circumstances and mechanisms that cause development to be delayed or arrested in subordinate individuals in the first place. For example, in adult male orang-utans, some individuals develop conspicuous, sexually-selected cheek flanges soon after reaching sexual maturity, whereas others may reach sexual maturity and remain unflanged for 20 years before developing this conspicuous secondary trait [57]; the causes of this individual variation in development are largely unknown. Societies with marked reproductive skew and suppression of subordinate reproduction provide an obvious place to investigate socially-responsive growth further, but as subordinate group members often enjoy a small share of reproduction in even heavily-skewed societies [58,59], it seems that if rank-related divergence in size and shape is identified in other mammals, it is unlikely to be of a magnitude comparable to the social mole-rats where subordinates never reproduce.

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Supplementary Material 1

Details of the study population

For: Reproduction triggers adaptive increases in body size in female mole-rats

In addressing the principle aims of the study we used information from several different sources, including X-ray data from both captive and wild Damaraland mole-rats. The general details of the study system and the X-ray methodology are presented, before we detail how specific data sources were used to meet our aims. The Damaraland mole-rat is a subterranean cooperative breeder that inhabits the red arenosols of the Kalahari in groups of 2-41 individuals (Jarvis and Bennett 1993). Our study population is located around the Kuruman River Reserve in the Northern Cape of South Africa (\$26.98706° E21.81229°). A captive population was founded at this location using animals sourced from the local population between February and September 2013, and these founding animals were either maintained in their original group or selected to create new groups, achieved through the pairing of a nonbreeding female with a nonbreeding male. Groups are housed in artificial tunnel systems constructed of PVC pipes and are provided ad libitum access to sweet potato and cucumber (additional details of the study system and animal husbandry can be found in Zöttl et al. 2016). The captive animals from which X-rays were taken therefore represent animals that were initially caught in the wild and were brought into captivity, or the lab-born offspring of wild-caught animals.

We also took X-rays from individuals living permanently in the local wild population where a long-term mark recapture study is ongoing. Groups were trapped periodically (6 or 12-month intervals) using modified Hickman traps that were positioned into tunnel systems by digging. Traps were baited with sweet potato. After trap setting, traps were checked every 2-3 hours throughout the day and night. On capture, animals were placed into a closed, sand-

filled box with other group members, and provided food and shelter. Intermittently, individuals were transported back to the laboratory where they were X-rayed, weighed, and had their total body length measured manually. Two people measured body length from the front of the snout to the tip of the tail to an accuracy of 1mm using a tape measure. Total body length was taken as the average of the two measures; the human measurement of body length is referred to as 'Total body length' to distinguish it from the 'Skeletal body length' measured from X-rays (below). When transporting animals from the field to the lab, traps were temporarily disabled to prevent individuals being kept in the traps for long periods. After sampling, groups were housed temporarily in tunnel systems in the laboratory (see methods below), and once a whole group was captured, as evidenced by an absence of activity for 24hrs, the animals were all returned to their natural burrow system.

Supplementary Figures and Tables

For: Reproduction triggers adaptive increases in body size in female mole-rats

Table S1. Proportional trait contributions (a) and trait loadings (b) to the first five principal components from a PCA of skeletal measures in captive adult female mole-rats (> 100g)

a) Proportional Contribution

Trait	PC1	PC2	PC3	PC4	PC5
Rostrum	3.4	0.42	11.94	0.98	20.6
Ulna	6.99	1.86	2.69	8.86	8.14
L5 Vertebra	15.74	26.35	1.08	12.66	27.97
Pelvic Girdle	10.75	22	18.39	24.72	20.21
Pelvis Length	6.02	0.04	40.93	39.57	11.19
Femur Length	31.43	41.89	16.31	2.12	3.81
Tibia Length	6.54	2.3	1.55	5.45	0.18
Skeletal Body Length	10.41	4.83	1.99	4.31	0
Skull Width	8.72	0.3	5.14	1.33	7.88

b) Trait Loadings

Trait	PC1	PC2	PC3	PC4	PC5
Rostrum	0.0271	-0.0058	0.0252	0.0065	0.0247
Ulna	0.0389	-0.0123	0.0119	0.0195	0.0153
L5 Vertebra	0.0584	0.0463	-0.0075	0.0232	-0.0285
Pelvic Girdle	0.0483	0.0423	-0.0313	-0.0325	0.0242
Pelvis Length	0.0361	0.0019	0.0466	-0.0411	-0.018
Femur Length	0.0825	-0.0583	-0.0294	-0.0095	-0.0105
Tibia Length	0.0377	-0.0137	0.0091	0.0153	0.0023
Skeletal Body Length	0.0475	0.0198	0.0102	0.0135	0.0005
Skull Width	0.0435	0.0049	0.0166	0.0076	0.0146

Table S2. Bivariate scaling relationships of skeletal size measures in captive female mole-rats; SBL = Skeletal Body Length. SW = Skull Width. All linear traits were log-transformed. Bold, underlined terms represent slopes that differ significantly at $\alpha = 0.05$ (*), 0.01 (**), 0.001 (*). Significantly different slopes were determined by the interaction between Trait₂ and Class. Significantly different intercepts are taken from the model including the interaction if it was significant, otherwise they derive from a simpler model in which the interaction term was removed. Note that the difference in intercepts for relationships with an interaction are estimated at length zero, so they lie outside the bounds of the data. Difference in intercepts represent an in-group nonbreeder relative to the queen. Note that total head width here refers to the skull width as measured on the anaesthetised animal with digital callipers (i.e. not from X-rays).

$Trait_1$	$Trait_2$	β Queen	β nonbreeder	diff Slopes	diff Intercept
Rostrum Length	SBL	0.561 ± 0.161	0.594 ± 0.189	0.033, n.s.	0.013, n.s.
<u>Ulna</u>	SBL	0.314 ± 0.154	0.774 ± 0.181	0.460, *	-2.26, *
L5 Vertebra	SBL	0.586 ± 0.160	0.912 ± 0.189	0.326, n.s.	-0.112, ***
Pelvic Girdle Width	SBL	0.205 ± 0.238	0.566 ± 0.281	0.362, n.s.	-0.076, ***
Pelvis Length	SBL	0.619 ± 0.234	0.627 ± 0.276	0.008, n.s.	0.018, n.s.
Femur Length	SBL	0.490 ± 0.334	1.039 ± 0.394	0.549, n.s.	0.017, n.s.
Tibia Length	SBL	0.449 ± 0.148	0.625 ± 0.174	0.176, n.s.	0.001, n.s.
Skull Arch	SBL	0.510 ± 0.116	0.914 ± 0.137	0.404, **	-1.99, **
Head Width	SBL	0.402 ± 0.216	1.281 ± 0.254	0.880, ***	-4.31, ***
Weight	SBL	0.424 ± 0.120	0.951 ± 0.014	0.527, ***	-2.60, ***
Rostrum Length	SW	0.566 ± 0.172	0.557 ± 0.196	-0.009, n.s.	0.012, n.s.
Ulna	SW	0.730 ± 0.144	0.689 ± 0.163	0.041, n.s.	0.012, n.s.
L5 Vertebra	SW	0.228 ± 0.193	0.756 ± 0.219	0.530, *	-1.89, *
Pelvic Girdle Width	SW	0.420 ± 0.251	0.548 ± 0.285	0.129, n.s.	-0.076, ***
Pelvis Length	SW	0.381 ± 0.248	0.681 ± 0.282	0.300, n.s.	0.018, n.s.
Femur Length	SW	0.457 ± 0.354	1.015 ± 0.402	0.559, n.s.	0.017, n.s.
Tibia Length	SW	0.482 ± 0.155	0.601 ± 0.177	0.119, n.s.	0.001, n.s.
Head Width	SW	1.257 ± 0.187	1.261 ± 0.212	0.004, n.s	0.056, ***
Weight	SW	0.749 ± 0.115	0.883 ± 0.131	0.134, n.s.	0.001, n.s.

Table S3. Full model outputs for linear mixed effects models exploring the influence of body length on three fitness: a) Litter size, Poisson errors) b) Total neonate mass, normal errors c) Individual pup mass, normal errors. Significance of fixed covariates were estimated from likelihood ratio tests comparing models with and without the fixed effect of interest.

	Fixed Terms				Random Terms			
	Term	Estimate	Standard Error	LRT (χ^2_1)	p value	Term	Variance	Standard Deviation
a) Litter Size	Intercept	-1.06	0.92			Mother	0	
	Total Body Length	0.12	0.05	5.81	0.016			
	Primiparity (YES)	-0.3	0.17	3.57	0.059			
b) Total Neonate								
Mass	Intercept	-39.52	23.02			Mother	37.7	6.1
	Total Body Length	3.83	1.23	8.89	0.003	Residual	82.49	9.1
	Primiparity (YES)	-5.65	2.52	5.05	0.025			
c) Individual Pup							0.01	0.0
Mass	Intercept	5.25	3.17			Mother	0.81	0.9
	Total Body Length	0.36	0.17	4.12	0.040	Litter	0.84	0.92
	Primiparity (Y)	0.45	0.33	1.77	0.183	Residual	1.24	1.11
	Litter Size	-0.6	0.08	47.76	< 0.001			

Figure S1 – Bimonthly change in a) L5 lumbar vertebra and b) Skull width of captive females experimentally manipulated to follow different social trajectories.

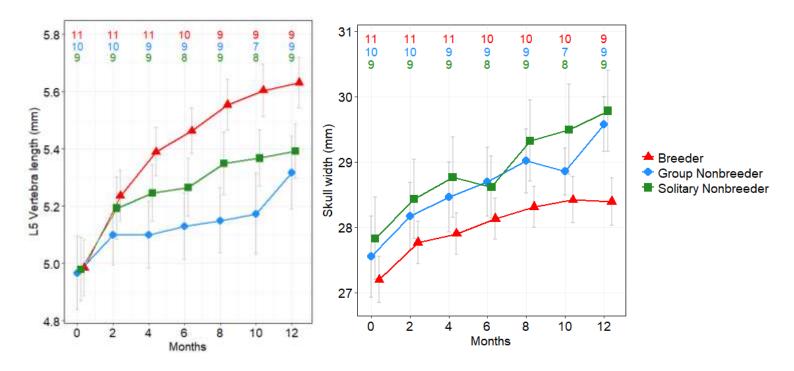


Figure S2 – Growth of L4 (a) and L6 (b) vertebrae of captive female mole-rats experimentally manipulated to follow different social trajectories.

