

1 Shaped from an early age: behavioural and hormonal phenotypes in juvenile  
2 male guinea pigs living in distinct social environments

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## Abstract

Behavioural plasticity enables individuals to vary their behaviour in response to different environmental conditions. As the social environment can change at any time, individuals need to be able to adjust throughout their lives. Our goal was therefore to elucidate when and how behavioural and hormonal adjustments in guinea pigs occur. We focused on juvenility, an important developmental phase characterized by prominent changes of the social environment, since the focus on social interactions shifts from parents to peers. For this approach, juvenile male guinea pigs (*Cavia aperea* f. *porcellus*) lived in two distinct social environments: while males of both groups lived in heterosexual pairs, males of one group were socially stimulated (e.g., an unfamiliar individual is introduced into the focus males home enclosure for 10 minutes) regularly whereas males of the other group were not. This procedure increased the number of social interactions. Socially stimulated males showed different adjustments to their social environment in comparison to non-socially stimulated males. They displayed an initially increased stress response, enabling them to adequately react to the unpredictable social encounters. Over time, males then adjusted to this challenging environment and displayed a decrease in stress response again. Moreover, only socially stimulated males showed a significant increase of courtship and sexual behaviour with age. Taken together, these findings demonstrate that already in juvenility the social environment induced hormonal adjustments and behavioural changes in male guinea pigs, thereby highlighting how early-life social experiences can shape individuals' phenotypes.

## Keywords

Behavior, behavioral development, behavioral plasticity, cortisol responsiveness, juvenility, niche conformance, social interactions, social niche, testosterone

## 1. Introduction

Behavioural plasticity enables individuals to vary their behaviour in response to different environmental conditions, so that they can adjust to changing social environments [1]. Such adjustments can result in an optimized phenotype-environment match and thus influence the fitness of an individual [2]. The effects of the social environment on behavioural phenotypes were demonstrated in several species. In birds, fish and mammals for example, individuals from social environments with many opportunities for social interaction reacted less aggressively towards potential competitors than individuals from social environments with only a few interaction partners [3–5]. These findings emphasize how interactions with conspecifics, especially potential mating partners and competitors, are thus an important driver for shaping behavioural phenotypes [6].

Such behavioural adjustments can happen through underlying endocrine mechanisms [7], for example through shaping of the principal neuroendocrine stress response system, namely the hypothalamic-pituitary-adrenocortical (HPA) axis. The HPA axis is regulating glucocorticoid secretion and thus stress response [8–10]. The causal links between social environment, stress responsiveness and behaviour were already demonstrated in several species. In guinea pigs for example, males raised in large mixed-sex groups frequently engaged in diverse social interactions, which triggered increased testosterone levels. Elevated testosterone in turn inhibited HPA activity, ergo reducing cortisol responsiveness [11, 12], which is the main glucocorticoid in guinea pigs [13]. This resulted in a low-aggression phenotype, facilitating integration into unfamiliar groups with several adult males and females. In contrast, males housed with only a same-age female experienced fewer social interactions, leading to lower testosterone levels, higher cortisol responsiveness and a high-aggression phenotype incompatible with unfamiliar males [12, 14]. Similar results were found in zebra finches: males raised in a group showed the lowest courtship behaviour and lowest aggressiveness, whereas males raised with a single female showed the most intense courtship behaviour and highest aggressiveness and were most attractive to females [15].

These examples show how the social environment can modulate the development of the males' social behaviour to be adaptive in their likely future environment [16]. This phenomenon is also described by the “predictive adaptive response hypothesis” [17] which describes how environmental cues can provide predictive information about the environment and alter the development to increase fitness during later life [18]. In principle, such shaping processes can occur in different life phases. Effects of environmental cues on the behavioural phenotype are facilitated by ongoing neural maturation. Thus, especially the early life phase (prenatal phase and early postnatal phase i.e., the time between birth and weaning in mammals) and adolescence (transition from infancy to adulthood) are referred to as sensitive windows of enhanced behavioural plasticity [18]. Already in the prenatal phase, stress

hormones secreted by the mother can directly affect intra-uterine development, because the offspring's HPA axis is susceptible to prenatal programming [19, 20]. During adolescence, developmental trajectories can be shaped directly by cues from the social environment. Prominent alterations occur in the endocrine system and neural circuitry [21], like activity and organization of the HPA axis [22]. Since this plasticity is assumed to decline with age [23], only a few studies examined behavioural plasticity during adulthood. In guinea pigs, however, evidence for an adaptive reshaping of hormonal phenotypes during adulthood was found, suggesting a greater role for behavioural plasticity in later life stages than commonly presumed [14]. These findings support that behavioural plasticity- and in consequence behavioural adjustments in response to the (social) environment- might be possible all throughout ontogeny. This is a plausible theory considering the social environment being able to change at any point during lifetime. To investigate behavioural plasticity holistically, it is therefore important to include every life stage. Research focussing on juvenility i.e., the time between weaning and adolescence, is however lacking. During this time, the social environment changes a lot since the focus on social interactions shifts from the parents to peers. There is also evidence that during juvenility the HPA axis displays a heightened sensitivity to stress from the environment and thus adjustment to the social environment could occur. In rats for example, stress experienced during juvenility affected behaviour in later life [24] and similar effects could be mimicked by applying corticosterone during juvenility [25].

Guinea pigs have a high flexibility regarding their social organization [26] and behavioural plasticity in other life phases has already been investigated in this species. Thus, they are a well-suited model organism for examining such processes in juvenility, too. Our goal in this study was therefore to investigate how juvenile male guinea pigs adjust to two different social environments. Thus, we repeatedly measured hormone concentrations and observed home-enclosure behaviour. We hypothesized that male guinea pigs living in different social environments differ in their behavioural and hormonal phenotypes. Since juvenile guinea pig males are sexually immature, direct fitness consequences in the form of reproductive success could not be measured. Instead, body weight as fitness proxy was assessed. To analyse the stability of hormone concentrations and body weight over time, a repeatability analysis was conducted.

## **2. Material and methods**

### **2.1 Animals and housing conditions**

All animals used for this study were bred from a breeding program of multi-coloured shorthaired guinea pigs (*Cavia aperea* f. *porcellus*) at the Department of Behavioural Biology at the University of Münster. They were born and reared in a total of six to eight harem groups within one breeding room, each consisting of one male, one to three females and their pre-weaned offspring. The offspring was routinely taken out of the harems after weaning at post-natal day (PND) 21 ( $\pm 1$ ) and adults were removed and

replaced at around 18-24 months of age. Each harem was kept in wooden enclosures with a base area of approximately 1.5 m<sup>2</sup> and a wall height of 0.5 m. The enclosures were filled with wood shavings (Tierwohl Super, J. Rettenmaier & Söhne GmbH + Co KG, Rosenberg, Germany) as bedding and enriched with red plastic shelters and wooden bridges.

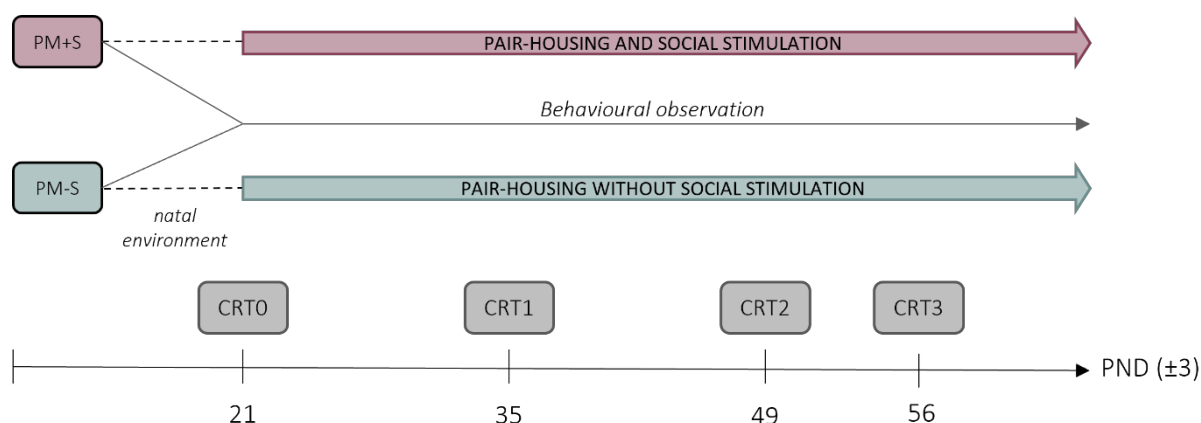
The experimental animals were transferred to enclosures in a different housing room after weaning at PND 21 ( $\pm 3$ ). These enclosures had a base area of 0.5 m<sup>2</sup>, a wall height of 0.5 m, were also filled with wood shavings and enriched with a big and a small red plastic shelter. Food (hasfit Cavia C pellets, EQUOVIS GmbH, Münster, Germany) and water mixed with vitamin C were available *ad libitum*. Additionally, hay was replenished daily and fresh fodder (carrots, cucumbers, apples) was fed regularly. All guinea pig housing rooms were kept under controlled conditions with a 12 h: 12 h light/ dark cycle (lights on at 07:00), temperature of approximately 22 °C and relative humidity of approximately 48 %.

## 2.2 Experimental design

For this study, twenty guinea pig males were used. The experimental phase started after weaning at PND 21 ( $\pm 3$ ) and lasted six weeks, meaning the animals were 60 ( $\pm 3$ ) days of age when the experiments ended. In guinea pig males, sexual maturity is usually reached around PND 70 [27]. Each male was paired with a female which was the same age. The male and his respective female partner stem from different harem groups, meaning they were neither half nor full siblings. To investigate the influence of distinct social environments on behavioural and hormonal phenotypes, they were randomly assigned to one of two treatment groups. Males of both groups lived in heterosexual pairs, but males of one group were socially stimulated (see 2.3) regularly (pair-housed male with social stimulation; PM+S group), while males of the other group were not (pair-housed male without social stimulation; PM-S group).

In total, four cortisol response tests (CRTs) to measure basal and reaction cortisol values (see *assessment of endocrine phenotypes*) were conducted within the six week long experimental phase (**Fig. 1**). The first CRT was conducted before the social stimulation treatment started and is thus referred to as CRT0. CRT0 was conducted in the first experimental week, CRT1 and CRT2 followed 14 ( $\pm 2$ ) days after the preceding one, while CRT3 was carried out 7 ( $\pm 2$ ) days after CRT2 (**Fig. 1**). Social stimulation and recording of home enclosure behaviour were each conducted three times per week during the whole experimental phase.

Please note: as part of another project, a battery of behavioural tests to further evaluate social and risk-taking behaviour plus fur swabbing with PMDS tubes to analyse chemical fingerprints was conducted in the last week.



**Figure 1:** Procedure of behavioural observations in the home enclosure and cortisol response tests (CRT). Focal males were housed with a female partner. One group (PM+S) was regularly stimulated by introducing other individuals into the home enclosure while the other group (PM-S) was not. This social stimulation started after CRT0 and lasted until the experimental phase was finished at post-natal day (PND) 60±3.

## 2.3 Social stimulation

The social stimulation procedure applied in the present study was adapted from Lürzel and colleagues [28, 29], where social stimulation successfully influenced hormonal profiles in adolescent guinea pig males. The social stimulation treatment for the stimulated males (PM+S) started after the first CRT. From then on, social stimulation was applied three times per week for the whole experimental phase. More in details, an unknown individual was introduced into the home enclosure of the focus male and his female partner for a maximum of ten minutes. In each week, two of these stimulations were done with another male and one with a female. In total, the focus males had a total of twelve social stimulation sessions with another male and six social stimulation sessions with a female. The female stimulation animals always came from the harems to ensure they were pregnant and thus in the same reproductive stadium, preventing a confounding influence of oestrus. While female stimulation animals were always adult, the age of male stimulation animals ranged from 44 to 994 days, however, they were always older than the focus male. The pool of stimulation males for each PM+S male included eight to twelve individuals and the pool of stimulation females four to six individuals. The stimulus animals were replaced at irregular intervals. If the focus male was stimulated more than once with the same stimulus animal, there was always a minimum interval of seven days between these stimulation sessions.

PM+S males never experienced more than one social stimulation session per day and the day and time of day at which social stimulation occurred was varied in order to avoid possible habituation effects by introducing unpredictability. Before the stimulation itself began, the red plastic shelters were temporarily removed from the home enclosure of the focus male and the video camera was turned on, since all stimulation sessions were recorded. After the stimulation animals were introduced into the home enclosure, a timer was started as the sessions had maximum length of ten minutes. When males displayed escalated aggressive behaviour, the sessions were aborted beforehand to minimise the risk

of injury. Out of a total of 120 stimulation sessions using males as stimulus animals, eight had to terminated because aggression escalated.

## 2.4 Assessment of behavioural parameters

To analyse how distinct social environments influence (social) behaviour, the home enclosure behaviour of the focus males in both treatment groups was observed by filming them 2-3 times per week for one hour each. For this purpose, a video camera (Panasonic HC-V785 or SONY HDR-CX405) was installed approximately 1.5 m above each experimental home enclosure. The day and time (usually between 09:00 and 15:00) at which the videos were recorded was randomized. In total, 12 h to 18 h of home enclosure behaviour was collected for each individual.

The subsequent analysis was done with the program Interact (Interact, Lab Suite Version 2022, Program version 20.8.3.0, Mangold International GmbH, Arnstorf, Germany). The videos were blinded and randomized, ensuring ID and treatment of the respective individual as well as the time of recording were unknown to the observer.

The observed behaviours were summarized into the following categories: courtship and sexual behaviour, sociopositive behaviour, agonistic behaviour, play and other (**Tab. 1**).

**Table 1:** Ethogram used for the observation of home enclosure behaviour. The abbreviation “FA” stands for “focus animal”, e.g., the experimental male.

Category	Behaviour	Description
Courtship and sexual behaviour	Ano-genital licking	The FA stretches its snout towards or touches another animals’ ano-genital region and lick or nuzzles the other animals’ genital region. The distance between the two animals is less than one snout-width.
Courtship and sexual behaviour	Chin-rest	The FA lays the bottom of its head on another animals’ torso.
Courtship and sexual behaviour	Mounting	The FA moves the forepart of its body onto the back of another animal from behind.
Courtship and sexual behaviour	Pelvic thrust	The FA mounts the other animal and moves the lower part of its body fast and rhythmically.
Courtship and sexual behaviour	Mating attempt	The FA puts at least one of its forepaws on another animal and tries to mate with the other animal, but the other animal prevents this.
Courtship and sexual behaviour	Rumba	The FA approaches the other animal slowly and visibly shifts its weight from one hind leg to the other and back, it can also move forward while doing so. This is often accompanied by a low purring noise. Behaviour ends when the FA stops for more than 3s.
Courtship and sexual behaviour	Flank	The FA walks parallel to another animal, touches its side with its own

		and slightly raises the hind leg on the side that is touching the other animal while moving forward.
Courtship and sexual behaviour	Chin-rump following	The FA walks or runs behind another animal with its nose towards the other animal's rear, trying to make contact with the chased animal. There is a maximum of 1 body length of distance between the two animals. Behaviour ends when the FA stops chasing for at least 3s.
Sociopositive behaviour	Naso-nasal sniffing	The FA stretches its nose towards another animal's nose or snout. The distance between the two animals is less than one snout-width.
Sociopositive behaviour	Naso-anal sniffing	The FA stretches its nose towards or touches another animals' anal region with its nose. The distance between the two animals is less than one snout-width.
Sociopositive behaviour	Social resting	The FA rests next to another animal at least 3s with a distance of less than a half a body length. Behaviour ends when not shown for at least 3s.
Play	Play	The FA makes one or a series of upward leaps and turns the head or foreparts sharply while in the air, or the FA starts with a short and fast run and then stops suddenly and changes the direction.
Agonistic behaviour	Displace	The FA approaches another animal or shows agonistic behaviour towards it, causing the other animal to move at least one body length away from the FA.
Agonistic behaviour	Evade	The FA moves at least one body length away from another animal that approached or interacted otherwise with it..
Agonistic behaviour	Head-thrust	The FA abruptly moves its head towards another animal, hitting or narrowly missing it, or biting it. The distance between the two animals is maximum half a body length.
Agonistic behaviour	Fight	A prolonged agonistic interaction of at least 3s between at least two animals. Head-thrusts, kicks and attack lunges can occur. The behaviour ends when one or both animals back away.
Agonistic behaviour	Kick	The FA abruptly moves one of its hind legs towards another animal.
Agonistic behaviour	Paw	The FA repeatedly moves one or both of its front paws across the bedding without moving in any direction.
Agonistic behaviour	Urine spray	The FA slightly arches its back and, with a small jolt, squirts urine behind it, usually towards another animal, which often reacts by stopping and cleaning itself. The urine squirt itself is not



		always directly or indirectly (wet spots on the enclosure wall) visible.
Agonistic behaviour	Curved body posture	The FA is standing within a distance of one body length in front of or sideways to another animal. Its body is usually curved with head and rump directed to the other animal, which is also displaying the same behaviour. This behaviour is often accompanied by growling and teeth chattering.
Agonistic behaviour	Head up	The FA is standing still but lifts its head up in such a way that the chin is facing upwards and towards another animal. The distance between both animals is maximum one body length.
Agonistic behaviour	Attack lunge	The FA jumps on or towards another animal, with the landing happening within one body length of the other animal.
Agonistic behaviour	Chase	The FA follows another animal over a distance of at least one body length. This happens with high velocity. During this interaction, the distance between both animals never exceeds two body lengths. Chasing is terminated, if the distance between the animals exceeds to body lengths for more than 3 s.
Other	Being under the house	The FA has moved under the small hideout with at least half of its body. Behaviour ends when the FA has moved at least half of its body out from under the hideout.
Other	Time-out	The FA is not visible.

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## 203 2.5 Assessment of hormone concentrations

204 Hormones were measured using blood samples obtained in cortisol response tests (CRTs), a  
205 standardized test used to measure the endocrine stress response to a challenge [30]. The male guinea  
206 pigs were exposed to the stressor of exposure to a novel environment [31] and stress responses at  
207 different time points were assessed by sampling blood. The test started between 12:30 and 13:30. Prior  
208 to that, the animals were undisturbed for one hour.

209 At the start of the CRT the male was taken out of his home enclosure and placed on the experimenter's  
210 lap outside of the housing room. To facilitate blood flow, a muscle salve (Finalgon® Wärmesalbe DUO,  
211 Zentiva Pharma GmbH, Frankfurt am Main, Germany) for expanding the blood vessels was applied to  
212 the guinea pig's ear and wiped off again. After that, the marginal ear vessel was punctured with a lancet  
213 (Solofix® Blutlanzetten, B. Braun Melsungen AG, Melsungen, Germany) and blood was collected in  
214 heparinized capillary tubes (Capillary tubes for microhaematocrits, 100 µl, Paul Marienfeld GmbH & Co  
215 KG, Lauda Königshofen, Germany) to later on determine basal cortisol (c0) and basal testosterone (t)  
216 levels. This procedure had to be completed within 3 minutes (cortisol) or 6 minutes (testosterone)

respectively after starting the test to avoid the sampling process from influencing the hormone values in the obtained sample itself [32]. Then, the guinea pig was singly placed into an unfamiliar enclosure in a different housing room where it stayed for a total of two hours. This enclosure had a size of 1 m<sup>2</sup>, wall height of 0.5 m and was equipped with wood shavings, food and water. Exactly one and two hours after the first one, blood sampling was repeated to determine first (c1) and second (c2) cortisol response values. The guinea pigs were weighed after each blood sampling and returned to their home enclosure after the last one.

To separate the blood plasma, the sample was centrifugated (13,000 × g for 5 min), transferred into a 1.5 mL Eppendorf tube and deep frozen at -20°C until assayed. Hormone concentrations were determined in duplicate using enzyme-linked immunosorbent assays (ELISA) (cortisol: RE52061, IBL International, Hamburg, Germany; antibody cross-reactivity: cortisol (100%), prednisolone (30%), 11-deoxycortisol (20%), cortisone (10.7%), prednisone (6.5%), 17 α-hydroxyprogesterone (5.4%), 6β-hydroxycortisol (4.4%), corticosterone 3.8%, desoxycorticosterone (1.8%); testosterone: RE52151, IBL International, Hamburg, Germany; antibody cross-reactivity: testosterone 100%, 11β-OH-testosterone 8.7%, 11α-OHtestosterone, 3.2%, dihydrotestosterone 1.9%). Intra- and inter-assay CVs were determined 2.09% and 3.98% for cortisol and 4.7% and 5.7% for testosterone.

In some cases, it was not possible to collect a sufficient amount of blood for the ELISA, resulting in a decreased sample size. For each CRT, the sample size per group ranged between n = 4 and n = 10.

## 2.6 Statistical analysis

Data analysis was carried out with RStudio version 2022.07.0 [33]. A priori sample-size calculation was conducted using the software G\*Power version 3.1.9.7 [34]. The calculations were based on baseline and response cortisol values. Previous studies showed that effects of the social environment on cortisol concentrations are large, with estimated effect size of  $f = 0.69$  [31, 35]. To detect effects with  $f = 0.69$  with an  $\alpha$  error probability of 0.05 and a power of 80% a total sample size of at least 19 animals would be needed. Thus, we decided to use a total sample size of n = 20 animals with n = 10 animals per treatment group.

Linear mixed-effect models were used to analyse the influence of the social stimulation treatment on hormone concentrations using the *lme4* [36] and *lmerTest* package [37]. In total, four models were fit with 1) baseline cortisol, 2) baseline testosterone, 3) cortisol responsiveness after 1 hour and 4) cortisol responsiveness after 2 hours as a respective response variable. To improve model fit, all response variables were square root transformed. Treatment (social stimulation versus no social stimulation) was added as a fixed effect. To investigate changes in hormone concentrations over time, we also included the variable CRT, representing the first, second and third CRT conducted after treatment, as a fixed

effect. We excluded data from the CRT conducted before the treatment (CRT0) from the analyses. However, hormone concentrations at CRT0 were still compared between the treatment groups using Wilcoxon rank-sum test to confirm there were indeed no differences between the groups prior to treatment. Furthermore, the continuous variable body weight was first mean-centered and then included as a fixed effect. We also added an interaction between treatment and CRT to determine whether effects of treatment varied across the three CRTs. Last, we fitted ID as a random effect. We used the *performance* [38] and *DHARMA* package [39] to check model assumptions. Marginal and conditional  $R^2$  values were calculated using the *performance* package [38], while partial  $R^2$  values for individual predictors were calculated using the *sensemakr* package [40]. Pair-wise comparisons for treatment, CRT and treatment\*CRT interaction were done by applying Tukey's adjustment for multiple comparison using the *emmeans* package [41].

Another linear-mixed effect model was fitted to analyse whether treatment affected body weight. Body weight measured after the first blood sampling in CRT1, CRT2 and CRT3 was modelled as a continuous response variable. The interaction between treatment and CRT was used as fixed effect to investigate the influence of treatment over time. ID was included as random effect. Pairwise comparison and  $R^2$  estimations were conducted as described for the hormone concentrations. Also, body weight at CRT0 was compared between the treatment groups using Wilcoxon rank-sum tests to confirm there were indeed no differences prior to treatment. Additionally, the relationship between body weight and cortisol responsiveness after 1 hour was examined by calculating Pearson's correlation coefficients for each treatment group separately. Body weight was mean-centered for each time point (CRT1, CRT2, CRT3) and the correlation coefficients were then calculated across all time points and for each time point separately. To determine whether the correlations for each time point differed significantly between treatment groups, Fisher's z-test was conducted using the *cocor* package [42].

Adjusted repeatability estimates of hormone concentrations and body weight were calculated for each of the treatment groups using the *rprR* package [43]. 95% confidence intervals were determined by parametric bootstrapping ( $N = 1000$ ), and likelihood ratio tests were used for significance testing. The models used to estimate adjusted repeatability were the same as mentioned before, with the only exception that treatment was removed as fixed effect.

For the analysis of the home enclosure behaviour, count data of behaviours from the coded videos was transformed into frequencies (occurrence per hour). Several behaviours were observed in only a few individuals, resulting in a zero-inflation of data which was detected using the *performance* package [38]. Therefore, we pooled behaviour into three categories: courtship and sexual behaviour, social behaviour and play, with individual behaviours being summed within each category. Agonistic behaviour was excluded from the analysis since it only occurred in a single individual. Generalized linear mixed-effect

models with negative binomial distribution accounting for the zero-inflated data were fit for each behavioural category using the *lme4* package [36]. Again, interaction between treatment and time was used as fixed effect in the models to investigate the influence of treatment over time. Time was categorized into “Phase 1” (1<sup>st</sup> and 2<sup>nd</sup> experimental week), “Phase 2” (3<sup>rd</sup> and 4<sup>th</sup> experimental week) and “Phase 3” (5<sup>th</sup> and 6<sup>th</sup> experimental week). ID was again fitted as a random effect. Model assumptions as well as the estimation of the different R<sup>2</sup> values were conducted in the same manner as for the analysis of hormone concentrations. Pair-wise comparisons for treatment, phase and treatment\*phase interaction were done by applying Tukey’s adjustment for multiple comparison using the *emmeans* package [41].

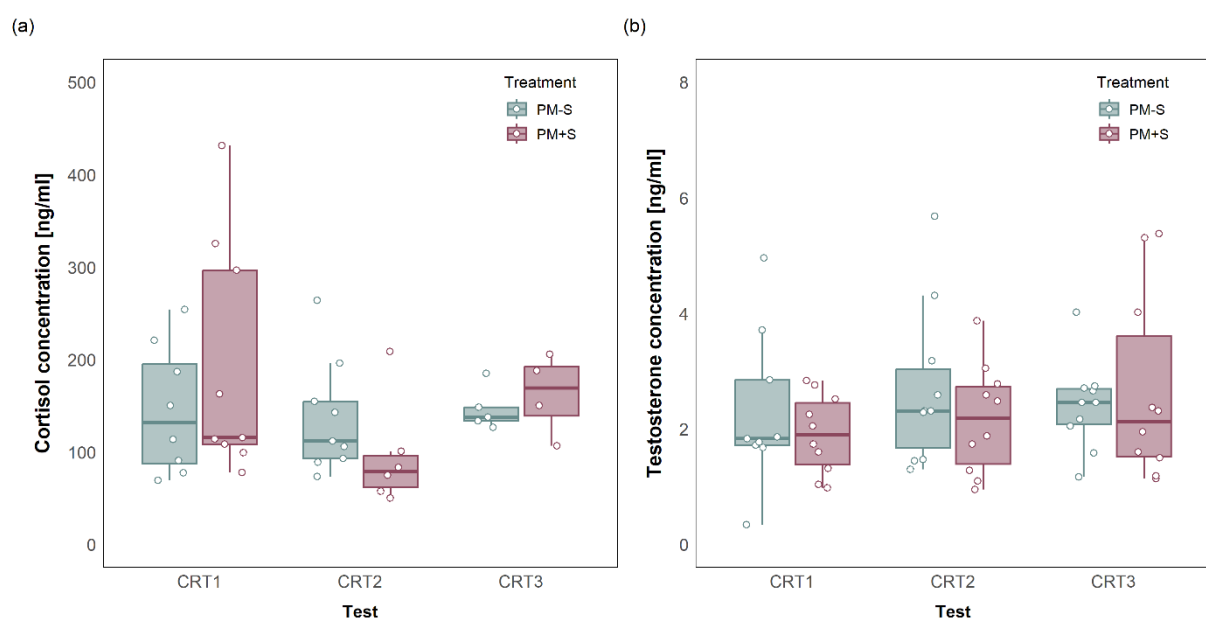
### 3. Results

Descriptive statistics for all hormone measurements, body weight and behaviour for each respective time point and over the entire time period can be found in the supplementary material (Tab. S1-3). Model summaries and detailed test statistics can be found in the supplementary material (Tab. S4-S22).

#### 3.1 Effects of social environment on hormone concentrations and body weight

The comparison of hormone concentrations (c0, c1, c2, t) at CRT0 using Wilcoxon rank-sum tests revealed no significant differences between the treatment groups prior to treatment.

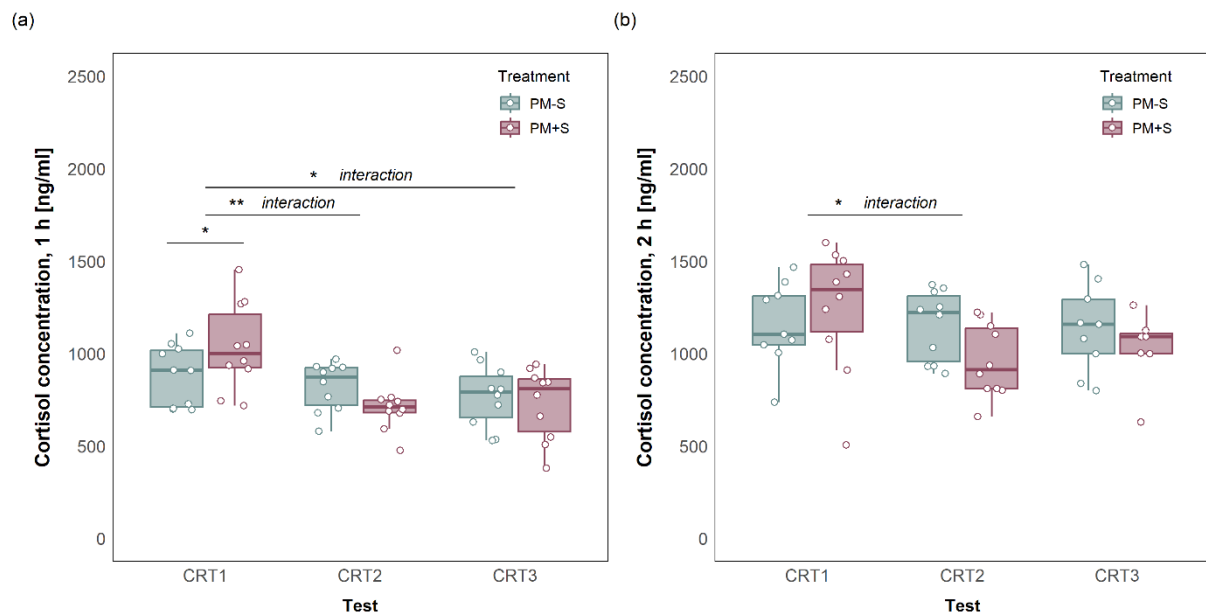
Regarding baseline testosterone and cortisol levels (Fig. 2), neither a significant effect of treatment or time (CRT), nor a significant treatment-by-time interaction effect was found.



**Figure 2:** Baseline cortisol (a) and testosterone (b) concentrations (ng ml<sup>-1</sup>) two weeks (CRT1), four weeks (CRT2) and five weeks (CRT3) after treatment start. Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted

are medians (horizontal marks), first to third quartiles (boxes), whiskers and all data points. Statistics: (a) Multiple comparisons of LMM; PM-S:  $n_{\text{CRT1}} = 8$ ,  $n_{\text{CRT2}} = 9$ ,  $n_{\text{CRT3}} = 5$ , PM+S:  $n_{\text{CRT0}} = 8$ ,  $n_{\text{CRT1}} = 9$ ,  $n_{\text{CRT2}} = 6$ ,  $n_{\text{CRT3}} = 4$ . (b) Multiple comparisons of LMM; PM-S:  $n_{\text{CRT1}} = 9$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ , PM+S:  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ .

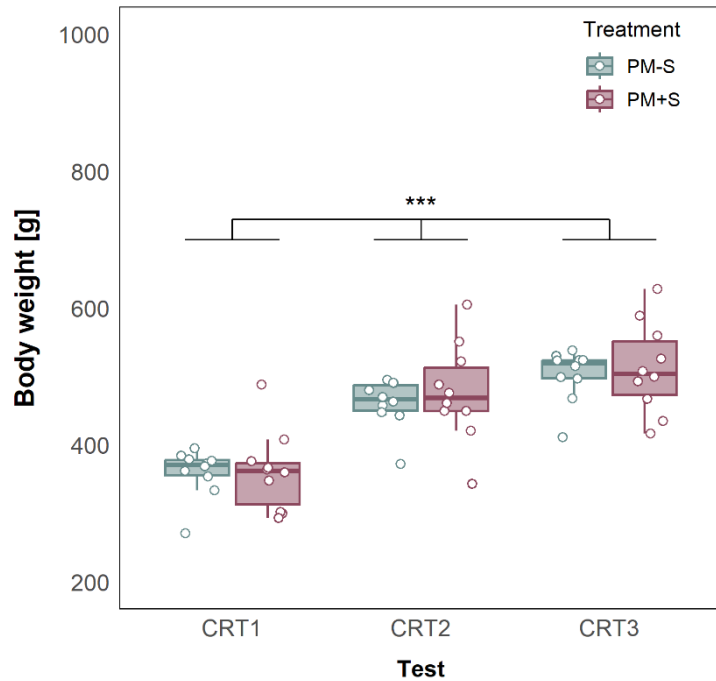
Regarding cortisol responsiveness at 1 hour (c1) of exposure to a novel environment, a significant treatment effect was found at CRT1 ( $\beta = -2.44 \pm 1.11$ ,  $t = -2.2$ ,  $p = 0.03$ ), where PM+S had significantly higher c1 values than PM-S (**Fig. 3a**). We also found a significant treatment-by-time interaction effect between CRT1 and CRT2 ( $\beta = 3.92 \pm 1.34$ ,  $t = 2.92$ ,  $p = 0.006$ ) as well as between CRT1 and CRT3 ( $\beta = 2.94 \pm 1.34$ ,  $t = 2.19$ ,  $p = 0.035$ ), where c1 values decreased for the PM+S group. Additionally, a significant effect of mass was found ( $\beta = -0.03 \pm -0.01$ ,  $t = -4.32$ ,  $p < 0.001$ ). For cortisol responsiveness at 2 hours (c2) of exposure to a novel environment (**Fig. 3b**), a significant treatment-by-time interaction effect between CRT1 and CRT2 ( $\beta = 4.2 \pm 1.7$ ,  $t = 2.46$ ,  $p = 0.019$ ) was found, with c2 values strongly decreasing for the PM+S group.



**Figure 3:** Cortisol concentrations ( $\text{ng ml}^{-1}$ ) at one hour (a) and two hours (b) of exposure to a novel environment two weeks (CRT1), four weeks (CRT2) and five weeks (CRT3) after treatment start. Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted are medians (horizontal marks), first to third quartiles (boxes), whiskers and all data points. Statistics: (a) Multiple comparisons of LMM;  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ , PM+S:  $n_{\text{CRT0}} = 10$ ,  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ ; \* $p < 0.05$  \*\* $p < 0.01$  (b) Multiple comparisons of LMM; PM-S:  $n_{\text{CRT1}} = 9$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 9$ , PM+S:  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 7$ ; \* $p < 0.05$ .

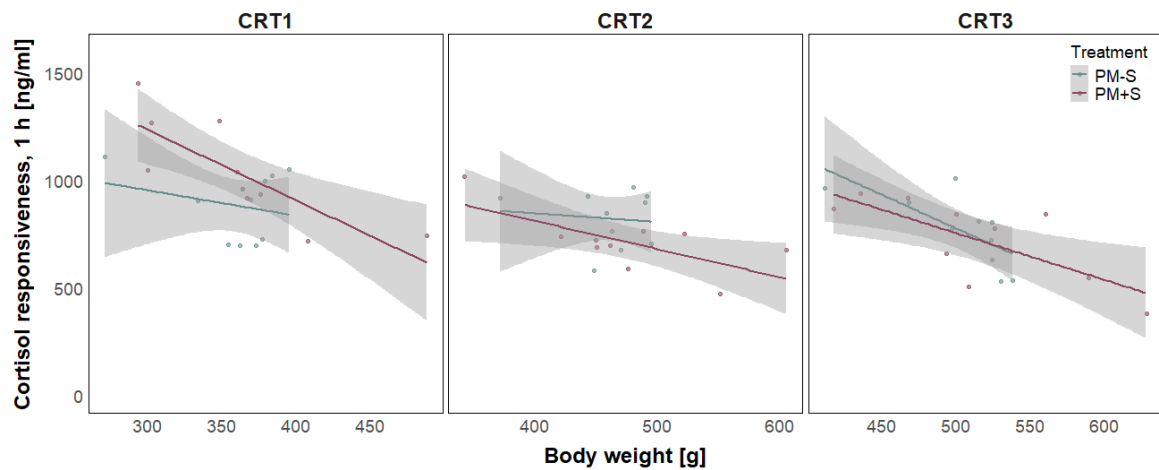
Regarding body weight, the comparison at CRT0 using a Wilcoxon rank-sum test revealed no significant differences between the treatment groups prior to treatment. A significant effect of time was found for the PM-S group from CRT1 to CRT2 ( $\beta = -101.2 \pm 6.62$ ,  $t = -15.28$ ,  $p < 0.001$ ), CRT1 to CRT3 ( $\beta = -143.1 \pm 6.62$ ,  $t = -21.61$ ,  $p < 0.001$ ) and CRT2 to CRT3 ( $\beta = -41.9 \pm 6.62$ ,  $t = -6.33$ ,  $p < 0.001$ ), as well as for the PM+S group from CRT1 to CRT ( $\beta = -116.1 \pm 6.62$ ,  $t = -17.53$ ,  $p < 0.001$ ), CRT1 to CRT3 ( $\beta = -151.7 \pm 6.62$ ,

$t = -22.91, p < 0.001$ ) and CRT2 to CRT3 ( $\beta = -35.6 \pm 6.62, t = -5.38, p < 0.001$ ) (Fig. 4). In each case, body weight was increasing.



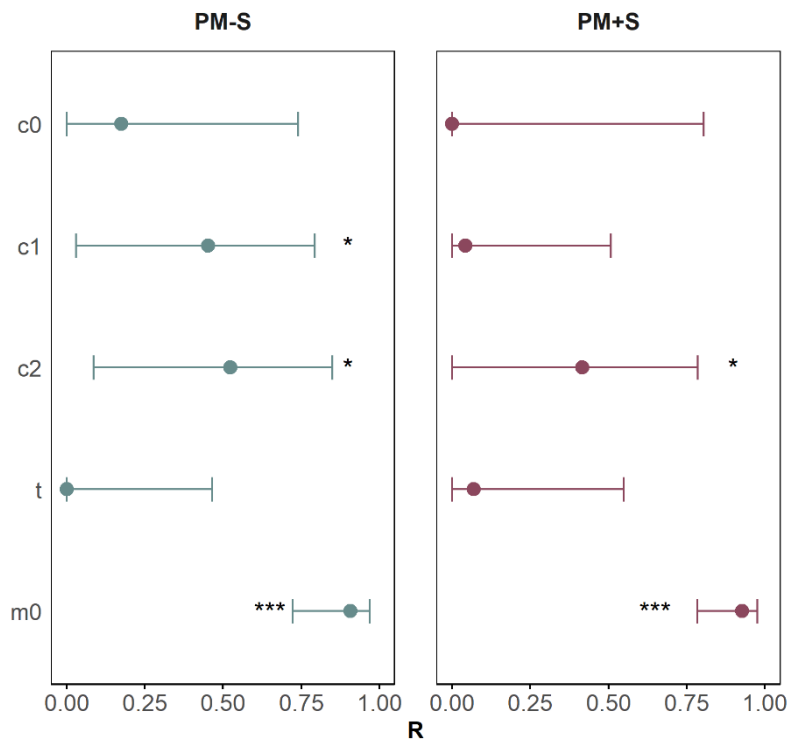
**Figure 4:** Body weight measured two weeks (CRT1), four weeks (CRT2) and five weeks (CRT3) after treatment start. Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted are medians (horizontal marks), first to third quartiles (boxes), whiskers and all data points. Statistics: Multiple comparisons of LMM; PM-S:  $n_{\text{CRT1}} = 10, n_{\text{CRT2}} = 10, n_{\text{CRT3}} = 10$ , PM+S:  $n_{\text{CRT1}} = 10, n_{\text{CRT2}} = 10, n_{\text{CRT3}} = 10$ ; \*\*\*  $p < 0.001$ .

The statistical analysis of hormone concentrations showed that c1 concentrations are significantly affected by body weight. At CRT1, body weight and c1 had a significant, strong negative correlation in the PM+S group ( $r = -0.81, t = -3.88, p = 0.005$ ) and a weak negative correlation in the PM-S group ( $r = -0.26, t = -0.76, p = 0.472$ ). At CRT2, body weight and c1 had a significant, moderate negative correlation in the PM+S group ( $r = -0.69, t = -2.67, p = 0.028$ ) and a weak, negative correlation in the PM-S group ( $r = -0.11, t = -0.31, p = 0.762$ ). At CRT3, body weight and c1 had a significant, strong correlation in the PM+S group ( $r = -0.74, t = -3.13, p = 0.014$ ) and a significant, strong correlation in the PM-S group ( $r = -0.71, t = -2.89, p = 0.02$ ). Comparisons between the correlations of the treatment groups were however not significant for any time point. These correlation between body weight and c1 concentrations over all timepoints (CRT1, CRT2, CRT3) are displayed in Figure 5.



**Figure 5:** Correlation between cortisol concentrations ( $\text{ng ml}^{-1}$ ) at one hour of exposure to a novel environment and body weight two weeks (CRT1), four weeks (CRT2) and five weeks (CRT3) after treatment start. Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted are regression lines, confidence intervals and all data points. PM-S:  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ , PM+S:  $n_{\text{CRT1}} = 10$ ,  $n_{\text{CRT2}} = 10$ ,  $n_{\text{CRT3}} = 10$ .

Adjusted repeatability was analysed for hormone concentrations (baseline cortisol, baseline testosterone, cortisol responsiveness after 1 and 2 hours) and body weight in both treatment groups (**Fig. 6**). Baseline cortisol ( $c_0$ ) was not repeatable in the PM+S group ( $R = 0$ ,  $\text{CI} = [0, 0.81]$ ,  $p = 1$ ). In the PM-S group, baseline cortisol had a low repeatability ( $R = 0.18$ ,  $\text{CI} = [0, 0.74]$ ,  $p = 0.331$ ). Baseline testosterone ( $t$ ) had a low repeatability in the PM+S group ( $R = 0.07$ ,  $\text{CI} = [0, 0.55]$ ,  $p = 0.44$ ) and was not repeatable in the PM-S group ( $R = 0$ ,  $\text{CI} = [0, 0.47]$ ,  $p = 1$ ). Cortisol responsiveness after 1 hour ( $c_1$ ) had a low repeatability in the PM+S group ( $R = 0.04$ ,  $\text{CI} = [0, 0.51]$ ,  $p = 0.495$ ) and a moderate repeatability in the PM-S group ( $R = 0.45$ ,  $\text{CI} = [0.03, 0.79]$ ,  $p = 0.014$ ). Cortisol responsiveness after 2 hours ( $c_2$ ) was moderately repeatable in the PM+S group ( $R = 0.42$ ,  $\text{CI} = [0, 0.55]$ ,  $p = 0.04$ ) and in the PM-S group ( $R = 0.52$ ,  $\text{CI} = [0.09, 0.85]$ ,  $p = 0.015$ ). Body weight ( $m_0$ ) was highly repeatable in the PM+S group ( $R = 0.93$ ,  $\text{CI} = [0.78, 0.98]$ ,  $p < 0.001$ ) and in the PM-S group ( $R = 0.91$ ,  $\text{CI} = [0.72, 0.97]$ ,  $p < 0.001$ ).

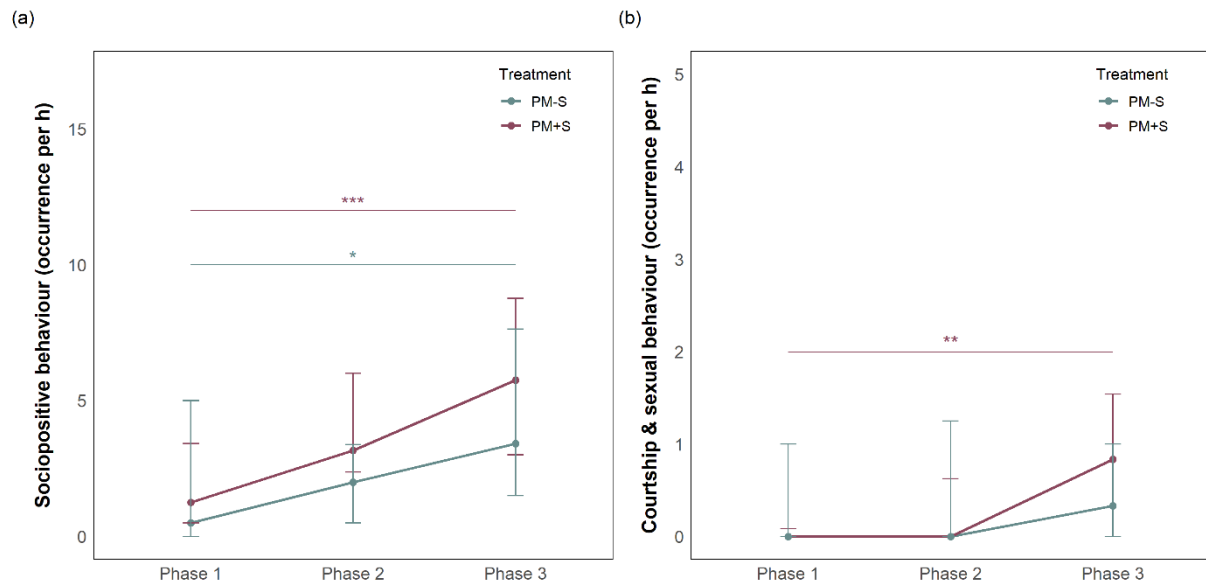


**Figure 6:** Repeatability (R) of baseline cortisol (c0), cortisol responsiveness after 1 (c1) and 2 hours (c2) of exposure to a novel environment, baseline testosterone (t) and body weight (m0). Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted are adjusted repeatability (data points) and confidence intervals (whisker). Statistics: repeatability analysis using permutation testing; PM-S:  $n_{c0} = 22$ ,  $n_{c1} = 30$ ,  $n_{c2} = 28$ ,  $n_t = 29$ ,  $n_{m0} = 30$ , PM+S:  $n_{c0} = 19$ ,  $n_{c1} = 30$ ,  $n_{c2} = 27$ ,  $n_t = 30$ ,  $n_{m0} = 30$ ; \* $p < 0.05$ , \*\*\*  $p < 0.001$ .

### 3.2 Effects of social environment on social behaviour

For sociopositive behaviour a significant effect of time (phase) was found for the PM-S group between phase 1 and phase 3 ( $\beta = -0.92 \pm 0.34$ ,  $z = -2.74$ ,  $p = 0.017$ ) and for the PM+S group between phase 1 and phase 3 ( $\beta = -1.29 \pm 0.33$ ,  $z = -3.9$ ,  $p < 0.001$ ). In both groups, the frequency of sociopositive behaviour increased over time (**Fig. 7a**). Furthermore, a significant increase of courtship and sexual behaviour was only found in the PM+S group ( $\beta = -2.38 \pm 0.76$ ,  $z = -3.13$ ,  $p = 0.005$ ) (**Fig. 7b**).





**Figure 7:** Frequency (occurrence per h) of **(a)** sociopositive behaviour and **(b)** courtship and sexual behaviour in the first (phase 1), second (phase 2) and third (phase 3) two weeks of treatment. Males in heterosexual pairs were either socially stimulated (PM+S) or not (PM-S). Plotted are medians (data points) and first to third quartiles (whiskers). Statistics: Multiple comparisons of GLMM. PM-S:  $n_{\text{Phase 1}} = 20$ ,  $n_{\text{Phase 2}} = 20$ ,  $n_{\text{Phase 3}} = 20$ , PM+S:  $n_{\text{Phase 1}} = 20$ ,  $n_{\text{Phase 2}} = 20$ ,  $n_{\text{Phase 3}} = 20$ ; \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ .

## 4. Discussion

In this study, we investigated how juvenile male guinea pigs adjust to distinct social environments through possible shaping of behavioural and hormonal phenotypes. By repeatedly analysing behavioural and hormonal parameters during juvenility, we aimed to explore when and how adjustments through behavioural plasticity occur in this early phase. For this purpose, male guinea pigs kept under pair-housing conditions with one female only (PM-S) were compared with males who lived with one female and received additional social stimulation by interactions with unfamiliar males and females (PM+S). Stimulated males showed an initially increased cortisol responsiveness which decreased again over time, as well as an increase in courtship and sexual behaviour over time. Moreover, cortisol responsiveness was significantly affected by body weight, this finding was however independent of treatment group.

### 4.1 Modulation of cortisol responsiveness by the social environment

Interestingly, cortisol responsiveness was different between treatment groups. More specifically, in the cortisol response test (CRT) conducted two weeks after start of social stimulation, cortisol responsiveness after one hour was higher in stimulated males than in non-stimulated males. Since baseline cortisol values did not differ between males of both treatment groups, social stimulation per se did not lead to prolonged higher stress levels. However, animals confronted with unpredictable interactions with unfamiliar conspecifics live in a much more challenging environment. Under such

conditions, a higher endocrine responsiveness to stressors in such a situation could be adaptive. This reactivity provides the organism with energy and shifts it into a state of heightened reactivity which is a prerequisite for responding to environmental challenges in an appropriate way. This has already been demonstrated in birds, where individuals with higher corticosterone responses are more successful in unpredictable conditions and thus better able to cope with environmental change [44, 45]. Consequently, the heightened stress response to this unpredictable environment presumably constitutes an adjustment process in stimulated males. This adjustment could also be interpreted as a process of *social niche conformance*. The concept of individualized social niches has recently gained prominence in behavioural biology and can be understood as “unit consisting of a focal individual and only those social interactions with other conspecific individuals that influence the focal individual’s inclusive fitness” [2]. Within this framework, social niche conformance describes the process where individuals adjust to an existing social environment, for example by adjustments of the behavioural or hormonal phenotype [2, 3, 7, 46]. In line with this interpretation, the significant decrease in cortisol responsiveness found in stimulated males in the subsequent CRTs could also reflect such a conformance process. At the end of the experimental phase cortisol responsiveness of stimulated and non-stimulated males almost converged, indicating juvenile males could adjust to the challenging situation. A stress-induced HPA activation is metabolically costly. Thus, it is adaptive for an organism to reduce HPA activity to stressors without harm [47].

Baseline cortisol levels did not differ between the treatment groups. Other studies have also reported no differences in baseline cortisol levels in guinea pigs living in different social environments [14, 29] or of different social status [48]. These findings suggest no influence of the social environment on baseline cortisol in guinea pigs, unlike in other species, such as mice, where plasma glucocorticoids can be affected by social interactions [49]. These differences may be due to differences in the social organization of these species. While male guinea pigs are able to integrate into unfamiliar groups with several adult males and females [12], male mice aggressively defend their territory and monopolize several females [50–52]. Another possible explanation is the sample size of baseline cortisol, which might have been too small to detect differences, since collecting a sufficient amount of blood was sometimes not possible.

Finally, the results from the repeatability analysis are in line with a meta-analysis, showing that repeatability estimates tend to be higher for peak hormone levels than for baseline levels [53]. The reason for this might elevated hormone responses (e.g., through stress) capturing a more defined aspect of endocrine function, while baseline hormone levels can represent multiple different biological functions [54, 55]. However, the results obtained here should be interpreted with caution, as the confidence intervals were wide and either close to or included zero [56, 57].

## **4.2 Social environment affected courtship and sexual behaviour, but not testosterone concentrations**

Sociopositive behaviour significantly increased from the beginning to the end of the experimental phase in both treatment groups, suggesting a social relationship has been established between the males and their respective female partners [26]. Sexual and courtship behaviour, however, only significantly increased over time in socially stimulated males. This finding leads to the consideration of socially stimulated males reaching sexual maturity earlier than non-stimulated males. Usually, sexual maturity is accompanied by a peak in testosterone concentration in male rodents [58, 59] and studies in Syrian hamsters have also shown most pronounced effects of testosterone on the organization of neural behavioural circuits and thus sexual behaviour are most pronounced during adolescence [60, 61]. Yet, we neither found differences in testosterone levels between stimulated- or non-stimulated males, nor did testosterone levels significantly increase over time in stimulated males in this study. Thus, an early onset of sexual maturity is unlikely to explain the significant increase in courtship and sexual behaviour. Instead, we favour a different explanation: stimulated males were able to observe such behaviour from adult stimulus males courting the focus male's female partner during the stimulation sessions. Immature guppies, for example, also learn courtship behaviour by observing experienced male conspecifics [62].

The lack of significant differences in testosterone levels between the treatment groups is also surprising for another reason: studies in adolescent male guinea pigs have demonstrated a causal relationship between the frequency of social interactions and increased testosterone concentrations [18, 29]. Yet, it is also known that for adolescent males specifically courting and agonistic encounters are responsible for increased testosterone levels [12, 63]. For juvenile male guinea pigs, however, it is unclear whether male-male interactions are really agonistic, and if male-female interactions are really sexual, when the male has not reached sexual maturity yet. In a study where baseline testosterone levels between colony-housed and individually-housed males were measured repeatedly from juvenility until adulthood, significant differences were also only found from an age of 90 days (i.e., adolescence), but not an age of 30 or 60 days (i.e., juvenility) [64].

## **4.3 Body weight as fitness proxy and its negative effect on cortisol responsiveness**

While reproductive success is a direct measurement of fitness, body weight as an index of body condition can be used as fitness proxy [65]. Body weight is related to many life history parameters, such as reproduction, survival and longevity. Animals with higher body weight have more body fat and in consequence more stored excess energy, which is beneficial for several reasons. They are better able to withstand harsher environmental conditions, and the development and expression of secondary

sexual traits are often dependent on body condition [66]. A larger body weight can also indirectly influence reproduction via a link to higher dominance status in social systems, as it was already demonstrated in guinea pigs and cavies [67, 68]. However, no differences regarding body weight were found between the treatment groups in this study. Furthermore, repeatability was very high in both stimulated and non-stimulated males, indicating that body weight is a stable individual trait independent of social environment.

More interestingly, body weight was significantly negatively correlated with cortisol responsiveness after 1 hour. The relationship between stress and body weight in animals has been studied a lot. The effects of acute stress on metabolic phenotypes can range from stress-induced anorexia [69] to increased food intake and thus obesity [70] and are influenced by factors like animal model and type of stress [71]. Regarding stress response, studies indicated high stress responsiveness is linked to obesity [72, 73]. However, the animals in this study were not only non-obese, but also a negative relationship was found between body weight and cortisol response, most presumably hinting at a different physiological process involved here. Even though no statistical differences between treatment groups could be found, this effect seemed to be more pronounced in socially stimulated males, since they had more negative and significant correlations for all time points. At the last time point, however, the correlation between body weight and cortisol responsiveness in non-stimulated males was almost as high as in stimulated males and also significant. This suggests an earlier onset of the effect that causes higher body weight to negatively influence cortisol responsiveness in socially stimulated males, potentially due to prior shaping of the HPA axis. This might also constitute a mechanism of the niche conformance process. However, further research is needed to determine underlying mechanisms.

Furthermore, it is particularly interesting that only cortisol responsiveness after 1 hour, but not after 2 hours, was affected by body weight. In guinea pigs, maximum cortisol responsiveness is usually reached after 2 hours, so cortisol responsiveness after 2 hours can be characterised as magnitude of stress response and cortisol responsiveness after 1 hour as speed of stress response [74, 75]. Speed and magnitude of stress response are correlated and especially speed of stress response is an important factor and possible target of selection [75], as it determines how quickly individuals can adjust to changes [76]. The observation of only cortisol responsiveness after 1 hour, but neither baseline cortisol levels nor cortisol responsiveness after 2 hours being negatively affected by body weight, indicates guinea pig males with higher body weights have a slower cortisol response. This would mean the maximum stress response might not be different between bigger and smaller individuals, but the time it takes to reach this maximum. Reasons for this could involve body weight dependent differences in the adrenal gland and availability or secretion of cortisol or cortisol binding globulins. Still, these hypotheses cannot yet be verified or explained, since studies investigating the exact physiological mechanisms involved in stress response in guinea pigs are lacking.

## 5. Conclusions

Socially stimulated males showed different adjustments to their social environment: at the beginning of the experimental phase, they displayed an increased stress response to be able to adequately react to the unpredictable social encounters. However, since such increases in stress are metabolically costly and social stimulation were not actually dangerous, the males then adjusted to this challenging environment and displayed a decrease in stress response again. Furthermore, body weight was found to have a significant, negative impact on speed of cortisol reactivity. These findings indicate the speed of cortisol reactivity is a flexible trait and able to adjust to external (social environment) and internal (body weight) parameters and thus forming the basis for individualised niches. Moreover, social stimulation did not only affect endocrine parameters, but also behaviour: while males of both treatment groups displayed a significant increase of sociopositive behaviour over time, only males with additional social stimulation also displayed a significant increase of courtship and sexual behaviour over time. Taken together, these findings demonstrate that already in juvenile guinea pigs the social environment induced hormonal adjustments and behavioural changes, thereby laying the grounds for social niche conformance. This process involves (behavioural) plasticity but goes beyond it by focusing on individual-by-environment interactions [77] and by emphasizing consequences for phenotype-environment-match and thus fitness [46]. For future studies repeating these experiments with adolescent males to investigate social niche conformance throughout ontogeny, we would expect the effects found here are further pronounced and persistent since social interactions become even more meaningful once the individuals reach sexual maturity.

## **Ethics**

All procedures complied with the regulations covering animal experimentation within Germany (Animal Welfare Act) and the EU (European Communities Council Directive 2010/ 63/ EU), and were approved by the local and federal authorities (Landesamt für Natur, Umwelt und Verbraucherschutz Nordrhein-Westfalen “LANUV NRW”, reference number: 81-02.04.2022.A080).

## **Funding**

This research was funded by the German Research Foundation (DFG) as part of the SFB TRR 212 (NC3), project numbers 316099922 and 396777165 (responsible PI: S.K.).

## **CRediT authorship contribution statement**

**Melanie Gleske:** Methodology; writing – original draft; investigation; formal analysis; visualization; data curation. **S. Helene Richter:** Conceptualization; Writing – review and editing. **Sylvia Kaiser:** Conceptualization; methodology; supervision, writing – review and editing; funding acquisition. **Carolin Munding:** Formal analysis. **All authors critically revised the manuscript and gave final approval for publication.**

## **Declaration of competing interests**

The authors declare no conflict of interests.

## **Data availability**

Open data/code are not available yet but will be accessible with peer-reviewed publication.

## **Acknowledgements**

We would like to thank Sabine Kruse for conducting the endocrine analysis and Raphael Beermann and Clarissa Lindemann for animal caretaking and assistance with the experiments. We are furthermore grateful to Maximilian Baldy for assistance with the experiments and for constructive comments and suggestions that improved the manuscript.

## **Appendix A. Supplementary data**

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## Supplementary material

Shaped from an early age: behavioural and hormonal phenotypes in juvenile  
male guinea pigs living in distinct social environments

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## Results: Descriptive statistics

**Table S1:** Descriptive statistics for baseline cortisol (c0), cortisol responsiveness after 1 hour (c1) and 2 hours (c2) of exposure to a novel environment and baseline testosterone (t).

Treatment	Hormone	Time point	n	mean	SD	min	max
PM+S	c0	CRT0	8	520.70	298.84	199.23	979.73
		CRT1	9	192.44	126.20	78.06	431.48
		CRT2	6	95.93	58.19	50.22	208.83
		CRT3	4	162.71	44.02	106.60	205.88
		Overall	27	263.85	244.88	50.22	979.73
	c1	CRT0	10	1583.90	365.44	968.58	2110.78
		CRT1	10	1036.95	236.58	718.33	1454.35
		CRT2	10	712.83	137.68	475.50	1017.43
		CRT3	10	729.12	193.55	380.30	942.00
		Overall	40	1015.70	429.22	380.30	2110.78
	c2	CRT0	9	1736.53	525.90	732.33	2296.40
		CRT1	10	1249.38	336.44	505.60	1599.15
		CRT2	10	959.57	197.91	660.11	1222.77
		CRT3	7	1029.25	196.66	630.45	1262.53
		Overall	36	1247.86	451.37	505.60	2296.40
	t	CRT0	7	1.00	0.44	0.66	1.87
		CRT1	10	1.91	0.68	0.98	2.84
		CRT2	10	2.17	0.94	0.95	3.87
		CRT3	10	2.68	1.63	1.14	5.38
		Overall	37	2.02	1.17	0.66	5.38
PM-S	c0	CRT0	9	414.35	184.21	141.28	712.00
		CRT1	8	145.56	69.33	69.57	254.20
		CRT2	9	136.86	61.10	73.58	264.23
		CRT3	5	146.43	23.05	126.85	185.23
		Overall	31	221.21	164.38	69.57	712.00
	c1	CRT0	9	1438.96	221.49	1032.23	1635.13
		CRT1	10	883.41	163.18	696.33	1110.35
		CRT2	10	822.30	131.59	580.20	969.28
		CRT3	10	768.61	166.59	530.18	1009.53
		Overall	39	966.51	312.68	530.18	1635.13
	c2	CRT0	9	1645.85	175.58	1370.85	1835.45
		CRT1	9	1158.78	227.61	736.08	1467.13
		CRT2	10	1155.06	189.08	892.17	1373.48
		CRT3	9	1136.01	234.90	799.63	1481.10
		Overall	37	1270.71	293.48	736.08	1835.45
	t	CRT0	9	1.18	0.87	0.25	3.22
		CRT1	9	2.30	1.35	0.34	4.96
		CRT2	10	2.69	1.38	1.30	5.68
		CRT3	10	2.40	0.76	1.17	4.02
		Overall	38	2.16	1.22	0.25	5.68

**Table S2:** Descriptive statistics for body weight.

Treatment	Time point	n	mean	SD	min	max
PM+S	CRT0	10	254.20	44.17	211	359
	CRT1	10	360.60	58.31	293	488
	CRT2	10	476.70	72.12	343	605
	CRT3	10	512.30	66.15	417	628
	Overall	40	400.95	118.47	211	628
PM-S	CRT0	10	263.50	32.76	193	313
	CRT1	10	359.80	35.49	271	395
	CRT2	10	461.00	36.16	372	495
	CRT3	10	502.90	38.18	411	538
	Overall	40	396.80	100.13	193	538

**Table S3:** Descriptive statistics for behaviour.

Treatment	Behaviour	Time point	n	mean	SD	min	max
PM+S	Sociopositive	Phase 1	20	2.12	2.02	0	6.50
		Phase 2	20	4.65	3.78	0.50	15.00
		Phase 3	20	7.78	7.27	0	26.00
		Overall	60	4.85	5.33	0	26.00
	Courtsip and sexual	Phase 1	20	0.15	0.29	0	1.00
		Phase 2	20	0.48	0.72	0	2.33
		Phase 3	20	1.70	3.16	0	13.50
		Overall	60	0.78	1.96	0	13.50
	Play	Phase 1	20	0.54	1.80	0	7.50
		Phase 2	20	0.48	1.27	0	5
		Phase 3	20	0.66	1.30	0	4.50
		Overall	60	0.56	1.45	0	7.50
PM-S	Sociopositive	Phase 1	20	2.52	3.75	0	10.67
		Phase 2	20	3.43	5.08	0	20.50
		Phase 3	20	5.98	8.06	0	36.33
		Overall	60	3.97	5.99	0	36.33
	Courtsip and sexual	Phase 1	20	0.60	1.05	0	3.33
		Phase 2	20	1.23	2.86	0	12.50
		Phase 3	20	1.72	4.41	0	19.67
		Overall	60	1.18	3.07	0	19.67
	Play	Phase 1	20	0.25	0.79	0	3
		Phase 2	20	0.08	0.18	0	0.50
		Phase 3	20	0.48	1.34	0	5
		Overall	60	0.27	0.90	0	5

## Results: Wilcoxon test for treatment comparisons of hormone concentrations and body weight at CRT0

**Table S4:** Wilcoxon rank sum test of hormone concentrations and body weight calculated for the first cortisol response test (CRT) conducted before treatment.

Wilcoxon rank sum test (CRT0)	W	r	p-value
Baseline cortisol	30	0.118	0.596
Cortisol responsiveness, 1h	31	0.246	0.270
Cortisol responsiveness, 2h	31	0.178	0.427
Baseline testosterone	35.5	0.083	0.711
Body weight	62.5	0.203	0.364

## Results: Model summaries of linear mixed effect models for hormone concentrations

**Table S5:** Model summary from mixed effect model (individual ID as random effect) to determine overall effect of time (CRT), treatment, time\*treatment interaction and body weight on baseline cortisol. CRT1 (time) and SM-S (treatment) were set as reference level by default.

Fixed effects		Estimate	Std. error	[95% CI]	t-value	p-value	R <sup>2</sup>
Baseline cortisol (Transformation: sqrt(x)) N = 41						Full model: Marginal R <sup>2</sup>	0.165
						Full model: Conditional R <sup>2</sup>	
	Intercept	11.298	1.106	[9.051, 13.546]	10.217	< 0.001	
Treatment		1.541	1.418	[-1.341, 4.423]	1.087	0.285	0.034
CRT1 - CRT2		0.982	1.793	[-2.662, 4.627]	0.548	0.587	0.009
CRT1 - CRT3		2.084	2.255	[-2.500, 6.667]	0.924	0.362	0.025
Body weight		-0.011	0.010	[-0.031, 0.008]	-1.164	0.253	0.038
Treatment*CRT1-CRT2		-3.117	2.115	[-7.416, 1.181]	-1.474	0.150	0.060
Treatment*CRT1-CRT3		-1.033	2.419	[-5.948, 3.883]	-0.427	0.672	0.005

**Table S6:** Model summary from mixed effect model (individual ID as random effect) to determine overall effect of time (CRT), treatment, time\*treatment interaction and body weight on cortisol responsiveness after 1 hour of exposure to a novel environment. CRT1 (time) and SM-S (treatment) were set as reference level by default. Significant ( $p < 0.05$ ) results are indicated in bold.

Fixed effects		Estimate	Std. error	[95% CI]	t-value	p-value	R <sup>2</sup>
Cortisol responsiveness 1h (Transformation: sqrt(x)) N = 60						Full model: Marginal R <sup>2</sup>	0.511
						Full model: Conditional R <sup>2</sup>	0.642
	Intercept	28.320	0.837	[26.631, 30.010]	33.836	< 0.001	
Treatment		2.435	1.106	[0.207, 4.662]	2.201	<b>0.033</b>	0.085
CRT1 - CRT2		2.312	1.221	[-0.137, 4.761]	1.894	0.064	0.071
CRT1 - CRT3		2.674	1.444	[-0.234, 5.583]	1.852	0.071	0.079
Body weight		-0.033	0.008	[-0.049, -0.017]	-4.315	<b>&lt; 0.001</b>	0.374
Treatment*CRT1-CRT2		-3.917	1.342	[-6.641, -1.194]	-2.918	<b>0.006</b>	0.105
Treatment*CRT1-CRT3		-2.936	1.339	[-5.654, -0.218]	-2.192	<b>0.035</b>	0.062

**Table S7:** Model summary from mixed effect model (individual ID as random effect) to determine overall effect of time (CRT), treatment, time\*treatment interaction and body weight on cortisol responsiveness after 2 hours of exposure to a novel environment. CRT1 (time) and SM-S (treatment) were set as reference level by default. Significant ( $p < 0.05$ ) results are indicated in bold.

Fixed effects		Estimate	Std. error	[95% CI]	t-value	p-value	R <sup>2</sup>
Cortisol responsiveness 2h (Transformation: sqrt(x)) N = 55						Full model: Marginal R <sup>2</sup>	0.175
						Full model: Conditional R <sup>2</sup>	0.543
	Intercept	32.914	1.245	[30.383, 35.444]	26.430	< 0.001	
Treatment		1.414	1.609	[-1.856, 4.684]	0.879	0.386	0.011
CRT1 - CRT2		2.031	1.707	[-1.402, 5.465]	1.190	0.240	0.045
CRT1 - CRT3		2.547	2.112	[-1.720, 6.813]	1.206	0.235	0.056
Body weight		-0.017	0.012	[-0.042, 0.008]	-1.431	0.167	0.149
Treatment*CRT1-CRT2		-4.198	1.702	[-7.675, -0.722]	-2.467	<b>0.020</b>	0.056
Treatment*CRT1-CRT3		-3.101	1.832	[-6.839, 0.636]	-1.693	0.101	0.032



**Table S8:** Model summary from mixed effect model (individual ID as random effect) to determine overall effect of time (CRT), treatment, time\*treatment interaction and body weight on baseline testosterone. CRT1 (time) and SM-S (treatment) were set as reference level by default.

Fixed effects		Estimate	Std. error	[95% CI]	t-value	p-value	R <sup>2</sup>
Baseline testosterone (Transformation: sqrt(x)) N = 59						Full model: Marginal R <sup>2</sup>	0.053
						Full model: Conditional R <sup>2</sup>	
	Intercept	1.475	0.130	[1.215, 1.736]	11.364	< 0.001	
Treatment		-0.094	0.171	[-0.437, 0.249]	-0.549	0.585	0.006
CRT1 - CRT2		0.087	0.197	[-0.307, 0.482]	0.445	0.658	0.004
CRT1 - CRT3		0.001	0.219	[-0.439, 0.441]	0.003	0.997	0.000
Mass		0.001	0.001	[-0.001, 0.002]	0.567	0.573	0.006
Treatment*CRT1-CRT2		-0.070	0.239	[-0.549, 0.409]	-0.293	0.771	0.002
Treatment*CRT1-CRT3		0.130	0.239	[-0.349, 0.609]	0.545	0.588	0.006

## Results: Multiple comparisons of linear mixed effect models of hormone concentrations

**Table S9:** Multiple comparisons (Tukey's) of linear mixed effect model to determine effects of time (CRT), treatment and time\*treatment interaction on baseline cortisol.

Baseline cortisol	Estimate	Std. error	df	[95% CI]	t-value	p-value
Pair-wise comparison (between treatment groups)						
CRT1	-1.541	1.438	33.951	[-4.463, 1.381]	-1.072	0.291
CRT2	1.577	1.605	33.909	[-1.686, 4.839]	0.982	0.333
CRT3	-0.508	2.044	33.965	[-4.663, 3.646]	-0.249	0.805
Pair-wise comparison (within treatment groups)						
CRT 1 - CRT 2 (PM-S)	-0.982	1.844	33.998	[-5.502, 3.537]	-0.533	0.856
CRT 1 - CRT 3 (PM-S)	-2.084	2.357	30.936	[-7.886, 3.719]	-0.884	0.654
CRT 2 - CRT 3 (PM-S)	-1.101	1.728	26.512	[-5.391, 3.188]	-0.637	0.801
CRT 1 - CRT 2 (PM+S)	2.135	2.161	33.034	[-3.168, 7.439]	0.988	0.590
CRT 1 - CRT 3 (PM+S)	-1.051	2.335	29.630	[-6.811, 4.709]	-0.450	0.895
CRT 2 - CRT 3 (PM+S)	-3.186	1.944	26.604	[-8.010, 1.637]	-1.639	0.247
Interaction contrasts (treatment*CRT)						
CRT1 - CRT2	3.117	2.152	24.259	[-1.321, 7.556]	1.449	0.160
CRT1 - CRT3	1.033	2.513	29.198	[-4.105, 6.170]	0.411	0.684
CRT2 - CRT3	-2.085	2.604	25.472	[-7.443, 3.273]	-0.801	0.431

**Table S10:** Multiple comparisons (Tukey's) of linear mixed effect model to determine effects of time (CRT), treatment and time\*treatment interaction on cortisol responsiveness after 1 hour of exposure to a novel environment. Significant ( $p < 0.05$ ) results are indicated in bold.

Cortisol responsiveness, 1h	Estimate	Std. error	df	[95% CI]	t-value	p-value
Pair-wise comparison (between treatment groups)						
CRT1	-2.435	1.106	45.448	[-4.662, -0.207]	-2.201	<b>0.033</b>
CRT2	1.483	1.113	45.143	[-0.758, 3.724]	1.333	0.189
CRT3	0.502	1.109	45.339	[-1.731, 2.734]	0.452	0.653
Pair-wise comparison (between CRTs)						
CRT 1 - CRT 2 (PM-S)	-2.312	1.229	52.686	[-5.275, 0.651]	-1.882	0.154
CRT 1 - CRT 3 (PM-S)	-2.674	1.457	45.455	[-6.205, 0.856]	-1.835	0.170
CRT 2 - CRT 3 (PM-S)	-0.362	1.000	42.549	[-2.791, 2.066]	-0.362	0.930
CRT 1 - CRT 2 (PM+S)	1.605	1.305	50.823	[-1.546, 4.757]	1.230	0.441
CRT 1 - CRT 3 (PM+S)	0.262	1.509	43.677	[-3.398, 3.922]	0.174	0.984
CRT 2 - CRT 3 (PM+S)	-1.344	0.985	40.817	[-3.740, 1.053]	-1.364	0.369
Interaction contrasts (treatment*CRT)						
CRT1 - CRT2	3.917	1.343	36.269	[1.195, 6.640]	2.918	<b>0.006</b>
CRT1 - CRT3	2.936	1.339	35.960	[0.220, 5.653]	2.192	<b>0.035</b>
CRT2 - CRT3	-0.981	1.339	35.888	[-3.696, 1.734]	-0.733	0.468

**Table S11:** Multiple comparisons (Tukey's) of linear mixed effect model to determine effects of time (CRT), treatment and time\*treatment interaction on cortisol responsiveness after 2 hours of exposure to a novel environment. Significant ( $p < 0.05$ ) results are indicated in bold.

Cortisol responsiveness, 2h	Estimate	Std. error	df	[95% CI]	t-value	p-value
Pair-wise comparison (between treatment groups)						
CRT1	-1.414	1.611	35.989	[-4.681, 1.853]	-0.878	0.386
CRT2	2.784	1.589	34.510	[-0.444, 6.013]	1.752	0.089
CRT3	1.687	1.731	40.019	[-1.812, 5.186]	0.975	0.336
Pair-wise comparison (between CRTs)						
CRT 1 - CRT 2 (PM-S)	-2.031	1.727	47.495	[-6.208, 2.145]	-1.177	0.473
CRT 1 - CRT 3 (PM-S)	-2.547	2.144	41.493	[-7.757, 2.664]	-1.188	0.467
CRT 2 - CRT 3 (PM-S)	-0.515	1.320	39.457	[-3.730, 2.699]	-0.390	0.920
CRT 1 - CRT 2 (PM+S)	2.167	1.844	44.515	[-2.303, 6.637]	1.175	0.474
CRT 1 - CRT 3 (PM+S)	0.555	2.225	37.425	[-4.875, 5.985]	0.249	0.966
CRT 2 - CRT 3 (PM+S)	-1.612	1.382	38.321	[-4.981, 1.756]	-1.167	0.480
Interaction contrasts (treatment*CRT)						
CRT1 - CRT2	4.198	1.704	31.886	[0.728, 7.669]	2.464	<b>0.019</b>
CRT1 - CRT3	3.101	1.838	32.550	[-0.640, 6.843]	1.687	0.101
CRT2 - CRT3	-1.097	1.813	31.909	[-4.790, 2.596]	-0.605	0.549

**Table S12:** Multiple comparisons (Tukey's) of linear mixed effect model to determine effects of time (CRT), treatment and time\*treatment interaction on baseline testosterone.

Baseline testosterone	Estimate	Std. error	df	[95% CI]	t-value	p-value
Pair-wise comparison (between treatment groups)						
CRT1	0.094	0.171	51.967	[-0.250, 0.437]	0.548	0.586
CRT2	0.164	0.167	51.932	[-0.171, 0.499]	0.980	0.332
CRT3	-0.036	0.167	51.955	[-0.370, 0.298]	-0.218	0.828
Pair-wise comparison (between CRTs)						
CRT 1 - CRT 2 (PM-S)	-0.087	0.198	50.284	[-0.565, 0.390]	-0.443	0.898
CRT 1 - CRT 3 (PM-S)	-0.001	0.220	51.403	[-0.533, 0.531]	-0.003	1.000
CRT 2 - CRT 3 (PM-S)	0.087	0.171	38.726	[-0.330, 0.504]	0.507	0.868
CRT 1 - CRT 2 (PM+S)	-0.018	0.200	51.749	[-0.501, 0.465]	-0.088	0.996
CRT 1 - CRT 3 (PM+S)	-0.131	0.221	50.309	[-0.664, 0.403]	-0.592	0.825
CRT 2 - CRT 3 (PM+S)	-0.113	0.170	37.724	[-0.527, 0.301]	-0.667	0.784
Interaction contrasts (treatment*CRT)						
CRT1 - CRT2	0.070	0.239	35.905	[-0.415, 0.555]	0.292	0.772
CRT1 - CRT3	-0.130	0.239	35.737	[-0.614, 0.354]	-0.545	0.589
CRT2 - CRT3	-0.200	0.235	35.050	[-0.677, 0.278]	-0.850	0.401

## Results: Model summary of linear mixed effect model for body weight

**Table S13:** Model summary from mixed effect model (individual ID as random effect) to determine overall effect of time (CRT), treatment and time\*treatment interaction on body weight. CRT1 (time) and SM-S (treatment) were set as reference level by default. Significant (p < 0.05) results are indicated in bold.

Fixed effects		Estimate	Std. error	[95% CI]	t-value	p-value	R <sup>2</sup>
Body weight N = 60						Full model: Marginal R <sup>2</sup>	0.585
						Full model: Conditional R <sup>2</sup>	0.968
	Intercept	359.8	16.834	[324.683, 394.917]	21.374	< 0.001	
Treatment		0.8	23.807	[-48.862, 50.462]	0.034	0.974	< 0.001
CRT1 - CRT2		101.2	6.623	[87.768, 114.632]	15.280	<b>&lt; 0.001</b>	0.251
CRT1 - CRT3		143.1	6.623	[129.668, 156.532]	21.606	<b>&lt; 0.001</b>	0.401
Treatment*CRT1-CRT2		14.9	9.366	[-4.096, 33.896]	1.591	0.120	0.004
Treatment*CRT1-CRT3		8.6	9.366	[-10.396, 27.596]	0.918	0.365	0.001

## Results: Multiple comparisons of linear mixed effect model of body weight

**Table S14:** Multiple comparisons (Tukey's) of linear mixed effect model to determine effects of time (CRT), treatment and time\*treatment interaction on body weight. Significant ( $p < 0.05$ ) results are indicated in bold.

Body weight	Estimate	Std. error	df	[95% CI]	t-value	p-value
Pair-wise comparison (between treatment groups)						
CRT1	-0.800	23.807	19.982	[-50.462, 48.862]	-0.034	0.974
CRT2	-15.700	23.807	19.982	[-65.362, 33.962]	-0.659	0.517
CRT3	-9.400	23.807	19.982	[-59.062, 40.262]	-0.395	0.697
Pair-wise comparison (within treatment groups)						
CRT 1 - CRT 2 (PM-S)	-101.200	6.623	36	[-117.389, -85.011]	-15.280	<b>&lt; 0.001</b>
CRT 1 - CRT 3 (PM-S)	-143.100	6.623	36	[-159.289, -126.911]	-21.606	<b>&lt; 0.001</b>
CRT 2 - CRT 3 (PM-S)	-41.900	6.623	36	[-58.089, -25.711]	-6.326	<b>&lt; 0.001</b>
CRT 1 - CRT 2 (PM+S)	-116.100	6.623	36	[-132.289, -99.911]	-17.530	<b>&lt; 0.001</b>
CRT 1 - CRT 3 (PM+S)	-151.700	6.623	36	[-167.889, -135.511]	-22.905	<b>&lt; 0.001</b>
CRT 2 - CRT 3 (PM+S)	-35.600	6.623	36	[-51.789, -19.411]	-5.375	<b>&lt; 0.001</b>
Interaction contrasts (treatment*CRT)						
CRT1 - CRT2	-14.900	9.366	36	[-33.896, 4.096]	-1.591	0.120
CRT1 - CRT3	-8.600	9.366	36	[-27.596, 10.396]	-0.918	0.365
CRT2 - CRT3	6.300	9.366	36	[-12.696, 25.296]	0.673	0.505

## Results: Correlation between body weight and cortisol responsiveness after 1 hour

**Table S15:** Calculation of correlation coefficient (Pearson) and significance testing for correlations (z-test) between body weight and cortisol responsiveness after 1 hour of exposure to a novel environment. Significant ( $p < 0.05$ ) results are indicated in bold.

Correlation between c1 and body weight	r	t-value	p-value
Within treatment groups			
CRT1 (PM+S)	-0.808	-3.883	<b>0.005</b>
CRT2 (PM+S)	-0.687	-2.671	<b>0.028</b>
CRT3 (PM+S)	-0.742	-3.131	<b>0.014</b>
Overall (PM+S)	-0.586	-3.829	<b>&lt; 0.001</b>
CRT1 (PM-S)	-0.258	-0.755	0.472
CRT2 (PM-S)	-0.110	-0.314	0.762
CRT3 (PM-S)	-0.714	-2.885	<b>0.020</b>
Overall (PM-S)	-0.350	-1.979	0.058
Comparison between treatment groups			
CRT1		-1.606	0.108
CRT2		-1.367	0.172
CRT3		-0.111	0.911
Overall		-1.125	0.261

## Results: Adjusted repeatability analysis of hormone concentrations and body weight

**Table S16:** Adjusted repeatability analysis of linear mixed effects models of baseline cortisol (c0), cortisol responsiveness after 1 hour of exposure to a novel environment (c1), cortisol responsiveness after 2 hours of exposure to a novel environment (c2), baseline testosterone (t) and body weight (m0). Significant ( $p < 0.05$ ) results are indicated in bold.

Repeatability	PM+S				PM-S			
	Std. error	[95% CI]	R	p-value	Std. error	[95% CI]	R	p-value
c0	0.254	[0, 0.805]	0	1	0.232	[0, 0.74]	0.175	0.331
c1	0.155	[0, 0.509]	0.042	0.495	0.196	[0.03, 0.793]	0.453	<b>0.014</b>
c2	0.210	[0, 0.786]	0.416	<b>0.040</b>	0.194	[0.085, 0.849]	0.523	<b>0.015</b>
t	0.164	[0, 0.549]	0.069	0.440	0.143	[0, 0.466]	0	1
m0	0.057	[0.784, 0.977]	0.927	<b>&lt; 0.001</b>	0.069	[0.723, 0.97]	0.908	<b>&lt; 0.001</b>

## Results: Model summaries of generalized linear mixed effect models for behaviour

**Table S17:** Model summary from generalized linear mixed effect model (individual ID as random effect) to determine overall effect of time (Phase), treatment and time\*treatment interaction on sociopositive behaviour. Phase 1 (time) and SM-S (treatment) were set as reference level by default. Significant ( $p < 0.05$ ) results are indicated in bold.

Fixed effects		Estimate	Std. error	[95% CI]	z-value	p-value	R <sup>2</sup>
Sociopositive behaviour N = 60						Full model: Marginal R <sup>2</sup>	0.213
						Full model: Conditional R <sup>2</sup>	0.376
	Intercept	0.726	0.294	[0.150, 1.303]	2.469	0.014	
Treatment		-0.004	0.407	[-0.801, 0.794]	-0.009	0.993	< 0.001
Phase 1 - Phase 2		0.311	0.350	[-0.374, 0.996]	0.890	0.374	0.002
Phase 1 - Phase 3		0.917	0.335	[0.261, 1.573]	2.741	<b>0.006</b>	0.035
Treatment*Phase 1 - Phase 2		0.430	0.484	[-0.518, 1.378]	0.890	0.374	0.004
Treatment*Phase 1 - Phase 3		0.376	0.471	[-0.548, 1.299]	0.797	0.426	0.007

**Table S18:** Model summary from generalized linear mixed effect model (individual ID as random effect) to determine overall effect of time (Phase), treatment and time\*treatment interaction on sexual and courtship behaviour. Phase 1 (time) and SM-S (treatment) were set as reference level by default.

Fixed effects		Estimate	Std. error	[95% CI]	z-value	p-value	R <sup>2</sup>
<b>Courtship and sexual behaviour N = 60</b>						<i>Full model:</i> <i>Marginal R<sup>2</sup></i>	0.304
						<i>Full model:</i> <i>Conditional R<sup>2</sup></i>	0.373
	Intercept	-0.607	0.472	[-1.532, 0.318]	-1.286	0.198	
Treatment		-1.344	0.811	[-2.933, 0.245]	-1.658	0.097	0.003
Phase 1 - Phase 2		0.622	0.598	[-0.551, 1.795]	1.040	0.298	0.005
Phase 1 - Phase 3		0.988	0.579	[-0.146, 2.122]	1.707	0.088	0.016
Treatment*Phase 1 - Phase 2		0.555	1.001	[-1.406, 2.516]	0.555	0.579	0.001
Treatment*Phase 1 - Phase 3		1.391	0.950	[-0.471, 3.254]	1.464	0.143	0.001

**Table S19:** Model summary from generalized linear mixed effect model (individual ID as random effect) to determine overall effect of time (Phase), treatment and time\*treatment interaction on play behaviour. Phase 1 (time) and SM-S (treatment) were set as reference level by default.

Fixed effects		Estimate	Std. error	[95% CI]	z-value	p-value	R <sup>2</sup>
<b>Play behaviour N = 60</b>						<i>Full model:</i> <i>Marginal R<sup>2</sup></i>	0.143
						<i>Full model:</i> <i>Conditional R<sup>2</sup></i>	0.277
	Intercept	-1.567	0.784	[-3.103, -0.031]	-2.000	0.045	
Treatment		0.444	1.048	[-1.611, 2.499]	0.424	0.672	0.005
Phase 1 - Phase 2		-1.176	1.209	[-3.545, 1.194]	-0.972	0.331	0.002
Phase 1 - Phase 3		0.590	0.980	[-1.331, 2.511]	0.602	0.547	0.003
Treatment*Phase 1 - Phase 2		1.203	1.509	[-1.754, 4.160]	0.797	0.425	< 0.001
Treatment*Phase 1 - Phase 3		-0.222	1.331	[-2.830, 2.387]	-0.167	0.868	< 0.001

## Results: Multiple comparisons of generalized linear mixed effect models of behaviour

**Table S20:** Multiple comparisons (Tukey's) of generalized linear mixed effect model to determine effects of time (Phase), treatment and time\*treatment interaction on sociopositive behaviour. Significant ( $p < 0.05$ ) results are indicated in bold.

Sociopositive behaviour	Estimate	Std. error	[95% CI]	z-ratio	p-value
<b>Pair-wise comparison (between treatment groups)</b>					
Phase 1	0.004	0.407	[-0.794, 0.801]	0.009	0.993
Phase 2	-0.427	0.381	[-1.174, 0.321]	-1.119	0.263
Phase 3	-0.372	0.362	[-1.081, 0.337]	-1.028	0.304
<b>Pair-wise comparison (within treatment groups)</b>					
Phase 1 - Phase 2 (PM-S)	-0.311	0.350	[-1.131, 0.508]	-0.890	0.647
Phase 1 - Phase 3 (PM-S)	-0.917	0.335	[-1.702, -0.133]	-2.741	<b>0.017</b>
Phase 2 - Phase 3 (PM-S)	-0.606	0.329	[-1.378, 0.165]	-1.842	0.156
Phase 1 - Phase 2 (PM+S)	-0.742	0.334	[-1.524, 0.041]	-2.220	0.068
Phase 1 - Phase 3 (PM+S)	-1.293	0.331	[-2.070, -0.516]	-3.900	<b>&lt; 0.001</b>
Phase 2 - Phase 3 (PM+S)	-0.551	0.307	[-1.271, 0.169]	-1.795	0.171
<b>Interaction contrasts (treatment*CRT)</b>					
Phase 1 - Phase 2	-0.430	0.484	[-1.378, 0.518]	-0.890	0.374
Phase 1 - Phase 3	-0.376	0.471	[-1.299, 0.548]	-0.797	0.426
Phase 2 - Phase 3	0.055	0.449	[-0.826, 0.936]	0.122	0.903

**Table S21:** Multiple comparisons (Tukey's) of generalized linear mixed effect model to determine effects of time (Phase), treatment and time\*treatment interaction on courtship and sexual behaviour. Significant ( $p < 0.05$ ) results are indicated in bold.

Courtship and sexual behaviour	Estimate	Std. error	[95% CI]	z-ratio	p-value
<b>Pair-wise comparison (between treatment groups)</b>					
Phase 1	1.344	0.811	[-0.245, 2.933]	1.658	0.097
Phase 2	0.789	0.653	[-0.491, 2.069]	1.208	0.227
Phase 3	-0.047	0.560	[-1.145, 1.050]	-0.084	0.933
<b>Pair-wise comparison (within treatment groups)</b>					
Phase 1 - Phase 2 (PM-S)	-0.622	0.598	[-2.025, 0.780]	-1.040	0.552
Phase 1 - Phase 3 (PM-S)	-0.988	0.579	[-2.344, 0.368]	-1.707	0.202
Phase 2 - Phase 3 (PM-S)	-0.366	0.557	[-1.672, 0.941]	-0.656	0.789
Phase 1 - Phase 2 (PM+S)	-1.177	0.800	[-3.053, 0.698]	-1.472	0.305
Phase 1 - Phase 3 (PM+S)	-2.379	0.758	[-4.156, -0.603]	-3.139	<b>0.005</b>
Phase 2 - Phase 3 (PM+S)	-1.202	0.592	[-2.588, 0.185]	-2.032	0.105
<b>Interaction contrasts (treatment*CRT)</b>					
Phase 1 - Phase 2	-0.555	1.001	[-2.516, 1.406]	-0.555	0.579
Phase 1 - Phase 3	-1.391	0.950	[-3.254, 0.471]	-1.464	0.143
Phase 2 - Phase 3	-0.836	0.818	[-2.440, 0.767]	-1.022	0.307

**Table S22:** Multiple comparisons (Tukey's) of generalized linear mixed effect model to determine effects of time (Phase), treatment and time\*treatment interaction on play behaviour.

Play behaviour	Estimate	Std. error	[95% CI]	z-ratio	p-value
<b>Pair-wise comparison (between treatment groups)</b>					
Phase 1	-0.444	1.048	[-2.499, 1.611]	-0.424	0.672
Phase 2	-1.647	1.209	[-4.016, 0.722]	-1.363	0.173
Phase 3	-0.223	0.946	[-2.076, 1.631]	-0.235	0.814
<b>Pair-wise comparison (within treatment groups)</b>					
Phase 1 - Phase 2 (PM-S)	1.176	1.209	[-1.658, 4.009]	0.972	0.594
Phase 1 - Phase 3 (PM-S)	-0.590	0.980	[-2.887, 1.707]	-0.602	0.819
Phase 2 - Phase 3 (PM-S)	-1.766	1.164	[-4.494, 0.963]	-1.517	0.283
Phase 1 - Phase 2 (PM+S)	-0.027	0.910	[-2.160, 2.106]	-0.030	1.000
Phase 1 - Phase 3 (PM+S)	-0.368	0.891	[-2.456, 1.719]	-0.413	0.910
Phase 2 - Phase 3 (PM+S)	-0.341	0.886	[-2.419, 1.737]	-0.385	0.922
<b>Interaction contrasts (treatment*CRT)</b>					
Phase 1 - Phase 2	-1.203	1.509	[-4.160, 1.754]	-0.797	0.425
Phase 1 - Phase 3	0.222	1.331	[-2.387, 2.830]	0.167	0.868
Phase 2 - Phase 3	1.425	1.461	[-1.439, 4.289]	0.975	0.330