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Intact GR dimerization is critical for restraining plasma ACTH levels during chronic psychosocial stress

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ABSTRACT

Male C57BL/6N mice exposed to the chronic subordinate colony housing (CSC; 19 days) paradigm, a preclinically validated model of chronic psychosocial stress, are characterized by unaffected basal morning plasma corticosterone (CORT) concentrations despite adrenal and pituitary hyperplasia and increased adrenocorticotropic hormone (ACTH) plasma concentrations, compared with single-housed control (SHC) mice. However, as CSC mice are still able to show an increased CORT secretion towards novel heterotypic stressors, these effects might reflect an adaptation rather than a functional breakdown of general hypothalamus-pituitary-adrenal (HPA) axis functionality. In the present study we used male mice of a genetically modified mouse line, to investigate whether genetically-driven ACTH overexpression compromises adaptational processes occurring at the level of the adrenals during CSC exposure. Experimental mice carried a point mutation in the DNA binding domain of the glucocorticoid (GC) receptor (GR), attenuating dimerization of GR (GR^{dim}), resulting in a congenially compromised negative feedback inhibition at the level of the pituitary.

In line with previous studies, CSC mice in both the wild type (WT; GR^{+/+}) and GR^{dim} group developed adrenal enlargement. Moreover, compared with respective SHC and WT mice, CSC GR^{dim} mice show increased basal morning plasma ACTH and CORT concentrations. Quantitative polymerase chain reaction (qPCR) analysis revealed neither a genotype effect, nor a CSC effect on pituitary mRNA expression of the ACTH precursor proopiomelanocortin (POMC). Finally, CSC increased anxiety-related behavior, active coping and splenocyte *in vitro* (re)activity in both WT and GR^{dim} mice, while a CSC-induced increase in adrenal lipid vesicles and splenic GC resistance was detectable only in WT mice. Of note, lipopolysaccharide (LPS)-stimulated splenocytes of GR^{dim} mice were resistant to the inhibitory effects of CORT.

Together our findings support the hypothesis that pituitary ACTH protein concentration is negatively controlled by GR dimerization under conditions of chronic psychosocial stress, while POMC gene transcription is not dependent on intact GR dimerization under both basal and chronic stress conditions. Finally, our data suggest that adrenal adaptations during chronic psychosocial stress (i.e., ACTH desensitization), aiming at the prevention of prolonged hypercorticism, are protective only to a certain threshold of plasma ACTH levels.

1. Introduction

While the underlying aetiology of most stress-associated diseases is

not fully understood, chronic stress-induced dysregulation of almost all psycho-neuro-immunological systems, including the hypothalamuspituitary-adrenal (HPA) axis, are likely to contribute to the complex,

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and multifactorial, aetiology of such disorders. Interestingly, both hyper- and hypoactivity of the HPA axis have been reported for patients diagnosed with stress-associated disorders, suggesting that stressinduced dysregulations in either direction are able to compromise mental and physical health (Langgartner et al., 2015; Raison and Miller, 2003; Holsboer, 2000, 2001; Heim et al., 2000; Fries et al., 2005; Pariante and Miller, 2001; Reber et al., 2016a). For instance, while major depression is most consistently reported to be associated with a hyperactive HPA axis, posttraumatic stress disorder (PTSD) is rather linked to a reduced HPA axis activity. In detail, patients with major depression exhibit increased concentrations of cortisol in plasma, urine, and cerebrospinal fluid (CSF), an exaggerated cortisol response to adrenocorticotropic hormone (ACTH) challenge and an enlargement of both the pituitary and the adrenal glands, all of which are likely mediated by a compromised glucocorticoid (GC) receptor (GR) sensitivity and negative feedback inhibition of the HPA axis (Holsboer, 1983, 2000, 2001; Pariante and Miller, 2001; Gold et al., 1988; Holsboer and Barden, 1996; Nemeroff, 1996: Owens and Nemeroff, 1993: Bardeleben and Holsboer, 1989). In contrast, PTSD patients typically show decreased baseline and 24h average plasma cortisol concentrations, probably mediated by an increased negative feedback sensitivity of the HPA axis (Lehrner et al., 2016; Morris et al., 2012; de Kloet et al., 2006). The latter is indicated by an enhanced cortisol suppression after administration of the synthetic GC dexamethasone (DEX) in PTSD patients versus healthy controls (de Kloet et al., 2006).

To investigate the psycho-neuro-immunological mechanisms underlying stress-related disorders, in particular PTSD, our laboratory employs the chronic subordinate colony housing (CSC) paradigm, which was extensively characterized over the past 20 years as animal model resembling a PTSD-like phenotype (Reber et al., 2016a). The CSC paradigm is based on chronic subordination (19 days) of four male mice towards a dominant male conspecific (Langgartner et al., 2015; Reber et al., 2016a). Compared to respective single-housed control (SHC) mice, CSC mice develop general- and social anxiety (Amoroso et al., 2020; Langgartner et al., 2017; Slattery et al., 2012), increased alcohol consumption/preference (Peters et al., 2013), hyperactivity (Slattery et al., 2012), spontaneous colitis (Langgartner et al., 2017; Reber et al., 2007, 2011, 2016b), and an aggravated dextran sulfate sodium (DSS)-induced colitis (Reber et al., 2008, 2016b; Amoroso et al., 2019). CSC mice further develop functional splenic in vitro GC resistance as well as splenomegaly when receiving a significant amount of bite wounds during CSC exposure or when undergoing abdominal surgery prior to CSC (Foertsch et al., 2017; Foertsch and Reber, 2020; Kempter et al., 2021).

In addition, CSC mice show a higher pituitary weight, accompanied by increased numbers of ACTH immunoreactive cells, which despite an increased DEX suppression of stress-induced ACTH, results in elevated basal and stress-induced plasma ACTH concentrations (Füchsl et al., 2013). Interestingly, as CSC is associated with unaffected basal plasma morning corticosterone (CORT) concentrations and basal evening hypocorticism, despite significant enlargement of the adrenal glands (Langgartner et al., 2017; Reber et al., 2007), increased basal plasma ACTH concentrations do not translate into increased plasma CORT levels. This is in line with the fact that adrenal explants of CSC mice show a decreased in vitro sensitivity to ACTH (Langgartner et al., 2017; Reber et al., 2007; Uschold-Schmidt et al., 2012). However, as CSC mice show an increased CORT secretion towards novel heterotypic stressors (Uschold-Schmidt et al., 2012), which is in line with what has been reported for PTSD patients (de Kloet et al., 2006), unaffected/lower basal plasma CORT levels in CSC mice and PTSD patients might reflect a beneficial adaptation in the face of adrenal enlargement and increased adrenal ACTH input rather than a functional breakdown of general HPA axis functionality. In support of the latter, CSC mice do not show any signs of depressive-like behavior/passive coping in the tail-suspension test (TST) and the forced-swim test (FST) (Slattery et al., 2012), typically seen in rats as a consequence of chronic CORT administrations (Donner et al., 2011). Importantly, it has not been investigated so far whether 1) the adaptive reduction of adrenal ACTH sensitivity in CSC mice showing a PTSD-like phenotype and no signs of depressive-like behavior is limited to a certain extent of ACTH overstimulation, and 2) whether overcoming these adaptational processes at the levels of the adrenal glands not only translates into increased plasma CORT levels but also development of depressive-like behavior/passive coping.

Glucocorticoids act through nuclear hormone receptors including the GR. The GR acts as a homodimer/monomer that either directly binds to specific DNA motives (Lim et al., 2015; Schiller et al., 2014) or interacts with other transcription factors at enhancer sites, resulting in tethering to other transcription factors (Vettorazzi et al., 2022). Based on an increased POMC mRNA expression and ACTH immunoreactivity in the anterior pituitary (Reichardt et al., 1998) in mice with an attenuated, albeit not completely abrogated GR dimerization (GR^{dim/dim} mice; GR^{dim}), it has been proposed that the negative feedback regulation at the level of the pituitary requires an intact GR dimerization interface (Reichardt et al., 1998).

In the current study we combine the GR^{dim} mouse model with the CSC paradigm to assess whether the CSC-induced increased of pituitary and plasma ACTH concentrations can be further exaggerated by a genetically compromised pituitary negative feedback inhibition, finally resulting in plasma ACTH levels exceeding the adaptive capacities at the level of the adrenal glands and, consequently, resulting in prolonged hypercorticism.

2. Material and methods

2.1. Animals

Male GRDim (Sv129Ev/B6 F1 hybrids) (GVO) mice generated by AG Tuckermann (Ulm University, Ulm, Germany) and characterized by a globally and congenitally attenuated, albeit not completely abrogated GR dimerization (GR^{dim/dim} mice; GR^{dim}) were used as experimental mice. Heterozygous GR^{dim/WT} mice were intercrossed to generate GR^{+/+} (WT) and GR^{dim/dim} (GR^{dim}) homozygous mutant mice and the offspring was genotyped by PCR on genomic DNA isolated from tail biopsies (as described in (Reichardt et al., 1998) with minor modifications). For the first five weeks after birth mice were kept under standard laboratory conditions (14h light/10h dark cycle, 22 °C, 60% humidity). Afterwards, experimental mice were transferred to an experimental room with a different type of standard laboratory conditions (12h light/12h dark cycle, 22 °C, 60% humidity) and kept single-housed until the start of the experiment. CD-1 mice (30-35g, Charles River, Sulzfeld, Germany) were used as dominant aggressors. All mice were kept in standard polycarbonate mouse cages (16 \times 22 \times 14 cm) and had free access to tap water and standard mouse diet. The animal study was carried out in accordance with the ARRIVE guidelines 2.0 (Percie du Sert et al., 2020) and was approved by the Federal Animal Care and Use Committee (Regierungspräsidium Tübingen, Germany). All efforts have been made to minimize the number of animals and their suffering.

2.2. Experimental procedures

Four weeks after arrival at the experimental room, WT and GR^{dim} mice were either exposed to the CSC paradigm or kept single-housed (SHC). All animals were tested for anxiety-related behavior in the open-field/novel object (OF/NO) test on day 15 and for depressive-like behavior in the forced swim test (FST) on day 19 of CSC between 0700 and 1000 a.m., respectively, before being sacrificed in the morning of day 20 between 0700 and 1000 a.m. following brief CO₂ anesthesia. Afterwards, the amount of bite wounds was assessed for bite score analysis. Trunk blood was collected for the assessment of plasma corticosterone (CORT) and adrenocorticotropic hormone (ACTH). The adrenal glands, the pituitary as well as the spleen were removed and weight. Adrenal glands were frozen for the assessment of lipid vesicles



Fig. 1. Experimental timeline

After arrival at the experimental room (Day –28), all mice were housed singly for four weeks. At day 1, experimental wild type (WT; GR^{+/+}) and GR^{dim} (GR^{dim/dim}) homozygous mice were weighed and assigned to either the single-housed control (SHC) or the chronic subordinate colony housing (CSC) group. In order to induce chronic psychosocial stress, a group of four CSC mice (two WT and two GR^{dim} mice per cage) was housed together with a dominant male CD-1 mouse for 19 consecutive days. To avoid habituation, CSC mice were exposed to a novel dominant male CD-1 mouse on days 8 and 15. All animals were tested for anxiety-related behavior in the open-field/novel object (OF/NO) test on day 15 and for depressive-like/coping behavior in the forced swim test (FST) on day 19 of CSC, before being euthanized in the morning of day 20. Afterwards, the severity of bite wounds was assessed for bite score analysis. Trunk blood was collected for the assessment of plasma corticosterone (CORT) and adrenocorticotropic hormone (ACTH). The adrenal glands, the pituitary as well as the spleen were removed and weight. Adrenal glands were frozen for the assessment of the adrenal lipid content by Oil Rred-O staining. The pituitary was frozen for quantitative polymerase chain reaction (qPCR) analysis of proopiomelanocortin (POMC). Moreover, splenocytes were isolated for an *in vitro* glucocorticoid (GC) sensitivity assay. Experimental timeline was partly created with BioRender.com.

by Oil Red-O staining. The pituitary was frozen for the analysis of qPCR analysis of proopiomelanocortin (POMC). Moreover, splenocytes were isolated for *in vitro* GC sensitivity assay. In total (n = 48) animals divided in four groups (SHC-WT: n = 13; CSC-WT: n = 11; SHC-GR^{dim}: n = 11; CSC-GR^{dim}: n = 13) were used in the present study. For details about the experimental timeline, see Fig. 1.

2.3. Chronic subordinate colony housing (CSC) procedure

The CSC paradigm was performed as described previously (Langgartner et al., 2015, 2017; Reber et al., 2007, 2008). Briefly, WT and GR^{dim} experimental mice were weighed and assigned to either the SHC or CSC group, matched according to their body weight. During CSC, experimental mice lived in chronic subordination to three different dominant CD-1 aggressor mice between days 1 and 20. In detail, four male mice (2 WT and 2 GR^{dim} mice per cage) were introduced into the home cage of a larger aggressor male CD-1 mouse on day 1 of CSC, resulting in immediate subordination of the four intruder mice. To disturb the established social hierarchy in each CSC cage and to avoid habituation, all four CSC mice of one cage were transferred into the home cage of a novel larger aggressor CD-1 male on days 8 and 15. All resident males were tested before CSC housing for their aggressive behavior and males that were not attacking their opponents or injure their opponents too excessively were not used. SHC mice remained undisturbed for the duration of the experiment, except for changing the bedding once a week and weighing twice a week. Besides being weighed twice a week, SHC and CSC mice were again weighed immediately before being euthanized at day 20.

2.4. Open-field/novel object (OF/NO) test

To assess the effects of CSC and an attenuation of GR dimerization on anxiety-related behavior, all mice were exposed to the OF/NO test on day 15 of CSC exposure. The OF/NO was performed as previously described (Langgartner et al., 2017, 2018). Briefly, the arena (45 cm length x 27 cm width x 27 cm height) was subdivided into an inner (27 cm \times 9 cm) and an outer zone (Fig. 2A). The arena was cleaned thoroughly before each trial. During each trial, the mouse was placed into the middle of the arena and allowed to explore the arena for 5 min. After 5 min of exploration, a plastic round object (diameter: 3.5 cm; height: 1.5 cm) was placed in the middle of the arena. The mouse was now allowed to explore the arena containing the unfamiliar/novel object for another 5 min (Fig. 2E). During OF exploration the overall distance moved, the number of inner zone entries and the time in the corners of the arena was assessed. During NO exploration the overall distance moved, the number of novel object explorations and the time in the corners of the arena was assessed. All parameters were analyzed using EthoVision XT (Version 11.5, Noldus Information Technology, Wageningen, Netherlands). The test was performed between 0700 and 1000 a. m. under white light conditions (350 lux).

2.5. Forced swim test (FST) test

To assess the effects of CSC and an attenuation of GR dimerization on depressive-like/coping behavior, all mice were exposed to the FST test on day 19 of CSC exposure. The FST was performed as previously described (Slattery et al., 2012). Briefly, each mouse was placed into a clear plastic cylinder (diameter: 12 cm; height: 40 cm) filled to 13 cm with 23 \pm 2 $^\circ C$ water for 6 min and mouse behavior was recorded (Fig. 2L). The cylinder was filled with fresh water following each trial. The test was performed between 0700 and 1000 a.m. under white light conditions (200 lux). During the last 4 min of the FST, the time spent mobile was analyzed as velocity >2.5 cm/s using EthoVision XT (Version 11.5, Noldus Information Technology, Wageningen, Netherlands). The time spent floating was calculated based on the following formula: 240s - time spent mobile [s]. Moreover, the time spent climbing was quantified by an experienced experimenter blind to treatment and the time spent swimming was calculated based on the following formula: time spent mobile [s] – time spent climbing [s].

2.6. Trunk blood sampling

Within 3 min after removing the cage from the animal room, mice were rapidly euthanized by decapitation following brief CO₂-exposure. Trunk blood was collected in ethylenediaminetetraacetic acid (EDTA)-coated tubes (Sarstedt, Nuembrecht, Germany) and stored on ice until centrifugation. Tubes were centrifuged at 4 °C (5000 g, 10 min). Plasma samples were stored at -20 °C until further analysis.

2.7. Assessment of adrenal-, pituitary and spleen weight

Adrenal glands were removed, pruned of fat and stored in ice-cold

D. Langgartner et al.

Neurobiology of Stress 24 (2023) 100541





Chronic subordinate colony housing (CSC; dark-grey bars; triangles) increased anxiety-related behavior during open-field (OF; Fig. 2A) and novel-object (NO; Fig. 2E) exposure in both wild type (WT; $GR^{+/+}$; unhatched bars) and GR^{dim} ($GR^{dim/dim}$; hatched bars) mice. In detail, CSC vs. single-housed control (SHC; light-grey bars, circles) mice showed an increased time in corners during NO (Fig. 2G) exposure as well as a decreased distance moved during NO exposure (Fig. 2F), respectively. GR^{dim} but not WT CSC mice showed an increased time in corners during OF exposure (Fig. 2C) and a decreased number of entries into the inner zone of the OF during OF exploration (Fig. 2D) as well as a decreased number of NO explorations (Fig. 2H) when compared to respective GR^{dim} SHC mice. The general locomotion (distance moved during OF exploration) was comparable between all experimental groups (Fig. 2B). Representative track- and time visualizations are shown in (Fig. 2I) and (Fig. 2K), respectively. To assess the effects of CSC and an attenuation of GR dimerization on depressive-like/coping behavior, all mice were exposed to the FST (Fig. 2L) in the morning of day 19 of CSC exposure. In WT and GR^{dim} mice, CSC decreased time of floating (Fig. 2M) and increased the time of swimming (Fig. 2N) during FST exposure, respectively. Moreover, CSC in WT mice decreased the time of climbing during FST exposure (Fig. 2O). Data are presented as bar graphs (mean +SEM) with individual dots. * $P \le 0.05$, ** $P \le 0.01$, *** $P \le 0.001$ vs. respective SHC. Illustrations (Fig. 2A, 2E and 2L) were created with BioRender.com.

Dulbecco's Modified Eagle Medium (DMEM/F-12, Life Technologies, Inc, Grand Island, NY, USA; supplemented with 0.1% bovine serum albumin (BSA; Biomol, Hamburg, Germany)), until being weight. Right adrenal glands were subsequently frozen at -80 °C until being used for Oilred-O staining. The pituitary was removed, immediately weight and frozen at -80 °C until being used for qPCR analysis. Spleens were removed, pruned from fat, weighed, and stored in ice-cold Hanks' Balanced Salt solution (HBSS; Sigma-Aldrich) until being used for the isolation of splenocytes.

2.8. Enzyme-linked immunosorbent assay (ELISA) for plasma CORT and ACTH

Singlets of plasma samples were analyzed using a commercially available ELISA for total CORT (intra-assay coefficient of variation (CV) \leq 7.7%; inter-assay CV \leq 10.8%; IBL International, Hamburg, Germany) and ACTH (intra-assay CV \leq 10.3%; inter-assay CV \leq 7.1%; IBL International, Hamburg, Germany).

2.9. Adrenal Oil Red-O staining

Right adrenal glands were embedded in freezing medium (Tissue-Tek, Sakura Finetek Europe, Zoeterwoude, Netherlands) and frozen at -80 °C. Five µm cryo-sections containing both, adrenal cortex and medulla, were cut using a cryostat (Leica Biosystems, IL, USA) and thawmounted onto precoated slides (SuperFrost ®Plus; VWR, Darmstadt, Germany). Adrenal sections were then fixed in paraformaldehyde (3.5–3.7%; Otto Fischar GmbH & Co. KG, Saarbruecken, Germany) for 72h at room temperature (RT). Subsequently, slides were washed in distilled water, rinsed in 60% isopropyl alcohol for 5 min and then stained in freshly filtered Oil Red-O solution (Certistain Oil red-O, Merck, Darmstadt, Germany) for 15 min. Afterwards, the sections were differentiated in 60% isopropyl alcohol and washed in distilled water before being preserved with Roti® Mount Aqua (ROTH, Karlsruhe, Germany) and a microscope cover glass (ROTH, Karlsruhe, Germany).

In order to quantify the amount of lipid vesicles in the adrenal cortex, four images (one image from each quarter of the section) were collected from each section using bright field microscopy (20x magnification (Leica DMI6000B; Leica Biosystems, Nussloch, Germany), camera (Olympus DP73; Olympus GmbH, Hamburg, Germany)). Image analysis was performed using the software Reimage (© Dr. Michael Noll-Hussong). Specifically, within the adrenal cortical zone of each image, a 300×300 pixel sized region of interest was defined and analyzed for the amount of Oil Red-O stained pixels. The percentage of stained pixels was calculated for each image. Within each section, the mean percentage of stained pixels was subsequently calculated over all four images. Afterwards, the mean percentage over all sections of an animal was calculated and used for statistics. For further statistical analysis, the mean percentage of stained pixels was multiplied with the weight of the right adrenal gland.

2.10. Quantitative reverse transcriptase PCR for POMC in pituitary mRNA

In order to determine effects of CSC and an attenuation of GR dimerization on pituitary POMC mRNA expression, the pituitary was stored at -80 °C until RNA isolation. RNA was isolated using the PAR-ISKit® (Thermo Fisher Scientific, Waltham, USA) according to the manufacturer's instructions. As Nanodrop 2000 technology (Thermo Fischer Scientific, Waltham, MA, USA) revealed a RNA concentration in all samples <100 ng/µl, a total volume of 10 µl and, thus, less than 1 µg RNA, per sample were used to generate cDNA employing random primers (High-capacity cDNA reverse transcription kit, Applied Biosystems, Thermo Fischer Scientific, Waltham, MA, USA) and additional RNase inhibitor "RNase out" to reduce RNase activity (Invitrogen,

Thermo Fischer Scientific, Waltham, MA, USA). Quantitative PCR was performed in triplets using a ViiA7 Real-Time PCR System (Applied Biosystems, Thermo Fischer Scientific, Waltham, MA, USA) and Platinum SYBR Green (Invitrogen, Thermo Fisher Scientific, Waltham, MA, USA). The primers (Sigma Aldrich, St. Louis, MO, USA) used were Ribosomal Proteins (RPL; Housekeeper) (forward: 5 CCTGCTGCTCTCAAGGTT 3'; reverse: 5' TGGCTGTCACTGCCTGG-TACTT 3') and POMC (forward: 5' CGGTGAAGGTGTACCCCAACGT 3'; reverse: 5' GGACCTGCTCCAAGCCTAATGGCC 3'). For the quantification of results, the corresponding Applied Biosystems Software was used and $\Delta\Delta$ Ct analysis was performed.

2.11. Isolation of splenocytes

The isolation of splenocytes was done as previously described (Foertsch et al., 2017). Briefly, spleens were mechanically disrupted using a nylon cell strainer (70 μ m; Corning, USA) and the plunger of a syringe to obtain a single cell suspension. Erythrocytes were lyzed by incubating the cell suspension for 2 min in ASK-lysis buffer (155 mM NH₄Cl, 10 mM KHCO₃, 10 mM EDTA) and stopped by addition of Hanks' Balanced Salt solution (HBSS; Sigma-Aldrich)/10% heat-inactivated fetal calf serum (FCS). Following one washing step with HBSS and filtration of the cell suspension through a 70 μ M nylon cell strainer, cells were resuspended in ice-cold Roswell Park Memorial Institute Medium (RPMI-1640, Sigma-Aldrich, St. Louis, MO, USA) containing 10% FCS, 50 U/ml penicillin and 50 μ g/ml streptomycin (RPMI+) and counted using a cell counter (TC-20, Bio-Rad Laboratories, Munich, Germany).

2.12. Assessment of functional in vitro GC sensitivity of isolated splenocytes

Splenocytes were adjusted to a concentration of 5×10^6 cells/ml in RPMI+. Afterwards all isolated cells were plated in singlets in flatbottom, 96-well plates in a final total volume of 100 µl/well (splenocytes: 2.5×10^5 cells/well). Cells either remained untreated or were stimulated with LPS (Escherichia coli O111:B4, final concentration 1μ g/ ml; Sigma-Aldrich, Deisenhofen, Germany). Moreover, corticosterone (CORT; Sigma-Aldrich, Deisenhofen, Germany; final concentration: 0 and 0.1 µM respectively), diluted in 95% ethanol, was added to the wells. Cells were incubated (37 °C, 5% CO₂) for 48 h and cell viability was determined using a commercially available colorimetric assay (CellTiter 96 Aqueous One Solution Assay, Promega, Madison, WI). Absorbance at 450 nm was measured using an enzyme-linked immunosorbent assay (ELISA) plate reader (Fluostar Optima, BMG Labtech, Offenburg, Germany). All values were blank-corrected. The absorbance (optical density (OD)) of unstimulated cells was subtracted from the corresponding LPS-stimulated cells and the resulting delta OD was set to 100% for the 0 µM CORT condition. A low cell viability at increasing CORT levels indicates high GC sensitivity and low GC resistance.

2.13. Assessment of bite wound severity

The severity of bite wounds in CSC mice was assessed using a bite score previously established by our group (Foertsch et al., 2017). Briefly, following decapitation, the skin (with fur attached) was removed so that skin and body were still connected at the tail and hind limbs. Afterwards, pictures were taken from each mouse and digitally overlaid with a standardized grid (GIMP 2.6.12), which contained 20 squares covering the skin (dermal wounds) and 20 squares covering the body (subdermal wounds). Each square on the skin was scored according to the affected area (score 0–4), intensity (score 0–4) and the degree of purulence (0–2), whereas each square on the body was scored for the affected area (score 0–3) and the intensity of injury (0–2). The bite score is then represented by the sum of all skin and body injuries and ranges from 0 to 300.



Fig. 3. Effects on HPA axis parameters

When compared to respective single-housed control (SHC; light-grey bars; circles) mice, chronic subordinate colony housing (CSC; dark-grey bars; triangles) resulted in an increased relative adrenal weight (Fig. 3A) in both wild type (WT; $GR^{+/+}$; unhatched bars) and GR^{\dim} ($GR^{\dim/\dim}$; hatched bars) mice. Relative adrenal weight of both GR^{\dim} SHC and CSC mice was decreased when compared to respective WT mice. Relative pituitary weight (Fig. 3B) was increased in WT but not GR^{\dim} CSC vs. SHC mice and decreased in GR^{\dim} vs. WT CSC mice. Plasma morning corticosterone (CORT; Fig. 3C) and adrenocorticotropic hormone (ACTH; Fig. 3D) concentrations were increased in GR^{\dim} CSC compared to GR^{\dim} SHC as well as WT CSC mice, respectively. Adrenal lipid content represented by the ratio of Oil Red-Ostained adrenal area to total adrenal area in sections of the right adrenal gland (Fig. 3E) as well as adrenal lipid content per adrenal weight (Fig. 3F) was increased in WT but not GR^{\dim} CSC vs. SHC mice and decreased in both GR^{\dim} vs. WT SHC and CSC mice, respectively. Representative Oil Red-O staining images of each experimental group are shown in (Fig. 3G). Data are presented as bar graphs (mean +SEM) with individual dots. *P# * $P \le 0.05$, ** $P \le 0.01$, *** $P \le 0.001$ vs. respective SHC. # $P \le 0.05$, ## $P \le 0.01$, ### $P \le 0.001$ vs. respective WT.

2.14. Statistics

For statistical analysis and graphical illustrations GraphPad Prism (version 9.3.1, GraphPad Software, LCC) was used. Kolmogorov-Smirnov test with Lilliefors' correction was employed to test for normal distribution. Outliers in normally distributed data sets were identified by Grubbs test and excluded from further analysis. Normally distributed data sets were analyzed by parametric statistics, i.e. two-way ANOVA (two factors, two or more independent samples) followed by post-hoc analysis using Bonferroni pairwise comparison, when a significant main effect was found. Not normally distributed data sets were analyzed by non-parametric statistics, i.e. Mann-Whitney *U* test (MWU; one factor, two independent samples) and Wilcoxon matched-pairs

signed rank test (one factor, two dependent samples). Data are presented as mean +SEM including individual values. The level of significance was set at $p\leq<0.05.$

3. Results

3.1. Effects on anxiety-related and depressive-like/coping behavior

To assess the effects of CSC and an attenuated GR dimerization on anxiety-related behavior, all mice were exposed to the OF/NO test (Fig. 2A–K) in the morning of day 15 of CSC exposure (Fig. 1). Both WT and $\mathrm{GR}^{\mathrm{dim}}$ mice displayed increased anxiety-related behavior in the CSC compared to SHC group. This was indicated by a decreased distance



Fig. 4. Effects on relative pituitary POMC mRNA expression The relative pituitary proopiomelanocortin (POMC) mRNA expression levels (Fig. 4A) did not differ between wild type (WT; GR^{+/+}; unhatched bars) and GR^{dim} (GR^{dim/dim}; hatched bars) mice exposed to chronic subordinate colony housing (CSC; dark-grey bars; triangles) or single-housing (SHC; light-grey bars; circles). As housekeeping gene, ribosomal proteins (RPL) was used. Data are presented as bar graphs (mean +SEM) with individual dots.

moved during NO exposure (Fig. 2F; WT: P = 0.016 (MWU); GR^{dim}: P = 0.006 (MWU)) and an increased time spent in the corners during NO (Fig. 2G; WT: P < 0.001 (MWU); GR^{dim}: P = 0.021 (MWU)) exposure in CSC vs. SHC mice of both the WT and GR^{dim} group, respectively. GR^{dim} but not WT CSC mice further showed an increased time spent in corners during OF exploration (Fig. 2C; GR^{dim}: P = 0.002 (MWU)), a decreased number of entries into the inner zone during OF exploration (Fig. 2D; P = 0.001 (MWU)), as well as a decreased number of NO exploration (Fig. 2H; P = 0.005 (MWU)) when compared to respective SHC mice. The general locomotion, represented by the distance moved during OF exploration, was comparable between all experimental groups (Fig. 2B). Representative track- and time visualizations are shown in (Fig. 2I) and (Fig. 2K), respectively.

To assess the effects of CSC and an attenuation of GR dimerization on depressive-like/coping behavior, all mice were exposed to the FST (Fig. 2L - O) in the morning of day 19 of CSC exposure. Compared with respective SHC mice, CSC significantly decreased the time floating (Fig. 2M, two-way ANOVA, Main effect for stress: $F_{1,40} = 28.37$; P < 0.001) in both, the WT (Bonferroni: P < 0.001) and GR^{dim} (Bonferroni: P = 0.02) group. Furthermore, CSC increased the time spent swimming (Fig. 2N, two-way ANOVA, Main effect for stress: $F_{1,40} = 36.23$; P < 0.001) in both the WT (Bonferroni: P < 0.001) and GR^{dim} (Bonferroni: P = 0.004) group compared with respective SHC mice. Finally, CSC significantly decreased the time spent climbing (Fig. 2O) during FST exposure compared with respective SHC mice in the WT (P = 0.014 (MWU)) group.

3.2. Effects on HPA axis parameters and body weight

Chronic subordinate colony housing resulted in an increased relative adrenal weight (Fig. 3A) in both the WT (P < 0.001 (MWU)) and GR^{dim} (P = 0.002 (MWU)) group when compared to respective SHC mice. Furthermore, relative adrenal weight of both GR^{dim} SHC (P < 0.001 (MWU)) and CSC (P < 0.001 (MWU)) mice was decreased when compared to respective WT mice. In contrast, relative pituitary weight (Fig. 3B) was only increased in CSC vs. SHC mice of the WT (P = 0.016 (MWU)) but not the GR^{dim} group. Moreover, relative pituitary weight was found to be significantly decreased in GR^{dim} vs. WT CSC mice (P = 0.005 (MWU)). Plasma morning CORT concentrations (Fig. 3C) were significantly increased in GR^{dim} SHC (P = 0.013 (MWU)) as well as WT CSC (P = 0.003 (MWU)) mice. Similarly, plasma morning ACTH concentrations (Fig. 3D) were significantly

increased in GR^{dim} CSC compared with GR^{dim} SHC (P = 0.002 (MWU)) as well as WT CSC (P = 0.015 (MWU)) mice. The availability of adrenal lipids represented by the ratio of Oil Red-O-stained adrenal area to total adrenal area in sections of the right adrenal gland (Fig. 3E, G) was increased in WT (two-way ANOVA: Interaction effect: $F_{1,43} = 6.981$; P =0.011; Main effect for genotype: $F_{1,43} = 53.62$; P < 0.001; Bonferroni: P= 0.004) but not GR^{dim} CSC vs. SHC mice. Moreover, adrenal lipid content was significantly lower in both GR^{dim} SHC (Bonferroni: P =0.003) and GR^{dim} CSC (Bonferroni: P < 0.001) mice compared with the respective WT group. When corrected for differences in adrenal weight (Fig. 3F), adrenal lipids were still increased in CSC vs. SHC mice in the WT (two-way ANOVA: Main effect for stress: $F_{1,42} = 34.03$; P < 0.001; Bonferroni: P < 0.001) but not GR^{dim} group, as well as significantly lower in both SHC (two-way ANOVA: Interaction effect: $F_{1.42} = 24.19$; P < 0.001; Main effect for genotype: F_{1,42} = 143.6; *P* < 0.001; Bonferroni: P < 0.001) and CSC (Bonferroni: P < 0.001) GR^{dim} mice compared with respective WT mice. Fig. 3G shows representative Oil Red staining images of each experimental group. Body weight was comparable between all experimental groups on day 20 of CSC (data not shown).

3.3. Effects on relative pituitary POMC mRNA expression

Neither CSC nor the attenuated GR dimerization had a significant effect on the relative POMC mRNA expression in the pituitary (Fig. 4A).

3.4. Effects on spleen weight and functional splenic in vitro GC sensitivity

In line with what has been reported by our group for a bite score above 20 (WT: 27.5; GR^{dim}: 44.2; Fig. 5B) (Foertsch et al., 2017), CSC increased relative spleen weight (Fig. 5A) in both WT (P = 0.009 (MWU)) and GR^{dim} (P < 0.001 (MWU)) mice when compared to respective SHC mice. Furthermore, relative spleen weight of both GR^{dim} SHC (P < 0.001 (MWU)) and GR^{dim} CSC (P < 0.001 (MWU)) mice was significantly increased compared to respective WT mice.

Compared to basal values, splenic cell viability in response to LPS (Fig. 5C) was increased in all groups (WT SHC: P < 0.001; WT CSC: P0.001; GR^{dim} SHC: P < 0.001; GR^{dim} CSC: P < 0.001 (all Wilcoxon)). Moreover, splenocytes from WT and GR^{dim} CSC mice showed a higher basal (WT: P < 0.001 (MWU); GR^{dim}: P < 0.001 (MWU)) and LPSinduced (WT: P = 0.002 (MWU); GR^{dim}: P = 0.007 (MWU)) cell viability compared to respective SHC mice. Basal cell viability was further increased in splenocytes from GR^{dim} vs. WT CSC mice (P = 0.047(MWU)). Delta (LPS - basal) cell viability (0 µM CORT set to 100%; Fig. 5D) in response to 0.1 μ M CORT was reduced in the SHC (P < 0.001(Wilcoxon)) and CSC (P < 0.001 (Wilcoxon)) mice of the WT group when compared to respective groups in the 0 µM condition with the effect being significantly less pronounced in CSC WT mice (CSC vs. SHC WT mice at 0.1 μ M CORT: *P* = 0.007 (MWU)). Of note, delta cell viability in response to 0.1 μ M CORT was significantly higher in GR^{dim} SHC (P < 0.001 (MWU)) and CSC (P < 0.001 (MWU)) vs. respective WT mice. Thus, GRdim splenocytes are less sensitive towards the immunosuppressive effects of GCs.

4. Discussion

In the present study we confirm own earlier findings in male C57Bl/ 6N mice, showing that Sv129Ev/C57Bl/6 (B6) F1 hybrid WT mice exposed to CSC are characterized by increased anxiety-related behavior and no signs of depressive-like/passive coping behavior. Moreover, CSC vs. SHC Sv129Ev/B6 WT mice show unaffected morning plasma CORT levels, despite an increased pituitary and adrenal mass as well as an increased Oil Red-O positive adrenal lipid content in the cortex. In extension, we show that male GR^{dim} mice expressing a GR with a point mutation (A465T) disrupting one of the GR dimerization interfaces and bred as F1 hybrids on a Sv129Ev/B6 background (Sv129Ev/B6 GR^{dim}) have unaffected basal morning plasma ACTH and CORT concentrations



Fig. 5. Effects on spleen weight and functional splenic in vitro GC sensitivity

When compared to respective single-housed control (SHC; light-grey bars; circles) mice, chronic subordinate colony housing (CSC; dark-grey bars; triangles) increased relative spleen weight (Fig. 5A) in both the wild type (WT; $GR^{+/+}$; unhatched bars) and GR^{\dim} ($GR^{\dim/\dim}$; hatched bars) group. Furthermore, relative spleen weight of GR^{\dim} SHC and CSC mice was increased compared to respective WT mice. Both, WT and GR^{\dim} CSC mice had a bite score above 20 (WT: 27.5; GR^{\dim} : 44.2; Fig. 5B). Splenic cell *in vitro* viability in response to lipopolysaccharide (LPS; Fig. 5C) was increased in all groups. Splenocytes from WT and GR^{\dim} CSC mice showed a higher basal and LPS-induced cell viability compared to respective SHC mice. Basal cell viability was increased in splenocytes from GR^{\dim} vs. WT CSC mice. Delta (LPS - basal) cell viability (0 μ M corticosterone (CORT) set to 100%; Fig. 5D) in response to 0.1 μ M CORT was reduced in WT SHC and CSC mice when compared to the 0 μ M condition. Delta cell viability in response to 0.1 μ M CORT was significantly increased in GR^{\dim} SHC and CSC vs. respective WT mice, as well as WT CSC vs. SHC mice. Data are presented as bar graphs (mean +SEM) with individual dots. ** $P \le 0.01$, *** $P \le 0.001$ vs. respective SHC. # $P \le 0.05$, ### $P \le 0.001$ vs. respective WT. ^{\$\$\$\$} P < 0.001 vs. respective basal or 0 μ M condition.

compared with respective WT mice. Together with earlier studies (Reichardt et al., 1998) suggesting a compromised negative feedback inhibition at the level of the pituitary in Sv129Ev GR^{dim} mice, these data support the hypothesis that the significant reduction in pituitary and adrenal weight in unstressed Sv129Ev/B6 GR^{dim} in the present study represent a beneficial adaptation to avoid basal hypercorticism and its well-known negative consequences. However, although Sv129Ev/B6 GR^{dim} mice do not develop pituitary enlargement when exposed to CSC, chronic psychosocial stress in Sv129Ev/B6 GR^{dim} mice seems to exceed their capacity of HPA axis adaptation, indexed by increased plasma morning ACTH and CORT levels and adrenal enlargement compared with respective SHC mice. As we neither found a genotype nor CSC effect on pituitary POMC mRNA concentrations, our data support the hypothesis that an affected GR signaling at least indirectly affects plasma ACTH concentrations during chronic psychosocial stress, most likely via regulating the expression of certain factors involved in POMC mRNA translation/post-translational modification.

In contrast to own previous studies in C57Bl/6N mice (Füchsl et al., 2013), CSC in Sv129Ev/B6 WT mice did not result in increased basal morning plasma ACTH concentrations, despite pituitary weight was significantly enhanced. However, given that only a view plasma ACTH

values were above the detection limit in both SHC and CSC mice of the Sv129Ev/B6 WT group, this missing CSC effect might be due to overall very low and, thus, undetectable ACTH levels in Sv129Ev/B6 compared with C57Bl/6N mice. Noteworthy, although in Sv129Ev/B6 GR^{dim} SHC mice only a view plasma ACTH concentrations were above the detection limit of the ELISA, comparable plasma ACTH concentrations between Sv129Ev/B6 WT SHC and Sv129Ev/B6 GR^{dim} SHC mice in the present study are in line with an earlier study finding no differences in plasma ACTH between Sv129Ev WT and Sv129Ev GR^{dim} mice (Reichardt et al., 1998). Further different between an earlier study done in mice from a Sv129Ev background (Reichardt et al., 1998) and the current study done in mice from a Sv129Ev/B6 background is the effect of the GR dimerization deficits on pituitary POMC mRNA levels. While non-quantitative in situ hybridization in the earlier study suggested an increased POMC mRNA expression in GR^{dim} vs. WT mice (Reichardt et al., 1998), which was further accompanied by an increased pituitary ACTH immunoreactivity (Reichardt et al., 1998), qPCR done in the current study failed to find a genotype effect. As qPCR analysis of POMC mRNA expression in the present study further failed to detect a CSC effect, we hypothesize that increased plasma ACTH levels in CSC vs. SHC Sv129Ev/B6 $\mbox{GR}^{\mbox{dim}}$ mice to be the consequence of an enhanced POMC mRNA



Fig. 6. Summary Figure

Upon binding of corticotrophin releasing hormone (CRH) to the CRH-receptor (R) 1 on corticotroph cells of the anterior pituitary, proopiomelanocortin (POMC) gene transcription is initiated. POMC mRNA is subsequently translated into POMC and post-transcriptionally cleaved into *inter alia* adrenocorticotropic hormone (ACTH). During chronic psychosocial stress exposure, glucocorticoid receptor (GR) dimerization is proposed to inhibit the translation of different enzymes/convertases involved in POMC mRNA translation or post-transcriptional processing into ACTH protein (Harno et al., 2018). Figure was created with BioRender.com.

translation/post-translational modification. At the level of the adrenal glands, CSC exposure in both WT and GR^{dim} mice of the Sv129Ev/B6 background caused adrenal enlargement, confirming that the stress procedure worked reliably in both genotypes. The latter is also supported by the CSC-induced increase in anxiety-related behavior in both WT and Sv129Ev/B6 GR^{dim} mice, an effect repeatedly reported for C57Bl/6N mice earlier (Reber et al., 2007, 2016a; Langgartner et al., 2017; Peters et al., 2013). However, while the CSC-induced increase in adrenal mass did not translate into increased plasma CORT concentrations in Sv129Ev/B6 WT mice, it did in Sv129Ev/B6 GR^{dim} mice. Interestingly, unaffected plasma morning CORT levels despite an increased adrenal weight and increased plasma ACTH were previously also reported for CSC-exposed C57Bl/6N mice (Langgartner et al., 2017; Reber et al., 2007; Uschold-Schmidt et al., 2012). Moreover, plasma ACTH levels in the current study were significantly higher in Sv129Ev/B6 GR^{dim} CSC vs. respective WT CSC mice. Therefore, our data supports the hypothesis that a reduction in adrenal ACTH sensitivity during times of chronic stress, adrenal enlargement and increased ACTH levels (Langgartner et al., 2017; Reber et al., 2007; Uschold-Schmidt et al., 2012) represents a beneficial adaptation, protecting the individual at least to a certain threshold of plasma ACTH levels from the negative consequences of stress-induced hypercorticism.

Interestingly, increased basal morning plasma ACTH concentrations in Sv129Ev/B6 GR^{dim} CSC mice resulted in increased plasma CORT concentrations, although the CSC-induced increase in adrenal lipids was only detectable in Sv129Ev/B6 WT but not GR^{dim} mice. The latter supports the hypothesis that while Sv129Ev/B6 GR^{dim} CSC mice are able to translate increased basal morning plasma ACTH concentrations adequately into increased plasma CORT concentrations, this might fail in response to a severe acute heterotypic stressor.

In line with previous studies done in C57Bl/6N mice (Slattery et al.,

2012), CSC in the present study did neither increase depressive-like behavior/passive coping in WT nor GR^{dim} mice of the Sv129Ev/B6 background. In contrast, while CSC in C57Bl/6N mice did not affect the time spent immobile during FST (Slattery et al., 2012), CSC in the present study reduced the time floating during FST exposure in both genotypes, suggesting a genotype-independent shift towards active stress coping or development of a hyperactive phenotype. The latter is rather unlikely, as earlier studies by our group revealed that home cage locomotor activity in CSC vs. SHC C57Bl/6N mice is not affected on day 20 of CSC (Slattery et al., 2012). Analysis of the time spent swimming and climbing during FST exposure indicated that the above described effect is predominantly mediated by an increase in swimming but not climbing behavior. One possible explanation for this discrepancy in FST immobility might be that C57Bl/6N mice in the Slattery et al., 2012 study were exposed to the FST 8 days following termination of CSC (Slattery et al., 2012), whereas in the current study the FST was performed on day 19 of CSC. Another possible explanation would be that in the Slattery et al., 2012 study mice exposed to the FST were not exposed to any other behavioral test before, whereas all mice in the present study were tested for their anxiety-related behavior in the OF/NO test on day 15 of CSC.

Strikingly, Sv129Ev/B6 GR^{dim} SHC and CSC mice were totally resistance to the anti-inflammatory effects of CORT on the LPS-induced increase in *in vitro* splenocyte viability. This is in line with findings showing that anti-inflammatory activity of GCs and the reduction of leukocyte abundance by GCs require intact GR activity, which is strongly hampered in GR^{dim} mice, as demonstrated during LPS-induced acute lung injury (Ichinose et al., 2001), sepsis (Barbee et al., 2010), contact hypersensitivity (Tuckermann et al., 2007), rheumatoid arthritis (Baschant et al., 2011) and TNF-induced inflammation (Vandevyver et al., 2012). Of note, in agreement with own previous findings in C57Bl/6N mice (Foertsch et al., 2017, 2020; Foertsch and Reber, 2020;

Kempter et al., 2021), 19 days of CSC in the present study caused an increased spleen weight in both WT and GR^{dim} mice of the Sv129Ev/B6 background, suggesting significant wounding during social stressor exposure in both genotypes (Foertsch et al., 2017, 2020; Foertsch and Reber, 2020). The latter was confirmed by a mean bite score of 27.5 in Sv129Ev/B6 WT and 44.2 in Sv129Ev/B6 GR^{dim} CSC mice. For comparison, C57Bl/6N CSC mice with a bite score above 20 reliably develop splenomegaly (Foertsch et al., 2017). Further in line with own previous data (Foertsch et al., 2017), isolated splenocytes of CSC WT and CSC GR^{dim} mice of the Sv129Ev/B6 background in the absence of CORT showed an increased cell viability when in vitro cultured both in the absence or presence of LPS. Of note, in contrast to $Sv129Ev/B6\ GR^{dim}$ mice, respective WT mice in general were sensitive to the anti-inflammatory effects of CORT, with CSC in combination with bite wounds facilitating development of functional splenic in vitro GC resistance, as shown in C57Bl/6N mice (Reber et al., 2007; Foertsch et al., 2017, 2020; Foertsch and Reber, 2020; Kempter et al., 2021).

Together our findings support the hypothesis, that POMC mRNA translation or posttranslational modification into ACTH protein is negatively controlled by intact GR dimerization under conditions of chronic psychosocial stress (summarized in Fig. 6). Moreover, our data suggest that adrenal adaptations during chronic psychosocial stress (i.e, ACTH desensitization), aiming at the prevention of prolonged hyper-corticism, are protective only to a certain threshold of plasma ACTH levels.

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CRediT authorship contribution statement

DL, MK, SK, MT, SV, JT and SOR, planned the study; DL, MK, SK, LG and LK and LGPR performed the experiments; MNH provided the Oil Red-O analysis software; DL and MK did the statistical analysis and created the figures; DL, MK, JT and SOR wrote the manuscript.

Declaration of competing interest

DL, MK, SK, LG, LK, MT, MNH, SV, JT and SOR have nothing to declare and do not have any conflict of interest.

Data availability

Data will be made available on request.

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D. Langgartner et al.

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