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## Impact of adolescent stress on the expression of stress-related receptors in the hippocampus of animals exposed to alcohol prenatally

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### Abstract

Many functions of the hippocampus are affected by prenatal alcohol exposure (PAE). In particular, dysregulation of the stress response is especially important because individuals with PAE carry increased risks for exposure to stressful environments throughout life. Little is known, though, about how adolescent stress in the context of PAE-related stress system dysregulation may further alter hippocampal development. Here we investigate the short- and long-term effects of adolescent chronic mild stress (CMS) on mRNA expression of stress-related mineralocorticoid (MR), glucocorticoid (GR), and type 1 CRH (CRHR1) receptors in the dorsal and ventral hippocampal formation of PAE and control rats. Our results indicate that PAE affects the expression of stress-related receptors in the hippocampus; however, PAE effects were more prominent during adolescence, as MR and CRHR1 mRNA expression were altered in both male and female PAE animals, with GR mRNA expression alterations observed only in PAE female. In adulthood, the effects of PAE were restricted to alterations in CRHR1 mRNA expression in females, while there were no effects in males. In contrast, the effects of adolescent CMS were more pronounced in adulthood, long after stress exposure termination. Importantly, PAE animals were less responsive to adolescent CMS, with effects only on CRHR1 in PAE animals compared to the altered MR, GR and CRHR1 mRNA expression observed in controls. Together, our results show that PAE and adolescent CMS induce dynamic alterations in the expression of stress-related receptors in the hippocampal formation that manifest differently depending on the age and sex of the animal.

### Keywords

prenatal alcohol exposure; chronic mild stress; mineralocorticoid receptor; glucocorticoid receptor; type 1 CRH receptor

### Introduction

Clinical and experimental studies have clearly demonstrated that alcohol consumption during pregnancy alters fetal brain development and can induce a wide range of cognitive,

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neurobehavioral, and physiological deficits (Drew and Kane, 2014; Hellemans et al., 2010; Riley et al., 2011; Schneider et al., 2011; Valenzuela et al., 2012; Weinberg et al., 2008). Although the structure and function of many brain regions are negatively affected by prenatal alcohol exposure (PAE), the hippocampus is particularly sensitive to the effects of PAE (Fontaine et al., 2016; Gil-Mohapel et al., 2010; Valenzuela et al., 2012). For example, several clinical studies indicate that PAE is associated with reduced hippocampal volume (Coles, et al., 2011; Donald et al., 2016; Gross et al., *in press*; Riikonen et al., 1999, Willoughby et al., 2008). Preclinical studies have considerably extended the clinical findings demonstrating that PAE results in hypertrophy of mossy fibers (West et al., 1981), reduced numbers of neurons (Burke et al., 2015) and dendritic spines (Abel et al., 1983), altered neural activity (Rainekei et al., 2014), impaired long-term potentiation (Christie et al., 2005; Pattern et al., 2013; Sutherland et al., 1997) and reduced adult neurogenesis (Gil-Mohapel et al., 2010; Redila et al., 2006; Sliwowska et al., 2010; Uban et al., 2010) in the hippocampus.

The hippocampus is a unique brain area as its structure is constantly adapting in response to environmental stimuli (McEwen, 1999). This plasticity is critical for the proper function of the hippocampus, playing a key role in a wide range of functions including learning and memory as well as emotional and stress regulation (Fanselow and Dong, 2010). This hippocampal plasticity is mediated, at least in part, by stress-related receptors (Maras and Baram, 2012; McEwen et al., 2015). Hippocampal neurons express high levels of the type 1 CRH receptor (CRHR1), glucocorticoid receptor (GR), and mineralocorticoid receptor (MR), which are major regulators of hippocampal development, structure, and function (Maras and Baram, 2012, McEwen, 2012; McEwen et al., 2015). PAE has been shown to reduce CRHR1 mRNA expression in the amygdala, medial prefrontal cortex (mPFC), and pituitary in adulthood (Glavas et al., 2007; Rainekei et al., 2016). Hippocampal CRHR1 expression following PAE has only been assessed in adolescent mice, and the data indicate that PAE reduces CRHR1 protein levels (Caldwell et al., 2015). Conversely, several studies have demonstrated that adult PAE male rats do not show changes in hippocampal MR and GR expression (Glavas et al., 2007; Kim et al., 1999; Uban et al., 2013). However, PAE adult females showed reduced hippocampal MR mRNA expression (Sliwowska et al., 2008; Uban et al., 2013). Moreover, PAE has been shown to reduce MR and to elevate GR protein levels in the nuclear fraction of hippocampal neurons in adolescent male mice, whereas levels of both receptors were not altered in the cytosolic fraction (Caldwell et al., 2014). Studies investigating PAE effects on MR, GR, and CRHR1 expression in the hippocampal formation have focused exclusively in the dorsal hippocampus (Glavas et al., 2007; Kim et al., 1999; Uban et al., 2013) or used the entire hippocampus (Caldwell et al., 2014, 2015). Importantly, the dorsal and ventral components of the hippocampal formation have different functions: the dorsal hippocampus is primarily involved in cognitive functions while the ventral hippocampus is more essential for stress and emotional regulation (Fanselow and Dong, 2010). Here, we expand this literature by evaluating the effects of PAE on the expression of MR, GR and CRHR1 in different subfields of the dorsal and ventral hippocampus.

Cortical and limbic areas, including the hippocampus, undergo extensive functional and structural reorganization during adolescence (Crews et al., 2007; Eiland and Romeo, 2013; Hueston et al., 2017; McCormick and Mathews, 2010; Spear, 2000). These maturation

processes occur in parallel with many changes in neuroendocrine function (Crews et al., 2007; Ojeda et al., 2006), including the unique hyperresponsivity of the hypothalamic-pituitary-adrenal (HPA) axis to both acute and chronic stressors (Doremus-Fitzwater et al., 2009; Eiland and Romeo, 2013; Green et al., 2016; Hollis et al., 2013; McCormick and Mathews, 2010; Romeo et al., 2004, 2006). The combination of extensive brain maturation and hyperresponsivity to stressors makes adolescence a period of increased vulnerability to adverse environmental stimuli. This is particularly relevant for individuals exposed to alcohol during gestation, as both human and rodent studies have demonstrated that PAE is associated with increased HPA activation and/or a delayed return to basal levels when faced with a wide range of stressful stimuli (Jacobson et al., 1999; Lee and Rivier, 1996; Nelson et al., 1986; Taylor et al., 1982; Weinberg et al., 2008). Notably, clinical data indicate that individuals prenatally exposed to alcohol are, in general, at a higher risk of being exposed to a more stressful environment throughout the lifespan (O'Connor and Kasari, 2000; O'Connor and Paley, 2006; Streissguth et al., 2004). Moreover, similar to PAE, exposure to chronic stress during adolescence can also lead to short- and long-term alterations in the expression of stress-related receptors in the hippocampus (Iredale et al., 1996; Isgor et al., 2004; Li et al., 2015; Sterlemann et al., 2008; Veenit et al., 2014). Nevertheless, little is known about how this higher risk for stress exposure affects ongoing hippocampal development during adolescence in the context of higher PAE-induced hyperresponsivity. To fill this gap in the literature, here we evaluate the short- and long-term effects of adolescent stress on stress-related receptor expression in PAE animals by performing a comprehensive assessment of CRHR1, MR and GR mRNA expression in specific subfields of the dorsal and ventral hippocampal formation.

## Methods

### Animals and Breeding

Male and female Sprague-Dawley rats were obtained from Charles River Laboratories (St. Constant, Canada). Rats were pair-housed by sex and maintained at a constant temperature ( $21 \pm 1^\circ\text{C}$ ) and on a 12 h light-dark cycle (lights on at 0700 h) with *ad libitum* access to water and standard lab chow (Harlan, Canada). After a 10-day acclimation period, male and female pairs were placed together for breeding. Vaginal smears were taken each morning, and the presence of sperm was used as an indicator of pregnancy (gestation day 1; G1). All experiments were performed in accordance with National Institutes of Health (NIH) guidelines for the care and use of laboratory animals, Canadian Council on Animal Care guidelines, and were approved by the University of British Columbia Animal Care Committee.

### Prenatal Alcohol Exposure

On G1, females were single-housed and randomly assigned to one of three treatment groups: Prenatal Alcohol Exposure (PAE), pair-feeding, or control. Dams in the PAE group were offered *ad libitum* liquid ethanol diet with 36% ethanol-derived calories (Weinberg-Keiver High Protein Control Diet #710109, Experimental Diet # 710324, Dyets Inc., Bethlehem, PA). The liquid ethanol diet was introduced gradually over the first 3 days with bottles containing: G1 - 66% control diet, 34% ethanol diet; G2 - 34% control diet, 66% ethanol

diet; G3-21 - 100% ethanol diet. This diet is formulated to provide adequate nutrition to pregnant rats regardless of ethanol intake (Lan et al., 2006). Blood alcohol levels were measured from tail blood samples collected on G15 approximately 4-6 h after lights off using Pointe Scientific Inc. Alcohol Reagent Set (Lincoln Park, USA; Workman et al., 2015). Alcohol-consuming dams showed an average of  $118.20 \pm 8.11$  mg/dl. Pair-fed dams were offered a liquid control diet with maltose-dextrin isocalorically substituted for ethanol, in an amount matched to the consumption of an alcohol-fed partner (g/Kg body weight/day of gestation). The control dams were offered *ad libitum* access to a pelleted form of the liquid control diet. All animals had *ad libitum* access to water, and those in the PAE and pair-fed groups were provided with fresh diet daily within 1 h prior to lights off to maintain the normal HPA circadian rhythm (Gallo and Weinberg, 1981; Krieger, 1974). Experimental diets were continued through G21. Beginning on G22, all animals were offered *ad libitum* access to standard laboratory chow and water, which they received throughout lactation. Pregnant dams were left undisturbed except for cage changing (G1, G7, and G14) and weighing (G1, G7, G14, and G21). On the day of birth (postnatal day 1, PN1) the litters were culled to 12 pups with an attempt to preserve an equal number of males and females per litter. Dams and pups were cage changed and weighed on PN1, PN8, PN15, and PN22. Dam and pup body weight data were published in Rainekei et al., 2016. On PN22 pups were weaned and group-housed by litter and sex.

Effects of adolescent CMS on stress-related receptors in PAE and *ad libitum*-fed control offspring is presented in the Results section below, whereas similar to Rainekei et al., 2017, specific pair-feeding effects are analyzed and presented separately in the Supplementary Materials. The adverse effects of alcohol *per se* on a vast number of outcomes, including hippocampal structure and function, are well established. Moreover, while the pair-fed group is used to account for the decreased food intake associated with chronic alcohol consumption, it can not separate alcohol effects from those of undernutrition, as it can never account for the nutritional effects associated with alcohol consumption, including changes in absorption and utilization of nutrients (Weinberg, 1984) and satiety (Lin et al., 1998). Moreover, pair-feeding is in many aspects a confounded “control” condition and in many ways is a treatment group in itself, as it leads to an abnormal feeding pattern: because pair-fed animals receive less food than they would consume if allowed to eat *ad libitum*, they consume their entire day’s ration within a few hours, and are thus essentially food deprived until the next feeding (Gallo and Weinberg, 1981; Weinberg 1984). This abnormal feeding pattern introduces a prenatal stress component, which in itself can have long-term impacts on the development of offspring neurobiological systems, including the neuroendocrine axis (Vieau et al., 2007).

### Adolescent Chronic Mild Stress (CMS)

One male and one female from each litter were randomly assigned to either the CMS or the non-CMS condition and were pair-housed with another animal of the same sex and prenatal group. Animals in the CMS condition were subjected to 10 consecutive days of chronic, unpredictable, mild stressors. To account for the sexually dimorphic time of pubertal onset (McCormick and Mathews, 2010; Vetter-O’Hagen and Spear, 2012), males and females were exposed to CMS at ages consistent with puberty onset – PN31-41 for females, PN

37-47 for males. On each CMS day, animals received two different stressors at random times: one in the morning (between 0800 and 1200 h) and one in the afternoon (between 1300 and 1800 h), with a minimum of 2 h between stressors. On day 1 of CMS and the day immediately following the end of CMS, within 2 h of lights on, basal blood samples were obtained from all animals (including those in the non-CMS group) via tail nick. After tail nick, all animals were weighed and placed in a new home cage. Pre- and post-CMS blood sample and body weight data were published in Raineke et al., 2016. Except for blood sampling and routine husbandry, animals in the non-CMS condition were left undisturbed during this period. CMS and non-CMS animals were housed in different colony rooms so that non-CMS animals were not exposed to the disturbance and stress odors of the CMS animals (Mackay-Sim and Laing, 1980). The order and type of stressor was randomized, but all animals received the same number of exposures to each stressor over the 10-day period. Stressors included: 1) Platform: exposure to an elevated Plexiglas platform (20 × 20 cm) mounted on 90 cm high post for 10 min; 2) Cage tilt: the home cage was tilted at a 30° angle for 2 h; 3) Novel cage: exposure to a novel cage without food and water, with a small amount of novel bedding for 1 h; 4) Soiled cage: exposure to a soiled cage of another animal of the same sex for 1 h; 5) Restraint: restraint in PVC tubes (tube size varied to ensure proper restraint/immobility of each animal) for 30 min; 6) Social isolation without food and water: overnight isolation in a smaller cage (28 × 17 cm with 12.5 cm high) for 12 h; and 7) Empty water bottle: animals given their empty water bottles for 1 h following the social isolation/food and water deprivation period.

### Behavioral Exposure and Brain Extraction

All animals were tested on the open field before and after CMS exposure. For the post-CMS test, half the animals were tested in adolescence (short-term effects of CMS) and the remaining animals were tested in adulthood (long-term effects of CMS). Following post-CMS open field tests, all animals were exposed to the FST (habituated for 15 min and then tested for 5 min the following day). Pre- and post-CMS behavioral data were published in Raineke et al., 2016. Animals were decapitated 30-min after the end of day 2 FST testing and brains were collected, quickly frozen on dry ice and stored at -80°C.

### Neural assessment of CRHR1, MR, and GR by in situ hybridization

**Probes**—CRHR1 (1.3kb fragment cloned into Bluescript SK plasmid) was provided by Dr. Victor Viau (source: Dr. Cyntia Donaldson, Perrin et al., 1993). Rat MR (550bp fragment in Bluescript SK) and GR (456bp fragment in pGem4) were provided by Dr. James Herman. All probes were transcribed using <sup>35</sup>S-UTP (Perkin-Elmer, Waltham, MA) and the Promega Riboprobe System (Promega Corp., Madison, WI) with polymerase T<sup>7</sup> for CRHR1 and GR antisense probes, and T<sup>3</sup> for MR antisense probe. Probes were purified using Micro Bio-Spin 30 Columns (Bio-Rad, CA, USA) and 0.1 M DTT was added to prevent oxidation.

**In situ hybridization**—Brains were sectioned coronally (20 μm) using a cryostat (-16°C) and stored at -80°C. Thawed sections were fixed in formalin for 30 min and pre-hybridized as follows: 1 × PBS for 10 min twice, proteinase K (0.1μg/L; 37°C) for 9 min, 0.1 M triethanolamine-hydrochloride (TEA) for 10 min, 0.1 M TEA with 0.25% acetic anhydride for 10 min, 2 × SSC for 10 min twice, dehydrated by a graded series of ethanol, chloroform,

and 100% ethanol and air-dried. Probes were applied at  $1 \times 10^6$  cpm/slide in 75% Hybridization buffer (75% formamide, 3  $\times$  SSC, 1 $\times$  Denhardt's solution, 200  $\mu$ g/mL yeast tRNA, 50 mM sodium phosphate buffer (pH 7.4), 10% dextran sulphate, 10 mM DTT) and covered with HybriSlips (Sigma-Aldrich, ON). Sections were incubated overnight at 55°C in chambers humidified with 75% formamide. HybriSlips were removed and slides were rinsed as follows: 2  $\times$  SSC twice for 20 min, 2  $\times$  SSC for 30 min, 50  $\mu$ g/L RNase A solution (37°C) for 60 min, 2  $\times$  SSC with 0.01 M DTT for 10 min, 1  $\times$  SSC for 10 min, 0.5  $\times$  SSC with 0.01 M DTT for 10 min, 0.1  $\times$  SSC with 0.01 M DTT (60°C) for 60 min, 0.1  $\times$  SSC for 5 min. Sections were dehydrated by a graded series of ethanol and air dried overnight. The hybridized slides were then exposed to Kodak BioMax MR film, and developed using Kodak GBX developer and fixer.

**Densitometric analysis**—The autoradiograph films were scanned and analyzed with ImageJ 1.48v (National Institutes of Health, USA). The left and right subregions of the dorsal (CA1, CA3, and DG) and ventral hippocampal formation (CA1, CA3, DG, and ventral subiculum) were traced free-hand in two sections per animal to determine mean grey density levels. Corrected grey levels were obtained by subtracting the background level (corpus callosum) from each of the four measurements. Left and right levels in each measured area were averaged together for analysis.

### Statistical analysis

All data are expressed as mean  $\pm$  SEM and were analyzed by two-way ANOVA (prenatal treatment and CMS as factors). When significant interactions between prenatal treatment and CMS were detected, ANOVAs were followed by Newman-Keuls *post hoc* tests. Sex was not used as a factor in the ANOVAs because females and males were exposed to CMS during different ages (PN31-41 and PN 37-47 respectively). Age was also not used as factor because *in situ* hybridizations for brains collected during adolescence and adulthood were run separately. Additionally, we utilized planned comparisons (Student's *t*-tests) to test the *a priori* hypotheses that: 1) PAE will alter animals' receptor expression compared to controls; and 2) CMS will differentially alter animals' receptor expression. In all cases, differences were considered significant when  $p < 0.05$ . Outliers were identified and removed using the Robust regression and Outlier removal (ROUT) method with  $Q=0.05$ .

## Results

### Mineralocorticoid receptor mRNA expression

**Dorsal hippocampal formation**—In adolescent males, PAE decreased MR mRNA expression in the CA1, CA3, and DG, independently of CMS exposure [Figure 1A,E,I; significant main effects of prenatal treatment for CA1 ( $F_{(1,28)}=9.63$ ,  $p=0.004$ ), CA3 ( $F_{(1,26)}=13.99$ ,  $p=0.0009$ ), and DG ( $F_{(1,28)}=4.85$ ,  $p=0.04$ )]. Neither PAE nor CMS affected MR mRNA expression in the dorsal hippocampal formation of adolescent females or in adult males or females (Figure 1).

**Ventral hippocampal formation**—In adolescent males, CMS decreased MR mRNA expression in the ventral subiculum in control but not PAE rats, while no changes were

observed in CA1, CA3, or DG [Figure 2A,E,I,M; *a priori* analysis for ventral subiculum comparing control non-CMS to control CMS ( $p=0.04$ )]. In adolescent females, however, CMS increased MR mRNA expression only in the CA1 of controls [Figure 2B; *a priori* analysis for CA1 comparing control non-CMS to control CMS ( $p=0.04$ )]. Moreover, adolescent PAE non-CMS females showed increased MR mRNA expression in the ventral subiculum compared to control non-CMS females [Figure 2N; *a priori* analysis for ventral subiculum comparing PAE non-CMS to PAE CMS ( $p=0.02$ )]. Neither PAE nor CMS affected MR mRNA expression in the ventral CA3 or DG of adolescent females (Figure 2F,J).

In adulthood, adolescent CMS reduced MR mRNA expression in the ventral subiculum of males independently of prenatal treatment [Figure 2O; significant main effect of CMS for ventral subiculum ( $F_{(1,30)}=5.46$ ,  $p=0.03$ )]. However, neither PAE nor CMS affected MR mRNA expression in the ventral CA1, CA3 and DG of adult males (Figure 2 C,G,K). In adult females, adolescent CMS increased MR mRNA expression in the CA3 and ventral subiculum of control but not PAE animals [Figure 2H,P; *a priori* analysis comparing control non-CMS to control CMS for CA3 ( $p=0.05$ ) and ventral subiculum ( $p=0.01$ )]. Neither PAE nor CMS affected MR mRNA expression in the ventral CA1 or DG of adult females (Figure 2D,L).

### Glucocorticoid receptor mRNA expression

**Dorsal hippocampal formation**—In adolescent males, CMS decreased GR mRNA expression in CA1 only in control rats, but no changes were observed in CA3 or DG [Figure 3A,E,I; *a priori* analysis for CA1 comparing control non-CMS to control CMS ( $p=0.05$ )]. In adolescent females, CMS increased GR mRNA expression in CA1, CA3, and DG in both groups independently of prenatal exposure (Figure 3B,F,J). Moreover, PAE reduced GR mRNA expression in CA1 and DG independently of CMS exposure [significant main effects of CMS for CA1 ( $F_{(1,28)}=7.78$ ,  $p=0.009$ ), CA3 ( $F_{(1,28)}=7.22$ ,  $p=0.01$ ), and DG ( $F_{(1,28)}=8.77$ ,  $p=0.006$ ) and significant main effects of prenatal treatment for CA1 ( $F_{(1,28)}=10.04$ ,  $p=0.003$ ), and DG ( $F_{(1,28)}=8.63$ ,  $p=0.006$ )].

In adult males, CMS during adolescence reduced GR mRNA expression in the CA1 and DG in both groups and in CA3 only in control animals [Figure 3C,G,K; significant main effects of CMS for CA1 ( $F_{(1,30)}=4.47$ ,  $p=0.04$ ), and DG ( $F_{(1,30)}=3.97$ ,  $p=0.05$ ); *a priori* analysis for CA3 comparing control non-CMS to control CMS ( $p=0.05$ )]. Neither PAE nor CMS affected GR mRNA expression in the dorsal CA1, CA3, or DG of adult females (Figure 3D,H,L).

**Ventral hippocampal formation**—Neither PAE nor CMS affected GR mRNA expression in the ventral hippocampal formation of adolescent males (Figure 4A,E,I,M). However, in adolescent females, PAE increased GR mRNA expression in the CA1 independently of CMS exposure, but no changes were observed in CA3, DG, or ventral subiculum [Figure 4B,F,J,N; *a priori* analysis for CA1 comparing control non-CMS to PAE non-CMS ( $p=0.04$ )].

In adulthood, adolescent CMS increased GR mRNA expression in the ventral subiculum of males independently of prenatal treatment [Figure 4O; significant main effect of CMS for

ventral subiculum ( $F_{(1,30)}=5.23$ ,  $p=0.03$ ). However, neither PAE nor CMS affected GR mRNA expression in the ventral CA1, CA3 or DG of adult males (Figure 4C,G,K). In adult females, adolescent CMS increased GR mRNA expression in the CA1 only in controls, but no changes were observed in CA3, DG, or ventral subiculum [Figure 4D,H,L,P; *a priori* analysis for CA1 comparing control non-CMS to control CMS ( $p=0.04$ )].

### Type 1 CRH receptors mRNA expression

**Dorsal hippocampal formation**—In adolescent males, PAE decreased CRHR1 mRNA expression in the CA1 independently of CMS exposure [Figure 5A; significant main effect of prenatal treatment for CA1 ( $F_{(1,28)}=7.51$ ,  $p=0.01$ )]. No changes were observed in CRHR1 mRNA expression in the CA3 and DG of adolescent males (Figure 5E,I). Additionally, neither PAE nor CMS affected CRHR1 mRNA expression in the dorsal hippocampal formation of adolescent females (Figure 5B,F,J).

In adult males, CMS decreased CRHR1 mRNA expression only in the CA3 of PAE rats, while no changes were observed in CA1 or DG [Figure 5C,G,K; *a priori* analysis for CA3 comparing PAE non-CMS to PAE CMS ( $p=0.02$ )]. In adult females, PAE reduced CRHR1 mRNA receptors in the CA1 and DG. In the CA1 the effect was only observed in the non-CMS animals and in the DG the effect was independent of CMS exposure [Figure 5D,L; *a priori* analysis for CA1 comparing control non-CMS to PAE non-CMS ( $p=0.05$ ); significant main effect of prenatal treatment for DG ( $F_{(1,28)}=5.75$ ,  $p=0.02$ )]. No changes were observed in CRHR1 mRNA expression in CA3 of adult females (Figure 5H).

**Ventral hippocampal formation**—In adolescent males, PAE decreased CRHR1 mRNA expression in CA1 independently of CMS exposure [Figure 6A; significant main effect of prenatal treatment for CA1 ( $F_{(1,27)}=12.01$ ,  $p=0.001$ )]. Moreover, CMS decreased CRHR1 mRNA expression in CA3 of PAE adolescent male rats [Figure 6E; significant interaction between prenatal treatment and CMS for CA3 ( $F_{(1,27)}=7.19$ ,  $p=0.01$ )]. Neither PAE nor CMS affected CRHR1 mRNA expression in DG or ventral subiculum of adolescent males (Figure 6I,M). In adolescent females, PAE decreased CRHR1 mRNA expression in DG independently of CMS exposure [Figure 6J; significant main effect of prenatal treatment for CA1 ( $F_{(1,28)}=4.18$ ,  $p=0.05$ )]. However, neither PAE nor CMS affected CRHR1 mRNA expression in CA1, CA3, or ventral subiculum of adolescent females (Figure 6B,F,N).

In adult males, CMS during adolescence increased CRHR1 mRNA expression only in CA1 of control animals [Figure 6C; *a priori* analysis for CA1 comparing control non-CMS to control CMS ( $p=0.02$ )]. Neither PAE nor CMS affected CRHR1 mRNA expression in the CA3, DG, or ventral subiculum of adult males (Figure 6G,K,O). In adult females, adolescent CMS increased CRHR1 mRNA receptors in CA3 of controls and reduced CRHR1 mRNA receptors in CA3 of PAE animals [Figure 6H; significant interaction between prenatal treatment and CMS for CA3 ( $F_{(1,28)}=8.23$ ,  $p=0.008$ ); *a priori* analysis for CA3 comparing control non-CMS to control CMS ( $p=0.04$ )]. Neither PAE nor CMS affected CRHR1 mRNA expression in CA1, DG, or ventral subiculum of adult females (Figure 6D,L,P).



## Discussion

Our ontogenetic approach investigating how PAE affects stress-related receptor mRNA expression in the hippocampus over the course of development revealed that the effects of PAE are more apparent during adolescence and decline by adulthood. Indeed, in adolescence, PAE resulted in altered MR, GR, and CRHR1 mRNA expression in several subfields of the dorsal and ventral hippocampal formation in a sexually dimorphic manner (Table 1). However, in adulthood, PAE only altered CRHR1 mRNA expression in the dorsal hippocampus of females. In contrast, the effects of adolescent CMS on stress-related receptor mRNA expression in the hippocampus showed the opposite pattern; that is, the effects of adolescent CMS on hippocampal expression of stress-related receptors were more pronounced in adulthood, long after the termination of the stress exposure. PAE animals also appeared to be less responsive to the effect of adolescent CMS, as CMS uniquely affected only CRHR1 in PAE animals compared to the altered MR, GR and CRHR1 mRNA expression observed in controls.

### PAE effects on stress-related receptors in the hippocampus are transient

Studies investigating the long-term effect of PAE on stress-related receptors expression in the dorsal hippocampus have demonstrated that, in adult males, hippocampal MR and GR expression is not altered (Glavas et al., 2007; Kim et al., 1999; Uban et al., 2013); however, in adult females, PAE has been shown to reduce hippocampal MR mRNA expression (Sliwowska et al., 2008; Uban et al., 2013). Furthermore, analysis of MR and GR protein levels in the nuclear and cytosolic fractions of whole hippocampus of adolescent male mice exposed to alcohol during gestation indicates that levels of both MR and GR were not altered in the cytosolic fraction, whereas PAE reduced MR and elevated GR protein levels in the nuclear fraction (Caldwell et al., 2014). Our evaluation of stress-related receptor mRNA expression in specific subfields of dorsal and ventral hippocampus in adolescent and adult rats considerably extends these previous findings by demonstrating that there are indeed effects of PAE on these receptors but they are transient; PAE resulted in altered MR and GR mRNA expression in the dorsal and/or ventral hippocampus during adolescence but not in adulthood. Moreover, the effects of PAE were sex and region dependent as adolescent PAE male rats showed reduced MR mRNA expression in dorsal CA1, CA3 and DG while adolescent PAE female showed reduced GR mRNA expression in dorsal CA1 and DG. In addition, adolescent PAE female rats showed increased MR in ventral subiculum and GR in ventral CA1. In contrast to previous studies from our laboratory where adult PAE females showed reduced MR mRNA in the hippocampus (Sliwowska et al., 2008; Uban et al., 2013), we observed no effects of PAE in adult females in the current study. This discrepancy is probably due methodological differences: previous studies reported measures of basal receptor expression while the current study evaluated receptor expression following an acute stressor (e.g. forced swim test).

MR and GR are both nuclear receptors that bind to corticosterone, though GR binds corticosterone with a 10-fold lower affinity. Because of this difference in affinity, MR is already almost fully occupied at basal corticosterone levels, while GR becomes saturated at stress corticosterone levels (Reul and de Kloet, 1985; Spencer et al., 1993). Importantly,

there is evidence that the nuclear type of MR can also be positioned in the membrane of hippocampal neurons (Karst et al., 2005). However, similar to GR, this membrane MR has a low affinity for corticosterone, responding only to high levels such as those present during stress-induced corticosterone release (Joëls et al., 2008). These findings suggest that both MR and GR in the hippocampus play a critical role in behavioral and hormonal stress responses, albeit in unique ways: whereas membrane MR is important for appraisal of the situation (i.e. how threatening is the stressor) and/or for choosing the appropriate response (i.e. determine the behavioral response to the stressor), GR is essential for returning brain activity to baseline after the stress response and for memory consolidation of the stressful event (de Kloet et al., 2005; Joëls et al., 2006; Oitzl and de Kloet, 1992; Schwabe et al., 2007, 2010, 2013). To this end, dysregulation of MR or GR expression in the hippocampus may affect different aspects of basal and/or stress response regulation. The current data show that PAE results in reduced MR in the dorsal hippocampus (CA1, CA3 and DG) of adolescent males and reduced GR in the dorsal hippocampus (CA1 and DG) of adolescent females. Taken together, these results suggest that the behavioral and hormonal alterations in response to acute and/or chronic stress observed in PAE animals (Hellemans et al., 2010; Weinberg et al., 2008) may be supported by different underlying mechanisms depending on the sex of the animal, with alterations in MR expression/regulation supporting the responses observed in PAE males, and alterations in GR expression/regulation supporting the responses observed in PAE females.

The current and previous data (Glavas et al., 2007; Kim et al., 1999; Uban et al., 2013) indicate that, in contrast to what is observed in adolescence, adult hippocampal MR and GR expression of PAE animals is not different from controls. It should be noted that the lack of PAE effects on the expression of stress-related receptor mRNA in adulthood does not necessarily preclude effects of PAE on protein levels, localization and/or function. Further investigation of underlying mechanisms, including epigenetic modifications will be important in future studies. Additionally, the PAE-related alterations in hippocampal MR and GR mRNA expression observed during adolescence in the current study may set the stage for the altered behavioral responses shown by PAE animals when faced with different concentrations of corticosterone in adulthood. Indeed, adrenalectomy in adulthood, which reduces corticosterone levels, resulted in an exacerbated increase in MR mRNA expression in dorsal CA3 of PAE females and in GR mRNA expression in dorsal CA3 of PAE males (Glavas et al., 2007). Additionally, corticosterone replacement in adrenalectomized animals was ineffective in normalizing MR mRNA expression in dorsal CA1 and DG of PAE males (Glavas et al., 2007). Together these data indicate that, even though the PAE effects of MR and GR mRNA expression in hippocampus appear to be transient, challenges to the system can uncover underlying alterations in the regulation of stress-related receptor expression in adulthood.

In addition to MR and GR, hippocampal CRHR1 also plays a role in emotional regulation. Indeed, abnormal activity of the central CRH system is associated with increased expression of many psychopathologies, including depression and anxiety (Binder and Nemeroff, 2010; Holsboer and Ising, 2008). Animal studies have confirmed this association by demonstrating that many of the behavioral abnormalities associated with depressive- and anxiety-like behaviors are mediated by dysregulated activity of CRHR1 in limbic areas, including the

hippocampus. Specifically, increased activity of the CRH system is associated with increased anxiety-like behavior as indicated by the finding that intracerebroventricular administration of CRH increases expression of anxiety-like behaviors (Dunn and File, 1987; Britton et al., 1982). In contrast, decreased activity of the CRH system is associated with reductions in anxiety-like behavior, as shown in studies using CRHR1-deficient mice (Smith et al., 1998; Timpl et al., 1998), conditional inactivation of the CRHR1 in limbic areas (Müller et al., 2003), and CRHR1 antagonism in rats (Keck et al., 2001; Sandi et al., 2008; Veenit et al., 2014). Previous studies using animal models of PAE have demonstrated that rats exposed to alcohol during gestation show reduced CRHR1 mRNA expression in the amygdala, mPFC, and pituitary in adulthood (Glavas et al., 2007; Raineki et al., 2016). Here, we extend those findings by demonstrating that PAE also reduced CRHR1 mRNA expression in the hippocampus. Nevertheless, and in contrast with previous findings, the current data indicate that PAE effects on CRHR1 mRNA expression in the hippocampus are more evident during adolescence than adulthood. PAE reduced the expression of CRHR1 in the dorsal and ventral CA1 of adolescent males and in the ventral DG of adolescent females. In adulthood, however, PAE effects were only observed in females, with a reduction of CRHR1 mRNA expression in dorsal CA1 and DG. Consistent with previous findings on central CRHR1 expression following PAE (Caldwell et al., 2015; Glavas et al., 2007; Raineki et al., 2016), we also observed a reduction in CRHR1 mRNA expression in the hippocampus. This overall reduction in central CRHR1 expression could be a response to increased CRH production; however, extended investigation of CRH mRNA expression following PAE indicated that PAE increased CRH only in the central amygdala (Lan et al., 2015), with no changes were observed in the paraventricular nucleus of hypothalamus, mPFC, or nucleus accumbens (Glavas et al., 2007; Lan et al., 2015; Uban et al., 2013). More work is necessary to characterize fully the effects of PAE on CRH expression in areas that are known to produce and release CRH in the brain, especially in the hippocampus and locus coeruleus.

Besides CRHR1 and GR, MR has increasingly been recognized as playing a crucial role in the pathophysiology of depression (Chen et al., 2016; Hinkelmann et al., 2016; Mostalac-Preciado et al., 2011; Wingenfeld et al., 2016). In our previous study using the same animals as in the current study (Raineki et al., 2016), we demonstrated that PAE males showed depressive-like behaviors only during adolescence and not in adulthood. Similar to the behavioral effects of PAE on depressive-like behaviors, here, we observed a temporary reduction in MR mRNA expression in dorsal CA1, CA3, and DG and in CRHR1 in dorsal and ventral CA1 during adolescence in PAE males. However, those reductions in MR and CRHR1 were also transient and disappeared in adulthood. This temporal alignment between behavioral and receptor expression suggests that the depressive-like behavior in PAE animals may be mediated, at least in part, by deficits in CRHR1 and MR expression in the hippocampus.

### **Effects of adolescent CMS on stress-related receptors in the hippocampus**

Exposure to chronic stress during adolescence has been shown to induce short- and long-term alterations to the structure and function of the hippocampus, including changes in the expression of stress-related receptors (Barha et al., 2011; Eiland and Romeo, 2013; Hueston

et al., 2017; Iredale et al., 1996; Isgor et al., 2004; Li et al., 2015; McCormick et al., 2012; Sterlemann et al., 2008, 2010; Veenit et al., 2014). Nevertheless, the direction of these effects can diverge among studies; this is not necessarily surprising given that different studies vary in their use of specific stress paradigms, duration of stress exposure, animal ages, and methodologies to measure outcomes. Despite limiting our ability to generalize conclusions, the use of different adolescent stress protocols has underscored the importance of how each variable within a stress model can differentially impact the outcome. Notably, very little is known about how adolescent stress affects the expression of stress-related receptors in the hippocampus of females (Wulsin et al., 2016) because, the literature has almost exclusively focused on assessing males (Iredale et al., 1996; Isgor et al., 2004; Li et al., 2015; Sterlemann et al., 2008; Veenit et al., 2014). The current dataset provides valuable information on the differential short- and long-term responses to chronic stress in adolescent males and females. Furthermore, as MR, GR, and CRHR1 in the hippocampus are critical regulators of cognition, stress and emotional responses, the current data may help us better understand the underlying mechanisms by which adolescent stress can lead to sexually dimorphic effects in: 1) learning and memory (Toledo-Rodriguez et al., 2012; Toledo-Rodriguez and Sandi, 2007); 2) anxiety- and depressive-like behaviors (Bourke and Neigh, 2011; McCormick et al., 2008; Raineki et al., 2016; Toledo-Rodriguez and Sandi, 2011); and 3) HPA axis function (Barha et al., 2011; Bourke and Neigh, 2011; Raineki et al., 2016).

It has been shown that adolescent stress exposure leads to similar short- and long-term effects on hippocampal GR expression but in a sexually dimorphic manner, with males showing decreased GR mRNA expression in dorsal CA1 and DG in both the short- and long-term (Isgor et al., 2004), while females show no observable changes (Wulsin et al., 2016). The current study replicates these findings in males, showing that adolescent CMS reduced GR mRNA expression in dorsal CA1 during adolescence and in dorsal CA1, CA3, and DG in adulthood (Table 1). Nevertheless, we also saw increased GR mRNA expression in the ventral subiculum of adult males as well as increased GR in dorsal CA1, CA3, and DG in adolescent females and in ventral CA1 in adult females following adolescent CMS.

In contrast to GR, less is known about the effects of adolescent stress on MR expression in the hippocampus. It has been shown that, in male mice, adolescent stress reduces the MR mRNA expression in dorsal CA1, CA3 and DG in adulthood (Sterlemann et al., 2008). Though our current results do not corroborate this previous study, our data do expand knowledge about how adolescent stress exposure can have profound impacts on MR expression in the hippocampus. Indeed, adolescent CMS reduces MR mRNA expression in ventral subiculum of males and increases MR mRNA expression in the ventral CA1 of females in the short term. However, the long-term effects of adolescence CMS on MR were only observed in females, where adolescent CMS increased MR in the ventral CA3 and subiculum in adulthood.

Previous studies found that in the short-term, adolescent stress increases CRHR1 mRNA expression in whole hippocampus in males (Iredale et al., 1996). However, we observed that adolescent CMS did not alter the expression of CRHR1 in the hippocampus immediately after the end of stress exposure. The literature on the long-term effects of adolescent stress on CRHR1 expression in the hippocampus is somewhat mixed, with data indicating that

adolescent stress can both increase CRHR1 mRNA expression (Veenit et al., 2014) and decrease CRHR1 protein expression (Li et al., 2015) in the hippocampus. This inconsistency is probably due to methodological differences between studies (as discussed above). Nevertheless, our data assessing the long-term effects of adolescent CMS indicate that both males and females show increased CRHR1 mRNA expression in the hippocampus in adulthood, although the effect was observed in different subfields of the hippocampus and was sex-dependent.

The effects of chronic stress during adolescence on brain and behavior can manifest differently depending on the age when observations are made. Certain effects are only present immediately after the end of the stress period, some are only observed long after the end of stress, and some are persistent and can be detected any time after stress exposure (Bourke and Neigh, 2011; McCormick and Green, 2013; Rainecki et al., 2016; Toledo-Rodriguez and Sandi, 2007; Tsoory et al., 2008). The current data indicate that exposure to CMS during adolescence induces both short- and long-term effects on the expression of stress-related receptors in the hippocampus; however, the effects of CMS were more pronounced in adulthood, long after the termination of stress exposure (Table 1).

Lastly, the hippocampus of PAE animals appears to be less responsive to the effect of adolescent CMS, as CMS uniquely affected only CRHR1 in PAE animals compared to the altered MR, GR and CRHR1 mRNA expression observed in controls (Table 1). This blunted responsiveness to adolescent CMS may indicate that the hippocampus of PAE animals shows reduced neuroplasticity, especially when facing chronic stress. Corroborating this suggestion are findings indicating that PAE is associated with long-term potentiation deficits in the hippocampus (Christie et al., 2005; Patten et al., 2013; Sutherland et al., 1997) and that chronic restraint stress in adulthood reduces hippocampal neurogenesis in control, but not PAE animals (Sliwowska et al., 2010). This PAE-induced blunted hippocampal responsiveness to environmental stimuli (e.g. stress) and reduced neuroplasticity observed in animal models has important implications for individuals exposed to alcohol during gestation. Indeed, appropriate responding to environmental stimulation (e.g. stress) is a required property for the hippocampus to play its critical role in learning and memory as well as in emotional and stress regulation, all of which show some level of dysregulation following PAE. Clinical studies support this suggestion by demonstrating that PAE-induced disruptions in hippocampal development is directly associated with problems in memory and learning (Coles, et al., 2011; Gross et al., in press; Willoughby et al., 2008).

## Conclusions

Together, the current results indicate that PAE and adolescent CMS induce dynamic alterations in the expression of stress-related receptors in the dorsal and ventral hippocampal formation, and that these alterations manifest differently depending on the age and sex of the animal. Due to the importance of hippocampal MR, GR, and CRHR1 for emotional and stress regulation as well as for learning and memory, the PAE- and adolescent stress-related alterations observed here may underlie some of the hippocampal-dependent behavioral deficits associated with those insults. Finally, the short-term impact of adolescent CMS on the expression of stress-related receptors in the hippocampus differs from the long-term

effects, suggesting a possible incubation period for some of the effects of adolescent stress in altering the developmental trajectory of the hippocampus.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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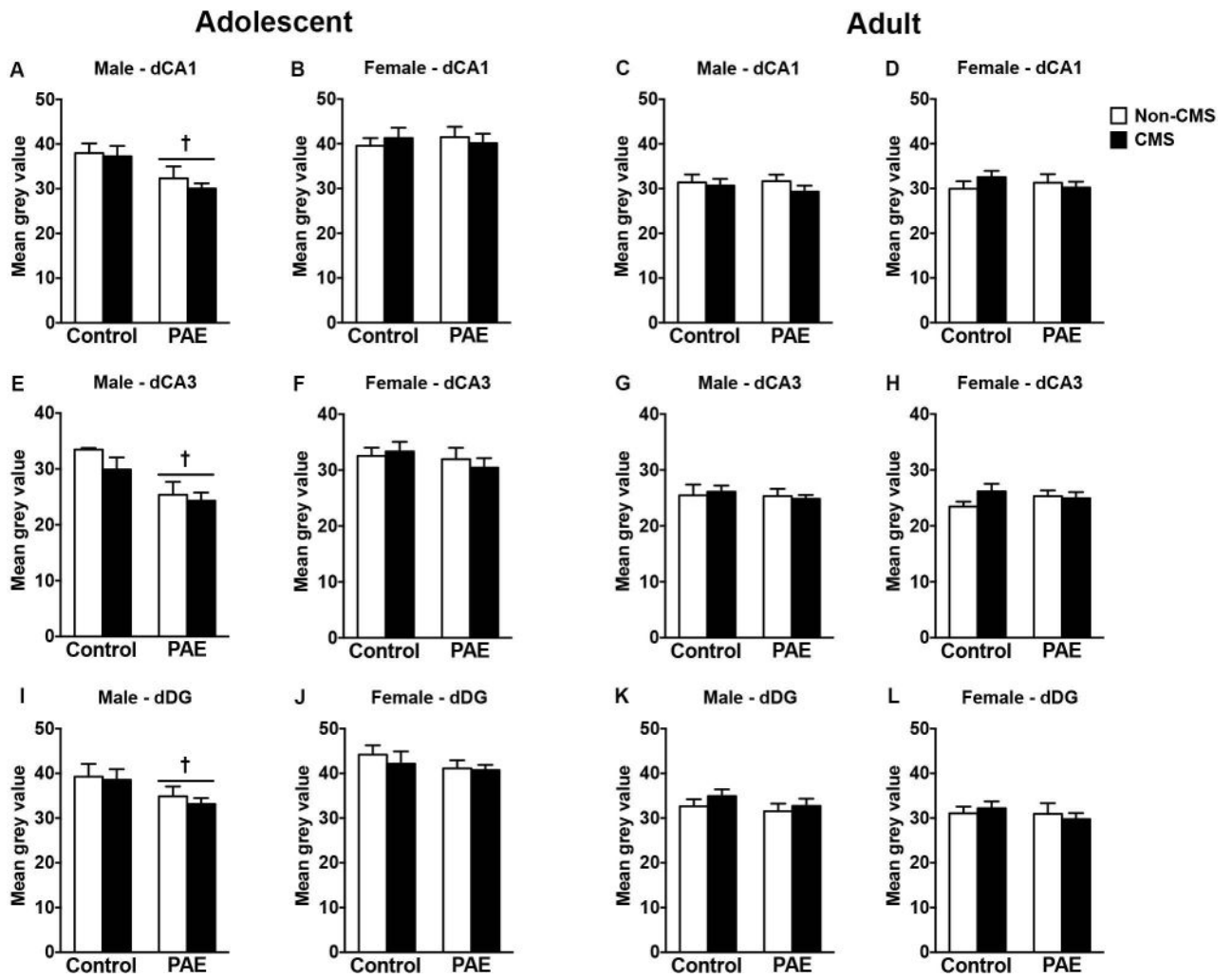
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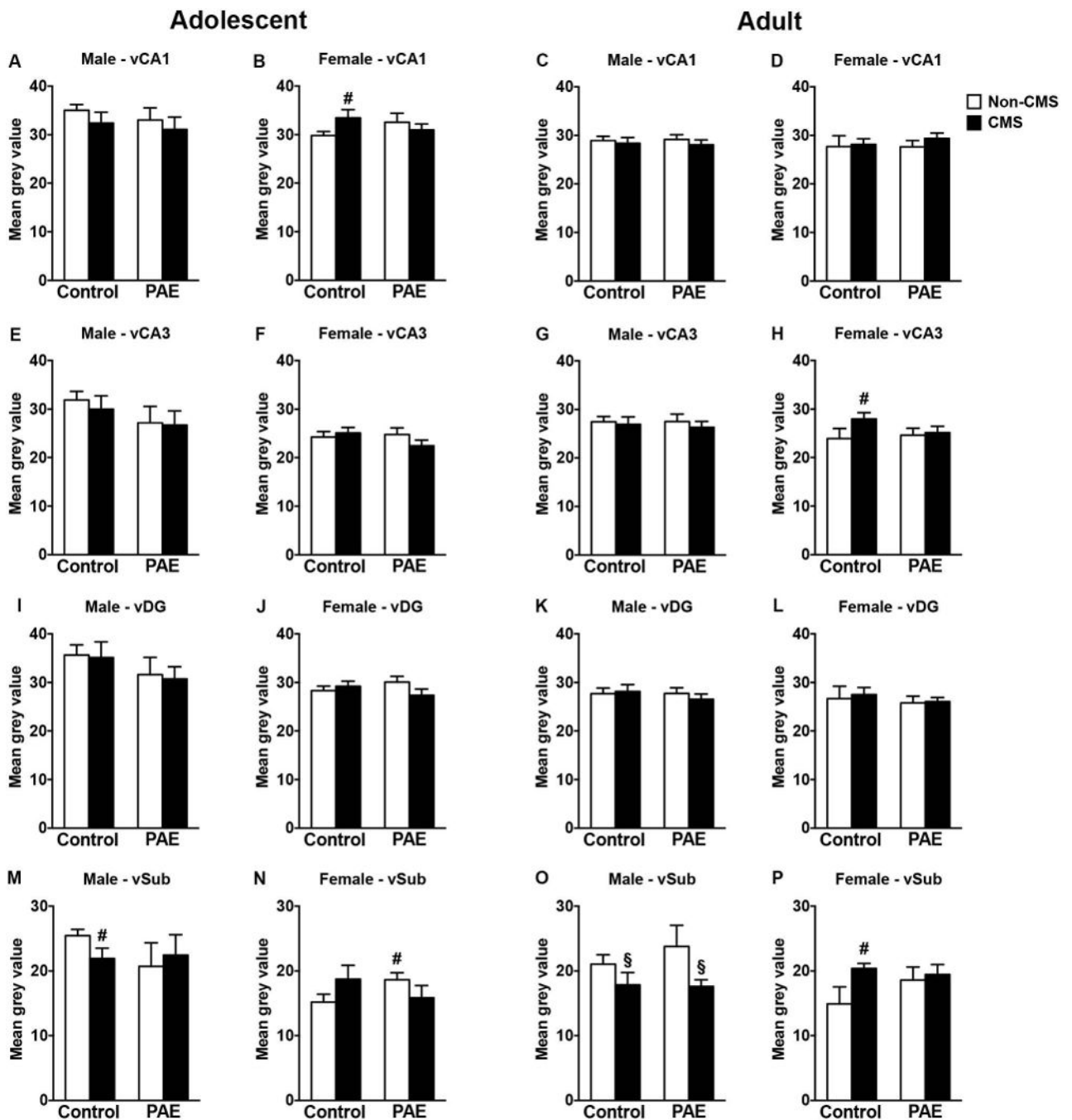
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**Figure 1.** Short- and long-term effects of adolescent CMS on dorsal hippocampus MR mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of MR mRNA expression in the CA1 (A-D), CA3 (E-H), and DG (I-L). † indicates a significant main effect of prenatal treatment, where all PAE animals are different from control animals (n = 6-10 for all groups).



**Figure 2.** Short- and long-term effects of adolescent CMS on ventral hippocampus MR mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of MR mRNA expression in the CA1 (A-D), CA3 (E-H), DG (I-L), and ventral subiculum (M-P). § indicates a significant main effect of CMS exposure, where all animals exposed to CMS are different from animals not exposed to CMS; for B, H, M, and P, # indicates that control CMS is different from control non-CMS based on *a priori* comparisons; for N, #

indicates that PAE non-CMS is different from control non-CMS based on *a priori* comparisons (n = 4-10 for all groups).

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Adolescent

Adult

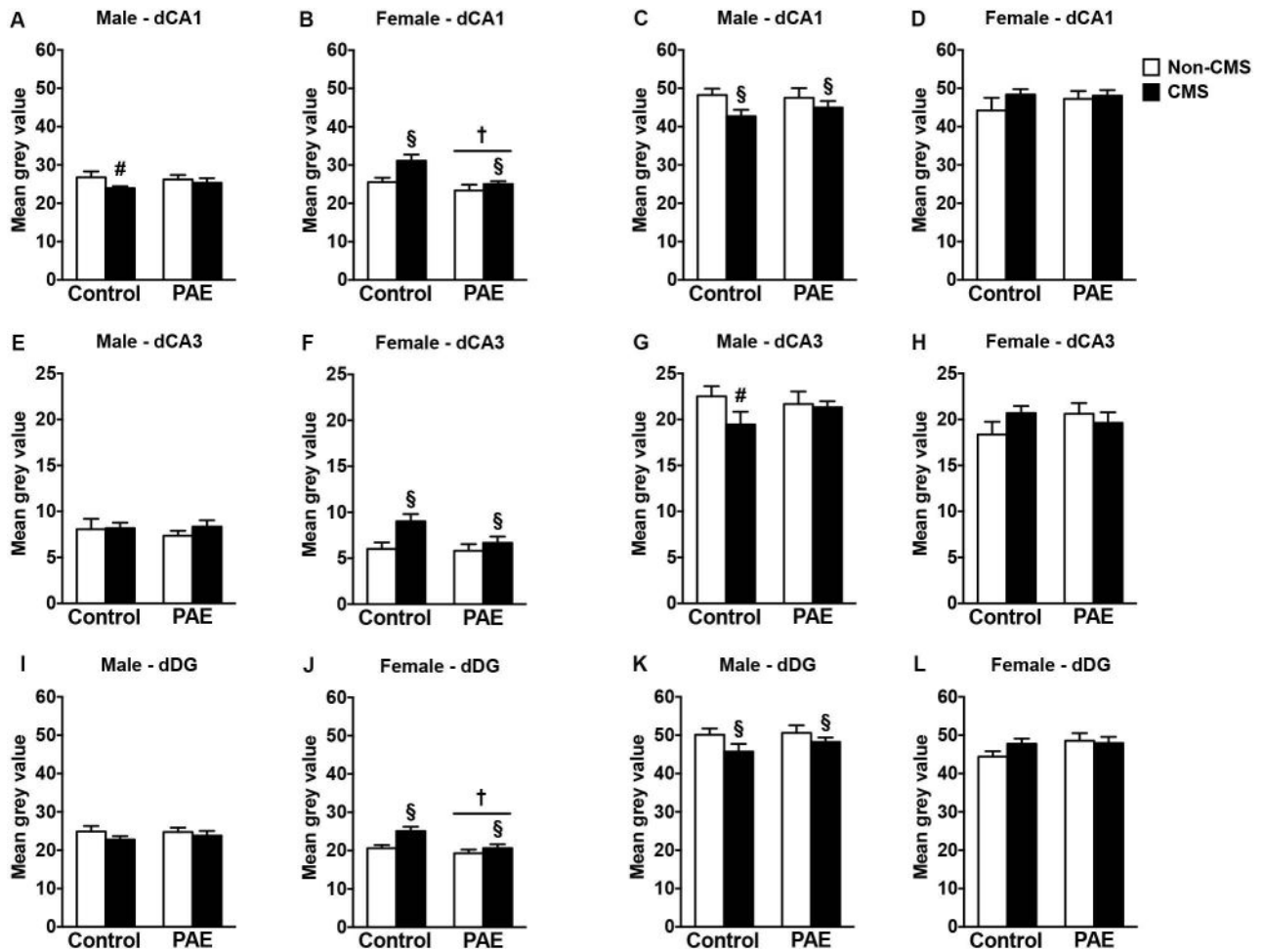
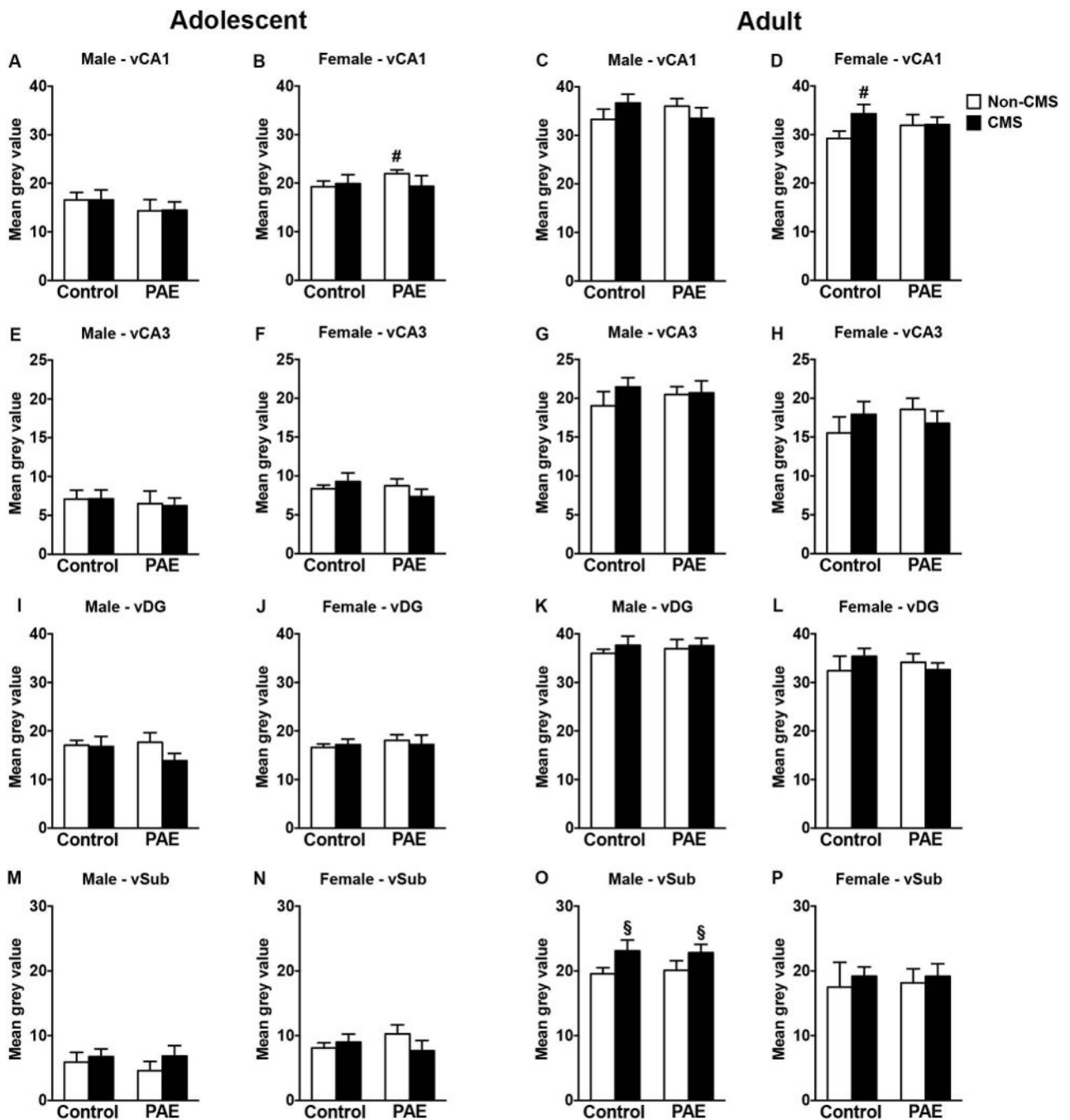


Figure 3.

Short- and long-term effects of adolescent CMS on dorsal hippocampus GR mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of GR mRNA expression in the CA1 (A-D), CA3 (E-H), and DG (I-L). † indicates a significant main effect of prenatal treatment, where all PAE animals are different from control animals; § indicates a significant main effect of CMS exposure, where all animals exposed to CMS are different from animals not exposed to CMS; for A and G, # indicates that control CMS is different from control non-CMS based on *a priori* comparisons (n = 6-10 for all groups).



**Figure 4.** Short- and long-term effects of adolescent CMS on ventral hippocampus GR mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of GR mRNA expression in the CA1 (A-D), CA3 (E-H), DG (I-L), and ventral subiculum (M-P). § indicates a significant main effect of CMS exposure, where all animals exposed to CMS are different from animals not exposed to CMS; for B, # indicates that PAE non-CMS is different from control non-CMS based on *a priori* comparisons; for D, # indicates that



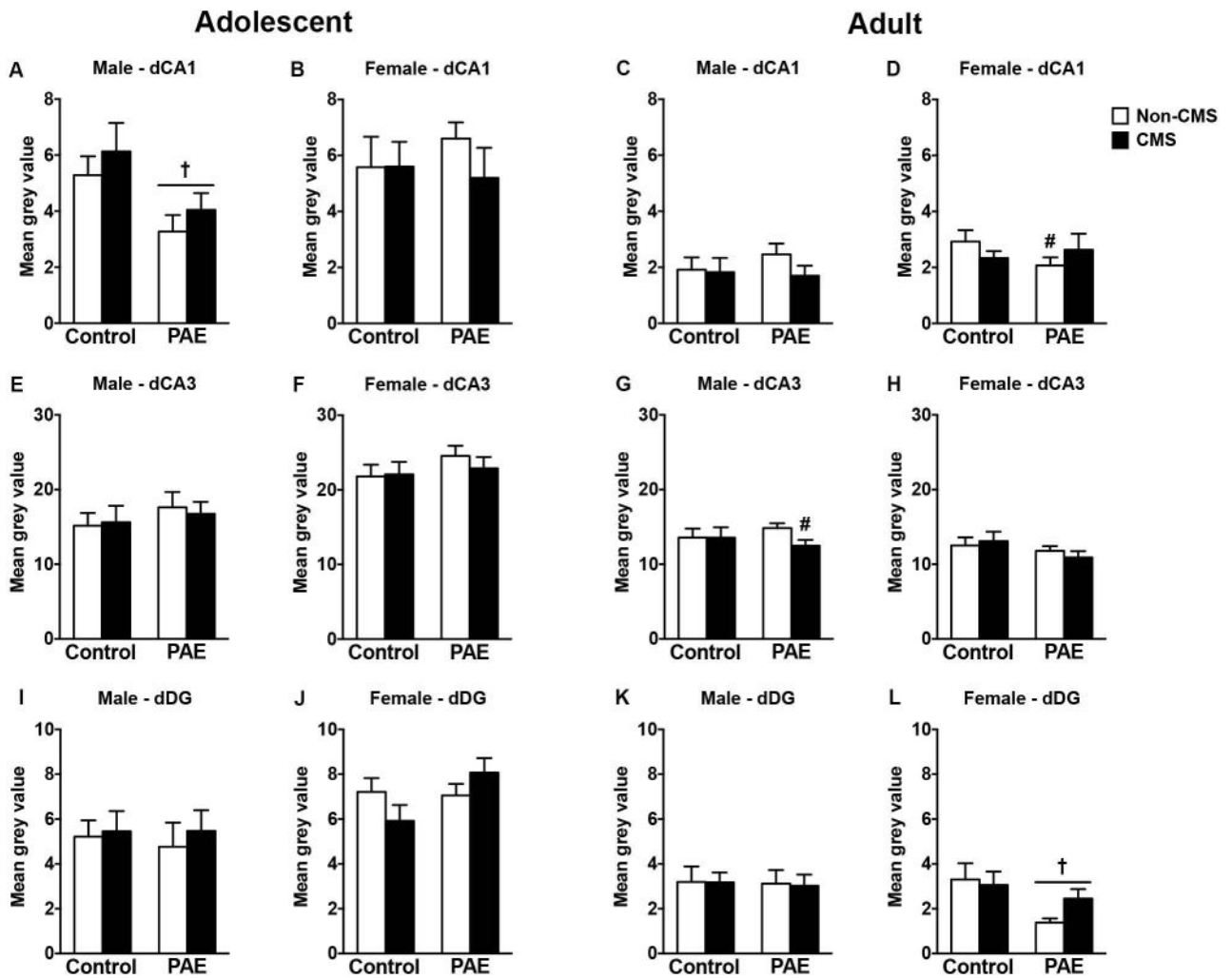
control CMS is different from control non-CMS based on *a priori* comparisons; (n = 4-10 for all groups).

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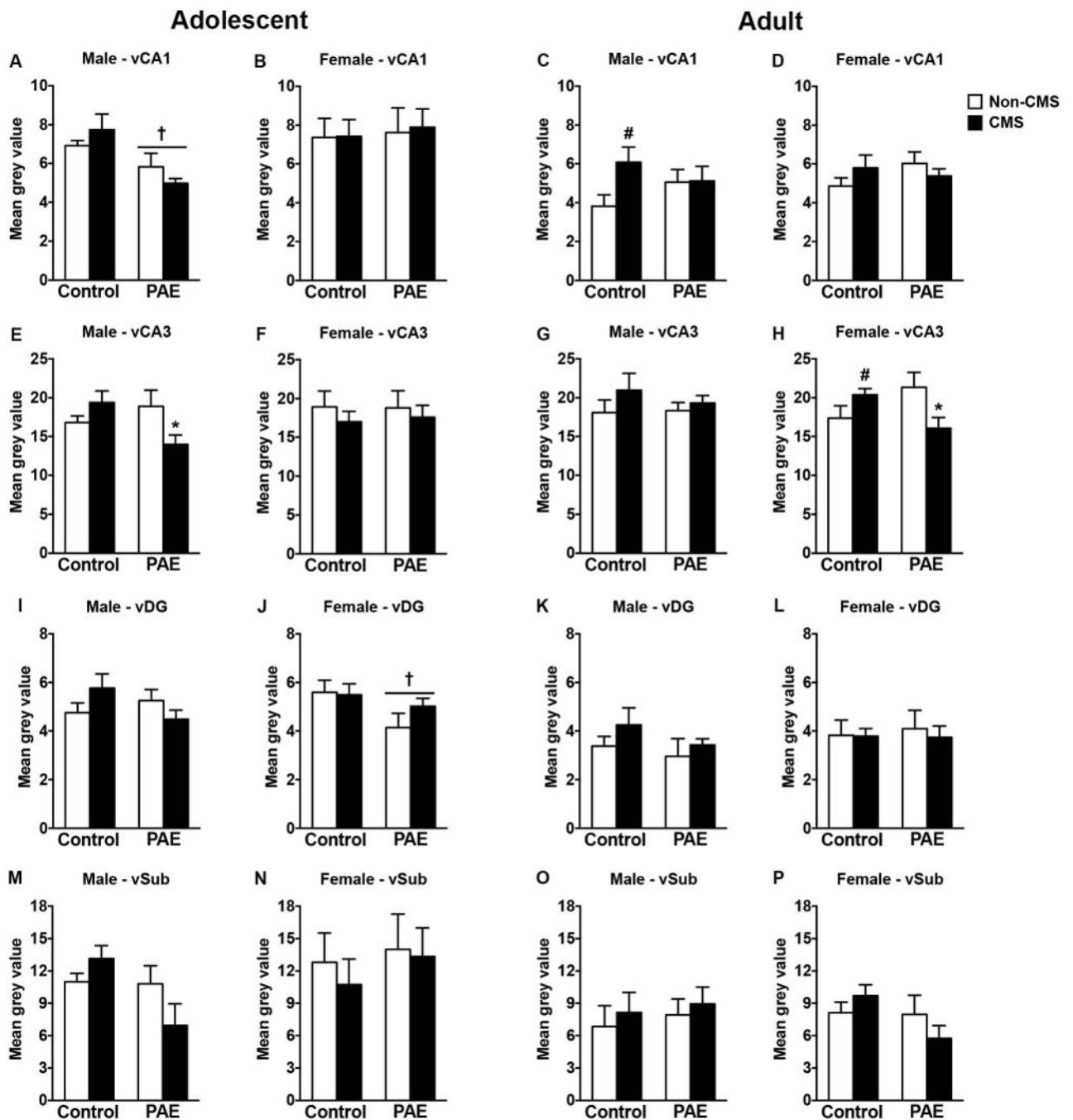
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**Figure 5.** Short- and long-term effects of adolescent CMS on dorsal hippocampus CRHR1 mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of CRHR1 mRNA expression in the CA1 (A-D), CA3 (E-H), and DG (I-L). † indicates a significant main effect of prenatal treatment, where all PAE animals are different from control animals; for D, # indicates that PAE non-CMS is different from control non-CMS based on *a priori* comparisons; for G, # indicates that PAE CMS is different from PAE non-CMS based on *a priori* comparisons (n = 6-10 for all groups).



**Figure 6.** Short- and long-term effects of adolescent CMS on ventral hippocampus CRHR1 mRNA expression in control and PAE rats. Bars represent the mean  $\pm$  SEM (mean gray value) of CRHR1 mRNA expression in the CA1 (A-D), CA3 (E-H), DG (I-L), and ventral subiculum (M-P). † indicates a significant main effect of prenatal treatment, where all PAE animals are different from control animals; for E and H, \* indicates that PAE CMS is different from PAE

non-CMS; for C and H, # indicates that control CMS is different from control non-CMS based on *a priori* comparisons (n = 3-10 for all groups).

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Summary of the short- and long-term effects of PAE and adolescent CMS on MR, GR, and CRHR1 mRNA expression in different subfields of the dorsal and ventral hippocampus

**Table 1**

<b>Adolescent</b>									
	PAE effects		CMS all groups		CMS control		CMS PAE		
	Male	Female	Male	Female	Male	Female	Male	Female	
MR	↓dCA1 ↓dCA3 ↓dDG	↑vSub	-	-	↓vSub	↑vCA1	-	-	-
GR	-	↓dCA1 ↓dDG ↑vCA1	-	↑dCA1 ↑dCA3 ↑dDG	↓dCA1	-	-	-	-
CRHR1	↓dCA1 ↑vCA1	↓vDG	-	-	-	-	↓vCA3	-	-
<b>Adulthood</b>									
	PAE effects		CMS all groups		CMS control		CMS PAE		
	Male	Female	Male	Female	Male	Female	Male	Female	
MR	-	-	↓vSub	-	-	↑vCA3 ↑vSub	-	-	-
GR	-	-	↓dCA1 ↓dDG ↑vSub	-	↓dCA3	↑vCA1	-	-	-
CRHR1	-	↓dCA1 ↓dDG	-	-	↑vCA1	↑vCA3	↓dCA3	↓vCA3	↓vCA3