

CTCAACCACCTG (470). Nucleotide sequences specific to GR messenger RNA (mRNA) are displayed in boldface type. We also obtained a vector control (lacking a shRNA insert) and a scrambled-sequence control virus (shRNA-Sc; proprietary sequence). All constructs contained a human U6 promoter to drive shRNA expression and contained a green fluorescent protein (GFP) cassette. For *in vitro* and *in vivo* studies, titers of 1×10^6 infection units (IU)/mL and of 1×10^9 IU/mL were used, respectively. All experimental procedures were approved by the University of Cincinnati Institutional BioSafety Committee.

In Vitro

Cell Culture and Transfection. The GR-expressing 4B cells (Dr. Toni Pak, Loyola University, Chicago, Illinois) were seeded in HyClone Dulbecco's modified Eagle's medium (DMEM)/high glucose media (with L-glutamine and L-glucose; Thermo Scientific, Waltham, Massachusetts) and 10% heat-inactivated fetal bovine serum (Atlanta Biologicals, Lawrenceville, Georgia). Cells were treated with trypsin (Invitrogen, Carlsbad, California) and subcultured and transferred to 6-well tissue-cultured plate such that they would reach 70% confluency overnight. Media was removed and replaced with 8×10^5 plaque forming units (PFUs) (or 800 μ L of 1×10^6 IU/mL), shRNA-GR 468, 8×10^5 PFUs shRNA-GR 469, 8×10^5 PFUs shRNA-GR 470, 800 μ L media, or 2 mL media (to control for transfection volume). After 16 hours, the contents of the wells were aspirated and media was replaced (2 mL). Cells were harvested 5 days later for quantification of GR mRNA.

Real Time Quantitative Polymerase Chain Reaction. The RNA was isolated with an RNeasy kit, according to manufacturer protocol

(Qiagen, Valencia, California). The RNA quantity and quality were determined with a NanoVue Plus spectrophotometer (General Electric Healthcare, Piscataway, New Jersey). The RNA was treated with Turbo DNA-free to remove genomic DNA (Ambion, Foster City, California) and reverse transcribed with an iScript complementary DNA synthesis kit according to manufacturer protocol (Bio-Rad, Hercules, California). Real time quantitative polymerase chain reaction (RT qPCR) analysis was performed in an iCycler iQ Multi-Color Real Time PCR Detection System (Bio-Rad). Primers for GR mRNA (10 μ mol/L) (forward: 5'- CCACTGCAGGAGTCTCACAA-3'; and reverse: 5'-ACTGCTGCAAT-CACCTTGACG-3') and the house-keeping gene L-32 (forward: 5'-CATCGTAGAAAGAGCAGCAC-3'; and reverse: 5'-GCACACAAGC-CATCTATTCAT-3') were used (Integrated DNA Technologies, Coralville, Iowa). Quantification of complementary DNA was determined with iQ SYBR Green Supermix (Bio-Rad). Values were calculated with L-32 as an internal standard, and GR mRNA expression is presented as a percentage of control GR expression. Threshold cycle readings for each of the unknown samples were used, and the results were calculated with the $\Delta\Delta$ Ct method (15). Negative RT samples were included to rule out genomic DNA contamination.

In Vivo

Subjects. Male Sprague Dawley rats from Harlan (Indianapolis, Indiana) weighing 250–275 g upon arrival were singly housed throughout the experiment in a temperature/humidity-controlled room on a 12-hour/12-hour light/dark cycle. Food (Teklad; Harlan) and water were available *ad libitum*. All experimental procedures were conducted in accordance with the National Institutes of Health Guidelines for the Care and Use of Animals and approved

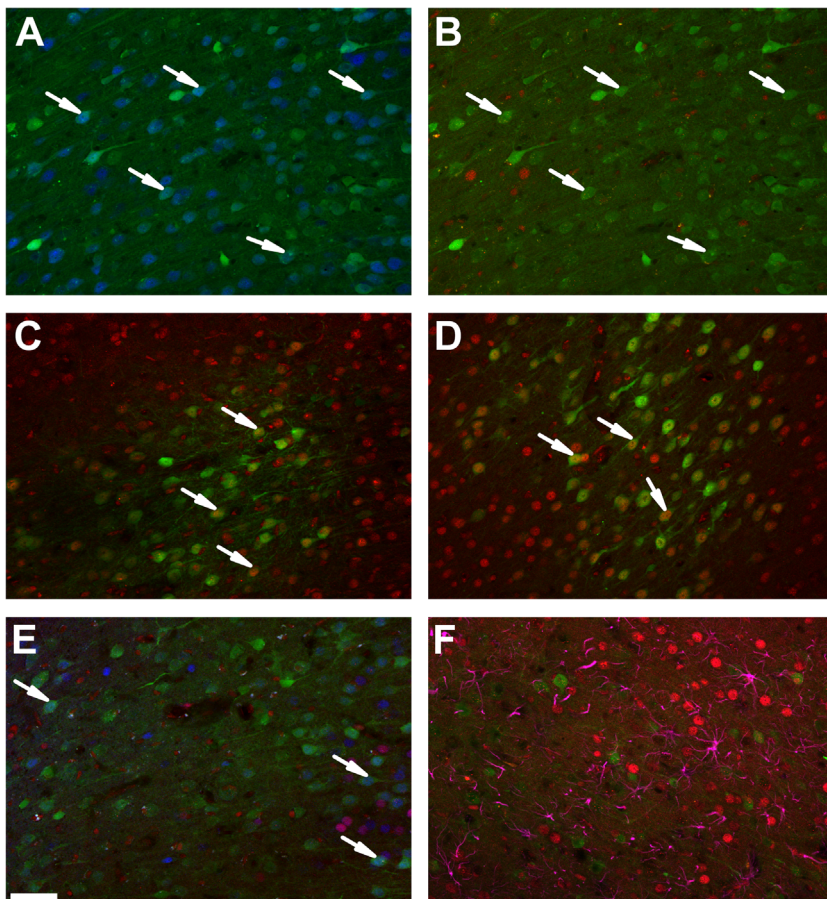


Figure 1. Verification and specificity of a short hairpin RNA targeting the glucocorticoid receptor (shRNA-GR). **(A)** Representative NeuN (blue) immunolabeled sections after microinjection with shRNA-GR (green). After microinjection of shRNA-GR, NeuN immunoreactivity remained intact in transduced neurons, indicating that neuronal viability was not affected (see arrows). **(B)** Representative GR (red) immunolabeled sections after microinjection with shRNA-GR (green), **(C)** shRNA-scrambled control (green), and **(D)** empty vector control (green). The shRNA-GR reduced GR in green fluorescent protein co-localized cells **(B)** relative to animals that received microinjections of shRNA-scrambled control **(C)** or empty vector control **(D)** (see arrows). **(E)** Representative dual GR (red) and mineralocorticoid receptor (MR) (blue) immunolabeled sections after intracranial microinjection with shRNA-GR. Mineralocorticoid receptor expression is intact in cells in which GR immunoreactivity is knocked down (as demonstrated in the superficial layers II/III on the left of the image and deep layers V/VI of the prelimbic prefrontal cortex on the right of the image, where MR is typically expressed. The agranular layer between layers III and V typically has little MR expression) (see arrows). **(F)** Representative glial fibrillary acidic protein (purple) and GR (red) immunolabeled sections after intracranial injection with shRNA-GR. The shRNA-GR did not transduce astrocytes (no green fluorescent protein and glial fibrillary acidic protein co-localization) and did not seem to knockdown astrocytic GR. Scale bar = 50 μ m.

by the University of Cincinnati Institutional Animal Care and Use Committee.

Stereotaxic Surgery. After 1 week of habituation, animals were anesthetized (90 mg/kg ketamine, 10 mg/kg xylazine), and preemptive analgesia (butorphanol) and antibiotic (gentamicin) were administered. Animals received 1- μ L bilateral microinjections into the iIPFC (anterior-posterior [AP] = +3.0, medial-lateral [ML] \pm .6, and dorsal-ventral [DV] = -4.3, Paxinos and Watson [16] coordinates) of shRNA-Sc ($n = 21$) or shRNA-GR ($n = 31$) or 2- μ L bilateral microinjections into the pIPFC (AP = +3.0, ML \pm .7, and DV = -3.3) of vehicle control (high glucose DMEM media with 4.5 g/L glucose, L-glutamine, and pyruvate [Mediatech, Manassas, Virginia]) and 10% heat-inactivated fetal calf serum (Invitrogen; $n = 21$), empty vector control ($n = 22$), or shRNA-GR (targeting position 1690 in the GR gene; $n = 22$) with a 25-gauge, 2- μ L Hamilton syringe (Reno, Nevada). To reduce tissue damage, each injection took place over 20 min. After the needle remained in place for 5 min, the virus was infused over 10 min with a microdriver (Model 5001; Kopf, Tujunga, California) and remained in place for 5 min to allow for complete diffusion. Animals recovered for at least 5–6 weeks before any experiments.

Chronic Variable Stress. Approximately half of the animals underwent chronic variable stress (CVS) for 14 days ($n = 11$ –16 from each microinjection group). The CVS was comprised of twice daily (AM and PM) repeated and unpredictable stressors, including cold swims (10 min, 16°–18°C), warm swims (20 min, 30°–32°C), cold room exposure (1 hour, 4°C), shaker stress (1 hour, 100 rpm), and hypoxia (30 min, 8% oxygen). Only animals undergoing CVS were used in the forced swim test (FST) and open field, and these tests were treated as morning stressors for the first 2 days of CVS.

FST. Approximately half of the animals from each microinjection group went through the modified FST, as described previously (17,18), to assess depression-like behavior. Animals were placed in a cylindrical container (46 cm in height \times 20 cm in diameter) filled with 30 cm of 29° \pm 2°C water for 10 min. Behavior was video recorded and scored every 5 sec for 10 min. Scoring was done by an observer blinded to the experimental condition. Mobility (swimming, climbing, headshakes, and diving) versus immobility was scored as previously described (18). Animals were not exposed to any swims before the FST, because the modified FST is a single exposure test.

Open Field Test. Animals were exposed to a novel open field to assess anxiety-like behavior and locomotor activity. Animals were placed in a 1-meter \times 1-meter black opaque acrylic glass box with 30.48-cm-tall white opaque walls surrounding each side for 5 min. A video recording of the behavior of the animal was scored and analyzed with Clever TopScan Software (CleverSys, Reston, Virginia). Time spent in the center versus the periphery of the open field was used as a measure of anxiety-like behavior (19).

Acute Restraint and Blood Collection. The morning after completion of CVS (at least 16 hours after last stress exposure), all animals were exposed to a novel 30-min restraint. Blood samples (approximately 250 μ L) were collected in tubes containing 10 μ L 100 mmol/L ethylenediamine tetraacetate by tail clip before (0 min) and 30, 60, and 120 min after onset of 30-min restraint and immediately placed on ice. Samples were collected in under 3 min before any rise in adrenocorticotropic hormone (ACTH) or corticosterone levels due to sampling (20). Blood samples were

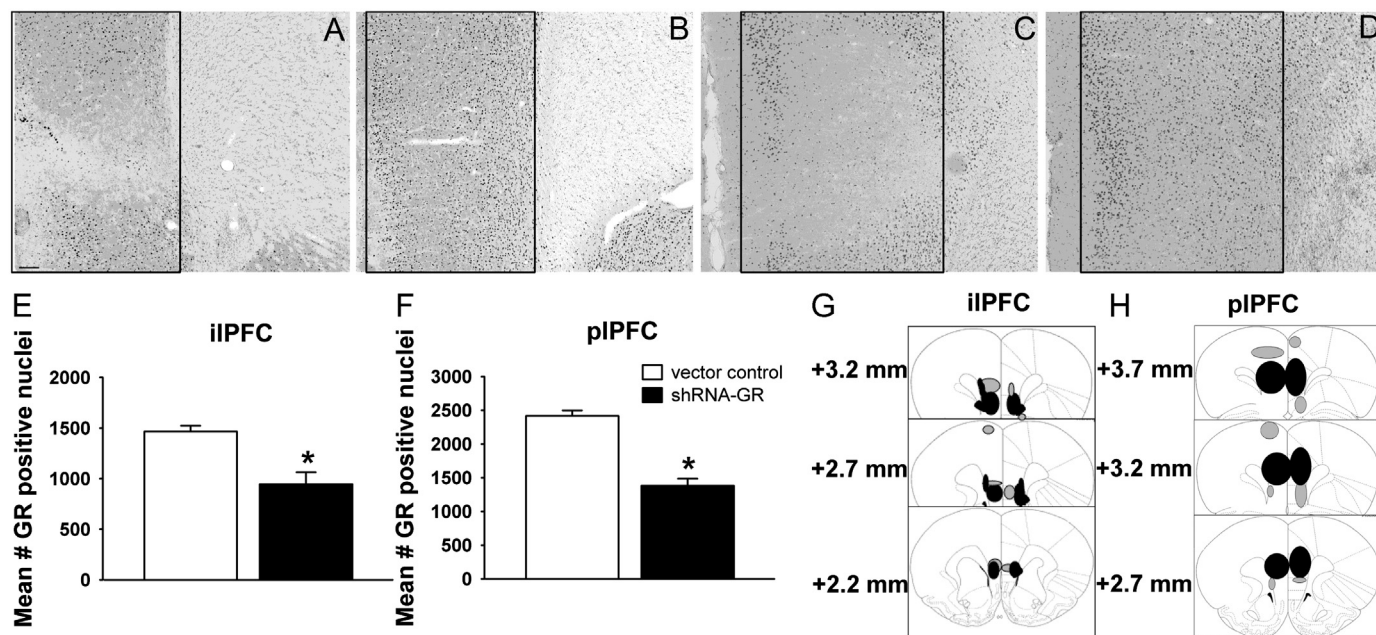


Figure 2. Selective decreases in glucocorticoid receptor (GR) immunoreactive neurons in the infralimbic prefrontal cortex (iIPFC) and prelimbic prefrontal cortex (pIPFC) after short hairpin RNA targeting the GR (shRNA-GR) microinjection. Representative GR-immunolabeled sections from vector control-microinjected animals in (B) the iIPFC and the (D) pIPFC and shRNA-GR-microinjected animals in (A) the iIPFC and (C) the pIPFC (representative areas of quantification are outlined in panels A–D). (E) Quantified GR expression from vector control-microinjected animals ($n = 10$) and shRNA-GR-microinjected animals ($n = 5$) in the iIPFC and (F) vector control-microinjected animals ($n = 6$) and shRNA-GR-microinjected animals ($n = 6$) in the pIPFC. The GR immunoreactivity was significantly reduced in animals that received shRNA-GR relative to vector control-microinjected animals ($p < .05$). (G) Extent of GR knockdown in the iIPFC of all shRNA-GR-microinjected animals that were considered “hits” ($n = 10$) or (H) in the pIPFC of all shRNA-GR-microinjected animals that were considered “hits” ($n = 16$) (reprinted from Paxinos and Watson [16] with permission from Elsevier, copyright 1998). Green fluorescent protein (GFP) expression throughout the pIPFC in each animal was traced onto stereotaxic images and compiled into one visual representation. Black circles indicate where GFP expression was most prominent ($n \geq 4$ in the iIPFC and $n \geq 8$ in the pIPFC), whereas gray circles represent areas where GFP was less prominent in animals that received shRNA-GR and were considered “hits” ($n \leq 3$ in the iIPFC or $n \leq 7$ in the pIPFC). Immunoreactive counts are mean \pm SEM. Scale bar = 100 μ m. * $p < .05$ vs. vector control-microinjected animals.

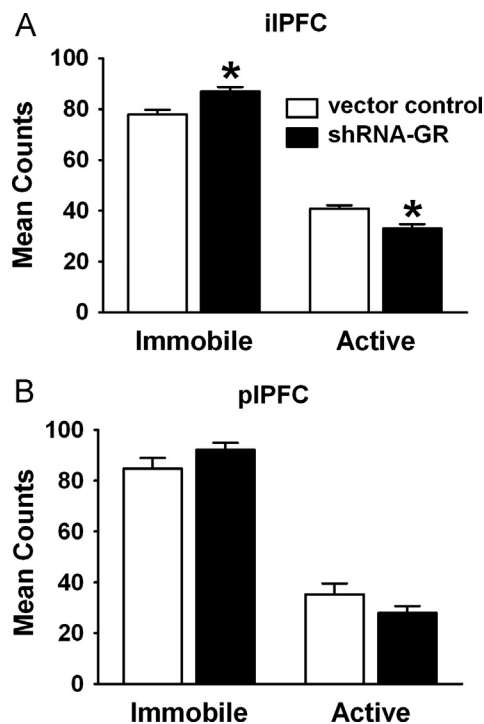


Figure 3. Increased helplessness behavior after GR knockdown in the iIPFC. **(A)** Immobility vs. activity in the modified forced swim test after vector control- or shRNA-GR-microinjection in the iIPFC ($n = 10$ or 5 , respectively) or **(B)** in the pIPFC ($n = 11$ or 8 , respectively). Animals receiving shRNA-GR in the iIPFC but not the pIPFC, exhibited increased immobility in the forced swim test relative to vector controls ($p < .05$). Data are mean \pm SEM. * $p < .05$ vs. vector control-microinjected animals. Abbreviations as in Figure 2.

centrifuged at $3000\times g$ for 15 min at 4°C , and plasma was stored at -20°C until time of radioimmunoassays (RIAs).

Tissue Collection. Animals were given an overdose of sodium pentobarbital and transcardially perfused with .9% saline followed by 4% sodium phosphate-buffered paraformaldehyde. Brains were postfixed in 4% sodium phosphate-buffered paraformaldehyde for 24 hours, then stored in 30% sucrose in diethylpyrocarbonate-treated water at 4°C . Brains were sectioned on a microtome in $30\text{-}\mu\text{m}$ coronal sections (Leica, Buffalo Grove, Illinois). Thymus and adrenal glands were dissected and weighed.

Immunohistochemistry. Sections were immunolabeled with primary antibodies against GR (M-20) (1:1000; Santa Cruz Biotech, Santa Cruz, California), neuronal nuclei (NeuN) (1:200; Millipore, Billerica, Massachusetts), MR (ID-5) (1:200 and 1:500; provided by Dr. Elise Gomez-Sanchez from University of Mississippi, Jackson, Mississippi) (21), or glial fibrillary acidic protein (1:2000; Dako, Carpinteria, California) with standard immunohistochemical procedures. For additional detail, see Supplement 1.

RIA. Plasma ACTH was determined by a RIA that used a specific antiserum (1:120,000 dilution; donated by Dr. William England University of Minnesota, Minneapolis, Minnesota) with ^{125}I ACTH (Amersham Biosciences, Piscataway, New Jersey) as labeled tracer. All samples were run in duplicate (when sample was sufficient) in the same assay. Plasma corticosterone levels were measured with an ^{125}I RIA kit (MP Biomedicals, Solon, Ohio). All samples were run in duplicate and each time point was run in the same assay. For additional detail, see Supplement 1.

Cell Counting. For analysis of GR-, NeuN-, or MR-positive immunoreactive nuclei, digital images of each side of the pIPFC or

iIPFC, as defined by the rat stereotaxic brain atlas of Paxinos and Watson, were captured at $5\times$ or $10\times$ magnification with a Carl Zeiss Imager Z.1 (Carl Zeiss Microimaging, Thornwood, New York). Quantitative analysis of cell counts was performed with the Automatic Measurement Program, Axiovision 4.4 (Carl Zeiss Microimaging). Images were captured on the same day with the same settings, and a uniform threshold was applied to all images in a given brain region.

Statistical Analysis

Data are expressed as mean \pm SEM. Behavioral data, body weight before CVS, and immunoreactive counts were analyzed with one-way analysis of variance (ANOVA). Body weight (after CVS), organ weights, and baseline corticosterone levels were analyzed with a two-way ANOVA (microinjection [vector control or shRNA-GR] \times stress [acute stress or CVS]). Fisher's least significant difference post hoc analyses were conducted. Hormonal data were analyzed with two-way repeated measures ANOVA (microinjection \times time [0, 15, 30, 60 or 120 min]) or three-way repeated measures ANOVA (microinjection \times stress \times time [0, 30, 60 or 120 min]), time being the repeated measure. Fisher's least significant difference was used for a priori planned comparisons across microinjection and stress at each time point. Data were analyzed with GBStat (version 6.5.4) software (Dynamic Microsystems, Silver Spring, Maryland), and statistical significance was set at $p \leq .05$. Where appropriate, behavioral data failing Levene's F , Hartley's F -max, Cochran's C , and Barlett's χ^2 homogeneity of variance tests were log transformed. Outliers were removed as outlined previously (22). Animals with unilateral or no GFP expression or injections outside the iIPFC or pIPFC were excluded ($n = 21$ or 6 , respectively). For simplicity of presentation, results are graphed by acute stress only (No CVS) or chronic stress (CVS), although data were part of the same statistical analysis. Experiments targeting the pIPFC or the iIPFC were conducted separately, and therefore statistical comparisons across experiments were not analyzed.

Results

shRNA Validation

We first performed in vitro studies to identify an shRNA sequence that can specifically knockdown GR expression. We transfected immortalized, GR-expressing hypothalamic 4B cells with several different lentiviral-packaged shRNAs predicted to target GR mRNA (shRNA-GR) (23). As determined by RT qPCR, GR mRNA expression was reduced after transfection with shRNA-GR 469 (99.8% reduction as shown by results from one PCR experiment) (Figure S1 in Supplement 1). We next validated the ability of this shRNA to knockdown expression in vivo. Immunofluorescence analysis revealed reduced GR immunoreactivity at the site of injection in animals that received shRNA-GR, without loss of neuronal viability (NeuN immunolabeling is intact in GFP-positive neurons) (Figure 1A). Reduced GR was observed as a loss of GR immunoreactivity in GFP-positive (i.e., virus-infected) neurons (Figure 1B). No reduction in GR was observed in animals that received a scrambled-sequence control (shRNA-Sc) (Figure 1C), empty vector control (Figure 1D), or vehicle control (data not shown). The MR immunoreactivity was also intact in transduced neurons that lack GR, demonstrating that the shRNA-GR does not downregulate expression of a closely related protein (Figure 1E). Furthermore, shRNA-GR microinjection did not produce recruitment of astrocytes to the region beyond that of a

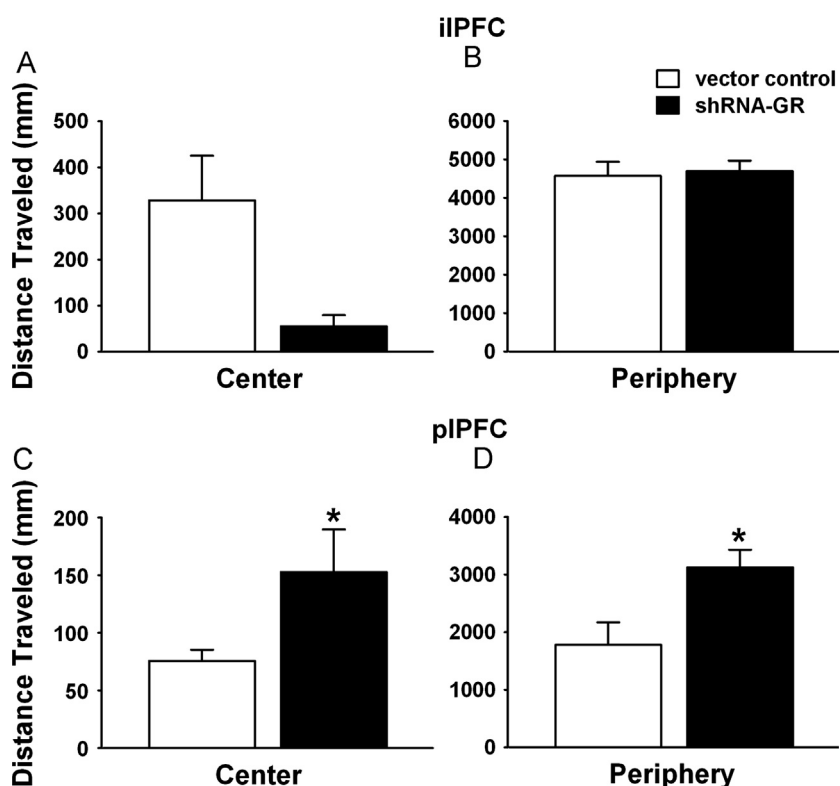


Figure 4. Increased locomotor activity after GR knock-down in the pLPFC relative to vector control-microinjected animals. **(A)** Locomotor activity in the center and **(B)** periphery after vector control- or shRNA-GR microinjections in the iLPFC ($n = 9$ – 10 or 4 , respectively). **(C)** Locomotor activity in the center and **(D)** periphery after microinjections of vector control or shRNA-GR in the pLPFC ($n = 11$ or 7 – 8 , respectively). Animals receiving shRNA-GR in the pLPFC traveled significantly more throughout the center **(C)** and periphery **(D)** than vector controls ($p < .05$). Data are mean \pm SEM. * $p < .05$ vs. vector control-microinjected animals. Abbreviations as in Figure 2.

control injection site. Importantly, astrocytes did not seem to incorporate the shRNA-GR (no co-localization of glial fibrillary acidic protein with GFP), and expression of the astrocytic GR was intact (Figure 1F).

To assess the extent of GR knockdown in the iLPFC and pLPFC, we quantified the number of GR-positive immunoreactive nuclei in the area of injection. The GR expression was selectively knocked down in shRNA-GR-microinjected animals in the iLPFC [$F_{1,13} = 20.33$, $p = .0006$] (Figure 2A,E) and in the pLPFC [$F_{1,10} = 59.48$, $p < .0001$] (Figure 2C,F) relative to vector control-microinjected animals (Figures 2B,D–F), without affecting the number of NeuN [$F_{1,10} = 1.06$, $p = .33$] or MR [$F_{1,11} = .168$, $p = .69$] immunoreactive cells at the site of injection (as quantified in the pLPFC) (Figure S2 in Supplement 1). Knockdown of GR expression is mostly confined to the iLPFC or the pLPFC (with minimal spread to adjacent areas) (Figure 2G,H).

Behavioral Testing

The medial PFC is thought to be an important mediator of depression-like behavior (24,25). To test the effect of GR knockdown on depression-like behavior, we first examined performance in the FST, commonly used as an assay for behavioral helplessness. Animals that received microinjection with shRNA-GR in the iLPFC had significantly increased immobility in the FST compared with animals receiving vector control ($F_{1,13} = 9.67$, $p = .008$), suggestive of a depression-like phenotype (Figure 3A). However, there were no significant differences in scored individual active behaviors (e.g., swimming [$F_{1,13} = .46$, $p = .51$], climbing [$F_{1,14} = .75$, $p = .40$], diving [$F_{1,14} = .23$, $p = .64$], or headshakes [$F_{1,14} = .55$, $p = .47$]). In contrast, knockdown of pLPFC GR did not affect immobility ($F_{1,17} = 1.81$, $p = .20$) or individually scored activities (e.g., swimming [$F_{1,17} = 3.18$, $p = .09$], climbing [$F_{1,17} = .04$, $p = .85$], diving [$F_{1,18} = .39$, $p = .54$], or headshakes [$F_{1,16} = .88$, $p = .36$]) (Table S1 in Supplement 1) in the FST (Figure 3B).

Previous studies indicate that electrolytic lesions of the iLPFC or the pLPFC decrease time spent in the center of the open field (26). Therefore, we tested anxiety-related behavior and locomotion in the open field test, with the same cohorts of animals used in the FST. Microinjection of shRNA-GR in the iLPFC did not precipitate an anxiety-like phenotype (no main effect of microinjection on time spent in the center of the open field) ($F_{1,11} = 1.69$, $p = .22$) (Table S2 in Supplement 1). Furthermore, there were no significant differences in overall locomotor activity ($F_{1,11} = .60$, $p = .46$) or locomotor activity in the center ($F_{1,11} = 2.98$, $p = .11$) (Figure 4A) and the periphery ($F_{1,12} = .04$, $p = .86$) (Figure 4B) relative to vector control-microinjected animals. Similarly, injection of shRNA-GR in the pLPFC was without effect on anxiety-related open field behavior ($F_{1,16} = .23$, $p = .64$) (Table S2 in Supplement 1) but did cause a substantial increase in total locomotor activity ($F_{1,17} = 7.59$, $p = .01$), distance traveled in the center ($F_{1,16} = 103.15$, $p < .0001$) (Figure 4C), and in the periphery ($F_{1,17} = 6.42$, $p = .02$) (Figure 4D) of the open field and a significant increase in rearing ($F_{1,17} = 7.69$, $p = .01$) (Table S2 in Supplement 1).

Body/Organ Weights

Rats were exposed to a 2-week CVS regimen to test the impact of GR signaling in prefrontal regions on physiological reactivity to prolonged adversity. As documented previously, attenuated weight gain, adrenal hypertrophy, and thymic involution are consistent attributes of chronically stressed rats (27). There was no main effect of microinjection on body weight gained before CVS in either the iLPFC ($F_{1,30} = .13$, $p = .72$) or pLPFC ($F_{1,35} = .99$, $p = .33$) of shRNA-GR-microinjected animals, indicating that PFC GR knockdown did not affect body weight. Gross somatic effects of chronic stress on adrenal hypertrophy were not affected by GR knockdown in either PFC subregion. However, thymic involution was selectively enhanced in the CVS-iLPFC group, consistent with greater cumulative exposure to glucocorticoids over the stress

regimen (main effect of stress [$F_{1,26} = 6.54, p = .02$]) (Table S3 in Supplement 1).

Hormonal Responses

We next tested the role of iLPFC and pLPFC in control of HPA axis responses to acute restraint stress. Both iLPFC and pLPFC injections of shRNA-GR enhanced acute activation of the HPA axis (increased peak corticosterone release; at 30 or 60 min post-stress time points, respectively) (Figure 5). Enhanced corticosterone release was accompanied by increased ACTH release 15 min after acute restraint (measured in a separate study after pLPFC GR knockdown [$F_{4,40} = 6.80, p = .0003$]) (Figure S3 in Supplement 1). After chronic stress, iLPFC GR knockdown potentiated the corticosterone response to a novel stressor, consistent with hypersensitization of the HPA axis ($F_{3,75} = 3.42, p = .02$). Knockdown of GR in the pLPFC did not affect the post-CVS peak HPA axis response to a novel stressor, and in fact corticosterone levels were significantly lower 60 min after restraint ($F_{3,102} = 4.43, p = .006$). Together, the data suggest differential roles of the iLPFC and pLPFC in chronic stress processing.

Finally, we determined the impact of pLPFC GR and iLPFC GR knockdown on baseline levels of stress hormones in unstressed animals or animals exposed to chronic stress. The shRNA-GR microinjection in the pLPFC increased baseline levels of

corticosterone in chronically stressed animals, reflected in a microinjection \times stress interaction ($F_{1,31} = 6.05, p = .02$) (Figure 6B), suggesting selective involvement in control of basal glucocorticoid homeostasis under chronic stress.

Discussion

Our study indicates that glucocorticoid control of stress responsiveness and emotional reactivity is mediated by distinct prefrontal cortical mechanisms, with the iLPFC particularly important for mediating chronic stress adaptation and emotional reactivity to stress. Loss of infralimbic GR caused increased helplessness behavior and hormonal hypersensitivity to chronic stress, consistent with a role in integrating glucocorticoid signals into appropriate behavioral and physiological responses to prolonged challenge. Importantly, in human, area 25 (iLPFC homolog) is linked to depression, a disease that is characterized by helplessness behavior and reduced central sensitivity to glucocorticoids (1,13,14). The current data suggest that local glucocorticoid signaling in this confined prefrontal locus might be critical for appropriate control of mood.

The prelimbic cortex seems to play a very different role in chronic stress adaptation. Like the iLPFC, the pLPFC participates in control of HPA responses to an acute stressful event. Our data

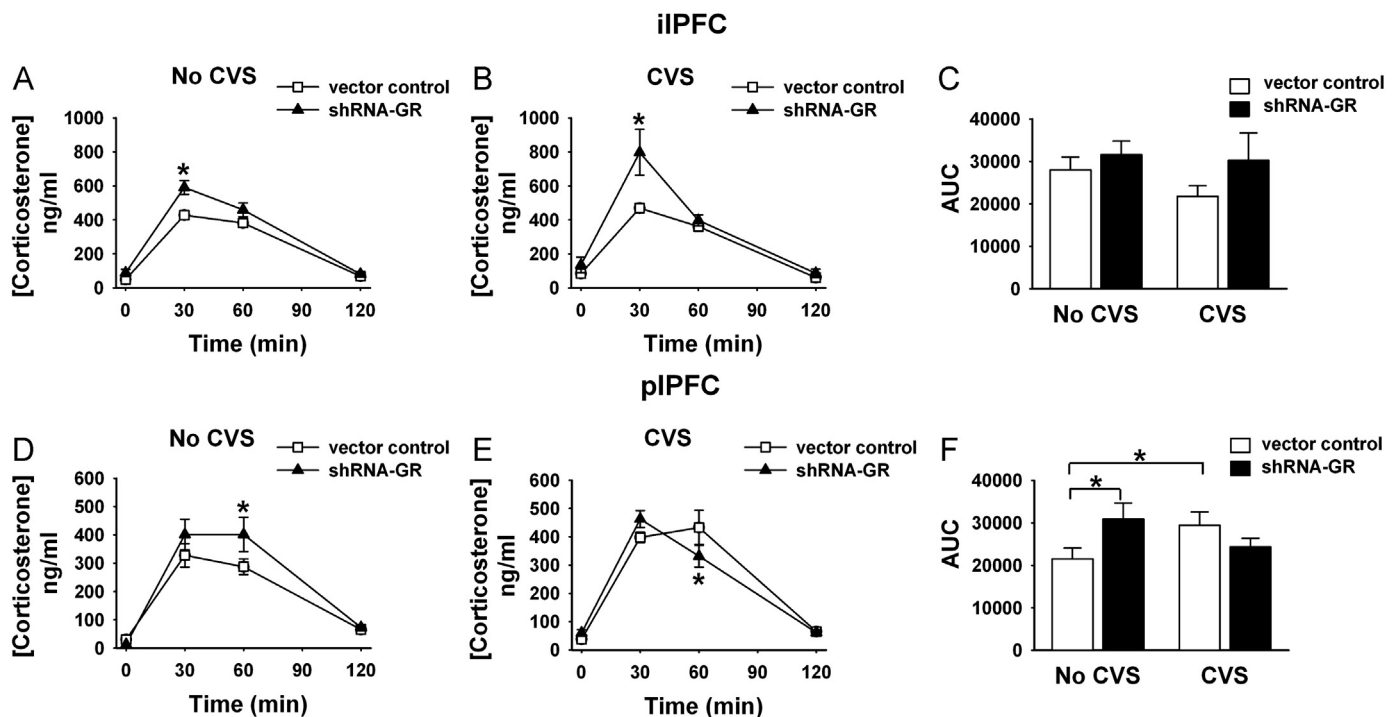


Figure 5. Differential impact of GR knockdown in the iLPFC vs. pLPFC on hypothalamic-pituitary-adrenal axis reactivity after acute and chronic stress. (A) Corticosterone responses after acute novel restraint in unstressed (no chronic variable stress [CVS]) and (B) CVS animals that received microinjections of vector control ($n = 9-11$ /unstressed or stressed group) or shRNA-GR ($n = 5$ /unstressed or stressed group) in the iLPFC. (C) Integrated area under the curve (AUC) for corticosterone responses (not including baseline values) after vector control-microinjections ($n = 9-11$ /group) or shRNA-GR ($n = 5$ /group) in the iLPFC. (D) Corticosterone responses after acute novel restraint in unstressed (No CVS) and (E) CVS animals that received microinjections of vector control ($n = 10-11$ /unstressed or stress group, respectively) or shRNA-GR ($n = 7-9$ /unstressed or stressed group) in the pLPFC. (F) Integrated AUC for corticosterone responses (not including baseline values) after vector control-microinjections ($n = 10-11$ /group) or shRNA-GR ($n = 7-9$ /group) or in the pLPFC. After iLPFC microinjection of shRNA-GR, acute stress caused a significant elevation in corticosterone at 30 min compared with vector controls, an effect that is exacerbated in chronically stressed animals relative to acutely stressed shRNA-GR-microinjected animals and controls ($p < .05$). After microinjection of shRNA-GR in the pLPFC, animals acutely stressed in the absence of CVS have significantly elevated corticosterone levels at 60 min compared with vector controls, whereas chronically stressed animals have significantly lower corticosterone responses at 60 min compared with vector control-microinjected animals ($p < .05$). Data are mean \pm SEM. * $p < .05$ vs. vector control-microinjected animals or between groups indicated by the brackets. Abbreviations as in Figure 2.

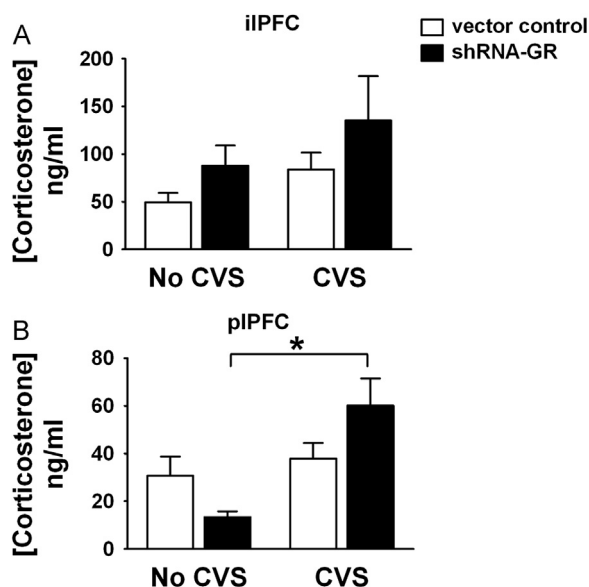


Figure 6. The GR knockdown in the pIPFC but not iIPFC of chronically stressed animals significantly increased baseline corticosterone levels. **(A)** Baseline corticosterone levels in unstressed and chronically stressed animals receiving vector control ($n = 10$ or 11 /group, respectively) or shRNA-GR ($n = 5$ /group) in the iIPFC and **(B)** in unstressed and chronically stressed animals receiving vector-control ($n = 10$ or 11 , respectively) or shRNA-GR ($n = 7$ or 9 , respectively) in the pIPFC. Baseline corticosterone levels were significantly different in chronically stressed animals receiving shRNA-GR in the pIPFC only (relative to acutely stressed animals that received shRNA-GR) ($p < .05$). Data are mean \pm SEM. * $p < .05$ vs. acutely stressed shRNA-GR-microinjected animals or between groups indicated by the brackets. Abbreviations as in Figures 2 and 5.

suggest that, unlike the iIPFC, the pIPFC GR does not seem to be involved in either regulation of HPA axis reactivity to chronic stress or control of initial emotional responses to stressors. However, pIPFC GR knockdown induces a significant increment in basal morning corticosterone release during chronic stress, suggesting that this region participates in setting the basal tone of the HPA axis under chronic stress. Thus, the prelimbic and infralimbic cortices, anatomical neighbors in the prefrontal region, are both involved in processing glucocorticoid information with regard to prolonged adversity but play very different roles in regulating behavioral and physiological responses.

Previous studies have used multiple techniques to assess the role of the PFC in regulation of the HPA axis in rats, including ibotenic lesions, acute activation, and corticosterone implants (6,7,28–31). The shRNA-mediated technique offers many advantages, because it allows for anatomical and molecular specificity through long-term knockdown confined to the region of shRNA expression. The shRNA-GR is packaged in a lentiviral construct and is therefore useful for targeting nondividing cells (i.e., neurons) without eliciting an immune response (32). Our data indicate that lentiviral delivery of shRNA-GR can effectively reduce GR immunoreactivity in virally transduced neurons locally in the region of the iIPFC or the pIPFC, affording the ability to use this method to query the role of the GR in defined neural populations. Our data suggest that the lentiviral knockdown spares medial PFC astrocytes, which play a role in depression-like behavior (33). Additional studies are required to determine whether the glial GR is also involved in modulation of mood.

In patients with MDD, area 25 is hyperactive, and deep brain stimulation ameliorates depressive symptoms in patients with treatment-resistant MDD (34). In rodent, the infralimbic cortex

projects to regions implicated in visceral/autonomic control (e.g., the nucleus of the solitary tract, lateral septum, bed nucleus of the stria terminalis, central amygdaloid nucleus, and posterior hypothalamus) (2), consistent with a role in mediating physical and emotional responses to chronic drive. Moreover, glucocorticoids are thought to inhibit limbic neuronal responses by reducing neural excitability and retracting dendritic trees (35–37). We hypothesize that the loss of a GR-mediated “brake” in the infralimbic cortex might allow prolonged activation of downstream targets, thereby promoting aberrant physical and emotional responses, as observed in the FST.

Despite its anatomic proximity to the infralimbic cortex, the prelimbic cortex has a markedly different efferent output, sending heavy projections to regions involved in limbic/cognitive processing, such as the nucleus accumbens, BLA, and raphe nuclei (2). These sites have polysynaptic input to stress regulatory systems, such as the paraventricular nucleus, and might buffer the impact of altered pIPFC GR signaling on HPA axis drive. For example, chronic stress causes morphological and functional neuroadaptation in regions such as the BLA, which might be sufficient to diminish the overall impact of pIPFC knockdown on HPA stress reactivity (38). The effect of pIPFC on basal morning corticosterone might either reflect modulation of processes relating to stress habituation (e.g., by the paraventricular thalamus) or energetic demands associated with stress-induced hyperactivity, as observed in the open field test (39,40).

Our work is consistent with dedicated roles of the GR within specific neurocircuits. Not surprisingly, total forebrain GR deletion (in mouse) has features in common with the lentiviral knockdown approach. Forebrain GR deletion (encompassing neocortex, hippocampus, and basolateral/cortical amygdala) produces glucocorticoid stress hyperresponsiveness, in response to acute restraint stress, characteristic of pIPFC and iIPFC GR knockdown in the rat (41). However, forebrain GR knockout does not cause chronic stress sensitization (like the pIPFC but not iIPFC knockdown). The latter data suggest that GR signaling in other forebrain regions (such as the hippocampus) might negate or magnify effects of local changes in the prefrontal cortex.

The present study indicates a distinct role of iIPFC versus pIPFC GR in acute versus chronic stress regulation. Human genetic studies link the GR to depression or posttraumatic stress disorder, either directly or via interactions with other proteins (e.g., FK506 binding protein 5, a chaperone protein that modulates GR function) (42). Given the overriding importance of area 25 (human iIPFC-equivalent) in affective disease states, our data suggest that stress-related GR signaling might be uniquely important in regulating resistance or resilience, with loss of function driving pathological behavioral and endocrine reactivity to situational adversity.

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