

Glucocorticoids desensitize hypothalamic CRH neurons to norepinephrine and somatic stress activation via rapid nitrosylation-dependent regulation of $\alpha 1$ adrenoreceptor trafficking

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eLife Assessment

This manuscript describes work that is **fundamental** to our understanding of the mechanism of how stress regulates the noradrenergic system and the CRH system. Using a combination of ex vivo physiology and in vitro methods, the study provides **compelling** evidence as to the signaling mechanisms of how glucocorticoids impact adrenergic GPCRs in CRH cells via receptor trafficking. While the ex vivo work is specific to the hypothalamus, the mechanisms here could be extrapolated and investigated in other brain regions that may have similar signaling regulation.

<https://doi.org/10.7554/eLife.102783.1.sa4>

Abstract

Summary

Noradrenergic afferents to hypothalamic corticotropin releasing hormone (CRH) neurons provide a major excitatory drive for somatic stress activation of the hypothalamic-pituitary-adrenal (HPA) axis. We showed that glucocorticoids rapidly desensitize CRH neurons to norepinephrine and suppress inflammation-induced HPA activation via a glucocorticoid receptor- and endocytosis-dependent mechanism. Here, we show that $\alpha 1$ adrenoreceptor (AR $\alpha 1$) trafficking is regulated by convergent glucocorticoid and nitric oxide synthase signaling mechanisms. Live-cell imaging of AR $\alpha 1$ b-eGFP-expressing hypothalamic cells revealed rapid corticosterone-stimulated redistribution of internalized AR $\alpha 1$ from rapid recycling endosomes to late endosomes and lysosomes via a nitrosylation-regulated

mechanism. Proximity assay demonstrated interaction of glucocorticoid receptors with AR α 1b and β -arrestin, and showed corticosterone blockade of norepinephrine-stimulated AR α 1b/ β -arrestin interaction, which may prevent AR α 1b from entering the rapid recycling endosomal pathway. These findings demonstrate a rapid glucocorticoid regulation of G protein-coupled receptor trafficking and provide a molecular mechanism for rapid glucocorticoid desensitization of noradrenergic signaling in CRH neurons.

Introduction

Increased circulating glucocorticoids induced by activation of the hypothalamic-pituitary-adrenal (HPA) axis play an important role in the metabolic, immunoregulatory, and cognitive effects of stress ^{1,2}. Rapid glucocorticoid-mediated negative feedback regulation of the corticotropin releasing hormone (CRH) neurons in the hypothalamic paraventricular nucleus (PVN) contributes to the termination of the HPA response ³. While the mechanism responsible for negative feedback regulation of the HPA axis has been classically attributed to transcriptional regulation by glucocorticoid receptors ^{4,5}, several studies have shown that glucocorticoids rapidly inhibit PVN CRH neurons (Di et al., 2003; Tasker et al., 2006) and pituitary corticotrophs ⁹ to mediate fast negative feedback (Evanson et al., 2010). Noradrenergic afferents to the CRH neurons provide a primary excitatory drive for the somatic stress activation of the HPA axis ¹¹ via activation of α 1 adrenergic receptors (AR α 1) ^{12–15}. We recently reported that glucocorticoid rapidly desensitizes PVN CRH neurons to AR α 1 activation by norepinephrine (NE), which suppresses the HPA response to somatic stress, but not psychological stress, due to the NE dependence of somatic stress activation of the HPA axis ^{11,16}. A main source of noradrenergic input to PVN CRH neurons is the brainstem nucleus of the solitary tract (NTS) ¹⁷. The NTS is activated by inputs from the vagal nerve and relays to the forebrain somatic interoceptive stress signals such as organ damage, metabolic challenge, and immune activation ¹⁸. Stimulation of an immune response by systemic lipopolysaccharide (LPS), for example, activates the HPA axis by way of the NTS noradrenergic system ^{19,20}. Our recent findings suggest that prior priming by acute stress exposure attenuates subsequent LPS activation of the HPA axis via a rapid glucocorticoid-induced desensitization of the CRH neurons to NE activation of AR α 1 ²¹, but the cellular and molecular mechanisms of this desensitization are not known.

The stress desensitization of PVN CRH neurons to NE is dependent on the ligand-induced internalization of AR α 1 ²¹. AR α 1 is internalized by clathrin-mediated endocytosis and is trafficked back to the membrane via recycling endosomes, a process that involves multiple trafficking proteins and that establishes a steady state of surface receptors and NE sensitivity ²². Initial steps of endocytosis following ligand binding and G protein activation include receptor phosphorylation, recruitment of β -arrestin to the phosphorylated receptor, and engagement of clathrin and dynamin to execute endocytosis. β -arrestin and the internalized AR α 1 undergo post-translational modifications, including nitrosylation and ubiquitination, which regulate trafficking through the cell's endosomal pathways ^{23,24}. Here, we tested the cellular mechanism by which glucocorticoids desensitize NE signaling. We found that corticosterone (CORT) desensitizes CRH neurons to NE in a process involving nitric oxide synthesis (NOS)-dependent protein S-nitrosylation that rapidly depletes the cells of surface adrenoreceptors by re-routing the AR α 1 out of the rapid membrane recycling endosomal pathway and into late endosomes and lysosomes.

Results

Corticosterone causes a rapid, long-lasting desensitization of AR α 1

Our previous study (Chen et al., 2019) showed that in *ex vivo* brain slices, PVN CRH neurons in the male mouse respond to NE with an increase in excitatory postsynaptic currents (EPSCs) that is mediated by postsynaptic AR α 1 activation and stimulation of retrograde glial-neuronal circuits by dendritic volume transmission (see diagram in **Fig. 1A**). A follow-up study showed that the PVN CRH neuron excitatory response to NE is suppressed by 5–10 min of preincubation of brain slices from unstressed mice in CORT (2 μ M) (**Fig. 1B**) and by prior restraint stress-induced endogenous glucocorticoid exposure *in vivo* (**Fig. 1C**). The rapid glucocorticoid-induced desensitization of CRH neurons to NE was blocked by a dynamin inhibitor, indicating that it is dependent on endocytosis (**Fig. 1C**).

Here, we tested the duration of the stress-induced desensitization of CRH neurons to NE by allowing the mice to recover from a 30-min restraint stress in their home cage for 4 h and 18 h prior to sacrifice and brain slice preparation. In animals allowed to recover from acute restraint for 4 h, the PVN CRH neurons remained insensitive to NE (100 μ M), whereas the EPSC response to NE was partially restored 18 h after the restraint stress (**Fig. 1D**). This indicated that the stress-induced desensitization of PVN CRH neurons to activation by NE takes longer than 18 h to fully reverse and suggests a surprisingly long-lasting diminished CRH neuron sensitivity to NE and to adrenoreceptor activation of the HPA axis following exposure to acute stress. Thus, acute glucocorticoid exposure causes a long-lasting desensitization of CRH neurons to NE via a mechanism with rapid onset, which is dependent on dynamin-dependent endocytosis of the AR α 1 (**Fig. 1E**).

AR α 1b has been reported to be the dominant α 1 adrenoreceptor subtype in CRH neurons, and we previously reported evidence of AR α 1b internalization in fixed N42 cell cultures. Here, we used live-cell imaging to follow the internalization and trafficking of the AR α 1b through the endosomal pathway. We used the N42 immortalized hypothalamic cell line, which we transiently transfected with an pEGFP-AR α 1b construct and imaged live by confocal microscopy to visualize AR α 1b receptor trafficking. A low NE concentration (1 μ M) caused AR α 1b internalization, which was seen by a decrease in membrane pEGFP-AR α 1b and an increase in cytosolic pEGFP-AR α 1b fluorescence, which reached a peak at $\sim$ 30 min, a time course that coincided approximately with that of the NE-induced increase in sEPSC frequency (see **Fig. 1**). Following 30 min of NE application, CORT (2 μ M) was co-applied with NE, which induced a rapid increase in the AR α 1b internalization over the NE-induced steady-state internalization. The onset of the CORT effect occurred within minutes and peaked within 10 min of its introduction into the chamber, which was consistent with a rapid CORT-induced desensitization of CRH neurons to NE. The combination of NE and CORT caused significantly more internalization than NE alone. The bulk of the NE- and CORT-induced increases in cytosolic AR α 1b appeared to be directed mainly to a single “hot-spot” in the cells (arrows in **Fig. 2A**), which we targeted for quantitation and used to map the trafficking of the AR α 1b receptor in the endocytic pathway with endosomal markers. The dynamic change in AR α 1b distribution within the cell during NE and NE+CORT application can be followed in real time in the supplemental video 1.

Cort redirects AR α 1b trafficking from rapid recycling endosomes to late endosomes

Receptor-mediated endocytosis forms early endosomes, which either cycle to the plasma membrane as recycling endosomes to replenish surface receptors or develop into late endosomes and are routed out of the recycling pathways. Specific Rab GTPases regulate distinct stages of endosomal trafficking. By co-transfecting the N42 cells with pEGFP-AR α 1b and constructs for different dsRed-tagged Rab proteins, we were able to use Förster resonance energy transfer (FRET) analysis to detect with 10-nm resolution the colocalization of the AR α 1b with the specific markers

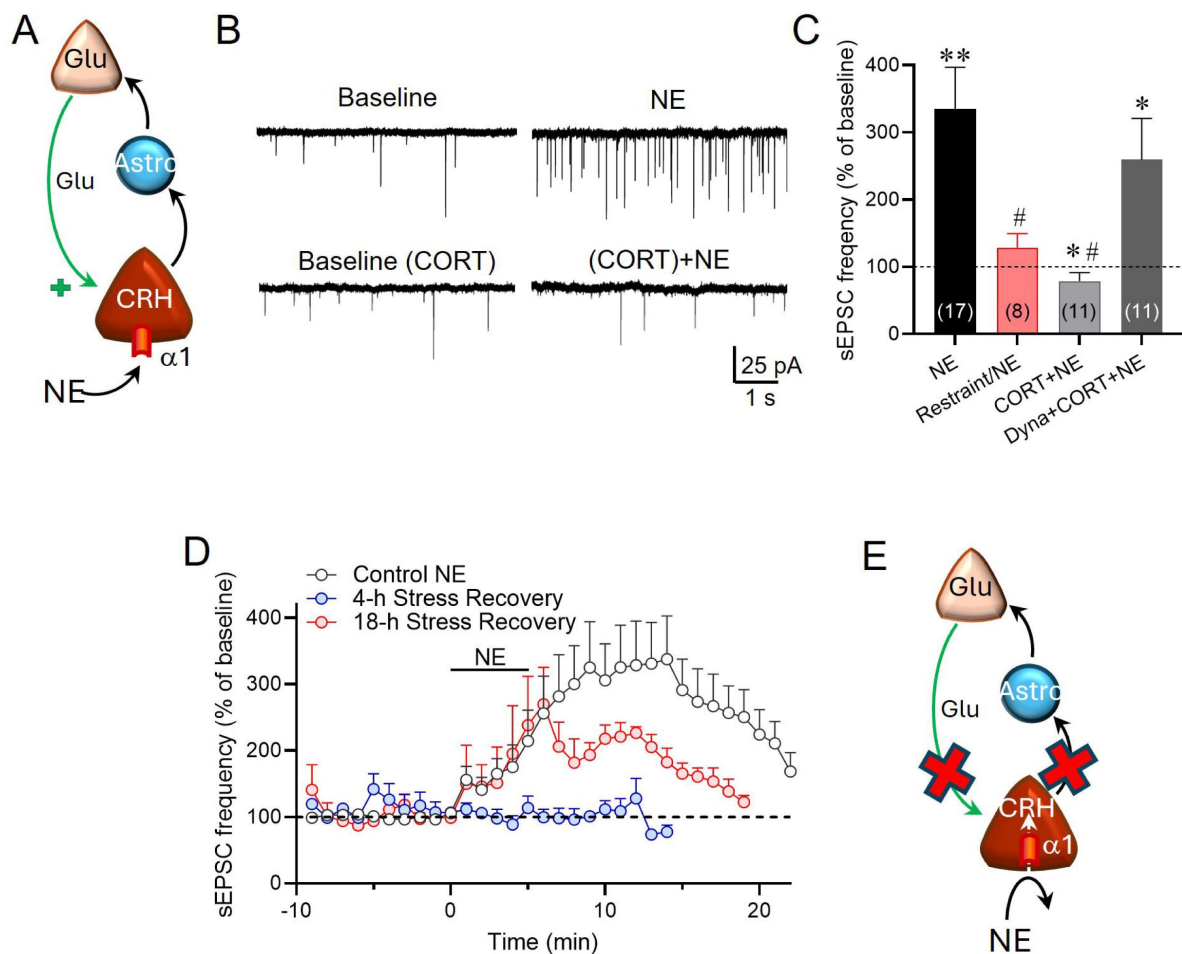


Figure 1.

Long-lasting glucocorticoid desensitization of NE AR α 1-induced increase in sEPSC frequency in CRH neurons via a dynamin-dependent mechanism.

A. Model of retrograde neuronal-glial signaling response triggered in CRH neurons by NE, based on Chen et al., 2019. **B.** Representative traces from our previous study of the sEPSC response in PVN CRH neurons to NE (100 μ M) without pretreatment (Baseline/NE) (upper trace) and in the presence of 1 μ M corticosterone (Baseline (CORT)/NE (CORT), 10 min) (lower trace). **C.** Summary graph of the stress (Restraint) and CORT (CORT) suppression of the NE-induced increase in sEPSC frequency and the rescue of the NE response by blocking endocytosis with the dynamin inhibitor dynasore (80 μ M, 10 min) (Dyna+CORT+NE). **D.** Time histogram of the NE-induced increase in sEPSC frequency in CRH neurons over time in slices after 4 h or 18 h recovery from a 30-min acute restraint stress (4-h and 8-h Stress Recovery), compared to the NE response following no prior stress (Control NE, taken from [21](#)). Complete desensitization of the NE response was seen after a recovery period of 4 h following restraint and the response was partially restored after 18 h of recovery. Panels B and C and the Control NE graph in D were modified from [21](#) with permission.

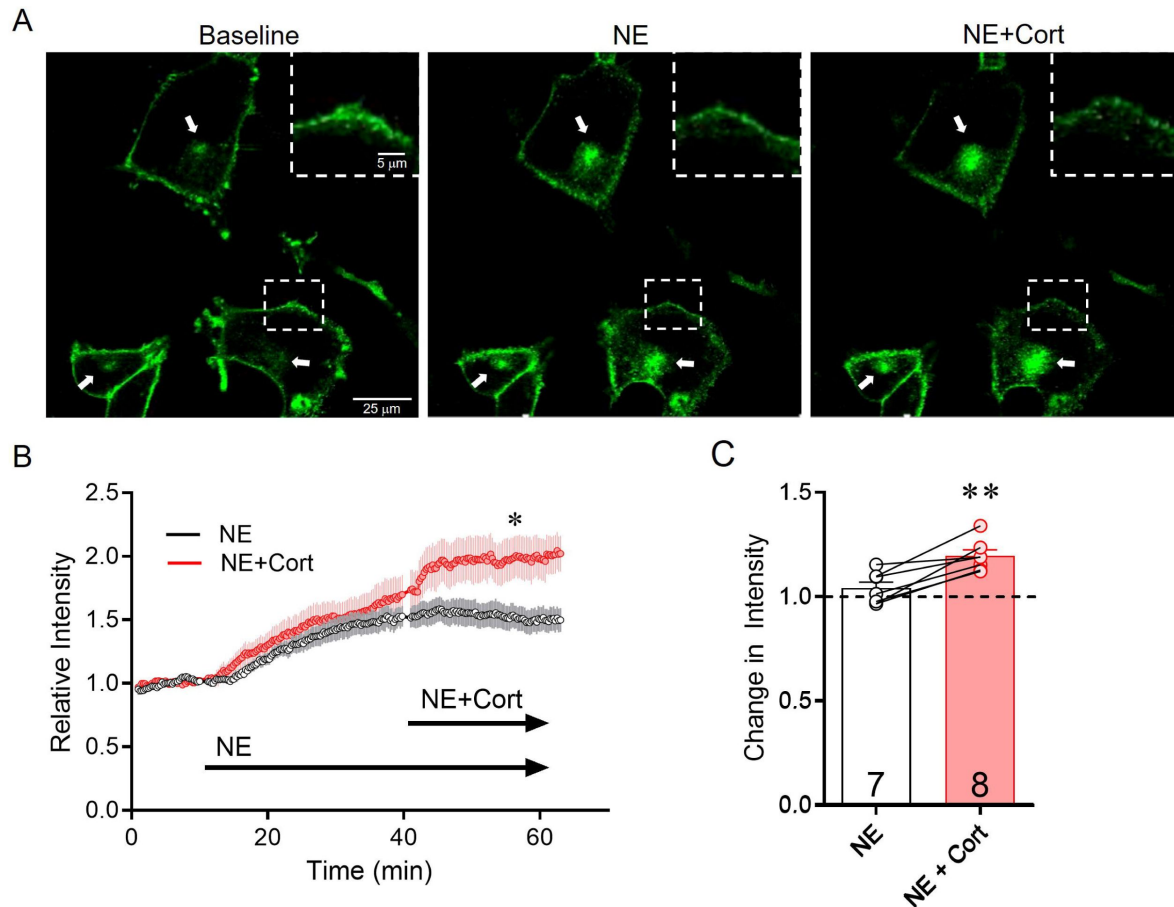


Figure 2.

Cort causes a rapid increase in cytosolic ARα1b accumulation in a distinct subcellular "hot spot".

A. In N42 cells transiently transfected with ARα1b-eGFP, ARα1b is mostly localized to the membrane at baseline, but redistributes from the membrane to internal 'hotspots' (arrows) after 30 min of NE application (NE) and further after 20 min of application of NE and CORT (NE+Cort). Membrane region in the dashed box is enlarged in the upper insets. **B.** Time course of increase in intracellular ARα1b-eGFP. Following plateau of the increase in the fluorescence signal by NE (Vehicle), the co-application of CORT caused a rapid, additional increase in cytosolic ARα1b that plateaued within 10 min (*, $p<0.05$, t-test vs NE). **C.** Relative increase in cytosolic intensity of ARα1b-eGFP compared to NE alone (NE) following the addition of CORT (NE+Cort). (**, $p<0.01$, paired t-test vs. NE).

of different endosomal compartments. We tested for AR α 1b association with the early endosomal marker Rab5, the late endosomal marker Rab 7, the rapid recycling endosomal marker Rab4a, and the slow recycling endosomal marker Rab 11 ²⁸.

Cells treated with NE (1 μ M) were first analyzed for an increase in AR α 1b FRET with dsRed-tagged Rab5, a marker of newly internalized cargo being transported to the early endosome. Norepinephrine caused the expected increase in AR α 1b localization in the early endosome, serving as a control for ligand-mediated endocytosis of the adrenoreceptor. However, subsequent CORT application failed to increase colocalization of AR α 1b with Rab5 compared to that caused by NE alone (**Fig. 3A**). This indicated that the CORT-induced increase in AR α 1b intracellular concentration was mediated by an accumulation of AR α 1b in the cytosol following standard ligand-dependent internalization and not by a CORT-induced increase in AR α 1b endocytosis. We next tested whether CORT caused an increase in the receptor trafficking to the late endosome using AR α 1b FRET with the late endosomal marker Rab7. Norepinephrine alone caused an increase in the AR α 1b-Rab7 FRET signal, and subsequent CORT treatment increased the FRET signal compared to that seen with NE alone (**Fig. 3B**). This indicated that CORT caused an increase in the NE-induced AR α 1b trafficking to the late endosome and suggests that the CORT-induced desensitization is not associated with an increase in the amount of AR α 1b undergoing internalization, but rather that it causes an increase in the receptor trafficking into the late endosomal pathway. A possible explanation for the accumulation of AR α 1b in the late endosome and loss of sensitivity to NE could be a decrease in the trafficking of the receptor back to the plasma membrane.

We tested this using FRET analysis with the rapid recycling endosomal marker Rab4a and with the slow recycling endosomal marker Rab11. Norepinephrine did not cause a change in the AR α 1b-Rab4a FRET signal, and the addition of CORT decreased the AR α 1b-Rab4a signal compared to NE alone (**Fig. 3C**), indicating reduced AR α 1b association with Rab4a and suggesting a reduction in trafficking of the AR α 1b through the rapid recycling endosome. Finally, we tested for changes in the AR α 1b interaction with the slow recycling endosomal marker Rab11. Neither NE nor NE+CORT had any effect on the AR α 1b-Rab 11 FRET signal (**Fig. 3D**), suggesting that neither treatment increases the slow recycling of AR α 1b to the membrane within the timeframe of the desensitization to NE. Together, these data indicate that CORT influences AR α 1b trafficking by redirecting the receptor away from rapidly recycling endosomes and into late endosomes. This would be expected to deplete the membrane of adrenoreceptors and provides, therefore, a molecular substrate for the glucocorticoid-induced desensitization to NE observed in *ex vivo* recordings.

Cort causes AR α 1b trafficking to the lysosome

To determine whether CORT causes AR α 1b to travel further down the endocytic pathway from late endosomes to lysosomes, we tested for AR α 1b trafficking to the lysosome by imaging the co-localization of pEGFP-AR α 1b with the lysosomal marker LAMP1 conjugated to red fluorescent protein (LAMP1-RFP). Application of NE (1 μ M) caused an increase in AR α 1b-LAMP1 co-staining (**Fig. 4A-C**), indicating an increase in the co-localization of the two proteins. After 30 min of NE application, CORT co-application (2 μ M) further increased the AR α 1b-LAMP1 co-staining (**Fig. 4A-C**), which revealed a CORT facilitation of AR α 1b trafficking to the lysosome for degradation. However, a large proportion of the AR α 1b was not co-stained with the LAMP1 marker, including the CORT-induced AR α 1b “hot spot”, resulting in a low Pearson’s correlation of AR α 1b and LAMP1 localization (maximum Pearson’s coefficient $\sim$ 0.5) (**Fig. 4B**), which suggests that, on this time scale ($\sim$ 50 min), most of the AR α 1b receptor was not trafficked to the lysosome for degradation, but rather was sequestered within the cell.

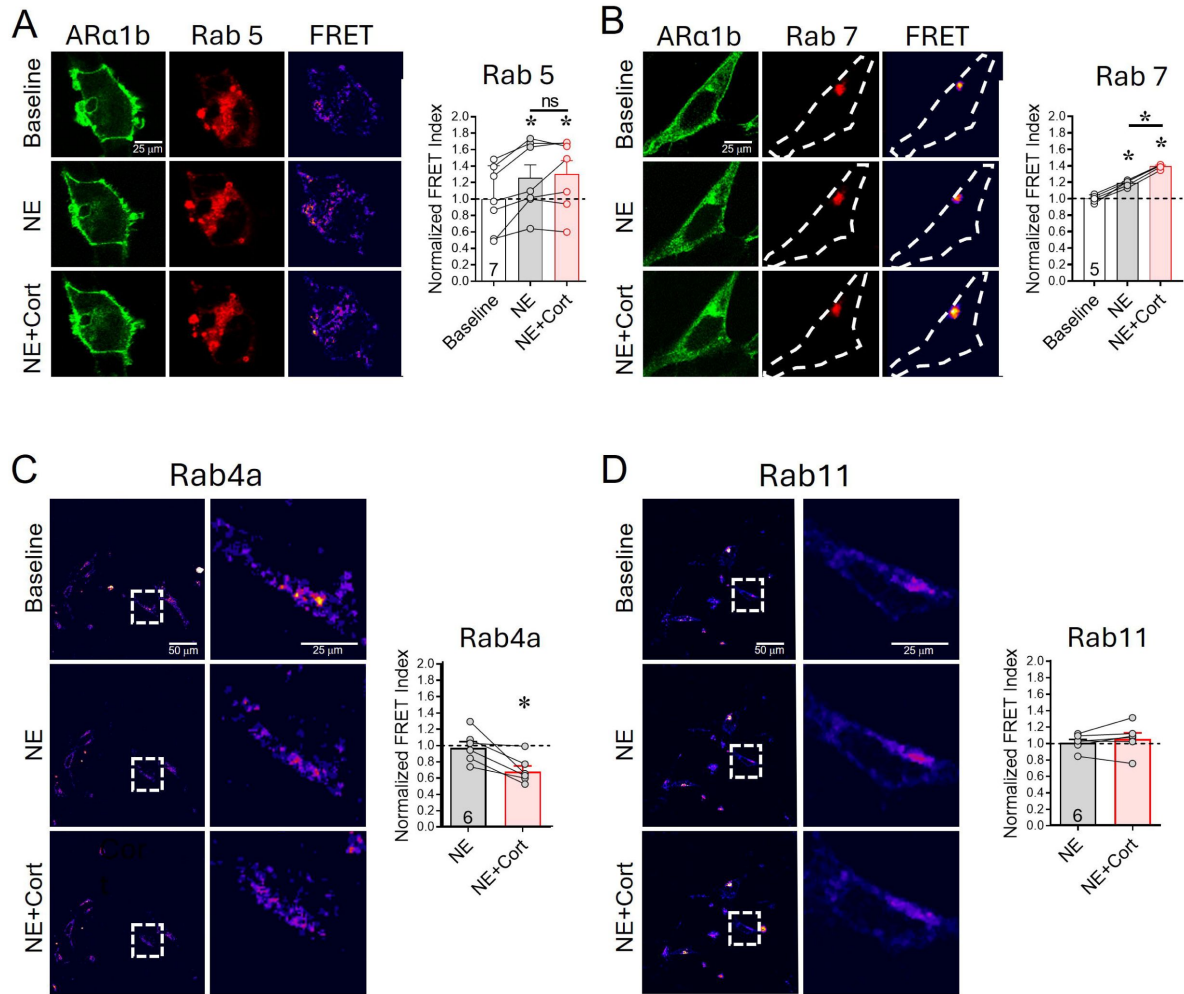


Figure 3.

Cort redirects ARA1b trafficking from the rapid recycling endosome to the late endosome.

A. N42 cells expressing ARA1b-eGFP (ARA1b) and Rab5-dsRed (Rab 5) and images of calculated FRET ratios (FRET) after treatments with NE and NE+Cort. **Right:** Mean (+/- SEM) of FRET signal in cells treated with NE (1 μ M) and NE+Cort showing increased FRET between ARA1b and Rab5 induced by NE, but no further increase after subsequent Cort application (2 μ M) (repeated measures ANOVA, Tukey's post-hoc, * $p < 0.05$ vs baseline; n.s., $p = 0.7423$). **B.** N42 cells expressing ARA1b-eGFP (ARA1b) and Rab7-dsRed (Rab 7) and images of calculated FRET ratios (FRET) after treatments with NE and NE+Cort. **Right:** Mean (+/- SEM) of FRET signal in cells treated with NE (1 μ M) and NE+Cort, showing increased FRET between ARA1b and Rab7 with NE, and a further increase in FRET ratio with subsequent Cort application (2 μ M) (repeated measures ANOVA, Tukey's post-hoc, * $p < 0.05$). **C.** Images of calculated FRET ratios of N42 cells expressing ARA1b-GFP and Rab4a-dsRed after NE and Cort treatments. Images in the right column show the areas in the squares at higher magnification. **Right:** Mean (+/- SEM) of FRET signal showing no significant change in ARA1b/Rab4a interaction after treatment with NE (1 μ M) and a decrease in FRET ratio with subsequent Cort application (2 μ M) (repeated measures ANOVA, * $p < 0.05$ vs NE). **D.** Images of calculated FRET ratios of N42 cells expressing ARA1b-GFP and Rab11-dsRed after NE and NE+Cort treatments. Images in the right column show the areas in the squares at higher magnification. **Right:** Mean (+/- SEM) of FRET signal in NE and NE+Cort, normalized to baseline. There were no significant changes to the ARA1b-Rab 11 FRET ratio after application of either NE (1 μ M) or NE+Cort (2 μ M) (repeated measures ANOVA, $p = 0.2895$).

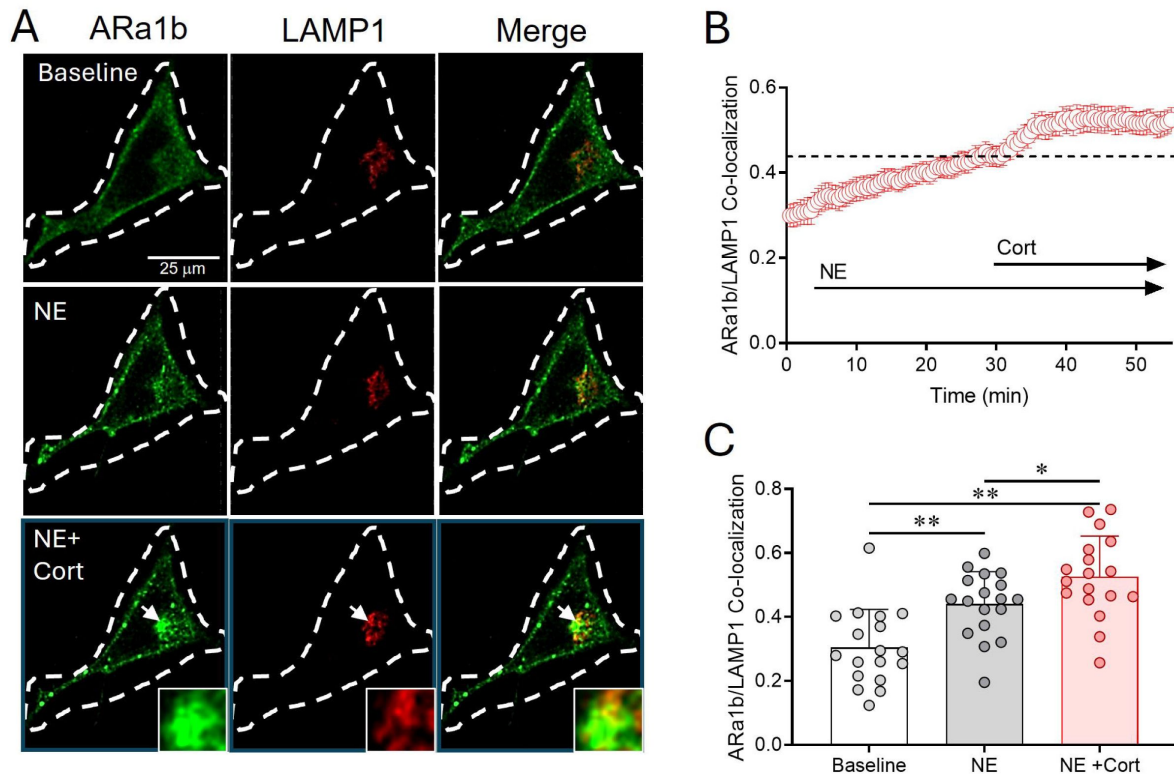


Figure 4.

ARA1b trafficking to lysosomes.

A. ARA1b-eGFP and LAMP1-RFP were co-transfected into N42 cells, and the cells were treated with NE for 30 min, followed by NE + CORT for 20 min. Arrow: CORT-facilitated internalized ARA1b co-localized with LAMP1, a lysosomal marker. **B.** Time lapse quantification of co-localization by Pearson's coefficient. **C.** Co-localization averages of baseline and the last 3 minutes of NE and NE +CORT treatment (Mean $\pm$ SEM, ANOVA, Tukey's post-hoc, * $p < 0.05$, ** $p < 0.01$).

Blocking AR α 1 reverses Cort-induced AR α 1b trafficking

We found previously that CORT alone without prior NE application has no effect on AR α 1b internalization in N42 cells and that inhibiting AR α 1 with prazosin during CORT application in hypothalamic slices blocked the glucocorticoid-induced desensitization of CRH neurons to NE ²¹, suggesting that glucocorticoid-induced desensitization to NE requires ligand (NE)-induced AR α 1 internalization. Here, we tested whether N42 cells in which CORT had already caused increased AR α 1b cytosolic accumulation would recover after blocking AR α 1 with prazosin. As described above (see **Fig. 1** and **Fig. 2A-C**), 1 μ M NE caused AR α 1b accumulation in the cytosol that was further increased by CORT (2 μ M). The addition of the AR α 1 antagonist prazosin (1 μ M) caused the internalized AR α 1b to return to near baseline levels within 10 min (**Fig. 5A, B**). These data suggest that blocking AR α 1 reverses glucocorticoid-induced desensitization to NE.

Nitrosylation dependence of the CRH neuron response to norepinephrine

Since GPCR internalization and trafficking are dependent on S-nitrosylation of proteins associated with receptor trafficking, including β -arrestin ²⁴, we next tested whether the CORT control of AR α 1 trafficking is regulated by nitric oxide synthase (NOS) activity. We first tested this with whole-cell recordings of the NE response in CRH neurons in *ex vivo* hypothalamic slices. Blocking NOS activity with the broad-spectrum NOS inhibitor N(ω)-nitro-L-arginine methyl ester (L-NAME, 50 μ M) inhibited the NE-induced increase in sEPSC frequency (**Fig. 6A, B**), which was similar to the glucocorticoid inhibition of the NE-induced increase in sEPSC frequency in CRH neurons (see **Fig. 1**). Since NOS induces both the production of the gaseous messenger NO and post-translational S-nitrosylation of target proteins, we tested for these two NOS-dependent mechanisms pharmacologically. We first blocked the NO receptor, soluble guanylyl cyclase, with 1H-[1,2,4]oxadiazolo[4,3-a]quinoxalin-1-one (ODQ) (100 μ M), but this failed to inhibit the NE-induced increase in sEPSC frequency in the CRH neurons (**Fig. 6A, B**). This confirmed our previous finding that NE does not induce an NO retrograde signal at glutamate synapses on the CRH neurons ¹⁵. We next tested the effect of the S-nitrosylation inhibitor N-ethylmaleimide (NEM) (50 μ M) on the NE response. NEM completely blocked the NE-induced increase in sEPSC frequency in the CRH neurons (**Fig. 6A, B**). These findings suggested that the NOS dependence of the NE activation of excitatory inputs to the CRH neurons is due to S-nitrosylation of target proteins in the CRH neurons, and not to the release of NO as a transmitter.

Several proteins involved in GPCR trafficking are regulated by nitrosylation, including G protein-coupled receptor kinase 2 (GRK2), β -arrestin, and clathrin ^{24,29}. We first tested for the nitrosylation dependence of intracellular trafficking of the AR α 1b with live-cell imaging in N42 cells. We found that N42 cells express endothelial NOS and inducible NOS, but not neuronal NOS (Supplemental Fig. 1). Co-application of the NOS inhibitor L-NAME (50 μ M) with NE (1 μ M) caused an increase in the AR α 1b internalization over that induced by NE application alone (**Fig. 6C, D**), suggesting that inhibiting S-nitrosylation enhanced ligand-dependent internalization/trafficking of the receptor. Co-application of CORT (2 μ M) with L-NAME failed to increase the AR α 1b cytosolic concentration compared to L-NAME alone (**Fig. 6C, D**), indicating that CORT did not cause any additional intracellular AR α 1b accumulation. This suggested, therefore, that blocking S-nitrosylation caused an occlusion of the CORT effect, revealing a convergence of the two signaling mechanisms and opposite effects of CORT (i.e., increased internal AR α 1b accumulation) and S-nitrosylation (i.e., decreased internal AR α 1b accumulation).

We next tested for changes in S-nitrosylation in the N42 cells with a biochemical approach using Western blot analysis and a proximity ligation assay following the application of a modified biotin-switch procedure. The biotin-switch procedure consists of cleaving the NO groups on cysteine

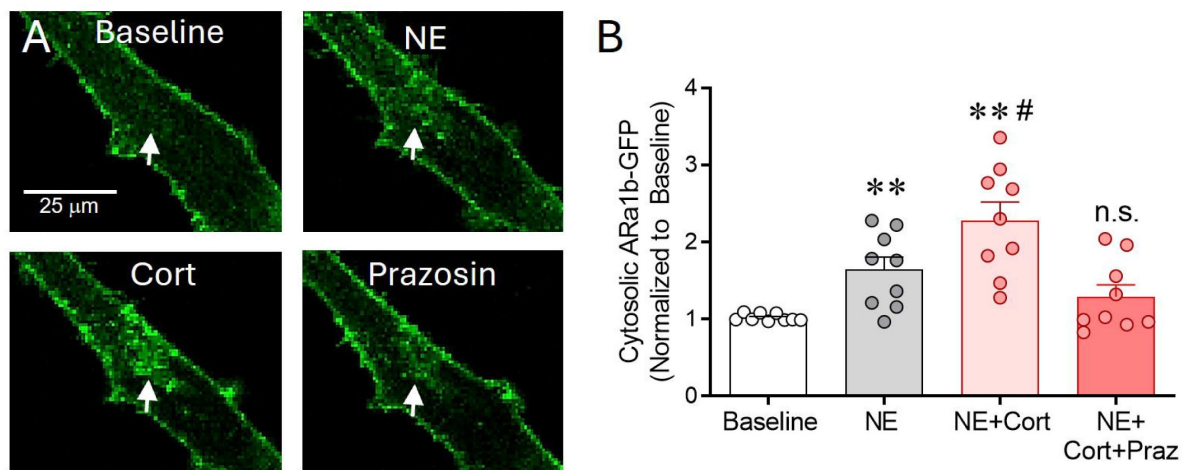


Figure 5.

Reversal of NE- and Cort-induced ARα1b trafficking.

A) ARα1b-EGFP-transfected N42 cells were treated with NE for 30 min, followed by the addition of Cort for 20 min and finally the addition of prazosin for 30 min. Arrows: ARα1b-EGFP hot spot. **B)** Normalized cytosolic fluorescence (ANOVA, Tukey's post-hoc, * $p < 0.05$ vs. baseline, # $p < 0.05$ vs. NE). The ARα1b hot spot returned to near baseline after 30 min prazosin co-application (n.s., $p = 0.35$ vs baseline).

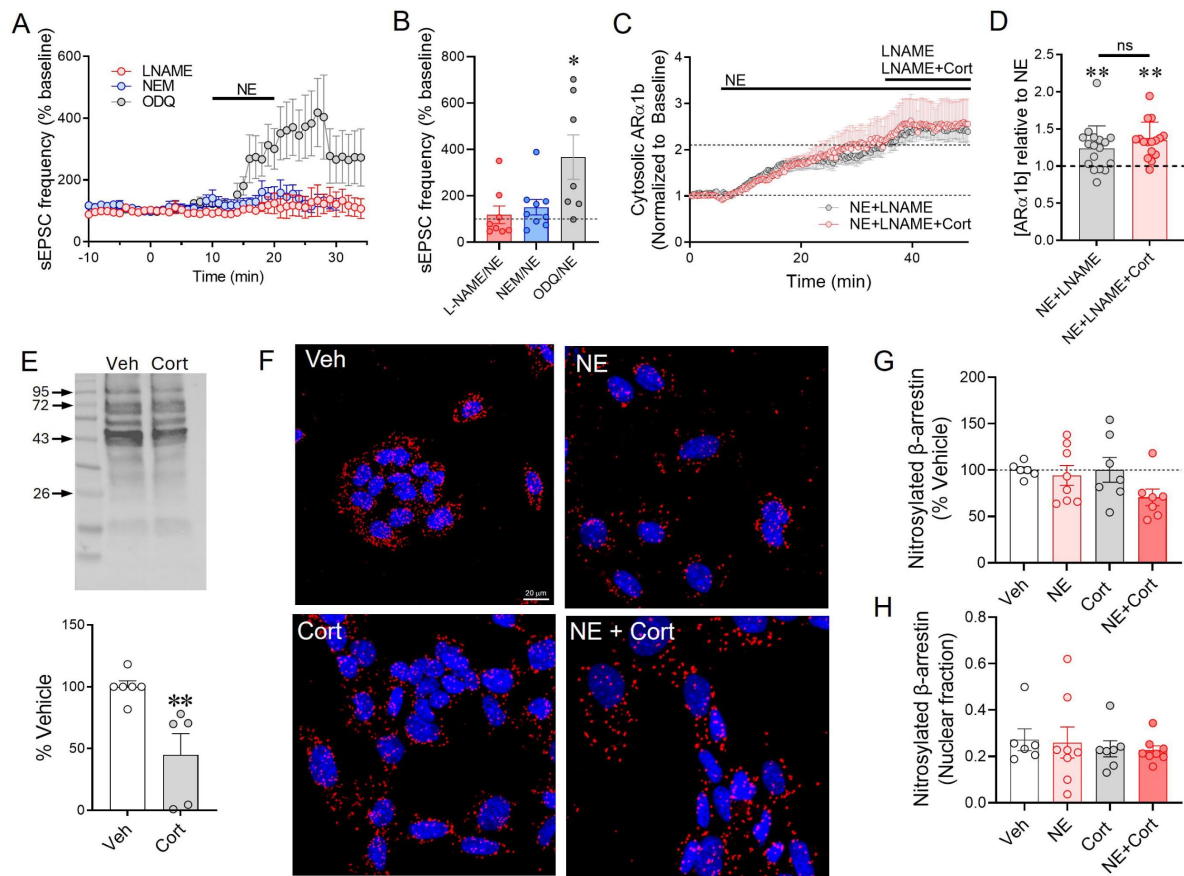


Figure 6.

CORT regulation of protein S-nitrosylation.

A. Nitrosylation dependence of the NE-induced increase in sEPSCs in PVN CRH neurons. The NE-induced increase in sEPSC frequency was blocked by the nitric oxide synthase (NOS) inhibitor L-NAME and the S-nitrosylation inhibitor NEM, but not by the soluble guanylyl cyclase inhibitor ODQ. **B.** Summary graph of the effects of L-NAME, NEM, and ODQ on the NE-induced increase in sEPSC frequency. **C.** Changes over time in the cytosolic intensity of ARα1b-eGFP in N42 cells after treatment with NE and L-NAME (NE+L-NAME) and NE, L-NAME and CORT (NE+L-NAME+Cort). **D.** Summary graph of the changes in ARα1b-EGFP fluorescence intensity indicative of internalization in NE+L-NAME and NE+L-NAME+Cort, normalized to the baseline fluorescence intensity (** $p < 0.01$ compared to baseline). Cort and L-NAME co-application was not significantly different from L-NAME alone ($p = 0.2009$, Student's t -test). Cytosolic ARα1b measurements in L-NAME and L-NAME+Cort were taken at 40-45 min and compared to measurements in NE alone at 30-35 min. **E.** Western blot analysis of whole-cell lysate (cytosolic fraction) from N42 cells treated with vehicle or 2 μ M Cort followed by TMT switch assay for total nitrosylated protein. Cort-treated cells showed a significant decrease in total nitrosylated protein [$t(9) = 3.368$, $p = 0.0083$, Student's t -test, $n = 5-6$]. **F.** Images of β-arrestin1/2 nitrosylation in N42 cells treated with NE, Cort, or NE+Cort for 20 min. The biotin-switch assay was performed, followed by proximity ligation assay (PLA) with antibodies to biotin and β-arrestin1/2. Maximum intensity projections of z-stacks. Red dots = PLA signal representing nitrosylated β-arrestin1/2; blue = DAPI. **G.** Nitrosylation of β-arrestin1/2 by NE, Cort, and NE+Cort relative to vehicle. There were no significant drug effects [One-Way ANOVA, $F(3, 24) = 1.861$; $p = 0.1632$, $n = 6-8$ fields from 4 separate assays]. **H.** None of the drug treatments altered the nuclear localization of nitrosylated β-arrestin1/2 [One-Way ANOVA, $F(3, 25) = 0.2101$; $p = 0.8885$, $n = 6-8$ fields from 4 separate assays].

residues of nitrosylated proteins and labeling these thiols with biotin or another tag such as a tandem mass tag (TMT) ³⁰. Western blot of TMT-tagged proteins in the cytosolic fraction of N42 cells showed a significant decrease in total S-nitrosylation with a 20-min application of CORT (2 μ M) (**Fig. 6E**). This was likely due to activation of the glucocorticoid receptor because we found a similar decrease in total S-nitrosylation in N42 cells with the synthetic glucocorticoid dexamethasone (1 μ M) (data not shown).

Western blot analysis revealed a strong nitrosylation signal at a molecular weight of about 50 kD, the approximate molecular weight of β -arrestin. Since β -arrestin has been shown to be regulated by S-nitrosylation ^{24,31}, we next performed a histological analysis of β -arrestin1/2 S-nitrosylation to test whether the glucocorticoid-induced reduction in total S-nitrosylation is due to changing the nitrosylation state of β -arrestin1/2. We combined the biotin-switch nitrosylation assay with a proximity ligation assay using a biotin antibody and a β -arrestin1/2 antibody to detect nitrosylated β -arrestin. Proximity ligation assay reveals protein-protein interactions with a 40-nm resolution ³². Cells were analyzed for the total number of biotin- β -arrestin interactions (i.e., nitrosylation association with β -arrestin) and for nuclear versus cytosolic localization of the interactions. Treatments with NE (1 μ M), CORT (2 μ M), and NE + CORT all failed to produce a statistically significant change in the PLA signal with respect to vehicle, suggesting they did not affect the nitrosylation of β -arrestin (**Fig. 6F, G**). The NE + CORT treatment caused a small decrease in the PLA signal, but this did not reach statistical significance ($p = 0.1470$, Dunnett's multiple comparisons test). Similarly, the treatments failed to change the nuclear fraction of nitrosylated β -arrestin (**Fig. 6H**). These findings suggest that the nitrosylation dependence of the NE and Cort effects on AR α 1b trafficking is not mediated by the regulation of β -arrestin1/2 S-nitrosylation and that nitrosylated β -arrestin1/2 is not trafficked to the nucleus in response to NE and/or Cort.

Ubiquitination is critical for endosomal processing and cellular trafficking. If glucocorticoid changes the internal trafficking of AR α 1b, then it may do so by changing the ubiquitination state of β -arrestin ³³. We tested for changes in the ubiquitination of β -arrestin in response to NE and CORT treatment (20 min). Both NE (1 μ M) and CORT (2 μ M) alone increased the ubiquitination of immunoprecipitated β -arrestin1/2 approximately two-fold (Supplemental Fig. 2). The increases in ubiquitinated β -arrestin1/2 caused by CORT and NE were additive when the cells were co-treated with the two drugs (Supplemental Fig. 2). Since the ubiquitination pattern of β -arrestin has been shown to affect trafficking of internalized cargo ³⁴, these data suggest that NE and CORT may alter AR α 1b receptor trafficking by increasing β -arrestin ubiquitination.

Glucocorticoid receptor interaction with the α 1 adrenoreceptor

To interrogate the mechanism by which NE and CORT exert cooperative effects on AR α 1b trafficking and β -arrestin ubiquitination, we tested for a direct interaction between AR α 1b and the glucocorticoid receptor (GR) with the proximity ligation assay. A human AR α 1b construct with a myc-DDK tag (hAR α 1b-myc-DDK) or an AR α 1b-eGFP construct was transiently transfected into N42 cells. Proximity ligation was achieved by using either a mouse anti-myc antibody or a mouse anti-GFP antibody to target AR α 1b and a rabbit anti-glucocorticoid receptor antibody (M-20, Santa Cruz) to target GR. Individual cells were analyzed for the number of protein-protein interactions and for nuclear versus cytosolic localization of the interacting proteins. A basal level of interaction between AR α 1b and GR was observed in vehicle-treated cells (**Fig. 7A**). Norepinephrine alone had no effect on AR α 1b-GR interaction, whereas CORT treatment caused a small but significant decrease in total AR α 1b-GR interaction compared to vehicle and NE treatment, which was unchanged by the combination of CORT + NE (**Fig. 7A, B**). We next analyzed the nuclear translocation of the AR α 1b-GR complex after treatment with NE and/or CORT. Norepinephrine alone had no effect on the nuclear fraction of the AR α 1b-GR complex, but CORT caused a significant increase in the nuclear AR α 1b-GR PLA signal compared to vehicle and to NE. The increase in nuclear localization of the AR α 1b-GR complex caused by CORT was reversed by co-application of NE (**Fig. 7A, C**). This suggested that NE activation of AR α 1b prevented the CORT-

induced nuclear translocation of the GR-AR α 1b complex. Thus, AR α 1b and GR can be found in a protein complex that is trafficked to the nucleus in response to CORT but not in the presence of NE, when the AR α 1b is presumably mostly bound to its ligand.

β -arrestin interactions with the α 1 adrenoceptor and glucocorticoid receptor

We next examined interactions of β -arrestin1/2 to determine whether the post-translational S-nitrosylation of β -arrestin is accompanied by changes in its interactions with AR α 1b and GR. As expected, NE treatment (1 μ M) of the N42 cells for 20 min caused an increase in AR α 1b- β -arrestin interaction (Fig. 8A, B). Treatment with CORT alone (2 μ M) had no effect on the number of AR α 1b- β -arrestin PLA profiles, but it blocked the NE-induced increase in AR α 1b- β -arrestin complex formation when co-applied with NE (Fig. 8A, B). CORT treatment alone also caused a significant increase in nuclear localization of the AR α 1b/ β -arrestin complex, which was blocked by NE when NE and CORT were co-applied (Fig. 8A, C).

Similar to what has been reported previously³⁵, we found a baseline interaction between GR and β -arrestin1/2. The drug treatments did not affect the total number of GR- β -arrestin1/2 interactions (Fig. 9A, B). CORT application caused the GR- β -arrestin1/2 complex to translocate to the nucleus, and co-application of NE with CORT did not prevent the CORT-induced nuclear translocation of the GR- β -arrestin1/2 complex (Fig. 9A, C).

Discussion

Acute stress desensitizes PVN CRH neurons in male mice to stimulation by noradrenergic afferents via a rapid glucocorticoid signaling mechanism. The glucocorticoid-induced NE desensitization suppresses the HPA response to somatic, but not psychological, stress and is dependent on the NE-mediated endocytosis of α 1 adrenoceptors²¹. Here, we tested for the cellular mechanisms of the α 1 adrenoceptor desensitization in slices from male mice (consistent with our previous studies, Chen et al., 2019; Jiang et al., 2022), and in the N42 hypothalamic cell line, which expresses both CRH²⁶ and a putative membrane receptor that mediates rapid glucocorticoid signaling^{21,36,37}.

Our findings reveal a rapid glucocorticoid regulation of the trafficking of α 1 adrenoceptors that diverts the receptors away from the rapid recycling endosomal pathway and into late endosomes. The glucocorticoid regulation of α 1 receptor trafficking requires internalization of the α 1 receptors by NE, is dependent on the overall nitrosylation state of the cells and involves ubiquitination of β -arrestin. The rapid glucocorticoid-induced re-routing of the α 1 adrenoceptors out of the rapid recycling endosomal pathway would be expected to deplete the cell of surface receptors and to cause the desensitization of CRH neurons to NE following acute stress exposure²¹.

Glucocorticoid secretion after immune activation of the HPA axis is essential for downregulation of the immune response³⁸. Left unchecked, inflammation and hypotension associated with immune system activation can cause problems beyond the immune challenge itself. One example is toxic shock syndrome, which occurs when bacteria-produced toxins cause over-activation of cytokines and inflammatory cells resulting in fever, rash, hypotension, and, in extreme cases, organ failure and death³⁹. Glucocorticoids constrain the immune response, and glucocorticoid therapy is often employed to inhibit the overactive immune system⁴⁰. For example, the cytokine storm occurring in some COVID-19 patients was treated with the synthetic glucocorticoid dexamethasone^{41,42}.

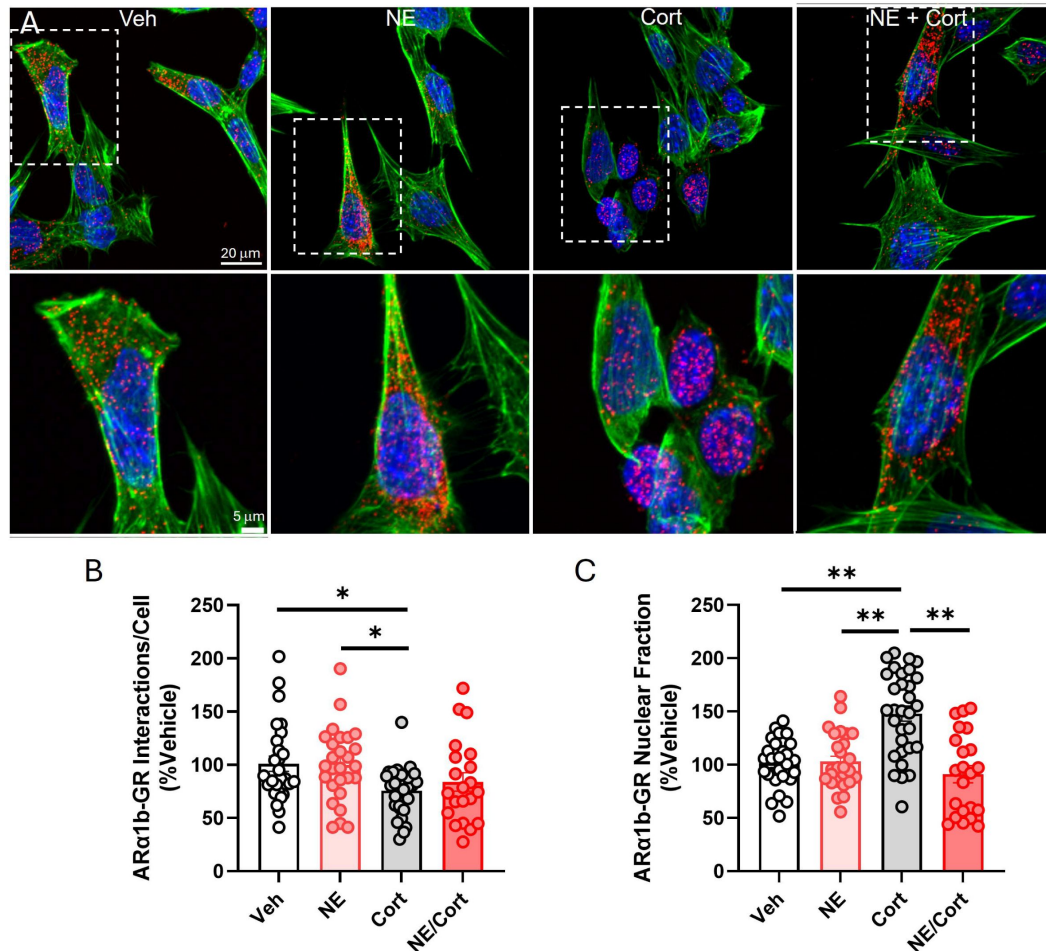


Figure 7.

Interaction between ARα1b and GR.

A. Representative images of ARα1b-myc-DDK-transfected N42 cells after the different treatments and following PLA. Lower images are high-magnification of boxes in upper images. Red = PLA signal, blue = DAPI staining of nuclei, green = phalloidin-FITC staining of F-actin. **B.** Summary graph of total numbers of ARα1b-GR interactions under the different drug conditions. There was an overall decrease in interactions with Cort treatment [F(3, 102) = 3.831 P=0.012]. **C.** Summary graph of the nuclear fraction of ARα1b-GR profiles under the different treatments. Cort treatment significantly increased nuclear localization of the receptor complex relative to vehicle and the other treatment groups [F(3, 103) = 17.30 P<0.0001]. n = 21-31 cells from 3 separate assays. Data were analyzed by one-way ANOVA followed by Tukey's multiple comparisons test. * p<0.05, ** p<0.0001.

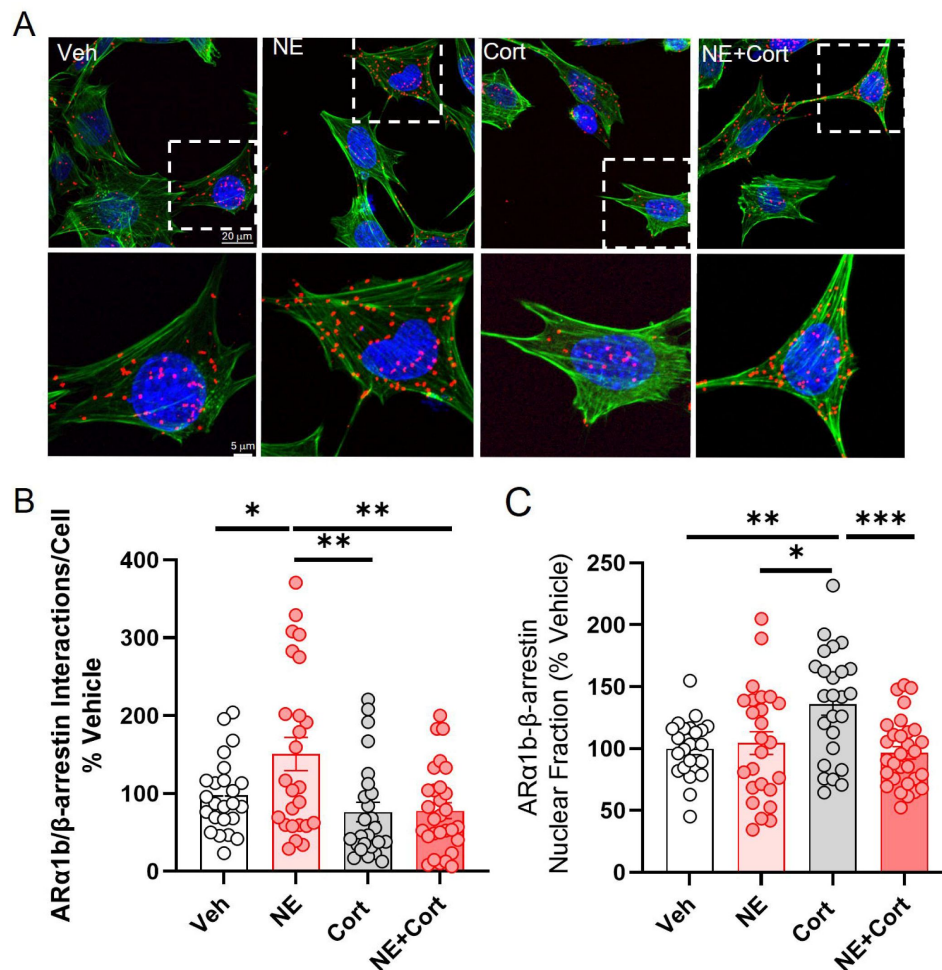


Figure 8.

Interaction between AR α 1b and β -arrestin.

Proximity ligation assay with antibodies to tagged AR α 1b and β -arrestin1/2. N42 cells were treated with drug for 20 min, followed by proximity ligation assay. **A**. Representative images. Blue = DAPI, red = PLA signal, green = phalloidin-FITC stain of F-actin. **B**. Treatments with NE, Cort, and NE+Cort. NE increased the number of AR α 1b- β -arrestin interactions per cell; Cort alone had no effect, but blocked the NE-induced increase in AR α 1b- β -arrestin interactions [F(3,99), $p = 6.157$, $p = 0.0007$]. **C**. Cort treatment increased the nuclear translocation of the AR α 1b- β -arrestin1/2 complex [F(3, 99) = 6.314, $p = 0.0006$], $n = 23$ -30 cells from 3 separate assays. * $p < 0.05$. ** $p < 0.01$, *** $p < 0.001$. Data were analyzed by one-way ANOVA Followed by Tukey's multiple comparisons test. Scale bar = 20 μ m (main), 5 μ m (inset).

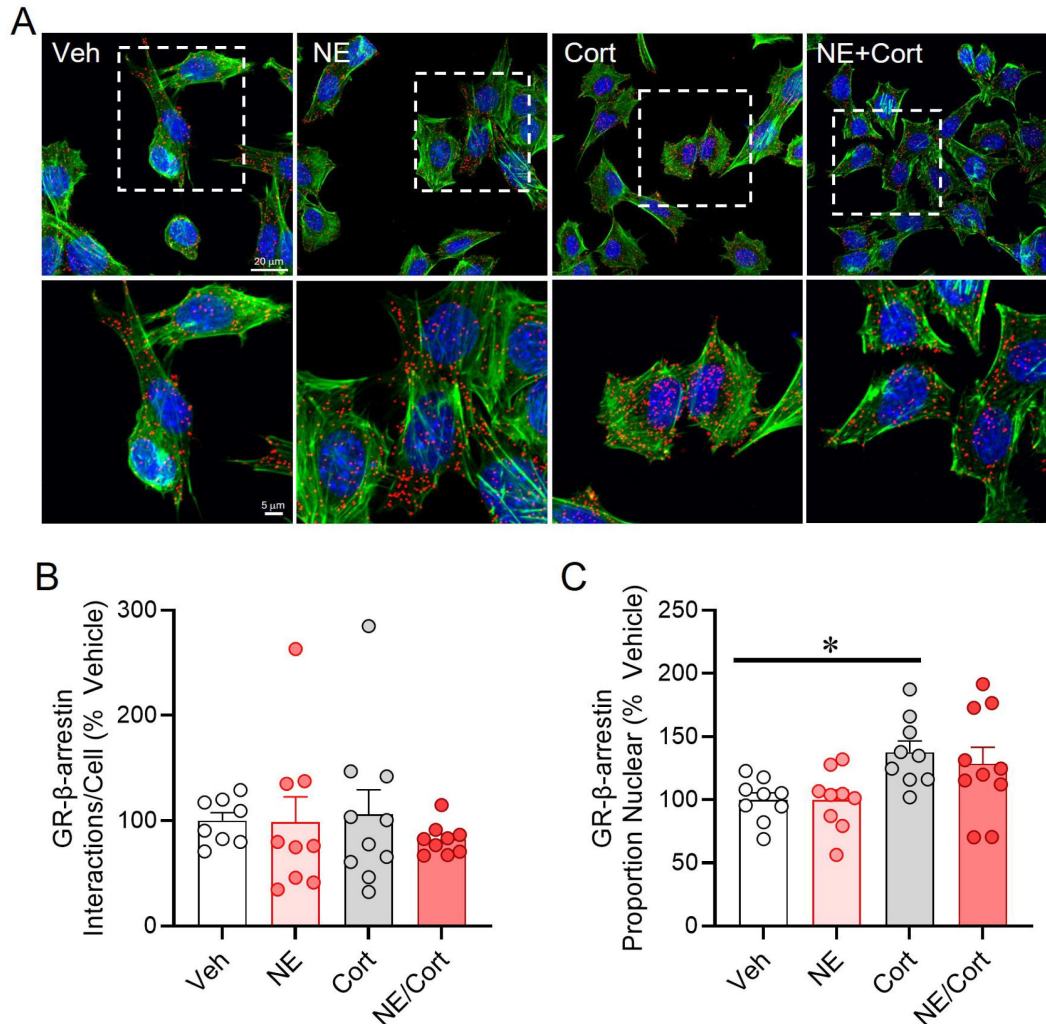


Figure 9.

Interaction between GR and β -arrestin.

Proximity ligation assay with antibodies to GR and β -arrestin1/2. **A.** Representative images of the PLA labeling with vehicle (Veh), NE, Cort, and NE + Cort. Lower images are high magnification of boxes in upper images. Blue = DAPI, red = PLA signal, green = phalloidin-FITC stain of F-actin. **B.** None of the treatments had an effect on the total number of GR- β -arrestin1/2 interactions per cell (one-way ANOVA, $[F(3, 32) = 0.3233, p = 0.8085]$). **C.** Cort treatment increased the nuclear fraction of the GR- β -arrestin1/2 complex (one-way ANOVA followed by Dunnett's multiple comparisons test, $[F(3, 33) = 4.058, p = 0.0147]$, $n = 9$ -10 cells from 3 separate assays, * $p < 0.05$).

Previously we reported that both acute restraint stress and CORT pre-exposure induced a desensitization of CRH neurons to the excitatory effect of NE that lasted for the entire period of brain slice recording (2-6 h) ²¹. However, whether *ex vivo* brain slice preparation prevented the CRH neurons from recovering from their desensitization to NE by placing them in a suspended desensitized state was not known, nor was it clear how long the desensitization to NE lasts following stress. Here we sacrificed the mice at two time points following a 30-min restraint, at 4 h and 18 h. A 4-h interval falls within the normal recording period following slice preparation and, we reasoned, would determine whether the observed desensitization of the CRH neurons sustained *ex vivo* is an accurate reflection of what occurs *in vivo*. The 18-h interval fell on the day after the stress exposure and tested whether the *in vivo* duration of the NE desensitization was sustained overnight. We found that the 4-h post-stress recovery period was insufficient for the CRH neurons to recover from glucocorticoid-induced NE desensitization, and that the NE response was partially, though not entirely, restored at 18 h after the acute stress. Thus, the desensitization of CRH neurons to noradrenergic activation is long-lasting, persisting for over 18 h.

We showed previously that the CRH neuron response and HPA activation to a somatic stress, LPS exposure, but not to a psychological stress, predator odor, are mediated by NE afferents and susceptible to desensitization by prior stress exposure ²¹. The long-lasting desensitization to NE appears, therefore, to be a negative feedback filter of the HPA axis that passes acute psychological stress activation while filtering out more long-lasting somatic stress activation (e.g., by immune challenge). This filtering of stress modality-specific stress responsiveness is adaptive in that it reduces the duration of stress exposure, since somatic stressors tend to be prolonged in nature, which would otherwise subject the organism to the damaging effects of sustained glucocorticoid exposure. However, this feature could be maladaptive in modern day, since deprioritizing somatic stressors in favor of psychological stressors may be beneficial in life-or-death situations, such as a looming predator, but those situations are less likely to occur in modern human life. Additionally, many modern stressors could be perceived as severe even though they may not be life threatening. Thus, acute stress-induced desensitization could be a maladaptive mechanism that needlessly deprioritizes somatic stressors and leaves the body susceptible to physiological distress. Further studies are needed to determine whether desensitization of the HPA axis generalizes to other somatic stressors, and to ascertain the role of noradrenergic desensitization of the HPA axis in diseases known to be caused by HPA dysregulation.

Glucocorticoid altered AR α 1b trafficking

Live-cell imaging of NE and CORT effects on trafficking of AR α 1b revealed that much of the trafficked receptor was directed to a single area of each cell, or a “hot spot”. This receptor-concentrated “hot-spot” co-localized well with the Rab7 marker of late endosomes. FRET analysis confirmed that Rab7 and AR α 1b are physically associated within 10 nm of each other, indicating that AR α 1b is likely located in the late endosome. The lack of increase in early endosomal AR α 1b in the presence of CORT indicates that the desensitization to NE by CORT is not caused by an increase in AR α 1b internalization, as we originally hypothesized based on its prevention by blocking dynamin-dependent endocytosis ²¹. The decrease in AR α 1b in rapidly recycling endosomes suggests that the increase in cytosolic receptor is due to a decrease in receptor trafficking back to the membrane. This rapid steroid-driven change in GPCR trafficking is not only novel *per se*, but also reveals a previously unobserved switch in the speed of receptor trafficking. Receptors of the Class A GPCRs, including the adrenoreceptors, are rapidly recycled back to the plasma membrane in minutes, whereas Class B receptors, which bind more stably to β -arrestin, take several hours ⁴³. The ubiquitination patterns of the β -arrestin that associates with these receptors during endocytosis appear to be specific for the receptor family type and ultimately control the speed of trafficking ³⁴. Our findings reveal a previously unobserved switch in GPCR trafficking kinetics and suggest that CORT could be altering the ubiquitination patterns of β -arrestin to switch GPCRs from rapidly trafficking Class A receptors to slowly trafficking Class B-like receptors. The overall increase in β -arrestin ubiquitination in the presence of CORT suggests that

other rapidly trafficking GPCRs may be similarly affected. Thus, this may represent a more universal mechanism of CORT desensitization of multiple neurotransmitters systems, a prospect that needs to be further explored.

While it remains unclear how glucocorticoids cause an increase in β -arrestin ubiquitination, changes in trafficking may also be achieved by S-nitrosylation of trafficking proteins. We found no change in the nitrosylation of β -arrestin 1/2 with NE or CORT treatment, indicating that the convergent regulation of AR α 1 by nitrosylation and glucocorticoid is not mediated by a change in the nitrosylation state of β -arrestin. S-nitrosylation regulates multiple other proteins involved in GPCR trafficking in addition to β -arrestin, including G protein-coupled receptor kinase 2, endophilin, and clathrin (Hayashi et al., 2018; Ozawa et al., 2008; Whalen et al., 2007). It is possible, therefore, that rapid CORT regulation of nitrosylation of these proteins may mediate the glucocorticoid modulation of AR α 1b intracellular trafficking, a possibility that remains to be determined.

The nuclear glucocorticoid receptor is one possible receptor candidate for the rapid CORT actions in this pathway. However, one or more other membrane receptors may also be candidates for the rapid glucocorticoid effects. Many studies have provided evidence for the existence of a glucocorticoid receptor on the membrane that signals via G protein or tyrosine kinase activity (Di et al., 2003; Rhen and Cidlowski, 2005), and a recent study shows glucocorticoid binding to a membrane adhesion Gai-coupled orphan receptor ⁴⁵. Our PLA analysis indicated a close physical association (<40 nm) of GR with AR α 1b, which suggests a possible direct interaction of the two receptors. We have found the M-20 GR antibody we used here to recognize proteins in the membrane, cytosolic, and nuclear fractions in Western blot analyses (**Figure 7**). The GR/AR α 1b complex detected by PLA responded to CORT by translocating to the nucleus, an effect that was blocked by concomitant activation of AR α 1b by NE. Thus, the receptor complex is differentially regulated depending on its component receptor activation. The interaction between the two receptors suggests this as a possible rapid mechanism whereby glucocorticoids could influence AR α 1b intracellular trafficking.

Based on the well-established role of β -arrestin in receptor trafficking, our finding of an additive effect of Cort and NE on β -arrestin ubiquitination suggests that β -arrestin is a locus of the Cort actions that regulate AR α 1b trafficking. The proximity ligation assay showed that both GR and AR α 1b can individually interact with β -arrestin, and further showed that GR and AR α 1b can interact with each other. However, the PLA did not allow for determination of whether GR, AR α 1b, and β -arrestin act together in a single complex. The ability of CORT to cause nuclear translocation of the AR α 1b- β -arrestin complex, the AR α 1b-GR complex, and the GR- β -arrestin complex suggests a trimeric complex that includes the GR, but we have not demonstrated the multimeric association of the three proteins by molecular assays. While adrenoreceptor localization in the nucleus has been previously reported in cardiomyocytes ⁴⁶ and astrocytes ⁴⁷, this phenomenon is still relatively novel. AR α 1b signaling could be further biased upon entering the nucleus, giving rise to more potential targets for synergistic NE-CORT signaling. Regardless, co-activation of AR α 1b and GR results in fundamental changes in the state of β -arrestin, increasing its ubiquitination, which are associated with profound alterations in the trafficking of the AR α 1b. The CORT regulation of adrenoreceptor trafficking is likely to be responsible for the stress desensitization of the CRH neurons to the noradrenergic excitatory synaptic input activated by somatic stress.

Finally, the concept of CORT-induced changes in endosomal trafficking is novel. Changes in receptor trafficking could have a broader impact on neuronal function including plasticity via regulation of ionotropic receptors ^{48,49}. Glucocorticoids have been shown to facilitate Rab4 cycling, in turn increasing AMPAR recycling ⁵⁰. These studies are crucial for understanding the full extent of rapid stress-induced glucocorticoid regulation of synaptic signaling beyond the classically studied transcriptional regulation.

Acknowledgements

We would like to thank Dr. Chris Hague of the University of Washington for his donation of the ARa1b-eGFP plasmid. This work was supported by NIH grant MH119283 and a Carol Lavin Bernick Faculty Grant from Tulane University.

Additional information

Author contributions

Conceptualization: G.L.W., L.M.H., and J.G.T.; methodology: G.L.W., L.M.H., Z.J., and J.G.T.; investigation: G.L.W., L.M.H., Z.J., A.M.N., M.S.F., S.N., P.S.T.; writing – original draft: G.L.W. and L.M.H.; writing – review & editing: J.G.T., G.L.W., L.M.H., Z.J., and G.L.W.; visualization: G.L.W., L.M.H., Z.J., and J.G.T.; supervision, J.G.T.; project administration, J.G.T.; funding acquisition, J.G.T.

Declaration of interest

The authors declare no competing interests.

Methods

Mice

CRH-eGFP transgenic mice were raised in-house from breeders provided by the Mutant Mouse Resource and Research Center at the University of California, Davis (MMRRC, stock: Tg(Crh-eGFP)HS57Gsat/Mm, RRID:MMRRC_017058-UCD). The mice were genotyped at 2-3 weeks using the primer set: CRH-F1 (CTG TCT TGT CGT GGG TGT CCG AT); GFP R2 (TAG CGG CTG AAG CAC TGC A). All animals and procedures were approved by the Tulane Institutional Animal Care and Use Committee (IACUC) and followed National Institutes of Health guidelines. Mice were housed in an AAALAC-accredited animal facility on a 12:12 light/dark cycle (lights on at 7:00 AM) under controlled temperature (20°C) and received food and water *ad libitum*.

Cell culture

We used an immortalized hypothalamic cell line, mHYPOE-N42 (N42) cells (Cellutions Biosystems), that expresses CRH ²⁶ as well as both the nuclear glucocorticoid receptor and a membrane-associated glucocorticoid receptor ^{36,37}. Cells were plated at a density of 4.5×10^4 cells/well on 12 mm glass coverslips (size 0, Carolina Biological, Burlington, NC) in 24-well plates (Corning, Corning, NY) in 1X Dulbecco's Modified Eagle's Medium (DMEM, Millipore Sigma, Burlington, MA) supplemented with 10% fetal bovine serum (FBS, Atlas Biologicals), 1% penicillin-streptomycin solution, and 0.2% Plasmocin Prophylactic® (Invivogen, San Diego, CA). Plates were incubated at 37°C with 5% CO₂ until cells were approximately 80% confluent. To remove endogenous hormones, the culture medium was then removed, cells were washed once with 1X Dulbecco's-phosphate buffered saline (PBS), and fresh DMEM without phenol red supplemented with 5% charcoal-stripped (CS) FBS was added for 16 h at 37°C with 5% CO₂. For proximity ligation assays, N42 cells were seeded at a density of 12,000/well in 16-well chamber slides and cultured as described above.

Alpha-1 adrenoreceptor overexpression

For live imaging assays, N42 cells were transiently transfected with a pEGFP-AR α 1b construct generously donated by Dr. Chris Hague of the University of Washington. Transfection was achieved by electroporation with two 10 ms, 1400 mV pulses using the NEON® Transfection System (Life Technologies, Carlsbad, CA). Transfected cells were plated onto No. 1 coverslip bottom plates for live-cell imaging with a Nikon A1 confocal microscope with incubator stage (Nikon USA, Melville, NY). For proximity ligation assays, cells were transfected with either pEGFP-AR α 1b or Myc-DDK-tagged human AR α 1b (Origene, Rockville, MD) using Lipofectamine 3000.

Cell imaging

Intracellular AR α 1b trafficking was tracked in live N42 cells using cytosolic intensity analysis, Förster resonance energy transfer (FRET), or pixel-based colocalization. For cytosolic intensity analysis, cells were transiently transfected as described above with pEGFP-AR α 1b. Images were captured every 15 seconds and intensity was measured within a manually defined cytosolic region of interest. For FRET, cells were transiently transfected with a combination of pEGFP-AR α 1b (the FRET donor) and Rab4a-dsRed, Rab5-dsRed, Rab7-dsRed, or Rab11-dsRed (FRET acceptors) and imaged repeatedly over time. Three images were taken at each time point: 1) donor excitation, donor emission, 2) acceptor excitation, acceptor emission, and 3) donor excitation, acceptor emission. The same set of images was also taken in cells only expressing either donor or acceptor fluorophores and used as bleed-through controls. Images were analyzed using the FRET analysis tool in Image J ⁵¹, which yielded images with a FRET intensity for each pixel. A region of interest around each cell expressing both constructs was defined, and the average FRET signal was calculated in the region to quantify the colocalization of AR α 1b with each Rab protein. For pixel-based colocalization, cells were transfected with pEGFP-AR α 1b and RFP-LAMP1 followed by colocalization analysis using Cell Profiler ⁵² to calculate each cell's mean Pearson's coefficient.

Ubiquitination assay

N42 cells were treated with drugs for 20 min at 37°C before lysis with NP40 lysis buffer containing 50 mM Tris-HCl (pH 7.4), 150 mM NaCl, 1% NP-40 and 5 mM EDTA for 1 h at 4°C. Immunoprecipitation (IP) of the protein of interest was performed using the Catch-and-Release kit (Millipore Sigma, Burlington, MA), eluting with the non-denaturing buffer. Eluates were run on a gradient SDS-PAGE gel (Bio-Rad, Hercules, CA) and transferred to PVDF membrane. Blots were blocked for 1 h in 5% dry milk in Tris-buffered saline with 0.05% tween (TBST). Primary antibodies were applied in 5% BSA in TBST overnight, followed by 4 washes in TBST. HRP-conjugated secondary antibody (Clean-Blot IP Detection Reagent, Thermo-Fisher, Waltham MA) was applied in 5% dry milk in TBST for 1 h at room temperature followed by 4 washes in TBST. Chemiluminescent substrate (Super Signal West Pico, Bio-Rad, Hercules, CA) was applied for 5 min followed by detection on a ChemiDoc densitometer (Bio-Rad). Densitometry scores for ubiquitin were quantified and normalized against the protein purified by immunoprecipitation. The antibodies used to target GFP, β -arrestin, and ubiquitin are described in Table 1.

Proximity ligation Assay

Proximity ligation assay (PLA) was carried out with the Duolink system (Millipore Sigma). N42 cells were seeded at a density of 12,000 cells/well on chamber slides (Thermo Scientific) and cultured as described above. For some assays, cells were transfected with human AR α 1b-myc-DDK (Origene) or pEGFP-AR α 1b using Lipofectamine 3000 (Thermo Scientific). Before assay, cells were cultured for 16h in phenol red-free DMEM supplemented with 10% CS FBS, pen-strep, and Plasmocin. Cells were treated for 20 min with the indicated drugs at 37°C, followed by three washes with PBS, fixation in 4% paraformaldehyde, and blocking with Duolink blocking solution. Incubation in primary antibody was carried out overnight at 4°C. For each assay, two primary antibodies were used, one for each protein or post-translational modification. One antibody of

each set was prepared in rabbit and the other in mouse in order to use the complementary secondary antibodies of the Duolink system. After washing with PBS, PLA was performed with the Duolink In Situ system according to the manufacturer's instructions (Millipore Sigma, Burlington, MA). Negative control wells were incubated with only one of the two primary antibodies and generally yielded less than 5 signals per cell (Supplemental Figure 3). For some assays, PLA was followed by incubation in phalloidin-FITC (1:750, Abcam, Boston, MA) to demarcate cells. Cells were imaged at 40x or 60x with a Nikon Eclipse Ti confocal microscope. For each treatment group, Z-stacks were taken from 2-4 fields. Image quantification was performed with NIH ImageJ/FIJI software. Protein-protein interaction signal was quantified as the number of "dots". For cells expressing the ARα1b-myc-DDK transgene, a threshold of 15 fluorescent dots per cell was used to distinguish the stain from background, and the phalloidin-FITC stain was used to demarcate the cell boundary. For assays employing pEGFP-ARα1b, transfected cells were determined by GFP signal, and these cells were individually analyzed for total number and nuclear fraction of interactions. See Table 1 for antibodies and dilutions.

Nitrosylation assay

For histological nitrosylation assays, cells were cultured on chamber slides as described above. A biotin switch assay was performed according to the manufacturer's instructions (Cayman Chemical Company, Ann Arbor, MI) to determine nitrosylation of endogenous β-arrestin1/2. After substitution of biotin for the NO group, PLA was performed with mouse anti-biotin and rabbit anti-β-arrestin1/2 antibodies (see Table 1 for antibody information). Phalloidin-FITC staining was not compatible with the biotin switch assay, so entire fields, rather than individual cells, were quantified for the post-translational modification signal. For each field, the total number of nuclei and the total number of PLA signals were determined in their individual channels. An average number of PLA (nitrosylation) signals per cell was determined by dividing the total number of PLA signals by the total number of nuclei. The nuclear outlines were superimposed onto the image of PLA signal, the PLA signal outside the nuclei was cleared, and the remaining PLA signal was determined to calculate the average nuclear fraction for each field.

For assays of nitrosylation in cell lysates, N42 cells were seeded at 0.5×10^6 in 6-well culture plates and grown to confluence. After treatment for 20 min with the indicated drugs at 37°C, cells were lysed, and NO post-translational modifications were substituted with a tandem mass tag (TMT) according to the manufacturer's instructions (Thermo Pierce). Lysates were subjected to Western blotting as described above and probed with an anti-TMT antibody. Densitometry was performed with Image Lab software (Bio-Rad), and results are presented as the sum density of all bands in each lane.

Electrophysiology

Ex vivo whole-cell patch clamp electrophysiological recordings were conducted in acutely prepared hypothalamic slices from 6-9 week-old male CRH-eGFP mice. On the mornings of experiments, a mouse was gently removed from its home cage to a transfer cage and transported to an adjacent room in the vivarium, where it was immobilized in a flexible plastic decapitation cone (DecapiCone, Braintree Scientific) and, within less than 2 min of removal from its home cage, decapitated without anesthesia using a rodent guillotine. Anesthetics activate the HPA axis and increase circulating levels of ACTH and CORT ⁵³, which otherwise remain low in our hands for ~3 min from the start of handling. Following decapitation, the brain was quickly removed and cooled in oxygenated, ice-cold artificial cerebrospinal fluid (aCSF) containing (in mM): 140 NaCl, 3 KCl, 1.3 MgSO₄, 11 Glucose, 5 HEPES, 1.4 NaH₂PO₄, 3.25 NaOH, 2.4 CaCl₂, pH 7.2-7.4, with an osmolarity of 290-300 mOsm. The forebrain was isolated, the ventral half was blocked, and the caudal surface of the block was glued to a vibratome chuck. Two or three 300 μm-thick coronal slices containing the hypothalamic PVN were then sectioned on a vibrating slicer (Leica) in cooled aCSF, bisected down the midline, and the hemi-slices were transferred to a holding chamber, where they were maintained in oxygenated aCSF at 20°C for at least 1 h to allow for recovery prior

to electrophysiological recordings. Single slices were then transferred and weighted with a silver wire to the bottom of a submerged recording chamber on a fixed-stage, upright microscope (Olympus BXW51) equipped with a long working distance, water immersion 40x objective, where they were perfused with fresh aCSF at a rate of ~2 ml/min at 20°C. eGFP-expressing CRH neurons in the PVN were first identified under epifluorescence illumination, which was then switched to infrared-differential interference contrast optics and the cells were visualized on a monitor using a near-infrared camera to target them for whole-cell patch clamp recordings. Patch pipettes were pulled from borosilicate glass (ID 1.2 mm, OD 1.65 mm) to a resistance of 3–6 MΩ on a horizontal puller (P-97, Sutter Instr.) and filled with an internal patch solution containing (in mM): 120 potassium gluconate, 10 KCl, 1 NaCl, 1 MgCl₂, 0.1 CaCl₂, 5.5 EGTA, 10 HEPES, 2 Mg-ATP, and 0.3 Na-GTP, with a pH adjusted to 7.3 with KOH and osmolarity adjusted to 300 mOsm with D-sorbitol. The GABA_A receptor antagonist picrotoxin (PTX, 50 μM) was applied via the bath perfusion to isolate excitatory postsynaptic currents (EPSCs). While multiple cells from the same mouse were sometimes recorded, only one cell was recorded per hemi-slice. The numbers of recorded CRH neurons are designated as the 'n' and numbers of mice used for each experiment are designated as the 'N'.

Statistical analyses

Data are presented as the mean ± standard error of the mean. Postsynaptic currents were selected and analyzed for changes in frequency, amplitude, and decay time with Minianalysis 6.0 (Synaptosoft Inc.). Baseline values were calculated from 3 min of recording just prior to drug application and compared with the 3 min of recording at the peak of drug responses using a within-cell analysis. For live cell imaging experiments, mean fluorescence values were calculated from the final 3 min of the baseline and of the drug application and were compared using a within-cell analysis. FRET values were taken from single time points at the end of both the baseline and drug applications. Statistical significance was determined with the two-tailed, paired Student's *t*-test for within-cell drug effects, or ANOVA with post-hoc Dunnett's or Tukey's analysis where indicated (GraphPad Prism 7-9 and SigmaPlot 11.0, Systat Software, Inc.).

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Reviewer #1 (Public review):

Summary:

In this manuscript, the authors investigate the cellular mechanism underlying suppression of adrenergic effects on excitatory transmission onto hypothalamic CRH neurons by stress. Experiments in ex-vivo slices show that this is a long-lasting effect that occurs through endocytosis of receptors. The authors then move into an immortalized hypothalamic cell line to enable investigation of the mechanism of changes in receptor trafficking. They use a series of immunohistochemistry, FRET, and biochemical experiments to show that application of corticosterone increases targeting of alpha1 adrenergic receptors to the late endosome and lysosome rather than the recycling endosome. Perhaps most interesting, they find that alpha1 receptors and glucocorticoid receptors form a complex that is ultimately transferred to the nucleus.

Strengths:

Overall, the studies in this manuscript are rigorous and well-conducted. The data supports their conclusions, and they've shown convincingly that glucocorticoid signaling affects trafficking of alpha1 receptors in the culture system they are using. These findings are important for the field of stress research, both in understanding how two components of the stress system (norepinephrine and HPA axis) interact with each other and in neuromodulatory modulation of hypothalamic CRH neurons. Their finding that alpha1 receptors and glucocorticoid receptors form a complex is particularly interesting and maybe impactful outside of the immediate application in the hypothalamus.

Weaknesses:

The study has two primary weaknesses. First, the majority of the experiments were conducted in an immortalized hypothalamic cell line. This was necessary to conduct the type of experiments needed to test the author's hypothesis, but it remains unclear how closely these cells resemble CRH neurons, or how the same mechanism may be preserved or altered in an intact circuit. Further discussion of these points would strengthen the manuscript. Second, while experiments are generally well-designed, the authors do not show that the effects of corticosterone can be blocked with a glucocorticoid receptor antagonist. This is fairly standard pharmacology and would strengthen confidence in the findings presented in the study.

<https://doi.org/10.7554/eLife.102783.1.sa3>

Reviewer #2 (Public review):

Summary:

In this manuscript, the authors report novel and exciting findings delineating a non-transcriptional mechanism whereby glucocorticoids desensitize CRH neurons to NE in response to somatic stress. The authors find that this desensitization induced by CORT 1. persists more than 18h, 2. reduces surface expression of AR1bR (NE receptors) by redirecting trafficking from rapid recycling to late endosomal pools and lysosomes, 3. is dependent on NE binding to the AR1bR, 4. involves cellular nitrosylation, 5. involves ubiquitination of beta-arrestin, and 5. involves interactions between glucocorticoid receptors and AR1bs, glucocorticoid receptors and ubiquitinated beta-arrestin, and AR1b and ubiquitinated beta-arrestin. While the authors do not directly provide evidence for a trimeric complex composed of these three proteins, their data that CORT causes translocation of these dimeric complexes to the cell nucleus suggests it is likely. Overall, these results are highly informative for understanding novel mechanisms mediating glucocorticoid regulation of GPCRs.

Strengths:

- Good rationale for each experiment, which describes many parts of the CORT-NE desensitization mechanism
- Great discussion of limitations of the approaches and the parts of the mechanism we do not fully understand yet
- Appropriate approaches for questions being answered
- Describes a highly novel CORT mechanism that non-transcriptionally switches GPCR trafficking dynamics, something that could have far reaching implications for other GPCRs involved in stress responses

Weaknesses:

- Unclear how this mechanism would generalize to other stressor modalities. Restraint stress is a somatic stressor, but can also be considered a psychological stressor (model of depression-like behavior). A purely somatic stressor might increase the robustness of this phenomenon.
- Remains unknown how nitrosylation plays into the mechanism in terms of specific proteins affected by CORT (GRK2, endophilin, clathrin possibilities)

<https://doi.org/10.7554/eLife.102783.1.sa2>

Reviewer #3 (Public review):

Summary:

In this manuscript, Weiss et al describe a mechanism through which glucocorticoids desensitize CRH neurons in the PVN to norepinephrine. This follows on from previous work from this lab showing rapid glucocorticoid suppression of adrenergic signaling in CRH neurons specific to somatic stress activation, and modality-selective glucocorticoid negative feedback.

Specifically, their previous work shows that:

- (1) NE increases glutamate drive to CRH neurons
- (2) CORT blunts the effects of NE through a dynamin-dependent mechanism
- (3) This contributes to loss of NE signalling after stress (specifically when the second stressor is a physiological one)

Here they extend this line of interrogation by showing that CORT redistributes Ara1b receptors from rapid recycling endosomes to late endosomes and lysosomes. They show a time window of CORT actions and provide additional mechanistic details implicating nitric oxide-dependent nitrosylation in receptor trafficking.

Strengths:

Builds on existing work to provide additional mechanistic details.
The experiments are well done and data are compelling.
The link to nitrosylation is novel (but see below)

Weaknesses:

(1) The link to nitrosylation is interesting, but a bit confusing. If I understand correctly, inhibiting the production of NO or using NEM increases receptor internalization, suggesting that NO-dependent nitrosylation prevents ligand-dependent internalization. What is unclear to me is how CORT is linked to this step. I note the authors show a decrease in nitrosylation with CORT. So, does CORT decrease the activity of NOS and, thus, the production of NO? If so, then exogenously activating this system in the presence of CORT should result in a recovery of NE-dependent increase in glutamate release. Or is the GCR directly decreasing nitrosylation? Linking these elements is critical in terms of furthering our mechanistic understanding of this process.

(2) It's not clear why/how blockade of Ara1 after CORT-induced cytosolic accumulation results in a reversal of effect. Unless I misunderstood something, this requires further explanation.

<https://doi.org/10.7554/eLife.102783.1.sa1>

Author response:

We appreciate the expression of enthusiasm for our paper by the editors and the three reviewers and the suggestions on how to improve the study. Here we outline how we will address the reviewers' concerns and suggestions in a planned revision of our manuscript.

Reviewer #1 listed two primary weaknesses:

(1) the need for discussion of the extent to which the cell line we used resembles CRH neurons and

(2) that we did not test for the effect of blockade of the glucocorticoid receptor.

(1) As the reviewer acknowledges, our experiments called for the use of a cell line to dissect intracellular trafficking of the $\alpha 1$ adrenoreceptor. We selected the N42 cell line for this purpose because it is an immortalized hypothalamic cell line (developed by Belsham and colleagues, Belsham et al., 2004) that expresses CRH. We have used this cell line successfully in the past to study transcriptional and rapid non-genomic actions of glucocorticoids, which indicated that, in addition to expressing CRH, these cells also express both the nuclear glucocorticoid receptor and a membrane-associated receptor that binds glucocorticoids (Rainville et al., 2019; Weiss et al., 2019). We believe that this hypothalamic cell line is the most closely related to native PVN CRH neurons of any cell line available. As requested, we will add to the Discussion of the manuscript to further justify our choice of cells.

(2) We agree that this experiment should be performed. We will test the classical GR (and progesterone) antagonist RU486 (mifepristone) for its effect on the cort regulation of $\alpha 1$ adrenoreceptor trafficking. Our ex vivo electrophysiology studies have indicated that the rapid glucocorticoid effect in native hypothalamic CRH neurons is not blocked by RU486 and is not, therefore, dependent on activation of the classical nuclear GR (Di et al., 2003; Di et al., 2016).

Reviewer #2 also listed two main weaknesses of the study:

(1) that we did not test whether the adrenoreceptor desensitization by restraint stress generalizes to other stress modalities and might be more robust with a pure somatic stressor, and

(2) the lack of identification of a target protein as a mechanism for the role of nitrosylation.

(1) We used restraint stress as a means to elicit corticosterone release, which desensitized the HPA response to a NE-dependent somatic stressor (lipopolysaccharide injection) but not to a NE-independent psychological stressor (predator odor) (Jiang et al., 2021). We got a near-complete loss of the sensitivity of CRH neurons to NE with restraint (i.e., near ceiling effect), such that a different stressor, including a more purely somatic stressor, should not increase the Cort-induced desensitization further. For that reason, we would argue that testing other stressors would not add value to the current study. That said, we plan and have received new funding to test in the future whether the Cort desensitization of the HPA response to LPS stress generalizes to other somatic stressors. We also have future plans to test for the Cort desensitization of other Gq-coupled receptors.

(2) We agree that finding the molecular target of nitrosylation as the mechanism for Cort desensitization of $\alpha 1$ adrenoreceptors would significantly improve the study, but this is a potentially enormous undertaking as it will require the screening and validation of multiple proteins involved in protein trafficking to find the one(s) targeted for nitrosylation by Cort. We tested β -arrestin as a possible target in the paper, but did not find Cort to regulate β -arrestin nitrosylation. We plan to undertake a general nitrosylation screen of proteins to identify multiple possible targets, but prefer to defer this and the validation of possible targets to a future, more thorough analysis.

Reviewer #3 also pointed out two main weaknesses of our study:

(1) that the glucocorticoid nitrosylation link was confusing, and

(2) that it was unclear how blocking $\alpha 1$ adrenoreceptors reversed the Cort-induced cytosolic accumulation of the receptor.

We appreciate the reviewer pointing out these deficiencies in our interpretation and explanation of our findings. We plan to address them directly in the revised version of the paper.

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<https://doi.org/10.7554/eLife.102783.1.sa0>