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Constitutive increases in corticotropin releasing factor and fatty acid amide hydrolase drive an anxious phenotype

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Abstract

Background—Corticotropin-releasing factor (CRF) mediates anxiogenic responses by activating CRF₁ receptors in limbic brain regions. Anxiety is further modulated by the endogenous cannabinoid (eCB) system that attenuates the synaptic effects of stress. In the amygdala, acute stress activates the enzymatic clearance of the eCB N-arachidonylethanolamine (anandamide; AEA) via fatty acid amide hydrolase (FAAH), although it is unclear whether chronic stress induces maladaptive changes in amygdalar eCB signaling to promote anxiety. Here, we used genetically-selected Marchigian Sardinian P (msP) rats carrying an innate overexpression of CRF₁ receptors to study the role of constitutive upregulation in CRF systems on amygdalar eCB function and persistent anxiety-like effects.

Methods—We applied behavioral, pharmacological, and biochemical methods to broadly characterize anxiety-like behaviors and amygdalar eCB clearance enzymes in msP versus non-selected Wistar rats. Subsequent studies examined the influence of dysregulated CRF and FAAH systems in altering excitatory transmission in the central amygdala (CeA).

Results—MsPs display an anxious phenotype accompanied by elevations in amygdalar FAAH activity and reduced dialysate AEA levels in the CeA. Elevations in CRF-CRF₁ signaling dysregulate FAAH activity, and this genotypic difference is normalized with pharmacological blockade of CRF₁ receptors. MsPs also exhibit elevated baseline glutamatergic transmission in the CeA, and dysregulated CRF-FAAH facilitates stress-induced increases in glutamatergic activity.

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Treatment with a FAAH inhibitor relieves sensitized glutamatergic responses in msPs and attenuates the anxiety-like phenotype.

Conclusions—Pathological anxiety and stress hyper-sensitivity are driven by constitutive increases in CRF₁ signaling that dysregulate AEA signaling mechanisms and disable neuronal constraint of CeA glutamatergic synapses.

Keywords

central amygdala (CeA); endocannabinoid; anxiety; CRF₁; glutamate; excitatory

Introduction

Acute stress mobilizes the peptide corticotropin-releasing factor (CRF) that orchestrates the endocrine, visceral, and behavioral responses to stress (1). Repeated induction of the CRF system contributes to maladaptive changes that promote negative affect and anxiety (2, 3). Although inextricably linked to the hypothalamic-pituitary-adrenal (HPA) axis, anxiety-like behavior is influenced by CRF type 1 (CRF₁) receptor signaling in the amygdala and other limbic regions (4, 5). The amygdala receives sensory information in the lateral and basolateral regions, which relay to the central nucleus of the amygdala (CeA) serving as the primary integrator and distributor of stress-related information (6, 7). Neuronal activation of the medial CeA exerts bidirectional control of anxiety-like function that is influenced by glutamatergic signaling pathways (8). In this regard, acute stress and CRF administration increase CeA glutamatergic activity (9–12), and this effect is sensitized with prior stress experience (13). Dysregulated CRF-CRF₁ signaling may therefore serve as a primer for inducing long-term neurological adaptations that influence amygdalar function and subsequent effects on stress and anxiety (3, 14–16).

Anxiety-like behavior is modulated by negative feedback systems that constrain the stress response. In this regard, the endogenous cannabinoid (eCB) system plays an important homeostatic role in the regulation of stress circuits. The primary eCBs, N-arachidonylethanolamine (anandamide, AEA) and 2-arachidonoylglycerol (2-AG), are synthesized “on demand” in the postsynaptic neuron and act on cannabinoid type 1 (CB₁) receptors in presynaptic terminals to suppress neurotransmitter release (17–20). Termination of eCB signaling occurs via enzymatic clearance by the serine hydrolases fatty acid amide hydrolase (FAAH) that degrades AEA, and monoacylglycerol lipase (MAGL) and α/β -hydrolase domain-containing 6 (ABHD6) that degrade 2-AG (21). Selective enzyme inhibitors reveal that eCB signaling by AEA and 2-AG produce distinct pharmacological profiles (22, 23) and mediate differential behavioral effects (24, 25) that may influence the development of anxiety-like disorders. In this regard, genetic polymorphisms in FAAH and MAGL are associated with disrupted limbic function (26, 27), and contribute to pathological changes in emotion- and reward-related processing (26, 28–30), post-traumatic stress disorder (31), and drug abuse (32, 33). Taken together, deficient eCB signaling may confer a critical loss in synaptic constraint of stress-related pathways that normally gate anxiety.

Stress exposure differentially alters eCB signaling in the amygdala that is associated with the HPA axis in distinct ways (16). For instance, acute stress rapidly increases amygdalar

FAAH activity that is observed with decreases in AEA content, and these effects are blocked with a CRF₁ antagonist (34). Although chronic stress induces similar responses (35), these effects vary with the nature of stress application that infer differences in amygdalar CRF-CRF₁ signaling. To this end, genetically-selected Marchigian Sardinian P (msP) rats, originally developed for high alcohol intake, have co-segregated an anxiogenic-like phenotype that is consistent with aberrations in brain stress signaling. Specifically, msPs display point mutations in the *Ctrh1* locus that elevate gene transcript levels and receptor binding densities in amygdalar regions (36), enhance sensitivity to CRF₁ antagonists (36, 37), and modulate stress responsivity and stress-coping performance (36, 38). Here, we utilized msP rats to explore the long-term consequences of dysregulated CRF₁ signaling on amygdalar eCB function and anxiety-like measures. We implemented a broad-scale biochemical approach evaluating multiple eCB clearance enzymes in the amygdalar proteome, and subsequently identified a genotypic dysregulation in FAAH. We then applied behavioral, neurochemical, and electrophysiological approaches to examine the hypothesis that dysregulated CRF-FAAH mechanisms in the amygdala contribute to AEA signaling deficiencies and prime stress-reactive glutamatergic systems in the CeA to facilitate anxiety.

Methods and Materials

Animals

Adult male msP rats (300–550g) were bred at The Scripps Research Institute (TSRI) from a colony obtained from the University of Camerino (Italy). For genotypic comparisons, we used adult male Wistar rats (300–550g; Charles River, Raleigh, NC) from which the msP line was generated. Rats were group-housed on a 12h reverse light/dark cycle (lights off at 8:00 am) with food and water available ad libitum, and were handled for 3–5 days before experimentation. We conducted all procedures in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals and with the Institutional Animal Care and Use Committee (IACUC) policies of TSRI.

Chemicals and Drug Preparation

A list of chemical compounds, and the preparation methods used for the selective FAAH inhibitor PF-3845, the peptide CRF (human/rat), and the selective CRF₁ receptor antagonist N,N-bis(2-methoxyethyl)-3-(4-methoxy-2-methylphenyl)-2,5-dimethylpyrazolo[1,5-a]pyrimidin-7-amine (MPZP) are provided in Supplementary Information.

Anxiety-like Behavioral Assessments

Elevated Plus Maze (EPM)—Rats were monitored for anxiety-like behavior using EPM procedures as previously described (39), but with minor modifications. Briefly, rats were habituated to a testing environment containing a standard EPM apparatus centered below a dimly-lit overhead lamp (see Supplemental Information). Rats were pretreated with doses of PF-3845 (3 or 10 mg/kg, i.p.), vehicle, or no injection and left undisturbed for 2h in an anteroom prior to EPM testing for open- and closed-arm activity (i.e., time/entries) for 5 min. A full assessment of performance is included in Supplemental Information (Tables S3, S4).

Novelty-induced Hypophagia—Rats were monitored for anxiety-like behavior using hypophagia procedures as previously described (40, 41). Briefly, rats were habituated to a quiet testing room illuminated by red light (see Supplemental Information). Bottles containing a palatable chocolate solution were presented for 30 min. Following acclimation, rats were monitored for baseline consummatory behaviors (i.e., latency to drink/overall intake). The next day, half of the rats from each genotype were pretreated with PF-3845 (10 mg/kg; i.p.) or vehicle, and left undisturbed for 2h prior to evaluation under novel testing conditions perceived to be stressful (i.e., lights on/new cages).

Biochemical Assessments of eCB Clearance Enzymes

Amygdalar Proteome—Amygdalar tissue was extracted bilaterally from a 2-mm coronal section with a 12 GA tissue extractor (Figure S1). Samples were snap-frozen in liquid nitrogen, and stored at -80°C . Membrane proteomes were processed in phosphate buffered saline as previously described (42).

Activity-based Protein Profiling (ABPP)—Amygdalar proteomes (1 mg/mL) were evaluated for eCB metabolic enzyme activity using ABPP as previously described (42). Details of the method are provided in Supplemental Information. Briefly, protein homogenates (50 μL) were incubated with a fluorophosphonate-rhodamine probe (1 μL , final concentration 1 μM) at 37°C for 30 min. Reactions were quenched with SDS-PAGE loading buffer (20 μL). Probe-labeled proteins were separated by SDS-PAGE (10% acrylamide) and visualized with a fluorescent scanner. Raw values of spectral counts were converted to reflect percentage change from Wistar controls.

FAAH Hydrolysis—A substrate hydrolysis assay was performed using liquid chromatography/triple quadrupole mass spectrometry as previously established (43). Briefly, amygdalar proteomes (50 $\mu\text{g}/\text{mL}$) were incubated with multiple concentrations of deuterated AEA standards (d4-AEA) to measure discrete timepoints of metabolic turnover into d4-ethanolamine (d4-EA; pmol/mg/min) (see Supplemental Information). The inverse of turnover values [$1/(\text{pmol}/\text{mg}/\text{min})$] were then used to calculate the maximum velocity of enzymatic reaction (V_{max}) and binding affinity (K_{m}).

Pharmacological studies examined the role of CRF systems in modulating FAAH hydrolysis. To assess the relevance of CRF₁ receptors across genotype, rats were given sub-chronic treatment with the selective CRF₁ antagonist MPZP (10 mg/kg, 2 mL/kg, s.c., b.i.d.) or vehicle for 3 days, and sacrificed ~2h after the final dose. To examine the effects of acute CRF administration, Wistar rats were implanted unilaterally with 23 GA cannulas (PlasticsOne, Roanoke, VA) aimed ~1.5 mm above the lateral ventricles (from bregma AP: -0.8 , ML: ± 1.6 , DV: -3.3 , skull). One week later, rats were acclimated to injection procedures, and received CRF (1 μg) or vehicle infusions via a syringe pump dispensing 2 μL of solution in 2 min. Animals were sacrificed 30 min after injections.

Neurochemical assessments of eCBs and neurotransmitters

In-vivo Microdialysis—Rats were implanted unilaterally with guide cannulas aimed 1 mm above the CeA (from bregma AP: -2.3 , ML: ± 4.0 , DV: -6.4 , dura). One week later,

microdialysis probes were inserted and secured into the guide cannulas. Dialysis and quantification of eCBs and neurotransmitters were performed in separate experiments as previously established (42, 44). Details are included in Supplemental Information.

Electrophysiological recordings

Slice Preparation—Coronal sections of the CeA (300–400 μm) were prepared using a Leica 1000S vibratome (Leica Biosystems, Buffalo Grove, IL) as previously described (12, 45). Slices were incubated with an interface configuration for 30–60 min, submerged, superfused, and equilibrated with 95% O_2 /5% CO_2 artificial cerebral spinal fluid (aCSF) of the following composition: (in mM) 130 NaCl, 3.5 KCl, 1.25 NaH_2PO_4 , 1.5 $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$, 2.0 CaCl_2 , 24 NaHCO_3 , and 10 glucose.

Whole-Cell Voltage-Clamp Recordings—Recordings of spontaneous excitatory postsynaptic currents (sEPSCs) in 31 medial CeA neurons ($n=16$ from 4 msPs, $n=15$ from 4 Wistars) were made with a Multiclamp 700B amplifier (Molecular Devices, Sunnyvale, CA), low-pass filtered at 2–5 kHz, digitized (Digidata 1440A; Axon Instruments), and stored on a computer using pClamp 10 software (Axon Instruments). The internal solution used was composed of (in mM): 145 K-gluconate; 5 EGTA; 5 MgCl_2 ; 10 HEPES; 2 Na-ATP; 0.2 Na-GTP. Recordings ($V_{\text{hold}} = -60\text{mV}$) were performed in the presence of bicuculline (30 μM) and CGP55845A (1 μM). Drugs were dissolved in aCSF and applied by bath superfusion. sEPSCs were analyzed and visually confirmed based on a minimum time interval of 3–5 min and a minimum of 60 events using semi-automated, threshold-based mini detection software (Mini Analysis, Synaptosoft Inc., Fort Lee, NJ).

Intracellular Recording of Evoked Glutamatergic Responses—We recorded evoked excitatory postsynaptic potentials (eEPSPs) from 48 medial CeA neurons ($n=24$ from 13 msPs, $n=24$ from 14 Wistars) with sharp micropipettes filled with 3M KCl using discontinuous current-clamp by stimulating the adjacent basolateral amygdala (BLA) through a bipolar stimulating electrode. We held neurons near their resting membrane potential [Wistar: -77.4 ± 1.0 , msP: -81.4 ± 0.9 mV]. Average input resistance by genotype was 157.1 ± 9.5 M Ω for Wistar and 166.1 ± 11.1 M Ω for msP. Data were acquired with an Axoclamp-2A preamplifier and stored for later analysis using pClamp software (Axon Instruments, Foster City, CA). Details of testing are included in Supplemental Information.

Statistics

For behavioral measures, genotype differences were examined using Student's t-tests (unpaired, two-tailed), repeated measures analysis of variance (RM-ANOVA), or two-way ANOVAs followed by Bonferroni-corrected *post hoc*s where appropriate. For enzymatic measures, genotype and/or drug differences were determined using non-linear regression analyses (d4-EA turnover), Student's t-test (unpaired, two-tailed) or two-way ANOVAs (V_{max} and K_m) followed by *post hoc* assessments. Analyses of d4-EA turnover at each concentration of d4-AEA are provided in Supplemental Information (Table S2). For microdialysis measures, mean baseline levels were analyzed with a multivariate ANOVA (MANOVA) followed by RM-ANOVAs assessing genotypic differences, drug effects across time, and changes relative to baseline levels (Fisher's PLSD). For electrophysiology studies,

genotype and drug treatment were evaluated using t-tests or ANOVA as reported in each experiment. $P < 0.05$ was considered statistically significant throughout.

Results

MsP rats display an innate anxiety-like phenotype

We characterized the anxious phenotype in msP rats using two models of anxiety-like behavior (Figure 1). In the EPM study, msPs displayed lower percentages of time in the open arms (Figure 1A) versus Wistars ($t_{(10)}=2.8$, $*p < 0.05$), albeit fewer closed-arm entries (Figure 1B), indicating reduced maze activity ($t_{(10)}=2.4$, $*p < 0.05$). In the hypophagia study, RM-ANOVA of latencies revealed a genotype x procedure interaction ($F_{(1,18)}=10.0$, $p < 0.01$). Relative to baseline, both groups exhibited increased latency (Figure 1C) during novelty stress ($+p < 0.05$), although msPs displayed higher latencies than Wistars in either procedure ($*p < 0.05$). Analysis of total consumption revealed a main effect of procedure ($F_{(1,18)}=38.9$, $+p < 0.001$), with decreased intake in both groups during novelty stress (Figure 1D), although msPs exhibited a further suppression of intake across procedures ($F_{(1,18)}=12.8$, $*p < 0.01$). The findings demonstrate an elevated anxiety-like phenotype in msPs.

Amygdalar FAAH activity is enhanced in msP rats

We examined genotypic differences in the functional state of amygdalar eCB signaling (Figure 2). ABPP labeling showed higher levels of active FAAH in msP (Figure 2B) than Wistars ($t_{(17)}=2.2$, $*p < 0.05$), with no difference in the 2-AG enzymes MAGL ($t_{(17)}=0.5$, N.S.) or ABHD6 ($t_{(17)}=1.3$, N.S.). Further assessment of FAAH activity using a substrate hydrolysis assay (Figure 2C) predicted higher d4-EA turnover ($F_{(2,92)}=3.8$, $*p < 0.05$) and Vmax rates (Figure 2D) in msPs than Wistars ($t_{(14)}=3.3$, $*p < 0.05$), whereas no differences in Km were observed (Wistar: 0.93 ± 0.06 , msP: 1.58 ± 0.39 μM , $t_{(14)}=1.6$, N.S.). To establish a functional role for genotypic differences in FAAH, *in vivo* microdialysis procedures examined eCB levels in the CeA, revealing lower baseline levels of AEA in msPs (Figure 2E) than Wistars ($F_{(1,17)}=4.6$, $*p < 0.05$). No differences were observed in CeA levels of 2-AG (Figure 2F) ($F_{(1,17)}=0.05$, N.S.), or the ethanolamides palmitoylethanolamine (PEA), or oleoylethanolamine (OEA) (Figure S2A, B). The findings show that msPs exhibit increased amygdalar FAAH activity and reduced AEA tone in the CeA.

Enhanced CRF₁ receptor signaling drives elevated amygdalar FAAH activity

We examined whether CRF-CRF₁ signaling contributes to genotypic differences in amygdalar FAAH activity (Figure 3). In the MPZP study, a two-way ANOVA of Vmax revealed a genotype x drug interaction ($F_{(1,26)}=7.9$, $p < 0.01$). While vehicle-treated msPs displayed increased FAAH hydrolysis versus Wistars ($*p < 0.05$), MPZP (10 mg/kg) abolished the genotypic difference in d4-EA turnover (Figure 3A) ($F_{(2,80)}=0.6$, N.S.), and reduced Vmax rates (Figure 3B) in msPs ($+p < 0.05$) without altering Wistar responses. No significant changes in Km were observed (Table S1). Conversely, CRF administration (1 $\mu\text{g}/2$ μL) induces FAAH activity in Wistars that predicts higher d4-EA turnover (Figure 3C) ($F_{(2,92)}=13.3$, $*p < 0.001$) and Vmax rates (Figure 3D) versus vehicle controls ($t_{(13)}=2.2$, $*p < 0.05$) without altering Km (Vehicle: 1.09 ± 0.07 , CRF: 1.02 ± 0.06 μM , $t_{(13)}=0.1$, N.S.).

Collectively, these findings establish a functional link between genotypic elevations in CRF-CRF₁ signaling and increased amygdalar FAAH activity.

Amygdalar glutamatergic transmission is enhanced in msP rats

Heightened activation of the amygdala facilitates anxiety-like behavior. To investigate the possibility of dysregulated amygdalar circuits, we characterized excitatory transmission in the rat CeA (Figure 4). Baseline levels of 10 different neurotransmitters were assessed using *in vivo* microdialysis. An omnibus MANOVA revealed a significant genotypic difference ($F_{(10,7)}=4.8$, $p<0.05$), with follow-up analyses confirming a robust increase in msP glutamate levels (Figure 4A) versus Wistars ($*p<0.05$), whereas levels of GABA (Figure 4B) and other transmitters were comparable across genotype (Table 1).

Given genotypic differences in CeA glutamate levels, we conducted whole-cell voltage-clamp recordings to characterize sEPSCs in 31 medial CeA neurons (n=16 from 4 msPs, n=15 from 4 Wistars). We found elevated baseline sEPSC frequency in msPs (Figure 4D, top panel) versus Wistars ($t_{(14)}=3.22$, $*p<0.05$), supporting enhanced glutamate release in the CeA. There were no genotypic differences in baseline amplitude (Figure 4D, bottom panel) ($t_{(29)}=0.09$, N.S.), rise (Wistar: 1.7 ± 0.1 , msP: 1.5 ± 0.1 msec, $t_{(29)}=2.0$, N.S.) or decay time (Wistar: 1.3 ± 0.1 , msP: 1.4 ± 0.1 msec, $t_{(29)}=0.7$, N.S.). The majority of glutamatergic terminals that innervate the CeA arise from the BLA (46, 47). To evaluate genotypic differences in glutamatergic signaling of this pathway, we recorded BLA-evoked EPSPs from 48 medial CeA neurons (n=24 from 13 msPs, n=24 from 14 Wistars). RM-ANOVA of eEPSP input-output curves (Figure S3A) revealed no significant differences in genotype, though there was a trend for higher responses in msPs. We did not observe genotypic differences in baseline paired-pulse facilitation ratios of eEPSPs (Figure S3B). These findings suggest that msPs display heightened glutamatergic activity in the CeA characterized by increased local spontaneous, but not BLA-evoked glutamate release.

FAAH inhibition alleviates stress-sensitive increases in glutamatergic transmission in msP rats

Enhanced glutamatergic tone in the amygdala may promote aberrant stress responsivity, and this may be influenced by FAAH. To examine the role of dysregulated FAAH in modulating stress-dependent changes in CeA glutamatergic transmission (Figure 5), the selective FAAH inhibitor PF-3845 was utilized for all *in vivo* and *ex vivo* studies. The specificity of PF-3845 is addressed in Supplemental Information.

To determine changes in CeA glutamatergic transmission during acute stress, glutamate release was assessed by *in vivo* microdialysis in restrained rats pretreated with PF-3845 (10 mg/kg) or vehicle. In msPs, RM-ANOVA of relative changes in glutamate levels revealed a drug x time interaction ($F_{(9,126)}=2.7$, $p<0.05$). While vehicle-treated msPs displayed a robust increase in glutamatergic transmission (Figure 5A) during restraint stress ($+p<0.05$), PF-3845 abolished this response ($*p<0.05$). Conversely, in Wistars, there were no significant changes in CeA glutamatergic transmission (Figure 5B) across the timeline of restraint. PF-3845 mildly reduced glutamate levels (Figure S5) prior to stress induction, although the

time-dependent decreases in this measure were comparable across genotype and did not produce baseline differences versus vehicle controls.

We evaluated the influence of CRF and FAAH on spontaneous glutamatergic responses in medial CeA neurons. Consistent with our previous work in msP neurons (12), CRF (100 nM) increased sEPSC frequency (Figure 5D, top panel) ($139.0 \pm 16.2\%$ of control, $n=7$ neurons from 4 rats). PF-3845 (1 μM) produced no changes in basal sEPSCs ($102.1 \pm 8.3\%$ of control, $n=9$ neurons from 4 rats), but significantly attenuated the CRF-induced increase ($95.1 \pm 3.2\%$ of control, $n=5$ neurons from 4 rats $t_{(10)}=2.2$, $*p<0.05$). Conversely, in Wistar neurons, CRF produced opposing (increased or decreased) effects on sEPSC frequency (Figure 5D, bottom) ($122.5 \pm 16.0\%$ of control, $n=8$ neurons from 4 rats). PF-3845 did not significantly alter basal sEPSCs ($101.4 \pm 3.1\%$ of control, $n=6$ neurons from 4 rats, $t_{(12)}=1.1$, N.S.), nor did the co-application of CRF alter this measure ($111.2 \pm 7.4\%$ of control, $n=6$ neurons from 4 rats). We next investigated the CRF-FAAH interaction on BLA-evoked CeA glutamatergic responses. CRF (100 nM) had no effect on eEPSP amplitude in msP neurons (Figure 5E) ($101.9 \pm 6.03\%$ of baseline, $n=8$ neurons from 7 rats, $t_{(7)}=0.32$, N.S.). However, PF-3845 (1 μM) significantly reduced basal eEPSPs (by $15.64 \pm 3.65\%$, $n=16$ neurons from 9 rats, $t_{(15)}=4.29$, $+p<0.05$) and this decrease persisted with the co-application of CRF ($89.23 \pm 5.51\%$ of baseline, 11/16 cells). Conversely, in Wistar neurons, CRF decreased eEPSPs (Figure 5F) ($83.32 \pm 3.36\%$ of baseline, $n=9$ neurons from 7 rats, $t_{(8)}=4.96$, $+p<0.05$), indicating that CRF inhibits BLA-evoked responses in the CeA. PF-3845 had no effect on basal eEPSPs ($92.85 \pm 3.54\%$ of baseline, $n=13$ neurons from 7 rats, $t_{(12)}=2.02$, N.S.), although the co-application of CRF no longer decreased this measure ($103.4 \pm 3.69\%$ of baseline, 10/13 neurons, $t_{(9)}=0.92$, N.S.). Our findings indicate that FAAH does not indiscriminately alter glutamatergic transmission, but rather modulates control of stress activation of these pathways. In this regard, msPs exhibit heightened sensitivity to stress-induced increases in glutamatergic transmission, and this effect is tempered with FAAH inhibition.

FAAH inhibition alleviates excessive anxiety-like behavior in msP rats

We examined whether FAAH inhibition alters the anxious phenotype in msPs (Figure 6). In the EPM study, a two-way ANOVA of open-arm time revealed a genotype x drug interaction ($F_{(2,83)}=3.2$, $p<0.05$). While vehicle-treated msPs displayed lower percentages of time spent on the open arm (Figure 6A) versus Wistars ($*p<0.01$), PF-3845 (10 mg/kg) abolished this genotypic difference, and restored open-arm exploration in msPs ($+p<0.01$) without altering Wistar behavior. Analysis of closed-arm entries (Figure 6B) revealed that drug treatment did not alter this measure in either genotype ($F_{(2,83)}=0.8$, N.S.), although msPs continued to display reduced maze activity ($F_{(1,83)}=29.3$, $p<0.001$). In the hypophagia study, a two-way ANOVA of latencies revealed a genotype x drug interaction ($F_{(1,33)}=4.2$, $p<0.05$) during novelty stress. Whereas vehicle-treated msPs displayed higher latencies (Figure 6C) than Wistars ($*p<0.001$), PF-3845 mitigated the genotypic difference ($*p<0.01$) and reduced latencies in msPs ($+p<0.05$) without altering Wistar behavior. Analysis of total consumption (Figure 6D) revealed similar findings in that drug treatment did not alter intake in either genotype ($F_{(1,33)}=0.02$, N.S.), although msPs continued to display suppressed intake

($F_{(1,33)}=10.4$, $*p<0.01$). Collectively, these data suggest that FAAH inhibition alleviates excessive anxiety-like responses in msPs.

Discussion

Here, we report that relative to non-selected Wistar rats, msPs exhibit lower dialysate concentrations of AEA in the CeA in association with increases in amygdalar FAAH activity. Our findings are specific to AEA signaling mechanisms, as those related to 2-AG processing were comparable across genotype. Upregulated FAAH activity appears to derive from innate increases in CRF₁ signaling, given that sub-chronic treatment with a CRF₁ antagonist reduces genotypic differences in this measure. MsPs also exhibit higher CeA glutamate levels, increased spontaneous excitatory signaling, and greater stress-induced activation of glutamatergic signaling. FAAH inhibition attenuates the stress- and CRF-induced elevations in glutamatergic transmission, and modestly diminishes evoked responses in msPs. FAAH inhibition also reduced the anxious phenotype of msPs in two distinct models. Collectively, we suggest that prolonged CRF₁ signaling induces long-term dysregulation of FAAH, the interaction of which primes stress-sensitive circuits in the CeA towards a hyper-excitabile state to induce persistent anxiety-like effects.

The well documented increase in CRF₁ signaling in msP rats (36, 48), together with the results showing that CRF₁ receptor antagonism blocks elevated AEA hydrolysis suggest a constitutive interaction between CRF₁ and FAAH that underlies anxiety-like pathologies. The findings are consistent with recent work showing that acute CRF₁ activation rapidly mobilizes FAAH activity in association with reductions in amygdalar AEA content (34). Importantly, the increases in FAAH activity are sustained in rats displaying upregulated CRF₁ systems that derived from phenotypic selection criteria of behavioral anxiety. Thus, we expand on the acute nature of the CRF₁-FAAH interaction to provide a direct link between dysregulation in these mechanisms and persistent anxiety-like effects. Our findings are consistent with studies showing a similar pattern of AEA signaling dysfunction induced by sustained glucocorticoid exposure that increases amygdalar CRF signaling, as well as in mutant mice exhibiting forebrain overexpression of the CRF gene (49). Finally, whereas prior studies characterized CRF-FAAH interactions in the BLA, we have extended these observations to the CeA that serves as an integrative hub for processing and converting stress-related information into behavioral and physiologic responses (6). The regional distinction is noteworthy given recent evidence that anxiety-like behavior is influenced by AEA signaling in the BLA under conditions that elicit low, but not high states of emotional arousal (50). Although the amygdalar dissections performed here did not exclusively isolate the CeA region, the neurochemical/electrophysiological data clearly establish an important role of the CeA in CRF-FAAH interactions and resulting effects on stress-reactive systems that influence anxiety.

Our present findings reveal that acute stressors facilitate CeA glutamatergic transmission, and sensitivity to this response is relieved with FAAH inhibitors. Restraint stress stimulated *in vivo* CeA glutamate levels in msPs, whereas Wistars displayed a tempered and non-significant increase in this measure. *Ex vivo* studies complemented these results by showing that CRF-induced increases in spontaneous glutamate release were more frequent in CeA

neurons from msP rats. Notably, our data support that dysregulated FAAH activity is a contributing factor to genotypic differences in CeA excitatory signaling (12). In this regard, FAAH co-localizes with CB₁ receptors that are primarily restricted to glutamatergic synapses on pyramidal neurons (51–54) found in the medial and lateral CeA (55). All FAAH-positive cells co-express CRF₁ labeling and gene expression, and mutant mice lacking the expression of CRF₁ receptors on forebrain glutamatergic neurons display blunted reductions in amygdalar AEA content following restraint (34). The influence of stress on CeA excitability likely contributes to the anxious phenotype observed in msPs. Acute stress increases CeA glutamate levels after prolonged periods of restraint (9, 10), and intra-CeA treatments that indirectly enhance extracellular glutamate also increase anxiety-like symptoms in EPM and fear conditioning studies (56). Moreover, prior stress exposure primes the neurochemical responses to restraint stress, since fear-conditioned rats display sensitized CRF-induced increases in CeA glutamate levels (13).

We observed robust anxiety-like symptoms in msPs, and genotypic differences in these measures were attenuated with FAAH inhibition. Our evaluations of EPM and novelty-induced hypophagia characterized the anxiety-like phenotype in distinct ways. The EPM studies examined the rats' willingness to actively explore open spaces, whereas hypophagia procedures examined consummatory behaviors under familiar versus novel contexts. PF-3845 selectively increased open arm exploration and reduced latencies to approach palatable substances in msPs, and these effects are closely related to the alleviation of anxiety-like function in rodents that are pre-stressed or exposed to aversive testing conditions (41, 57, 58). It is noteworthy that msPs continued to display suppressed motor activity and reduced chocolate intake despite the anxiolytic effects of PF-3845. These effects may relate to other known attributes of the msP line showing hypohedonic and depressive-like states (59), as well as innate tendencies for freezing that are resistant to pharmacological treatment (36, 38). The selectivity for which PF-3845 alleviates excessive anxiety may relate to the restoration of dysfunctional AEA-CB₁ receptor signaling. Disrupted CB₁ signaling is well known to facilitate anxiety (60–64), whereas CB₁ agonists generally decrease this measure (61, 65–67). FAAH inhibition increases brain AEA levels and is observed to dampen anxiety-like responses under acute (30, 41) and chronic (35, 68, 69) stress conditions. The anxiolytic effects of FAAH inhibitors are thought to derive from CB₁-mediated suppression of glutamatergic versus GABAergic signaling. Specifically, CB₁ deletion on glutamate, but not GABA forebrain neurons reverses the anxiolytic effects produced by low-dose administration of a CB₁ agonist (70). Relatedly, the anxiety-like effects of social defeat are reversed with a FAAH inhibitor and are associated with the dampening of spontaneous excitatory, but not inhibitory signaling in the mouse striatum (69). Taken together, our findings suggest that the anxiety-like phenotype in msPs results from increased FAAH activity allowing for enhanced stress activation of CeA glutamate systems that regulate anxiety.

The evidence of overactive FAAH in our studies represents a previously uncharacterized link between stress-promoting systems (i.e., CRF₁ receptors) and *in vivo* dysregulation of amygdalar glutamatergic mechanisms. Accordingly, msP rats appear to be in a heightened state of vulnerability to the activation of stress mechanisms. We propose that the stress-sensitive features in msP rats relate to dysfunctional AEA signaling elements that regulate

CeA glutamatergic transmission, and contribute to the etiopathology of anxiety. The evidence linking CRF₁-FAAH dysfunction in amygdaloid circuitry with negative affective symptoms has translational value for recent work establishing a parallel between clinical symptoms of aberrant stress reactivity, anxiety disorders, and genomic variations in CRF₁ and FAAH (26, 30, 71, 72). Moreover, our previous work relating the point mutations in the *Crhr1* gene in msPs to aberrant stress responses in fear conditioning studies (36, 38) suggest a possible link between innately dysregulated eCB systems and pathological symptoms of post-traumatic stress disorder. At present, the source of the CRF₁-FAAH interaction has not been determined. It is possible that second messenger systems linked to CRF₁ receptor activation, such as cAMP/PKA and/or MAPK (ERK1/2) pathways (73), may differentially activate FAAH in msPs. However, we cannot rule out other possibilities including genotypic differences in local CRF-CRF₁ neurocircuitry, synaptic function (12), AEA biosynthesis, and/or CRF₁-eCB-glutamatergic mechanisms.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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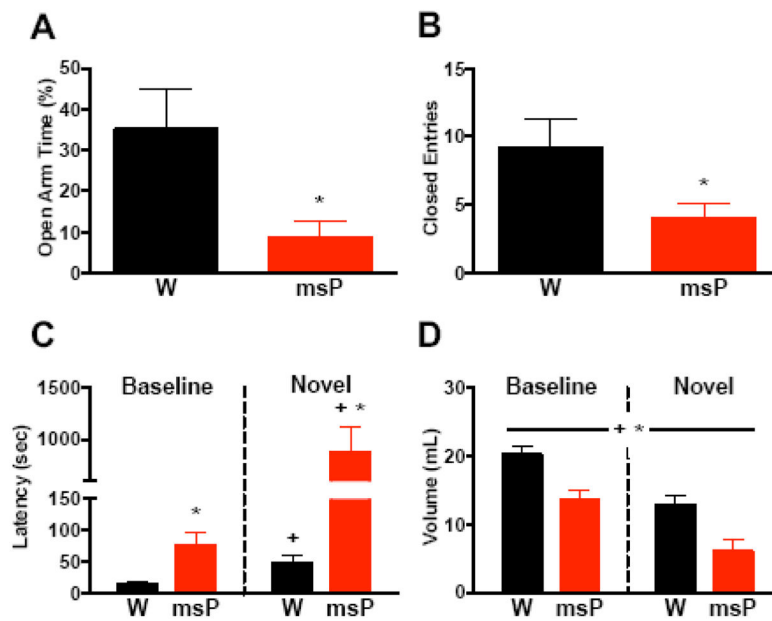


Figure 1. MsP rats display an innate anxiety-like phenotype

(A) Percentage of time spent on the open arm of the elevated plus maze (EPM) in non-selected Wistar (W, $n=5$) and msP ($n=7$) rats. (B) Total entries made into the closed arm of the EPM in rats from A. (C) Latency to drink from a palatable chocolate solution under baseline and novel environments in Wistars ($n=9$) and msPs ($n=11$). (D) Total volume of chocolate solution consumed in rats from C. Data expressed as mean \pm SEM. Asterisks (*) denote significant genotype differences, whereas plus signs (+) denote significant differences from baseline conditions ($p < 0.05$). A continuous line across groups denotes significant main effects.

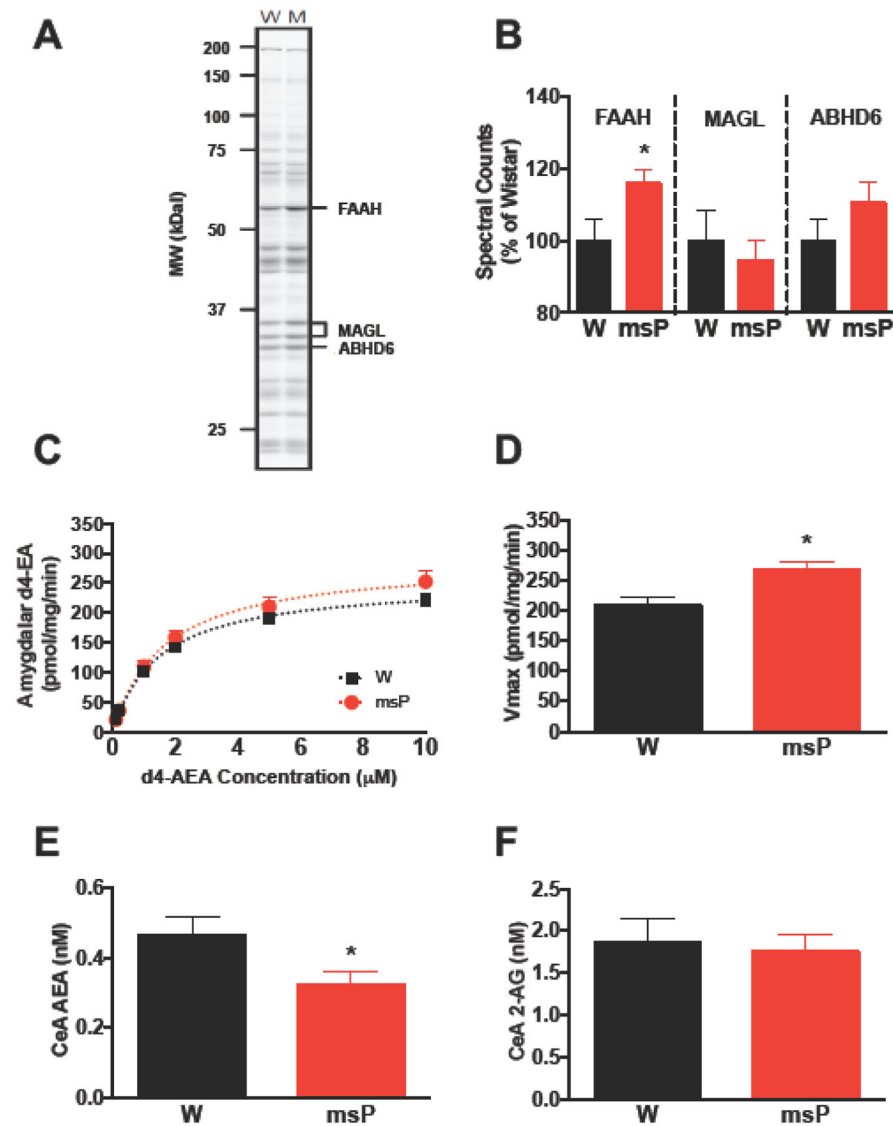


Figure 2. Amygdalar FAAH activity is enhanced in msP rats

(A) Representative gel image of serine hydrolase activity in amygdalar tissue collected from non-selected Wistar (W, n=10) and msP (M, n=9) rats. (B) Percentage of spectral counts in active site labeling of the endocannabinoid clearance enzymes fatty acid amide hydrolase (FAAH), monoacylglycerol lipase (MAGL), and α/β -hydrolase domain-containing 6 (ABHD6) in rats from A. (C) Substrate conversion of deuterated N-arachidonylethanolamine (d4-AEA) into ethanolamine (d4-EA) in amygdalar tissue collected from Wistars (n=8) and msPs (n=8). (D) Velocity of reaction (V_{max}) in rats from C. (E) Baseline dialysate concentration of N-arachidonylethanolamine (anandamide, AEA) in the central nucleus of the amygdala (CeA) of Wistars (n=9) and msPs (n=10). (F) Baseline dialysate concentration of 2-arachidonoylglycerol (2-AG) in the CeA of rats from E. Data expressed as mean \pm SEM. Asterisks (*) denote significant genotype differences ($p < 0.05$).

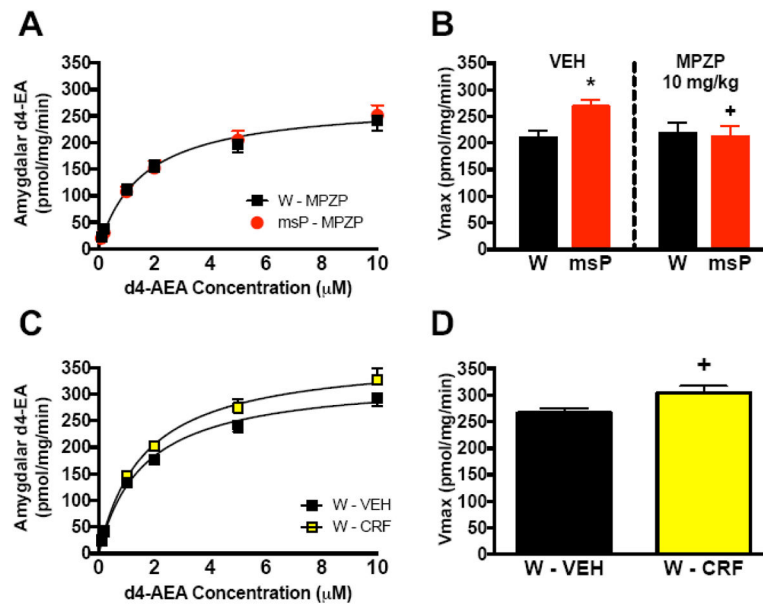


Figure 3. Enhanced CRF₁ receptor signaling drives elevated amygdalar FAAH activity
(A) Substrate conversion of deuterated N-arachidonylethanolamine (d4-AEA) into ethanolamine (d4-EA) in amygdalar tissue collected from non-selected Wistar and msP rats treated with the corticotropin-releasing factor type 1 (CRF₁) receptor blocker N,N-bis(2-methoxyethyl)-3-(4-methoxy-2-methylphenyl)-2,5-dimethylpyrazolo[1,5-a]pyrimidin-7-amine (MPZP, 10 mg/kg, 2 mL/kg, s.c., b.i.d., n=7 per genotype). **(B)** Velocity of reaction (V_{max}) in Wistars and msPs treated with either MPZP (n=7 per genotype) or vehicle (VEH, n=8 per genotype). **(C)** Substrate conversion of d4-AEA into d4-EA in amygdalar tissue collected from Wistars receiving administration of the peptide hormone CRF (1 μg/2 μL, i.c.v., n=7) or VEH (n=8). **(D)** V_{max} rates in rats from C. Data expressed as mean ± SEM. Asterisks (*) denote significant genotype differences ($p < 0.05$), whereas plus signs (+) denote significant differences relative to vehicle controls ($p < 0.05$).

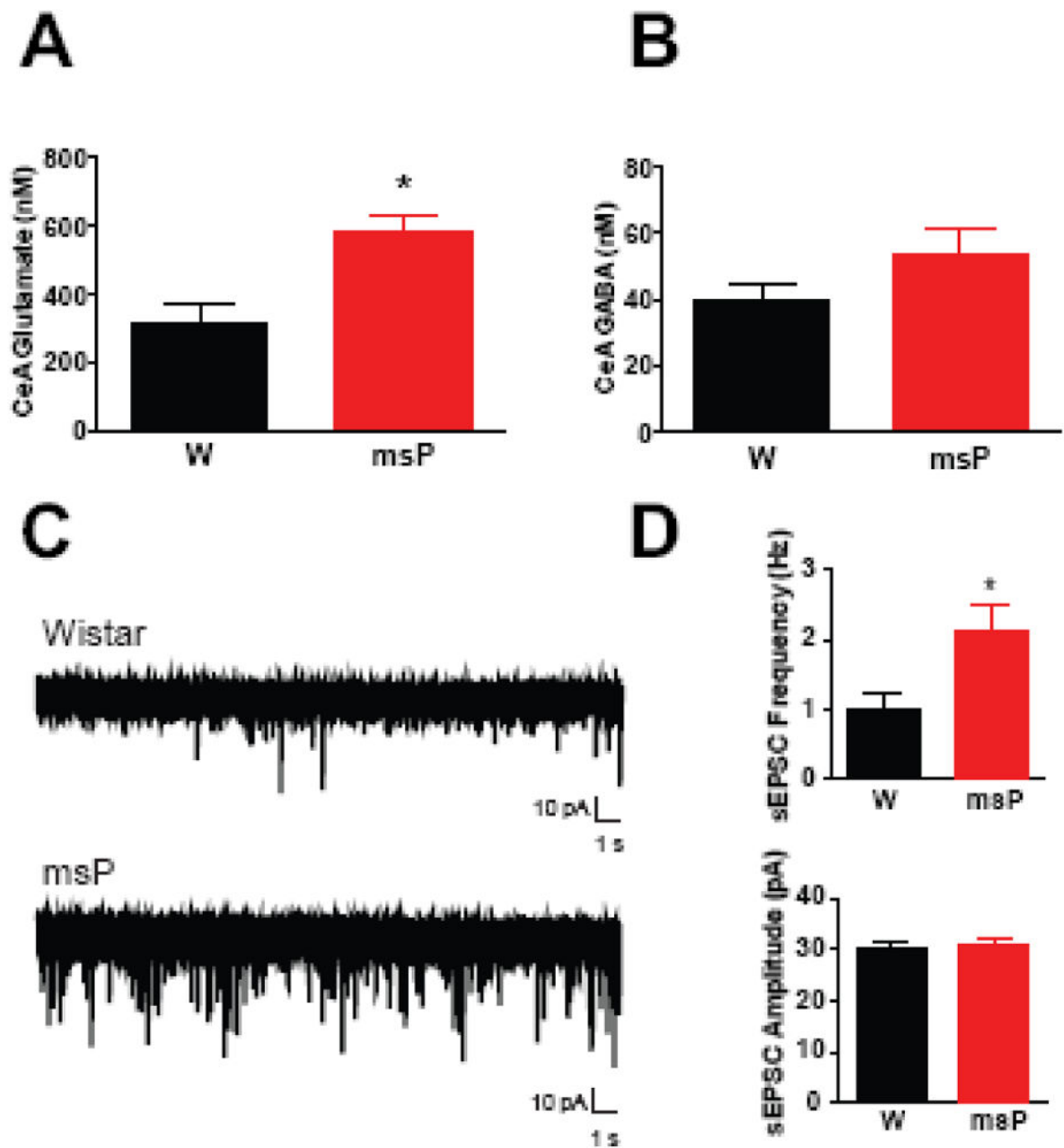


Figure 4. Amygdalar glutamatergic transmission is enhanced in msP rats

(A) Baseline dialysate concentrations of glutamate in the central nucleus of the amygdala (CeA) of non-selected Wistar (W, $n = 9$) and msP ($n = 9$) rats. (B) Baseline dialysate concentrations of gamma aminobutyric acid (GABA) in the CeA of rats from A. (C) Representative whole-cell patch-clamp recordings of spontaneous excitatory postsynaptic potentials (sEPSCs) from Wistars (top trace) and msPs (bottom trace). (D) Baseline frequency (top panel) and amplitude (bottom panel) of CeA sEPSCs in Wistars ($n=15$ neurons from 4 rats) and msPs ($n=16$ neurons from 4 rats). Data expressed as mean \pm SEM. Asterisks (*) denote significant genotype differences ($p < 0.05$).

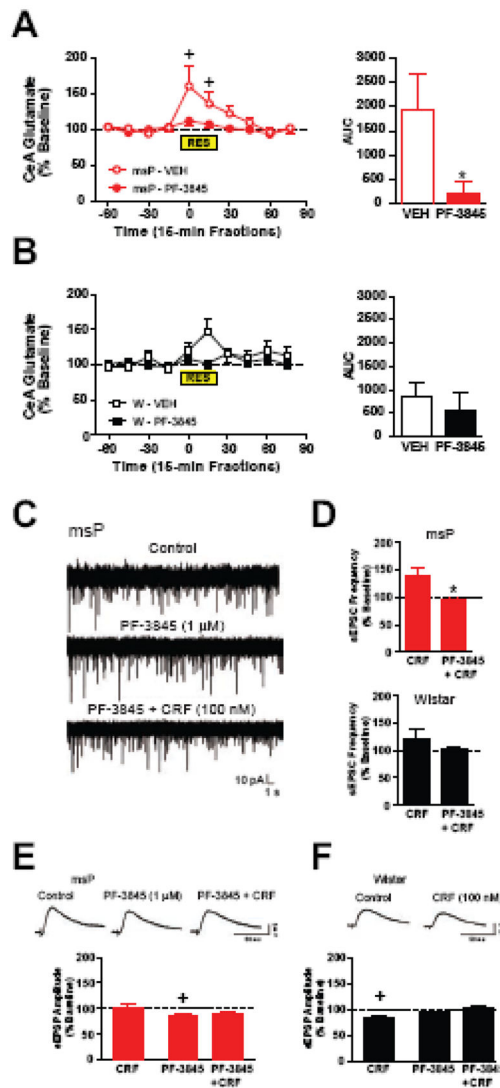


Figure 5. FAAH inhibition alleviates stress-sensitive increases in glutamatergic transmission in msP rats

(A) Percentage change in glutamatergic transmission in the central nucleus of the amygdala (CeA) across time (left panel) in which restraint stress (RES) was applied and measured for accumulative response (area under the curve, AUC, right panel) in msP rats pretreated with the fatty acid amide hydrolase (FAAH) inhibitor PF-3845 (10 mg/kg, i.p., n=9) or vehicle (VEH, n=7). (B) Percentage change in CeA glutamatergic transmission for similar procedures in non-selected Wistar (W) rats pretreated with PF-3845 (n=9) or VEH (n=7). (C) Representative whole-cell patch-clamp recordings of spontaneous excitatory postsynaptic potentials (sEPSCs) in msPs during control (top trace), superfusion of PF-3845 (1 μ M, middle trace), and PF-3845 in combination with the peptide hormone corticotropin-releasing factor (CRF, 100 nM, bottom trace). (D) Baseline frequency of CeA sEPSCs following superfusion of CRF or CRF in the presence of PF-3845 in msPs (n=5–9 neurons from 4 rats, top panel) and Wistars (n=6–8 neurons from 4 rats, bottom panel). (E) Representative BLA-evoked excitatory postsynaptic potentials (EPSPs) from an msP CeA

neuron (top panel) during control, PF-3845 application, and CRF co-applied with PF-3845. Histograms plotting the effects of CRF alone (n=8 neurons from 7 rats), PF-3845 alone (n=16 neurons from 9 rats) and CRF in the presence of PF-3845 (bottom panel, 11/16 neurons) in CeA neurons from msPs. (F) Representative BLA-evoked EPSPs from a Wistar CeA neuron (top panel) during control and CRF application. Histograms plotting the effects of CRF alone (n=9 neurons from 7 rats), PF-3845 alone (n=13 neurons from 7 rats) and CRF in the presence of PF-3845 (bottom panel, 10/13 neurons) in CeA neurons from Wistars. Data expressed as mean \pm SEM. Asterisks (*) denote significant differences relative to PF-3845 treatment, whereas plus signs (+) denote significant differences relative to baseline levels ($p < 0.05$).

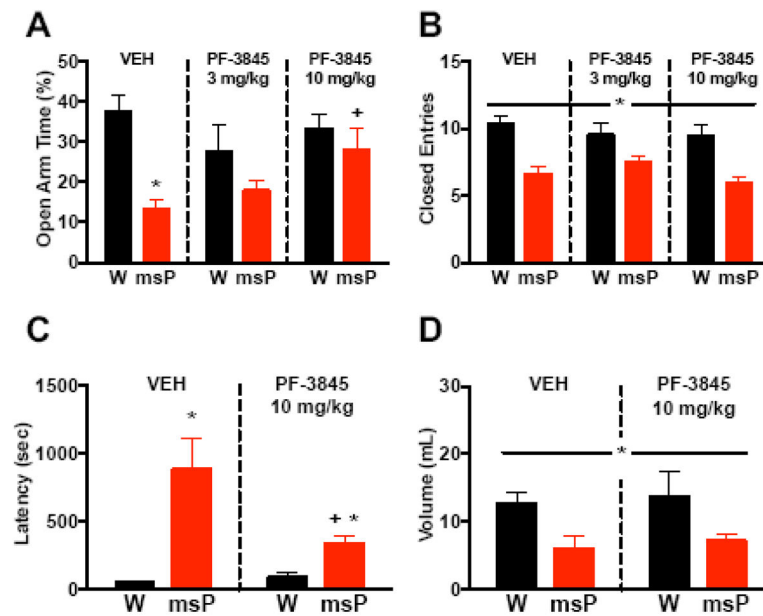


Figure 6. FAAH inhibition alleviates excessive anxiety-like behavior in msP rats

(A) Percentage of time spent on the open arm of the elevated plus maze (EPM) in non-selected Wistar (W) and msP rats pretreated with doses of the fatty acid amide hydrolase (FAAH) inhibitor PF-3845 (3 or 10 mg/kg, i.p., $n=12-15$ per dose and genotype) or vehicle (VEH, $n=19-20$ per genotype). (B) Total entries made into the closed arm of the EPM in rats from A. (C) Latency to drink from a palatable chocolate solution during novelty stress in Wistars and msPs pretreated with PF-3845 (10 mg/kg, i.p., $n=8-9$ per genotype) or VEH ($n=9-11$ per genotype). (D) Total volume of chocolate solution consumed in rats from C. Data expressed as mean \pm SEM. Asterisks (*) denote significant genotype differences, whereas plus signs (+) denote significant differences relative to vehicle controls ($p < 0.05$). A continuous line across groups denotes significant main effects.

Table 1
Neurotransmitter levels in the rat central amygdala

Values reflect mean \pm SEM baseline dialysate concentrations of a broad-scale analysis of neurotransmitters in the central nucleus of the amygdala of non-selected Wistar (n = 9) and msP (n = 9) rats. Asterisks (*) denote significant genotype differences ($p < 0.05$).

Baseline Dialysate (nM \pm SEM)	Wistar	msP	Genotype Comparisons
Aspartate	66.75 \pm 11.54	121.8 \pm 27.9	F(1,16) = 3.3
Dopamine	0.27 \pm 0.14	0.24 \pm 0.04	F(1,16) = 0.04
GABA	40.08 \pm 4.53	53.48 \pm 8.19	F(1,16) = 2.1
Glutamate	317.8 \pm 56.1	579.4 \pm 50.2*	F(1,16) = 12.1*
Glutamine	49.52 \pm 2.72	49.25 \pm 3.33	F(1,16) = 0.01
Glycine	3419 \pm 401	3611 \pm 414	F(1,16) = 0.1
Norepinephrine	0.60 \pm 0.16	0.47 \pm 0.06	F(1,16) = 0.6
Serine	8056 \pm 649	7872 \pm 536	F(1,16) = 0.1
Serotonin	0.09 \pm 0.01	0.07 \pm 0.01	F(1,16) = 3.3
Taurine	2370 \pm 212	2850 \pm 389	F(1,16) = 1.2