Specific and converging modulation of the CRH and noradrenergic systems across drug classes and natural rewards in an incubation of seeking paradigm.

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ABSTRACT

Stress is known to play a critical role in relapse to drug use as well as in food craving. Craving itself is a key determinant of relapse, and cue-induced drug craving has been shown to increase, or 'incubate', over time for certain drugs such as cocaine and nicotine, though this effect is less consistent for others such as opiates. However, the contribution of stress-related biochemical systems to the incubation of seeking has not yet been systematically examined in animal models, nor has the specificity of these mechanisms been tested across different drug classes or reinforcers. To address this gap, we trained rats to self-administer cocaine (0.75 mg/kg, i.v.), heroin (0.075 mg/kg, i.v.), or saline, and subsequently assessed changes in plasma corticosterone, ornithine and other stress-related amines, alongside central gene and protein expression (CRH, CRH2 receptor, and α- and β-adrenergic receptor subunits) after 1 or 30 days of withdrawal. A parallel experiment was conducted using sucrose as a reinforcer. Our findings indicate that although most effects were substance-specific, convergent adaptations were also observed, particularly within noradrenergic systems, suggesting that these mechanisms may represent a common substrate of incubation and a potential target for pharmacological interventions in relapse.

Keywords: alpha adrenergic receptors; beta adrenergic receptors; corticosterone; ornithine; incubation of seeking.

INTRODUCTION

Addiction, or substance use disorder according to the DSM-5-TR terminology, has been frequently defined as a chronic relapsing disorder [1–3]. As such, it is a crucial research goal to obtain a deep and comprehensive understanding of the mechanisms involved in relapse. Among other factors, stress plays a crucial role in relapse into drug use [4–6]. There is ample preclinical literature which indicates that rodents will reinstate their drug seeking behaviour after exposure to different stressors [7]. Moreover, a previous stressful experience can facilitate reinstatement of drug-seeking behaviour [8]. In humans, early clinical reports showed that stress facilitates relapse [6,9], and numerous preclinical studies have examined the neural and endocrine bases of the phenomenon [7]. An important factor driving relapse is craving [10] which is a time-dependent phenomenon. Indeed, while basal craving diminishes over time, cue-induced craving increases ('incubates') over time in cocaine users [11]. Importantly, this phenomenon seems to be absent in opioid users [12] which underscores the importance of comparing across drug classes in incubation studies. While a tremendous effort has been made to search for neurobiological mechanisms of this incubation phenomenon [13] with potential targets for treatment being identified [14], the role of stress is still poorly understood.

In addition to comparing a psychostimulant drug (cocaine) with an opioid drug (heroin) in this work we also introduced a third rewarding substance, sucrose-sweetened water. An incubation phenomenon similar to the one observed for cocaine has also been described for sucrose-sweetened water [15,16] and human studies have also reported incubation of food craving [17,18] which underscores the importance of including this third reward as a comparator and also due to its own significance. We have focused on the corticotrophin releasing hormone system and noradrenergic systems as the main mechanisms of the stress response and due to their implications in craving in humans [19] and the lack of data on the role of these systems in the incubation of seeking across drug classes and different rewards. In addition to measuring central parameters in different brain regions and study their potential correlations as we have done in previous works [20,21] in this work we have also measured peripheral mediators such as corticosterone and different monoacids relevant to metabolic processes involved in the stress response.

We show that incubation of cocaine, heroin, and sucrose seeking involves early, late, and persistent stress-related alterations, largely substance-specific but converging on *Adra2a* gene in the nucleus accumbens core and *Adrb1* in the basolateral amygdala. This

is the first report using the incubation of seeking paradigm focusing on the CRH and noradrenergic systems, filling an important gap in the literature.

MATERIALS AND METHODS

Animals. We used 84 Lewis male rats (300-320 g at the beginning of the experiments) purchased from Harlan International Ibérica. They belonged to a subset of animals from a previous study published by us [21]. Regarding the self-administration pattern, they behaved as a parallel group of animals expressing incubation of seeking with no significant differences. The behaviour of these animals has been described in a previous publication (see Roura-Martínez et al [21] for more details).

Experimental design. One batch of rats were submitted to jugular catheter surgery (for drug or saline self-administration) and the other left intact (sucrose/water self-administration). After ten days of drug/saline or sucrose/water self-administration, the rats underwent 1 or 30 days of forced withdrawal with regular handling. The rats that received intravenous administration of cocaine, heroin, or saline (iv control) were segregated into six groups (3 substances × 2 withdrawal periods; n=8 rats per group), while the rats that received sucrose or water (oral control) were distributed in four groups (2 substances × 2 withdrawal periods; n=9 rats per group).

Surgical procedure. An intravenous polyvinylchloride tubing (0.064 mm i.d.) catheter was implanted into the right jugular vein at approximately the level of the atrium and passed subcutaneously to exit the midscapular region [22]. Surgical procedures were performed under isoflurane gas anaesthesia (5% for induction and 2% for maintenance) and buprenorphine analgesia, and the catheters were flushed daily with 0.5 mL of gentamicin (40 mg/mL) dissolved in heparinized saline in order to prevent infection and to maintain patency (see [21]).

Self-administration. Self-administration sessions were performed in Skinner boxes (Coulbourn Instruments or Med-Associates) and two levers (active and inactive) monitored with Med-PC software. The house light was off during the sessions, and the door of the sound-attenuating cubicle left ajar, allowing some environmental light from the room. Each time the active lever was pressed (fixed-ratio 1), a pump outside the box was switched on for 5 seconds and either the drug or saline solution was infused through the catheter, or

the sucrose solution or water was dispensed into a receptacle placed in between the levers. A cue-light over the active lever switched on for 10 seconds at the same time. Subsequently, there was a time-out period of 40 seconds with no programmed consequences. Cocaine, heroin, or saline self-administration sessions lasted 6 hours per day, as described previously. [21]. Rats orally self-administering sucrose or water were subjected to shorter sessions (2 h/d; [23]). The doses per injection used in the experiments were 0.075 mg/kg of heroin; 0.75 mg/kg of cocaine-HCl; and 10% w/v sucrose (Sigma-Aldrich S1888), diluted in 0.1 mL of saline (0.9%, NaCl physiological saline. To facilitate the acquisition of this behaviour, we placed two sucrose pellets on the active lever in the first two self-administration sessions.

Animal sacrifice and tissue processing. We weighed and sacrificed the rats after 1 or 30 days of abstinence by decapitation, between 11:00 and 13:00. Immediately after, the brain was removed in about one minute, frozen by ten seconds of immersion in isopentane (VWR 24872.298) cooled with dry ice, and stored at -70°C. We collected trunk blood in a heparinized tube and extracted plasma by centrifugation at 1000g for 10 min at 4 °C, then stored at -70°C. Adrenal glands were removed and weighed.

Brain dissection. Brain was tempered 1 h at -20°C in the cryostat chamber (Microm, Cryostat HM 500O) embedded in TissueTek (Sakura, 4583). We collected 300μm thickness slices on a sterile cold surface and dissected them with a lancet, according to a rat neuroanatomical atlas (Paxinos and Watson, 2007). We stored the tissue during dissection in sterile tubes of 0.2 mL in dry ice and then at -70°C.

Tissue homogenization. We weighed and homogenized the samples in HEPES buffer (50 mM, pH 7.5) prepared in DEPC-treated water, containing sucrose (320 mM), protease and phosphatase inhibitors, and sodium butyrate (20 mM) using a motorized mortar (Sigma-Aldrich, pellet pestle Z359971). After being kept on ice for 10 min they were centrifuged at 1000 g for 10 min at 4°C. A volume of supernatant equivalent to 3-4 mg of original tissue (≤80 μL) was added to 800 μL of QIAzol (Qiagen 79306) for extraction of cytoplasmic RNA. Another volume was added to loading buffer (Tris 62.5 mM pH 6.8, glycerol 20% v/v, SDS 2% w/v, dithiothreitol 50 mM) for Western blot. Then, we stored the samples at -70°C.

Level of plasmatic corticosterone by radioimmunoassay. The plasmas were thawed, centrifuged at 10,000 g for 10 min at 4°C to discard precipitates, tempered at room

temperature (RT), and then analysed using the Coat-a-Count® Rat Corticosterone kit (Siemens). Briefly, 50µL of sample or calibrator was added to tubes covered with anti-corticosterone antibody. Then, 1 mL of I125-corticosterone solution was added, vortexed and incubated at RT for 2 h. The content was then decanted and the signal analysed (1 min/tube, in counts per minute, cpm) in a gamma counter (Wallac, Automatic Gamma Counter 1470 Wizard). To estimate the non-specific binding, samples or calibrators in non-covered tubes were measured. The maximum binding value was estimated by measuring a calibrated tube without labelled corticosterone. The corticosterone values were interpolated in a semi logarithmic line facing the logarithm of the concentration with the binding value and expressed in ng/mL.

Level of plasmatic amines by capillary electrophoresis. The content of amines was analysed according to the method described in Lorenzo et al. (2013) [23]. Samples were passed through a 0.22 µm filter (VWR 28145-491 13 mm syringe filter 0.2 µm PTFE membrane) and then stored at -70°C until use. In parallel, cocktails of different amines were prepared to be used as a calibration curve (from Sigma: L-glutamic acid (49449), glycine (G7126), L-glutamine (49419), taurine (T0625), L-serine (S4500), D-serine (S4250), L-proline (81709), L-isoleucine (W527602), L-ornithine (O2375), L-threonine (T8625)). L-2-aminodipidic acid (Sigma, A7275) was used as internal standard (IS). In each cocktail, the concentration of each amine was different, but the total concentration of amines was maintained constant. The derivatization (20 µL sample, 20 µL 200 mM IS, 25 µL 40 mM 4-fluoro-7-nitro-benzofurazan (Alfa-Aesar J61336), 150 µL 10 mM borate buffer pH 10.0) was carried out at 60°C for 15 min in darkness. Then, the samples were stored in the chamber of the capillary electrophoresis apparatus (Beckman Coulter PA 800 plus) at 7°C for at least 30 min before the start of the electrophoresis, carried out at 17°C (electrophoresis buffer: 175 mM borate buffer pH 10.25, 12.5 mM β-cyclodextrin (SAFC W40,282-6)). Before its first use, the capillary column of silica (length: 60cm; inner diameter: 75µm) was conditioned with 1 M NaOH (15 min) and water (15 min), and between electrophoresis with 0.1 M HCl (3 min), water (5 min) and electrophoresis buffer (5 min). The sample was injected at the anode at 0.5 psi (33 mbar) for 10 seconds. A potential difference of 21 kV was applied, observing currents of 120 µA. The molecules were detected at the cathode by means of laser induced fluorescence (LIF), exciting at 488 nm (argon lamp) and detecting the emission at 522 nm. To obtain reproducible electropherograms, the electrophoresis buffer was renewed every six analyses. In each session, a calibration curve was also injected. The electropherograms were analyzed with

the 32 KaratTM software, using the corrected area normalized to the internal standard. The values were expressed as pmol/mL.

Isolation of RNA. The RNA was extracted using a protocol adapted from Chomczynski and Sacchi [24,25] . Briefly, the sample (stored in QIAzol) was thawed and left at RT for at least 5 min. 160 µL of chloroform (Merck 1.02445.2500) were added, mixed for 15 seconds and incubated at RT for 2-3 min. It was then centrifuged at 12 000 g for 15 min at 4°C. The upper (aqueous) phase was transferred to a new sterile 1.5 mL tube, with 400 µL of isopropanol (Fischer Scientific, BP2618) and 10 µg of glycogen (Sigma G1767), mixed and incubated at RT for 10 min. It was centrifuged at 12000 g for 10 min at 4°C and the supernatant discarded. The precipitate was washed twice in 1 mL of cold (-20°C) 75% v/v ethanol and centrifuged at 7500g for 5 min at 4°C, discarding the supernatants. Finally, the precipitate was allowed to air-dry, resuspended in nuclease-free water and stored at -70°C. The RNA was quantified in a Bioanalizer 2100 system (Agilent, RNA Nano Chips, 5067-1511), with values of RIN≥7.0. Unless exceptions, absorbance ratios 260nm/280nm ≥1.8 were obtained. An isolation efficiency of 0.7-1.0 µg RNA/mg tissue was obtained.

RT-qPCR. About 250-500ng RNA (depending on the region) was treated with DNaseI (Invitrogen, 18068-015), retrotranscribed (Biorad, iScript cDNA Synthesis kit, 170-8891), diluted 1:10 in nuclease-free water and stored at -70°C. Gene expression was analyzed using SYBR Green (Biorad, SsoAdvanced Universal SYBR Green Supermix), in a CFX96 C1000 Touch of Biorad system. The total reaction volume was 10µL, and the primers were in the range of 500-750nM. The thermal protocol was as follows: 5 min at 95°C, 40 cycles of 5 min at 95°C, and 30 seconds at 60°C, followed by a melting curve. We calculated the Ct values and the efficiencies with the free software LinRegPCR [26] and then calculated the fold change for each sample and gene according to Pfaffl [27], using *Gapdh* as housekeeping gene. Because adrenoceptors have different (and sometimes opposite) effects in different brain regions, we also calculated ratios between receptors for each region and ratios between regions for each receptor.

Primer list

α1 adrenoceptor (*Adra1).*

S: 5'-ctcgagagaagaaagctgcca-3' A: 5'-aaaacggtttccgaaggcttg-3'

α2A Adrenoceptor (*Adra2a*).

S: 5'-tcctgagagggaagggattt-3' A: 5'-agttactggggcaagtggtg-3'

β1 adrenoceptor (*Adrb1*).

S: 5'-gctctggacttcggtagacg-3' A: '-acttggggtcgttgtagcag-3'

Protein quantification. Protein content was quantified in microplates using the colorimetric method described by Bradford [28], with minor modifications. Briefly, 250 µL of the reagent (a 1:5 dilution of the Bio-Rad Protein Assay Dye Reagent Concentrate 500-0006) was added to 5 µL of the sample. A calibration curve of BSA was made. The 595nm/450nm absorbance ratio (quantified in an Asys Hitech DigiScan) was used to interpolate unknown values.

Western blot. 5 µg of protein from each sample was resolved on 12% SDS-PAGE gels in a MiniProtean® Tetra System (Bio-Rad). To quantify the total protein, the gels were exposed to UV light for about 1 min to activate the reaction between trichloroethanol and tryptophan [29], and they were then recorded on an Amersham Imager 600 (General Electrics). The proteins were transferred to 0.2 µm PVDF membranes (Trans-Blot® TurboTM transfer pack, Bio-Rad) in a Trans-Blot® Turbo transfer system (Biorad), blocked with 5% w/v BSA in TBST for 1 h at RT and then probed overnight in TBST at 4 ºC with the primary anti- antibody anti-CRH (Abcam, ab184238) or anti- CRH receptor 2 (Abcam, ab104368) diluted 1:3.000. Antibody binding was recognized for one hour at RT with an HRP-conjugated anti-rabbit IgG secondary antibody (Abcam ab6721) diluted 1:5,000, and the signal was developed using ECL2 and recorded on an Amersham Imager 600. Proteins were analysed by densitometry using ImageJ free software and the data were normalized by gel according to the criterion described by Degasperi [30].

Missing samples. Due to storage problems, two brains were lost (water wd1 group).

Univariate analysis. We carried out two-way ANOVA (IBM Statistics) to compare specific biochemical parameters (log or inverse transformation when necessary) between groups due to the treatment and/or withdrawal, followed by simple effects analysis (Sydak correction). Following a significant effect of the treatment, we performed the Ryan post hoc procedure (REGWQ) to allocate the differences between groups [31].

Bivariate statistics. To detect correlated variables that could have a psychobiological meaning, a screening of Pearson correlations (IBM Statistics) followed by their respective comparisons between groups was performed. The Fisher transformation was applied to the r-value, and then the Z-value and its p-value were calculated to compare the

correlations between groups. We made two types of comparisons for each substance: a) the wd1 group versus its wd1 control and its wd30 group, and b) the wd30 group versus its wd30 control and its wd1 group. To select the correlations with differences between groups, no multiple comparisons correction was applied but we filtered them on the basis of three criteria: 1) n>5 in the three compared groups; 2) a significant r (p<.05) in the group to be tested; and 3) a significant Z (p<.05) for comparisons.

RESULTS

For all variables, descriptive statistics (N, mean, and SD per group) and inferential results are presented in Tables 1 (saline, cocaine, heroin) and 2 (water, sucrose). Table 3 shows the parameters from the correlation analyses.

A. Stress-related peripheral parameters

Body weight. Differences in body weight were only attributable to the passage of time (WD1 vs WD30), with no effects of drug or sucrose self-administration, nor of their interaction with withdrawal time (Tables 1 and 2).

Adrenal gland index. In parallel with weight gain, the adrenal gland index decreased over time. In addition, this index was higher in the cocaine and heroin groups during early abstinence (treatment × withdrawal day: $F_{(2,42)}=5.68$, p=.007, $\eta^2p=.21$; saline WD1 vs cocaine WD1, p=.001; saline WD1 vs heroin WD1, p<.001) (Fig. 1A and B; See Tables 1 and 2 for further details).

Corticosterone. Plasma corticosterone levels were significantly higher in the heroin group compared with the saline group during early withdrawal. Moreover, a significant effect of withdrawal was observed in the heroin group (Fig. 1C), with a reduction in corticosterone levels over time (treatment × withdrawal day: $F_{(2,42)}=3.32$, p=.046, $\eta^2p=.14$; saline WD1 vs heroin WD1, p=.033; heroin WD1 vs heroin WD30, p=.004). There were no effects in the sucrose/water experiment (Fig. 1D).

Plasma amino acid levels and L-ornithine metabolism. Among the amines analysed, effects were found in those related to L-ornithine metabolism. This amino acid is metabolically linked to L-glutamate, L-glutamine, and L-proline [32], and is also associated with stress (see Discussion). We analysed both the concentrations of these amines individually and the ratios of L-ornithine to each of the others. Cocaine, heroin, or sucrose self-administration did not alter absolute concentrations. However, when examining ratios, we

found reductions in the L-ornithine/L-glutamine ratio for cocaine and heroin at early withdrawal compared with saline (treatment × withdrawal day: $F_{(2,41)}=3.77$, p=.031, η²p=.16; saline WD1 vs heroin WD1, p=.024; cocaine WD1 vs cocaine WD30, p=.008; heroin WD1 vs heroin WD30, p<.001) (Fig. 1E). Similarly, the L-ornithine/L-proline ratio decreased after one month of withdrawal (treatment × withdrawal day: $F_{(2,41)}=3.71$, p=.033, η²p=.15; saline WD1 vs heroin WD1, p=.048; cocaine WD1 vs cocaine WD30, p=.088; heroin WD1 vs heroin WD30, p=.032), with heroin also showing increases at early withdrawal (Fig. 1G). No significant effects were observed in the sucrose/water experiment (Fig. 1F and 1H).

B. Stress-related central parameters

Self-administration effects persisting after withdrawal.

All three rewarding substances tested produced long-lasting effects in the nucleus accumbens. In terms of gene expression, cocaine self-administration was associated with a trend towards decreased *Adra2a* expression levels in the shell (Fig. 2C; treatment: $F_{(2,38)}=3.55$, p=.039, η²p=.16; saline vs cocaine, p=.068) and an increase in the *Adra2a* core/shell ratio (Fig. 2G; treatment: $F_{(2,36)}=3.65$, p=.036, η²p=.17; saline vs cocaine, p=.033). This ratio also increased after sucrose consumption (Fig. 2H; treatment: $F_{(1,19)}=8.18$, p=.010, η²p=.30), accompanied by additional changes in the core: decreases in *Adra1* (Fig. 2B; treatment: $F_{(1,30)}=9.28$, p=.005, η²p=.24) and increases in the *Adra2a/Adra1* ratio (Fig. 2F; treatment: $F_{(1,21)}=7.82$, p=.011, η²p=.27).

We then computed correlations between measured parameters and examined their modulation by treatment and withdrawal (Table 3). In the core, CRH and *Adra2a* levels were negatively correlated in saline groups (saline WD1, r=−.949, p=.001; saline WD30, r=−.779, p=.023). This correlation was abolished following cocaine (WD1: Z=−2.533, p=.011; WD30: Z=−3.079, p=.002) and heroin self-administration (WD1: Z=−2.411, p=.016; WD30: Z=−1.983, p=.047), with a reversal to a positive correlation after one month of cocaine withdrawal (cocaine WD30, r=.771, p=.042; WD1 vs WD30, Z=−1.707, p=.088) (Fig. 5).

In the prefrontal cortex, sucrose self-administration also produced lasting changes in the dmPFC: decreases in *Adrb1* (Fig. 3F; treatment: $F_{(1,30)}=21.40$, p<.001, η²p=.42), *Adra1* (Fig. 3B; treatment: $F_{(1,30)}=9.96$, p=.004, η²p=.25), *Adra2a* (Fig. 3D; treatment: $F_{(1,30)}=8.48$, p=.007, η²p=.22), *Adra1/Adrb1* ratio (Fig. 4F; treatment: $F_{(1,30)}=24.37$, p<.001, η²p=.45), and *Adra1* dmPFC/vmPFC ratio (Fig. 4H; treatment: $F_{(1,30)}=10.14$,

p=.003, η²p=.26); and increases in the *Adra2a/Adra1* (Fig. 4B; treatment: F(1,30)=29.38, p<.001, η²p=.50), *Adra2a* BLA/dmPFC ratio (Fig. 4J; treatment: F(1,22)=8.94, p=.007, η²p=.29), and dmPFC *Adra2a/Adrb1* ratio (Fig. 4D; treatment: F(1,30)=4.53, p=.042, η²p=.13).

Modulations occurring during withdrawal.

Apart from the cocaine WD30 correlation between CRH and *Adra2a* in the accumbens core, treatment × withdrawal interaction effects were mainly observed in the mPFC and BLA. A key common finding across all reinforcers was a central role of *Adrb1* expression in the BLA. For example, sucrose self-administration modified correlations with CRHR2 in the vmPFC (sucrose WD1, r=.907, p=.005; WD1, water vs sucrose, Z=−1.983, p=.047; sucrose WD1 vs WD30, Z=2.697, p=.007), cocaine modified correlations in the dmPFC (cocaine WD30, r=.859, p=.006; WD30, saline vs cocaine, Z=−2.038, p=.042; cocaine WD1 vs WD30, Z=−2.662, p=.008) and in the BLA (cocaine WD30, r=.944, p<.001; WD30, saline vs cocaine, Z=−2.270, p=.023; cocaine WD1 vs WD30, Z=−3.060, p=.002), while heroin modified correlations in the BLA (heroin WD30, r=−.905, p=.005; WD30, saline vs heroin, Z=2.471, p=.013; heroin WD1 vs WD30, Z=1.910, p=.056). *Adrb1* correlations with *Adrb1* in the dmPFC and *Adra2a* in the vmPFC also shifted during withdrawal from both drugs of abuse (Fig. 5; Detailed statistics are provided in Table 3).

After one month of sucrose withdrawal, some dmPFC alterations already evident after treatment became more pronounced: decreases in *Adrb1* (Fig. 3F; treatment × withdrawal day: F(1,30)=5.93, p=.021, η²p=.17; water WD30 vs sucrose WD30, p<.001; sucrose WD1 vs sucrose WD30, p<.001), *Adra1* dmPFC/vmPFC ratio (Fig. 4H; treatment × withdrawal day: F(1,29)=3.90, p=.058, η²p=.12; water WD30 vs sucrose WD30, p=.001; sucrose WD1 vs sucrose WD30, p=.002), and *Adra1/Adrb1* (Fig. 4F; treatment × withdrawal day: F(1,30)=4.68, p=.039, η²p=.14; water WD1 vs sucrose WD1, p=.051; water WD30 vs sucrose WD30, p<.001; sucrose WD1 vs sucrose WD30, p<.001); as well as increases in *Adra2a/Adra1* (Fig. 4B; treatment × withdrawal day: F(1,30)=11.69, p=.002, η²p=.28; water WD30 vs sucrose WD30, p<.001; sucrose WD1 vs sucrose WD30, p<.001). The ratio *Adra1/Adrb1* increased after one month of heroin withdrawal (Fig. 4E; treatment × withdrawal day: F(2,42)=3.34, p=.045, η²p=.14; saline WD30 vs heroin WD30, p=.016).

DISCUSSION

Our aim in this study was to characterise the processes occurring during the incubation of seeking for several rewarding substances (cocaine, heroin, and sucrose) in relation to peripheral and central stress-related parameters. Using a rat self-administration model followed by one month of forced abstinence, we identified changes emerging both at early and later stages of withdrawal, as well as persistent effects. Some of these alterations were shared between two substances, but most were substance-specific. For example, during early drug withdrawal (cocaine and heroin), we observed adrenal hypertrophy and disruptions in ornithine metabolism. Nevertheless, common nodes emerged across all reinforcers: *Adra2a* expression in the nucleus accumbens core (long-lasting effects) and *Adrb1* expression in the basolateral amygdala (late effects). This study addresses a gap in the literature, as few investigations on incubation of seeking have examined peripheral or central stress-related parameters, despite the well-established role of stress in modulating drug seeking.

As early as 1986, Gawin and Kleber hypothesised that induced craving could exhibit an incubation phenomenon [as cited in 34]. However, it was not until the 2010s—approximately a decade after its discovery in animal models—that systematic studies of incubation of craving in humans began. Several publications have examined the evolution of cue-induced craving throughout abstinence in individuals addicted to tobacco [34], methamphetamine [35], heroin [35], alcohol [36], and cocaine [37]. Overall, basal craving and/or withdrawal symptoms decrease as abstinence progresses, whereas cue-induced craving increases during abstinence before subsiding after several months (incubation of craving). Heroin represents an exception, although the only study began assessments after one month of abstinence rather than within the first week. In another investigation, Nava (2006) reported incubation of craving without differentiating between basal and cue-induced forms [38]. Incubation of basal anxiety and/or depressive symptoms has only been observed in methamphetamine and tobacco users, and no studies have investigated cue-induced anxiety.

With regard to non-drug reinforcers, at least three studies reported that basal craving diminishes during abstinence from restrictive diets [39–41], whereas cue-induced food craving tends to be higher among individuals undergoing more restrictive regimes [42]. A related phenomenon, incubation of cue-induced fear responses, has been recognised since the early 20th century [43] and is considered necessary for the manifestation of trauma-related symptoms.

Although this gap in human research concerning the relationship between stress and incubation of craving has not yet been bridged using animal models, some studies have addressed the issue. For instance, stress-induced seeking incubates when footshocks are applied following heroin consumption, but not following sucrose intake [44], and fails to incubate when yohimbine (an ADRA2 antagonist) is used after methamphetamine administration [45]. Resistance to punishment has been shown to incubate after histamine administration in cocaine-exposed animals [46] and after lithium chloride in sucrose-consuming animals [47]. Regarding cue-induced incubation of seeking, a negative state appears to enhance seeking and a positive state to reduce it, but only when the state is temporally proximal to the extinction test—indicating a transient effect. Such findings have been reported with caloric restriction (heroin: [48]), immobilisation (cocaine: [49]), and environmental enrichment (cocaine: [50–52]; sucrose: [53–56]).

Voluntary abstinence protocols have further demonstrated that footshock-induced abstinence prevents incubation of food seeking but not methamphetamine seeking [57–59], and food-induced abstinence prevents incubation of heroin seeking [60] but not methamphetamine seeking [60,61]. The general limitation of these studies is that treatments are rarely compared across both early and late withdrawal. More recently, Nicolas (2022) [62] investigated whether anxiety-like symptoms incubate in a rat model. In the novelty-induced hypophagia test—the only test performed and only in males—no effect was detected. Notably, each group comprised a pool of animals that had or had not undergone extinction, conflating basal and cue-induced effects. Thus, the relationship between stress and cue-induced incubation of seeking remains unresolved.

If behavioural data leave this relationship uncertain, what insights are gained from physiological and biochemical parameters? In drug users, consumption-related cues activate both the autonomic nervous system (ANS) and the hypothalamic–pituitary–adrenal (HPA) axis, with stronger responses in heavy users [63]. Both human and animal studies have shown that drug use itself alters ANS and HPA axis activity [64,65]. For example, heroin consumption suppresses HPA axis activity, whereas cocaine consumption enhances it. Abstinence from both substances induces HPA hyperactivity [66]. Similarly, rats with extended access to cocaine show higher plasma corticosterone levels than those with restricted access (Mantsch, 2014) . In humans, drug users often exhibit a dysfunctional HPA axis, characterised by elevated basal plasma cortisol and exaggerated cortisol responses [64].

By contrast, contingent consumption of natural reinforcers reduces ANS and HPA activation, unless cues are presented in the absence of the reinforcer, in which case both systems are activated [67]. Notably, none of the aforementioned human studies on incubation reported parallel incubation of cue-induced cardiovascular stress. In our rat studies, we observed early adrenal hypertrophy after cocaine and heroin, but not sucrose, self-administration—an effect typically associated with chronic stress [67]—which normalised by one month. Plasma corticosterone levels were transiently elevated following heroin self-administration, consistent with earlier findings for cocaine [68]. Neither we nor Grimm [69] observed such effects after sucrose self-administration (in their case, animals were anaesthetised before blood collection). These findings suggest that, unlike sucrose, drug self-administration under conditions that produce incubation of seeking is capable of activating the HPA axis in the short term.

Additionally, we observed apparent metabolic increases in plasma L-ornithine coinciding with adrenal hypertrophy after drug consumption. This is noteworthy because a homeostatic loop may exist between glucocorticoids and L-ornithine: oral administration of L-ornithine reduces plasma glucocorticoid levels in stressed mice [70] and humans [71], and corticosterone elevation in mice is negatively regulated via L-ornithine metabolism [72]. Our findings on L-ornithine in the cocaine and heroin groups may therefore represent a compensatory response to HPA axis activation.

Further evidence for HPA activation derives from endocannabinoid (eCB) alterations in the same animals [21]. Both chronic stress and corticosterone administration are known to induce changes in central eCB signalling [73–77], typically decreasing anandamide (AEA) and increasing 2-AG. For example, in the basolateral amygdala (BLA), such manipulations produce long-lasting decreases in AEA tone through FAAH upregulation and transient increases in 2-AG [78]. In our experiments, rats showed persistent eCB alterations detectable from the first day of abstinence and lasting at least one month. Both drugs of abuse reduced C*nr1* expression in the dorsomedial prefrontal cortex (dmPFC). Cocaine additionally decreased *Napepld/Faah* expression in the accumbens shell, whereas heroin increased *Faah* in the ventromedial PFC, consistent with reduced AEA signalling. Surprisingly, sucrose consumption also induced changes in the same direction: increased *Dagla/Mgll* in the dmPFC, reduced *Napepld* in the accumbens core, and increased *Faah* and *Cnr1* in the accumbens shell. These findings suggest increased 2-AG and reduced AEA.

Regarding the central noradrenergic system, Hamed and Boguszewski (2018), using a morphine-conditioned place preference model, reported reduced MHPG (the main noradrenaline metabolite) in the nucleus accumbens, but not in the medial prefrontal cortex or amygdala, after prolonged abstinence. In another study, Shepard [79] showed that yohimbine, an ADRA2 antagonist, potentiated methamphetamine seeking but did not produce incubation-like effects across withdrawal days 1, 21, and 51. In our experiments, we observed long-lasting noradrenergic alterations in the nucleus accumbens following all three reinforcers, and late effects after cocaine exposure, primarily involving *Adra2a*. Alterations were also detected in other regions, involving both noradrenergic and corticotropin-releasing hormone receptors. Across all reinforcers, these changes converged on *Adrb1* expression in the basolateral amygdala, linked to distinct receptor networks in this region and the medial prefrontal cortex. Both the prefrontal cortex and BLA receive noradrenaline from the locus coeruleus via the dorsal bundle, yet receptor dynamics differ: ADRA2A, a high-affinity receptor active under low arousal, exerts excitatory effects in the PFC but inhibitory effects in the BLA; conversely, ADRB1, a low-affinity receptor active during stress, is inhibitory in the PFC but excitatory in the BLA [80–83]. The potential role of ADRB1, rather than ADRA2A, in incubation is intriguing given that the late positive potential—a neurophysiological marker of incubating craving in abstinent cocaine users [84]—depends on β-receptor-mediated activation of the amygdala [85].

CONCLUSION

In summary, our findings demonstrate that incubation of drug seeking is accompanied by peripheral and central stress-related alterations that are largely substance-specific but converge on common noradrenergic nodes. Short-term activation of the HPA axis, adrenal hypertrophy, and changes in L-ornithine metabolism highlight the interplay between stress and craving. Persistent adaptations in the endocannabinoid and noradrenergic systems further implicate these pathways in the incubation process. Overall, these results underscore stress-related mechanisms as critical modulators of drug seeking and potential targets for intervention.

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Figure 1

Figure 1. Changes in stress-related peripheral parameters during the incubation of seeking. Levels of adrenal glands index (A, B), plasma corticosterone (C, D), and L-ornithine metabolism ratios (E-H) during cocaine and heroin (A, C, E, G) or sucrose (B, D, F, H) withdrawal. Individual values are presented as well as the mean ± SEM. Differences relative to the control group, *p<0.05, **p<0.01, ***p<0.001; differences relative to the same treatment on different withdrawal days, *f* p<0.05, *ff* p<0.01, *fff* p<0.001; general differences between the two withdrawal days, Δ p<0.05, ΔΔ p<0.01, ΔΔΔ p<0.001.

Figure 2. Changes in gene expression of adrenoceptors in the nucleus accumbens during the incubation of seeking. Levels of *Adra1* NAc core (A, B), *Adra2a* NAc core (C, D), *Adra2a*/*Adra1* NAc core (E, F), and *Adra2a* NAc core/shell (G, H) during cocaine and heroin (A, C, E, G) or sucrose (B, D, F, H) withdrawal. Individual values are presented as well as the mean ± SEM. Differences relative to the control group, *p<0.05, **p<0.01, ***p<0.001.

Figure 3. Changes in gene expression of adrenoceptors in the dorsomedial prefrontal cortex during the incubation of seeking. Levels of *Adra1* (A, B), *Adra2a* (C, D), and *Adrb1* (E, F) during cocaine and heroin (A, C, E) or sucrose (B, D, F) withdrawal. Individual values are presented as well as the mean ± SEM. Differences relative to the control group, *p<0.05, **p<0.01, ***p<0.001; differences relative to the same treatment on different withdrawal days, *f* p<0.05, *ff* p<0.01, *fff* p<0.001; general differences between the two withdrawal days, Δ p<0.05, ΔΔ p<0.01, ΔΔΔ p<0.001.

Figure 4

Figure 4. Changes in gene expression of adrenoceptors in the dorsomedial prefrontal cortex during the incubation of seeking. Levels of *Adra2a*/*Adra1* dmPFC (A, B), *Adra2a*/*Adrb1* dmPFC (C, D), *Adra1*/*Adrb1* dmPFC (E, F), *Adra1* dmPFC/vmPFC (G, H), and *Adra2a* BLA/dmPFC (I, J) during cocaine and heroin (A, C, E, G, I) or sucrose (B, D, F, H, J) withdrawal. Individual values are presented as well as the mean ± SEM. Differences

relative to the control group, *p<0.05, **p<0.01, ***p<0.001; differences relative to the same treatment on different withdrawal days, *f* p<0.05, *ff* p<0.01, *fff* p<0.001; general differences between the two withdrawal days, Δ p<0.05, ΔΔ p<0.01, ΔΔΔ p<0.001.

Figure 5. Heatmap of correlations between different parameters. For each pair of variables that experienced changes, we report the r value of the correlation for each experimental group. A bold value indicates a significant r value; a boxed value indicates a significant change relative to their control groups (control treatment and abstinence day).

Figure 6. General scheme summarizing the perdurable (A, D), early (B, E), and late (C, F) effects in central (A-C) and peripheral (D-F) parameters after the incubation of cocaine (red), heroin (green), and sucrose (yellow) seeking. Individual molecules are depicted with a circle; ratios are depicted with a double headed arrow linking both parameters; correlations are depicted with a line connected both parameters. A grey line indicates a direct metabolic pathway between metabolites (Q: L-glutamine; E: L-glutamate; P: L-proline; O: L-ornithine).

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Figure 1

Figure 2

Figure 3

Figure 4

Figure 5

	wd1			wd30			wd1		wd30	
	saline	cocaine	heroin	saline	cocaine	heroin	water	sucrose	water	sucrose
Adrb1 BLA x CRHR2 vmPFC	0,233	0,069	0,466	0,059	0,474	0,176	0,107	**0,907**	0,153	-0,500
Adrb1 BLA x CRHR2 dmPFC	0,041	-0,375	-0,434	0,001	**0,859**	0,262	0,398	-0,009	0,477	-0,234
Adrb1 BLA x CRHR2 BLA	-0,147	-0,432	-0,212	0,245	**0,944**	**-0,905**	-0,267	0,054	0,001	-0,554
Adrb1 BLA x *Adrb1* dmPFC	0,182	0,628	-0,198	0,166	**-0,939**	**0,696**	0,139	-0,367	0,711	-0,309
Adrb1 BLA x *Adra2a* vmPFC	-0,524	0,281	**-0,642**	0,290	**-0,753**	0,323	0,067	-0,205	-0,342	0,141
CRHR2 BLA x CRHR2 dmPFC	0,235	**0,873**	0,339	0,110	**0,826**	0,029	-0,210	**0,915**	0,244	0,321
CRH core x *Adra2a* core	**-0,949**	-0,122	-0,202	**-0,779**	**0,771**	0,208	0,809	0,378	-0,175	-0,122