

Extinction of Cocaine Self-Administration Produces a Differential Time-Related Regulation of Proenkephalin Gene Expression in Rat Brain

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The purpose of this study was to examine the time course effects of extinction of cocaine self-administration behavior on proenkephalin (PENK) gene expression in caudate-putamen nucleus (ST), nucleus accumbens (Acc), olfactory tubercle (Tu), piriform cortex (Pir), ventromedial hypothalamic nucleus (VMN), and central amygdala (Ce) as measured by in situ hybridization histochemistry. Seventy-two littermate male Lewis rats were randomly assigned in triads to one of three conditions: (1) contingent intravenous self-administration of 1 mg/kg/injection of cocaine (CONT); (2) noncontingent injections of either 1 mg/kg/injection of cocaine (NONCONT); or (3) saline yoked (SALINE) to the intake of the self-administering subject. The self-administering rats were trained to self-administer cocaine under a FR5 schedule of reinforcement for a minimum of 3 weeks. After stable baseline levels of drug intake had been reached, saline was substituted for drug. Following this first extinction period, cocaine self-administration was reinstated for an additional period of 2

weeks. Immediately after cessation of the last session of cocaine self-administration (day 0) and 1-, 5-, and 10-day after the second extinction period, animal brains in each triad were removed to be processed for in situ hybridization. PENK mRNA levels were significantly higher in the cocaine groups when compared with SALINE group in the ST, Acc, Pir, and Tu regions on days 0, 1, 5, and 10 of the extinction and lower in the Ce region of CONT group when compared to NONCONT and SALINE groups on days 1, 5, and 10 of the extinction period. In the VMN nucleus, PENK mRNA content in CONT group versus NONCONT and SALINE groups was also lower, but there were statistically significant differences only on day 5. These results suggest that changes in PENK gene expression after contingent cocaine administration might be involved in cocaine withdrawal states.

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Cocaine abuse and dependence remain major public health and social problems. Cocaine abuse results from a complex interplay of behavioral, pharmacological, and neurobiological determinants. The main pharmacological effect of cocaine is to inhibit the reuptake of monoamines dopamine, norepinephrine, and serotonin at presynaptic terminals. As a consequence of these ac-

tions, cocaine potentiates neurotransmission of all three monoamines (Hadfield et al. 1980; Heikkila et al. 1975; Ross and Renyi 1969). The major behavioral effect of cocaine is a psychomotor stimulant action with reinforcing addictive properties. This behavioral effect is produced primarily by inhibition of dopamine uptake, thereby increasing extracellular concentrations of released dopamine (Ritz et al. 1987; for a review see Kuhar et al. 1991; Spanagel and Weiss 1999). However, a recent study using mice lacking dopamine transporter proposed other mechanisms, such as increases in serotonin transmission, may mediate the reinforcing effects of cocaine (Rocha et al. 1998).

Several studies have shown that cocaine also affects the expression of opioid peptides and opioid receptors. For example, chronic cocaine administration leads to increases in circulating β -endorphin levels (Moldow and Fischman 1987), striatal preprodynorphin mRNA levels (Hurd et al. 1992; Daunais et al. 1993; Daunais and McGinty, 1995; Spangler et al. 1993, 1996, 1997), and striatonigral dynorphin content (Sivam 1989; Smiley et al. 1990) in rats and to decreases in preproenkephalin mRNA levels in rhesus monkeys (Daunais et al. 1997) and humans (Hurd and Herkenham 1993). Repeated administration of cocaine can also regulate the activity of opioid receptors in discrete brain regions of rats. Indeed, it has been shown that 2 weeks of either continuous administration of cocaine via subcutaneously implanted osmotic mini-pumps (Hammer 1989) or repeated daily injections (Unterwald et al. 1992, 1994) lead to increase in μ and κ -opioid receptor density in terminal fields of the nigrostriatal and mesolimbic-mesocortical dopamine systems (for a review see Izenwasser 1998).

Identification of the alterations in brain function by chronic cocaine administration has been a focus of many research efforts over the last two decades (for review see Nestler 1993; White et al. 1995; Pierce and Kalivas 1997). This effort has consistently revealed the induction of many changes in cellular processes, although the data sometimes appear contradictory. Some of the discrepancies found on neuroadaptations to cocaine administration may be related to differences in the duration of the treatment, dosage of cocaine administered, time after the last drug administration, or the animal model employed. In this respect, the majority of studies investigating cocaine-induced cellular neuroadaptations use repeated noncontingent administration of this drug; whereas, fewer studies employ drug self-administration. However, long-term neuroadaptations to chronic cocaine administration could arise not only from the pharmacological actions of cocaine on neurotransmitter systems but also by an interaction between this pharmacological action and the animal's environment (Dworkin et al. 1992, 1995; Wilson et al. 1994; Hemby et al. 1997). Indeed, it has been suggested that in humans the classical conditioning of the pharmaco-

logical actions of cocaine with environmental stimuli plays an important role in long-term addictive potential of this drug and in the liability to relapse in recovering cocaine addicts (Ehrman et al. 1992).

The purpose of the present study was to examine the time course effects in proenkephalin (PENK) gene expression after cocaine self-administration and its extinction using a yoked-box procedure. This behavioral experimental design involves the use of triads to assess relative differences related to contingent versus noncontingent presentations of environmental events (Smith and Dworkin 1986). PENK mRNA levels were measured by *in situ* hybridization histochemistry in several rat brain regions immediately after the last day of cocaine self-administration and 1, 5, and 10 days after extinction.

MATERIALS AND METHODS

Animals

Adult male Lewis rats (CRIFFA, France) weighing approximately 300–350 g at the beginning of their training were used. All animals were experimentally naive, housed individually in a temperature-controlled room (23°C) with a 12-h light–dark cycle (08:00–20:00 lights on), and given free access to Purina laboratory chow and tap water before initiation of the experiments. Animals used in this study were maintained in facilities according to European Union Laboratory Animal Care Rules.

Surgery

Experimentally naive subjects were surgically prepared with an IV catheter placed in the jugular vein. Polyvinylchloride tubing (0.064 i.d.) was implanted in the right jugular vein approximately at the level of the atrium under ketamine and diazepam anesthesia. The catheter was passed SC and exited in the midscapular region. The catheter then passed through a spring tether system (Alice King, Chatham, MA) mounted to the skull of the rat with dental cement. All subjects were housed individually following surgery and given at least 7 days to recover.

Apparatus

Twelve operant chambers (Coulbourn Instruments, Allentown, PA, USA) were used for cocaine self-administration studies. Two levers designed to register a response when 3.0 g of force was applied were placed 14 cm apart on the front wall of the chamber. A microliter injection pump (Harvard 22) was used to deliver IV saline or drug injections to the rat. Drug delivery, operant data acquisition, and storage were accomplished on IBM computers (Med Associates, USA).

Experimental Procedure

Cocaine-reinforced behavior was studied according to a procedure described previously (Ambrosio et al. 1996). Briefly, before surgical implantation of the IV catheter, animals were trained to lever press for food reinforcement under a fixed ratio (FR) 5 schedule of reinforcement. Initially, a single lever press on the left-hand lever resulted in delivery of a food pellet (45 mg, P.J. Noyes Company, Lancaster, NH, USA) and turned on a stimulus light above the lever. After responding was initiated, the response requirement to food delivery was raised in increments to 5, and a programmed 30-s time-out (TO) period in which responses had no programmed consequences followed each food pellet delivery (FR5:TO 30 s). When behavior was maintained under the FR5 schedule of food-reinforced behavior, the catheter was surgically implanted as described.

After the post-operative period, 72 litter mate male Lewis rats were randomly assigned in triads to one of three conditions: (1) contingent IV self-administration of 1 mg/kg/injection of cocaine (CONT); (2) noncontingent IV injections of either 1 mg/kg of cocaine (NONCONT); or (3) saline (SALINE) yoked to the intake of the self-administering subject. Initially, substitution of food delivery by cocaine began under a FR1 schedule of reinforcement and was subsequently raised to FR5. In this case, a programmed 30-s time out in which responses had no programmed consequences followed each cocaine injection (FR5:TO 30 s). Animals were allowed to self-administer cocaine in daily 2-h sessions between 9:00 and 14:00 h, 7 days a week for a minimum of 3 weeks. After stable behavior was established (less than 10% of variability in the number of injections for 5 sequential days), saline was substituted for a minimum of 3 days in the operant chambers. After this first extinction period, cocaine self-administration behavior was reinstated and maintained for an additional minimum period of 2 weeks. Saline substitution was carried out again for 1, 5, and 10 days in the operant chambers. Immediately after the last day of cocaine self-administration in which the animals reached the stability criterion (DAY 0) and after 1-day (DAY 1), 5-day (DAY 5), and 10-day (DAY 10) period of extinction, animal brains of each triad were removed to be processed for ISHH. Subjects in each group of triads was 18 (CONT = 6; NONCONT = 6; SALINE = 6), and the total amount of brains removed was 72 [4 groups of triads (DAY 0, DAY 1, DAY 5, and DAY 10) \times 18].

In Situ Hybridization Histochemistry (ISHH)

Brain sections were cut at 20 μ m at three different levels (six slides/level; two sections/slide) containing the brain regions of interest (ROIs). The first level includes nucleus accumbens (Acc), caudate putamen (ST), piriform cortex (Pir), and olfactory tubercle (Tu). The second level contains central amygdala (Ce), and the third level contains ventromedial hypothalamic nucleus (VMN). All these sections were obtained according to Paxinos and Watson Atlas (Paxinos and Watson 1986) [RREFOL], mounted onto gelatin-coated slides and stored at -80° C until the day of the assay.

ISHH was performed as described previously (Young et al. 1986) using synthetic 48-bases oligonucleotide probes complementary to PENK (NEN-Dupont, NEP-502, Madrid, Spain). The oligonucleotide probe was labeled using terminal deoxytransferase (Boehringer, Madrid, Spain) to add a 35 S-labeled deoxyATP (1000 Ci mmol $^{-1}$; Amersham, Madrid, Spain) tail to the 3' end of the probes. The probe (in 50 μ l of hybridization buffer) was applied to each section and left overnight at 37°C for hybridization. Following hybridization, sections were washed four times for 15 min each in 0.15 M NaCl, 0.015 M sodium citrate, pH 7.2 (1X saline sodium citrate, SSC) at 55°C, followed by two 30-min washes in 1X SSC at room temperature, one brief water dip, and were then blown dry with air. Slides from all experimental groups were classified in three sets (each one corresponding to a different level of brain sectioning and containing the areas of interest). To control for imaging enhancement variables, each set of slides was apposed to the same film (Kodak BioMax MR-1, Amersham, Madrid, Spain) in individualized cassettes 3 days for the level containing Acc, ST, Pir, and Tu 7 days for the level containing VMN and 14 days for the level containing Ce.

Two slides per level (two slices/slide; two measurements/slice) in each animal were analyzed with a Macintosh computer using the public domain NIH Image program (developed at the U.S. National Institutes of Health and available on the Internet at <http://rsb.info.nih.gov/nih-image>). Brain regions positive for PENK mRNA hybridization were identified according to the previous reported PENK mRNA distribution in the rat brain (Harlan et al. 1987). Previous experiments in our laboratory have found that the selected times of exposure to film, in these brain regions, and under our experimental conditions (oligonucleotide probe, radioactivity added to each slide, incubation conditions, type of film selected), renders a hybridization signal whose gray levels are linear with the optical density, according to the NIH Image Program. Therefore, optical densities were calculated from the uncalibrated mode of the Image Program by subtracting from each measurement its corresponding background and expressed in gray-scale values. The background measurement was taken from an area of the slice with the lowest nonspecific hybridization signal and subtracted from the hybridization signal measurement in the same slice. Measurements were pooled from brain sections, and the values were averaged. Results were presented considering mean control values as 100%.

Additional brain sections were cohybridized with a 100-fold excess of cold probes or with RNase to assert the specificity of the signal. As expected, no hybridization signal was detected in these sections (data not shown).

Statistical Analyses

The analysis of the PENK gene expression data was performed using a two-way analysis of variance (ANOVA) with the type of cocaine administration (CONT, NONCONT, or SALINE) serving as one independent variable and the brain area serving as the other independent variable. Subsequently, a one-way ANOVA followed by Student–Newman–Keul's test was performed for each brain region and day of extinction (0, 1, 5, and 10), with the type of cocaine administration (CONT, NONCONT, or SALINE) serving as independent variables. Differences were considered significant if the probability of error was less than 5%.

RESULTS

In Figure 1, we present the behavioral data of the contingent, noncontingent, and saline groups used in our study. These data really represent responding of the contingent group only, because the noncontingent and saline groups did not have the possibility of responding; in both groups the subjects received cocaine (1 mg/kg/injection) or saline passively; whereas, the contingent animals were able to self-administer the drug. In Figure 1, the behavioral data presented are divided in four consecutive phases. Phase 1 corresponds to acquisition and maintenance of cocaine IV self-administration behavior; phase 2 to first extinction of cocaine IV self-administration behavior; phase 3 to re-acquisition; and phase 4 to second extinction. The data depicted in Figure 1 clearly show that cocaine maintained robust responding, as compared to the responding observed with saline (first and second extinction phases). The last 15 days of phase 1 (acquisition and maintenance) were very similar to the 15 days of phase 3 (re-acquisition). In

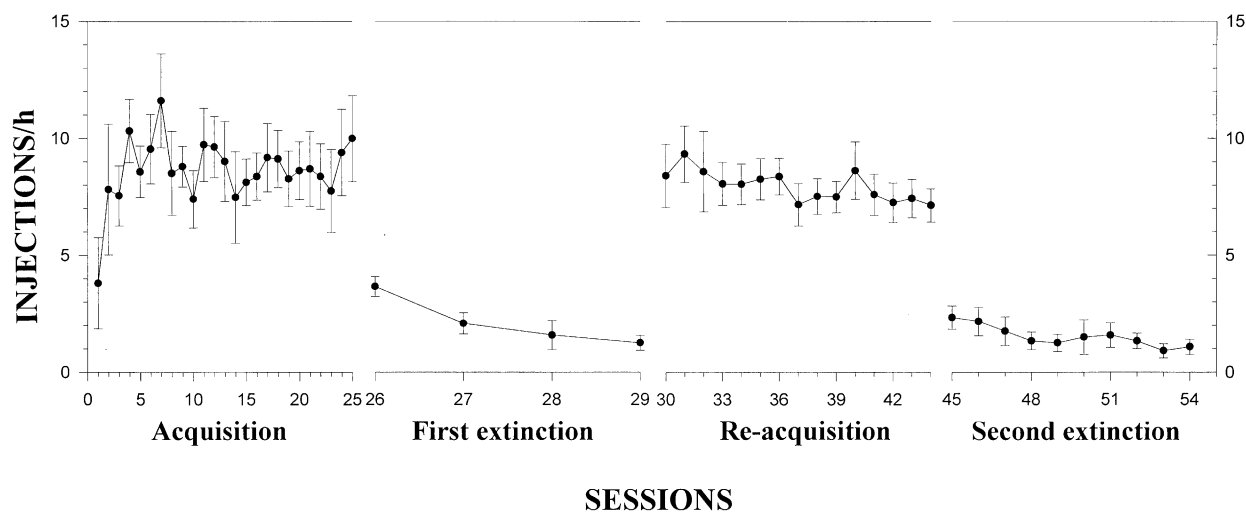


Figure 1. Cocaine self-administration behavior and its extinction in contingent, noncontingent, and saline groups. The behavioral data correspond to the responding of cocaine (1 mg/kg/infusion) contingent self-administering animals, because noncontingent and saline subjects received passively cocaine (1 mg/kg/infusion) or saline when the contingent animals self-administer. The behavioral responses are presented divided in four consecutive phases: phase 1 corresponds to the acquisition and maintenance of cocaine IV self-administration behavior; phase 2 to the first extinction of cocaine IV self-administration behavior; phase 3 to the reacquisition; and phase 4 to the second extinction. When animals reached a stable cocaine self-administration behavior (less than 10% of variability in the number of injections for 5 sequential days) during acquisition and maintenance phase, saline substitution of the first extinction period was carried out. Subsequently, subjects reacquired cocaine self-administration behavior and were maintained self-administering it for 2 weeks. When the animals again reached the stability criterion, a second extinction period started for 10 days. Cocaine maintained robust responding as compared to the responding under saline (first and second extinction phases). The responding during the last 15 days of acquisition phase is very similar to that of the 15 days of reacquisition phase. In both acquisition phases, the number of cocaine injections/h was clearly higher than the number of saline injections/h. In addition, both extinction phases were similar, and the extinction of cocaine self-administration behavior occurred in all the subjects in a very consistent manner. In the days 0 (last day of cocaine IV self-administration), 1, 5, and 10 of the second extinction phase brain animals of the triads ($n = 18$ in each triad; contingent, $n = 6$; noncontingent, $n = 6$; saline, $n = 6$) were removed to be processed for ISHH. See details in the text. *Ordinates:* Number of cocaine or saline injections per hour in 2-h sessions of cocaine or saline IV self-administration behavior. *Abscissas:* Consecutive phases and sessions of cocaine or saline IV self-administration behavior.

both phases of acquisition, the number of cocaine injections/hour was higher than the number of saline injections/hour. Both extinction phases were very similar too, and it is clear from the figure, that extinction occurred in all the subjects in a very consistent manner. This behavioral responding pattern suggests that cocaine was serving as a positive reinforcer in our experimental conditions.

A two-way ANOVA was performed to analyze differences in PENK gene expression during extinction of cocaine self-administration behavior considering the type of cocaine administration and brain region as independent variables. There was a significant main effect of type of administration ($F(2,17) = 107.69, p < .00001$), brain region ($F(5,17) = 461.90, p < .00001$), and type of administration \times brain region interaction ($F(10,17) = 18.71, p < .00001$) on PENK gene expression during extinction of cocaine self-administration behavior.

An over-all view of time course effects of extinction of cocaine self-administration on PENK mRNA levels is shown in Figure 2. Extinction from cocaine self-admin-

istration produces a significant increase of PENK mRNA levels in CONT and NONCONT animals in the ST, Acc, TU, and Pir when compared with each corresponding saline group. By contrast, in the VMN and Ce, extinction from cocaine self-administration causes a reduction of PENK mRNA levels in the CONT group with respect to saline or to NONCONT group. No statistical differences were found between the different time points of extinction of cocaine self-administration in the saline group of all brain regions examined.

The detailed one-way ANOVA analysis revealed that PENK mRNA levels in the ST are different in the last day of self-administration session (DAY 0) and during the days of extinction period (DAY 1, DAY 5, and DAY 10). PENK mRNA levels in both cocaine groups significantly increased when compared to SALINE group in DAY 0 ($F(2,12) = 12.8, p < .0011$), DAY 1 ($F(2,13) = 21.12, p < .0001$), DAY 5 ($F(2,13) = 45.2, p = .0001$), and DAY 10 ($F(2,13) = 48.0, p < .0001$) of our study. No statistical differences were found between CONT and NONCONT groups.

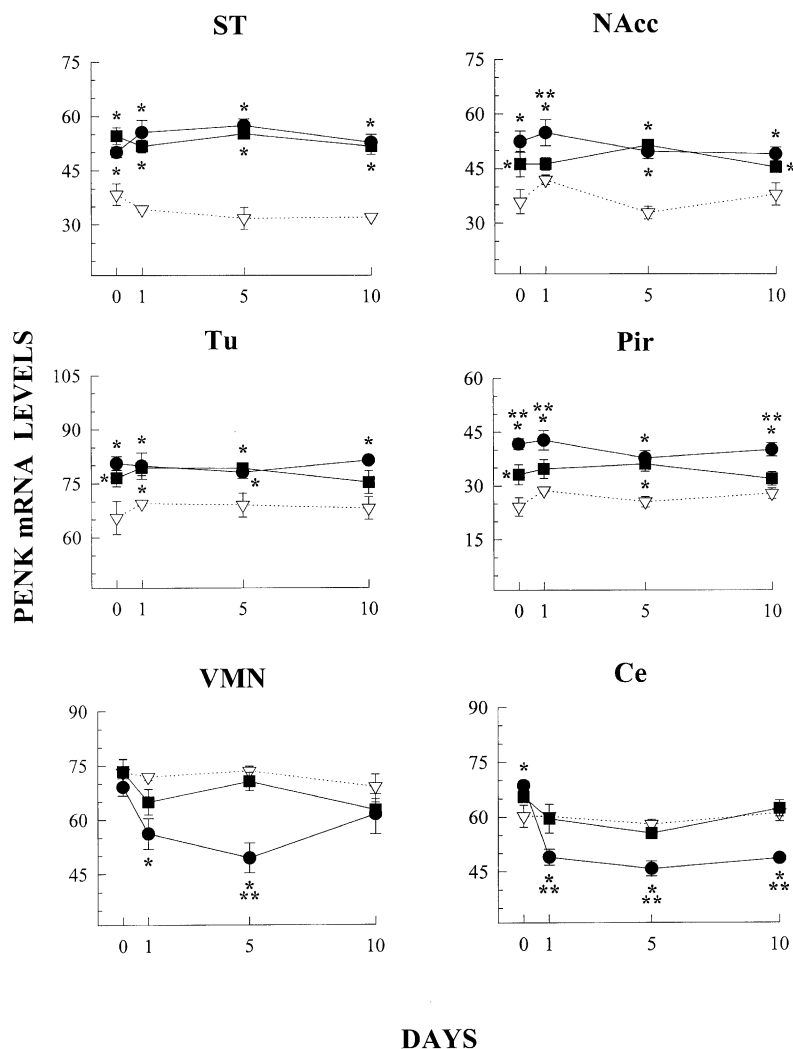


Figure 2. Time course effects of extinction of cocaine self-administration on PENK mRNA levels in caudate putamen (ST), nucleus accumbens (Acc), olfactory tubercle (Tu), piriform cortex (Pir), ventromedial hypothalamic nucleus (VMN), and central nucleus of the amygdala (Ce) in the rat brain of SALINE, CONTINGENT, and NONCONTINGENT groups. Experimental design has been described in Figure 1 and in the Material and Methods section. Symbols represent the means and vertical lines ± 1 SEM of PENK mRNA levels (arbitrary units of optical density) in 4–6 rats. *, values from CONTINGENT or NONCONTINGENT animals that are significantly different ($p < .05$) from SALINE group. **, values from CONTINGENT animals that are significantly different ($p < .05$) from NONCONTINGENT group.

The levels of PENK mRNA in Acc of the three groups differed during the extinction of cocaine self-administration. PENK mRNA levels in the CONT group were significantly higher than NONCONT and SALINE groups on DAY 1 ($F(2,11) = 8.40, p < .0061$) of extinction. Both cocaine groups also showed higher PENK mRNA levels than SALINE on DAY 0 ($F(2,12) = 6.52, p < .0121$), DAY 5 ($F(2,13) = 31.7, p < .0001$), and DAY 10 of extinction period ($F(2,13) = 6.87, p < .0092$). No statistical differences were found between NONCONT and SALINE groups on DAY 1 of extinction.

The levels of PENK mRNA in the Tu region of both cocaine groups were significantly higher than SALINE group across all the days of extinction DAY 0 ($F(2,12) = 6.01, p < .0155$), DAY 1 ($F(2,13) = 4.74, p < .0284$), DAY 5 ($F(2,14) = 6.43, p < .0105$), DAY 10 ($F(2,11) = 6.69, p < .0126$), although no differences were found between cocaine groups.

In the Pir region, PENK mRNA levels in both cocaine groups were higher than in SALINE group DAY 0 ($F(2,13) = 12.10, p < .0011$), DAY 1 ($F(2,13) = 9.64, p < .0027$), DAY 5 ($F(2,15) = 12.35, p < .0007$), and DAY 10 ($F(2,15) = 12.44, p < .0007$). In addition, the levels of PENK mRNA in the CONT group were significantly

higher than those found in the NONCONT group on all days of extinction in this brain region.

As mentioned previously, PENK mRNA levels in ST, Acc, Tu, and Pir increased during the extinction of cocaine self-administration. However, in the VMN and Ce, the results revealed a significant decrease in PENK mRNA levels in CONT group when compared to NONCONT and SALINE groups. Indeed, PENK mRNA levels in the VMN of the hypothalamus were lower on DAY 1 ($F(2,9) = 5.12, p < .032$) and DAY 5 ($F(2,10) = 20.03, p < .0003$) of extinction of CONT group, as compared to NONCONT and SALINE groups. No differences were found between CONT on DAY 0 or DAY 10 and NONCONT or SALINE groups, respectively. Similarly, in the Ce region mRNA levels markedly decreased in the CONT group across all the days of extinction, as compared to the levels of NONCONT and SALINE groups DAY 0 ($F(2,15) = 4.23, p < .034$), DAY 1 ($F(2,12) = 5.77, p < .017$), DAY 5 ($F(2,12) = 18.78, p < .0002$), and DAY 10 ($F(2,12) = 16.32, p < .0004$). There were not statistically significant differences between NONCONT and SALINE groups.

Figure 3 shows representative autoradiograms of the groups SALINE, CONT, and NONCONT used to ana-

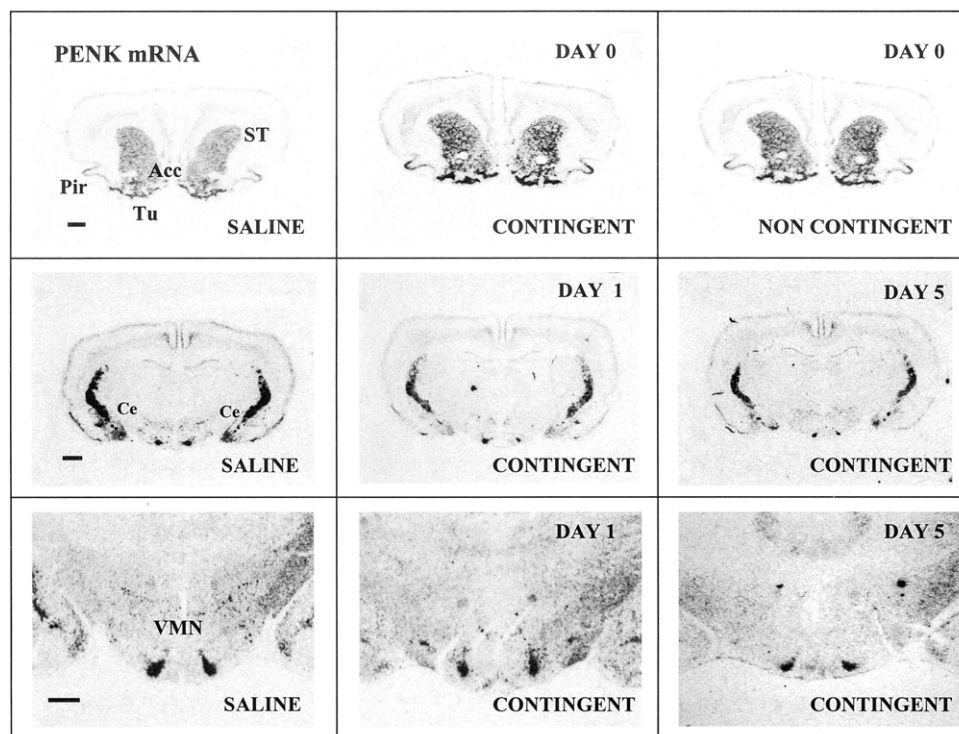


Figure 3. Representative autoradiograms of coronal brain sections at the levels caudate-putamen (ST), nucleus accumbens (Acc), olfactory tubercle (Tu), piriform cortex (Pir), ventromedial hypothalamic nucleus (VMN), and central nucleus of the amygdala (Ce) of the time course effects of extinction of cocaine self-administration on PENK mRNA levels in SALINE, CONTINGENT, and NONCONTINGENT groups. Slides were apposed to film (Kodak, Biomax, MR-1) 3 days for ST, Acc, Tu, and Pir, 7 days for VMN, and 14 days for Ce. Experimental design was as indicated in Figure 1 and in the Material and Methods section. Bar represents 1 mm.

lyze PENK mRNA levels in the ST, Acc, Pir, Tu, Ce, and VMN after the extinction of cocaine self-administration. In the top row, autoradiograms corresponding to CONT and NONCONT animals in DAY 0 presented a higher intensity in the hybridization signal in ST, Acc, TU, and Pir than the one corresponding to the SALINE group. In contrast, in the middle and bottom rows of Figure 3, it is clear that the hybridization signal in Ce and VMN in CONT animals of DAY 5 is lower than in the SALINE group. The optical densities of the hybridization signal in these autoradiograms have been measured, and the values are similar to mean values of their respective experimental groups.

DISCUSSION

In this work, *in situ* hybridization was used to study the effect of long-term cocaine self-administration and its extinction on PENK gene expression in caudate putamen (ST), nucleus accumbens (Acc), olfactory tubercle (Tu), piriform cortex (Pir), central nucleus of amygdala (Ce), and ventromedial hypothalamic nucleus (VMN). The magnitude and significance of the changes in PENK gene expression found in this study depend upon the brain region examined and the type of drug administration (contingent, noncontingent, and saline). Both types of cocaine administration, contingent and noncontingent, increased the expression level of PENK mRNA either on the last day of cocaine self-administration or during the extinction in all the brain regions examined when compared to saline group, except in VMN and Ce nuclei, where a significant decrease was found in contingent animals, as compared to noncontingent and saline groups. In the VMN, the decrease in PENK gene expression returned to noncontingent and saline values 10 days after the extinction of cocaine self-administration behavior; whereas, in the Ce, the decrease is maintained for the whole 10-day period of extinction. The maintained effects found in PENK gene expression in our study do not suggest a response to short-term compensatory adaptations in brain function after cessation of cocaine self-administration. Interestingly, our results showed an important additional difference in PENK gene expression in some brain regions, depending upon the type of cocaine administration. Although PENK mRNA levels in both cocaine groups were different from those found in the saline group in most of the brain regions examined, with the exception of VMN, the changes persisted mainly in the cocaine contingent administration group 10 days after extinction. These results suggest that an interaction between environmental stimuli and the pharmacological action of cocaine during drug self-administration and its extinction may represent an important variable in the regulation of cocaine action on PENK gene expression.

Our data are not in agreement with those found by others (Branch et al. 1992; Spangler et al. 1997), reporting that 14 days of repeated noncontingent cocaine administration in a bingeing model produced no alterations in PENK mRNA levels. The differences in the results found between both studies underscore the importance of the pattern, dose, route, and time course of cocaine administration on PENK gene expression. In contrast, different modes of cocaine administration or self-administration (Branch et al. 1992; Spangler et al. 1993, 1997) result in similar increases in prodynorphin gene expression in the brain, suggesting that cocaine actions on prodynorphin systems seem more independent of environmental conditions or type of drug administration.

Several of the brain regions included in this study have been proposed as elements of a rewarding circuitry in the mesocorticolimbic dopaminergic system (Koob and Bloom 1988). Ascending monoaminergic projections of the A₁₀ dopamine neurons include the nucleus accumbens, the prefrontal cortex, the cingulate, entorhinal, and piriform cortices, the medial part of the lateral septum, the interstitial nucleus of the stria terminalis, the olfactory tubercle, the anterior medial part of the caudate nucleus, and the amygdaloid complex (Versteeg et al. 1976; Fallon and Moore 1978; Moore and Bloom 1978; Simon et al. 1979). With the exception of the hypothalamic VMN, all the forebrain regions examined in the present study belong to the hypothesized rewarding circuitry and exhibit modifications in PENK gene expression during cocaine self-administration and its extinction. In addition, the decrease of PENK gene expression found in the present study may play a relevant role in cocaine withdrawal. It has been shown that cocaine self-administered animals exhibit behavioral symptoms upon withdrawal similar to the anxiogenic effects upon exposure to stress (Carroll and Lac 1987; Kleven and Woolverton 1991). Furthermore, PENK mRNA levels in the VMN may be altered after the effects of acute and chronic stress (García-García et al. 1998). Thus, it is possible that the decrease in PENK mRNA content in VMN found in our study represents a response to stress during early cocaine withdrawal phases.

It is particularly interesting to note the persistent decrease in PENK gene expression found in Ce. This nucleus receives the densest dopaminergic innervation arising from the ventral tegmental area to the amygdaloid complex (Fallon and Ciofi 1992), and it is thought to play a role in the modulation of the reinforcing and discriminative stimulus properties of cocaine (Caine et al. 1995; Hurd et al. 1997; Callahan et al. 1995). In addition, it represents an important part of the "extended amygdala," a proposed forebrain continuum bordered by the central nucleus of the amygdala and the shell of the nucleus accumbens, which includes portions of the substantia innominata and bed nucleus of stria terminalis (Alheid and Heimer 1988). Several animal studies

have suggested the "extended amygdala" as a neural substrate for the behavioral effects of self-administered cocaine (Robledo and Koob 1993; Caine and Koob 1994; Hurd et al. 1997; McGregor and Roberts 1993; Robbins et al. 1989; Robledo et al. 1996). In humans, recent evidence from brain-imaging studies suggest a role for dopamine-rich brain regions, including the nucleus accumbens and amygdala, in cocaine craving (Grant et al. 1996; Breiter et al. 1997; Childress et al. 1999). Along this line, considering the particular conditions of this work, it is tempting to speculate that, in addition to previously documented inhibition of mesocorticolimbic activity (Rossetti et al. 1992; Kuhar and Pilotte 1996), a decrease in proenkephalin gene expression in the Ce may represent a possible mechanism mediating in cocaine withdrawal states.

In conclusion, our data provide evidence of long-lasting neuroadaptative changes in proenkephalin gene expression after long-term contingent cocaine administration and its extinction in forebrain regions involved in reinforcing properties of psychostimulants. Although the relevance of changes in PENK mRNA levels elicited by cocaine self-administration and its extinction awaits further analysis of complementary changes in receptor function, the present results are in accordance with other studies showing that chronic cocaine administration produces long-lasting neuroadaptations in gene expression, especially in dopamine terminal fields (Hope et al. 1994; Moratalla et al. 1996; Cha et al. 1997). Long-lasting behavioral alterations exist associated with cocaine addiction, including the development of paranoia and drug craving (Satel and Edell 1991; Childress et al. 1999). In this sense, research on alterations in gene expression produced by long-term cocaine self-administration and its extinction could be a useful approach to determine neuroadaptative changes involved in cocaine withdrawal states.

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