

cAMP Response Element-Mediated Gene Transcription Is Upregulated by Chronic Antidepressant Treatment

J. Thome,¹ N. Sakai,¹ K.-H. Shin,¹ C. Steffen,¹ Y.-J. Zhang,¹ S. Impey,² D. Storm,² and R. S. Duman¹

¹Division of Molecular Psychiatry, Yale University School of Medicine, New Haven, Connecticut 06508, and

²Department of Pharmacology, University of Washington, Seattle, Washington 98195

Regulation of gene transcription via the cAMP-mediated second messenger pathway has been implicated in the actions of antidepressant drugs, but studies to date have not demonstrated such an effect *in vivo*. To directly study the regulation of cAMP response element (CRE)-mediated gene transcription by antidepressants, transgenic mice with a CRE-LacZ reporter gene construct were administered one of three different classes of antidepressants: a norepinephrine selective reuptake inhibitor (desipramine), a serotonin selective reuptake inhibitor (fluoxetine), or a monoamine oxidase inhibitor (tranylcypromine). Chronic, but not acute, administration of these antidepressants

significantly increased CRE-mediated gene transcription, as well as the phosphorylation of CRE binding protein (CREB), in several limbic brain regions thought to mediate the action of antidepressants, including the cerebral cortex, hippocampus, amygdala, and hypothalamus. These results demonstrate that chronic antidepressant treatment induces CRE-mediated gene expression in a neuroanatomically differentiated pattern and further elucidate the molecular mechanisms underlying the actions of these widely used therapeutic agents.

Key words: CRE enhancer; β -galactosidase; gene transcription; phosphorylation; fluoxetine; desipramine; tranylcypromine

Although the acute action of antidepressant treatment is known to be mediated by blockade of the reuptake or breakdown of serotonin (5-HT) and norepinephrine (NE), the molecular adaptations underlying the therapeutic action of these agents has not been determined. One signal transduction pathway that has been implicated in the action of antidepressant treatment is the cAMP second messenger cascade. Recent studies have demonstrated that different classes of antidepressants upregulate the cAMP system at several sites, including increased Gs activation of adenylyl cyclase (Ozawa and Rasenick, 1991) and upregulation of cAMP-dependent protein kinase (PKA) (Nestler et al., 1989; Perez et al., 1989). In addition, inhibition of cAMP metabolism produces antidepressant-like effects in behavioral models of depression (Wachtel, 1983; Griebel et al., 1991; O'Donnell, 1993) as well as therapeutic responses in depressed patients (Horowski and Sastre-Y-Hernandez, 1985; Bobon et al., 1988; Fleischhacker et al., 1992; Malison et al., 1997).

One potential target of the cAMP system that could mediate the action of antidepressants is the cAMP response element binding protein (CREB) (Montminy, 1990; Meyer and Habener, 1993). CREB is a transcription factor that mediates the actions of cAMP on gene expression and could thereby underlie some of the long-term effects of antidepressant treatment. A role for CREB in the action of antidepressants is supported by studies demonstrating that chronic antidepressant treatment increases the expression of CREB in limbic regions of rat brain (Nibuya et al., 1996). CREB regulates gene transcription by binding to a cAMP response element (CRE), a *cis*-acting enhancer element in the

regulatory region of various genes. The function of CREB is regulated largely by its state of phosphorylation at Ser¹³³, which results in activation of gene transcription (Montminy, 1990; Meyer and Habener, 1993). Phosphorylation of CREB at Ser¹³³ can occur via activation of the cAMP cascade and PKA, but also via activation of calcium-dependent protein kinases (i.e., protein kinase C and calcium/calmodulin-dependent protein kinase) (Duman et al., 1997, 1999). This raises the possibility that CREB could act as a common downstream target of different classes of antidepressants that influence 5-HT and/or NE (Duman et al., 1997, 1999).

The focus of the present study is to determine the influence of antidepressant treatment on the function of CREB. This is a critical issue because although CREB expression is upregulated by antidepressant treatment, the function of this transcription factor could be unchanged without a corresponding increase in phosphorylation. To address this issue, the influence of antidepressant administration on CRE-mediated gene expression and CREB phosphorylation were examined in CRE-LacZ transgenic mice. The transgene in these mice is a CRE-LacZ reporter gene construct, and stimulation of the CRE site leads to increased expression of the LacZ gene product, β -galactosidase. These mice have been used to establish the role of CRE-mediated gene expression in cellular and behavioral models of learning and memory and in circadian rhythm (Impey et al., 1996, 1998; Obrietan et al., 1998; Pham et al., 1999).

MATERIALS AND METHODS

CRE-LacZ transgenic mice. All experiments were conducted in mice heterozygous for the CRE-LacZ reporter gene construct. This construct consists of six tandem CREs upstream of a minimal HSV promoter driving the expression of β -galactosidase (Meinkoth et al., 1990). The mice were generated using C57BL6/SJL F2 blastocytes for microinjection; founders were bred to C57BL6 mice (Impey et al., 1996). The animals were bred and maintained under standard conditions (12 hr light/dark cycle, food and water *ad libitum*). PCR was used to determine the genotype of individual mice. All animal use procedures were in strict

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Correspondence should be addressed to R. S. Duman, 34 Park Street, New Haven, CT 06508. E-mail: ronald.duman@yale.edu.

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accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals and were approved by the Yale Animal Care and Use Committee.

Drug administration paradigms. For chronic paradigms, fluoxetine (10 mg/kg, i.p.), desipramine (15 mg/kg, i.p.), or tranylcypromine (10 mg/kg, i.p.) was administered daily for 14 d. For acute paradigms, mice received 0.9% saline (i.p.) daily for 13 d and a single injection of tranylcypromine, desipramine, or fluoxetine (i.p.) on day 14 (same dose as above). Control animals were administered 0.9% saline (i.p.) for 14 d. Haloperidol (1 mg/kg, i.p.) or cocaine (10 mg/kg, i.p.) were administered daily for 14 d.

β -Galactosidase staining. Mice were killed 6 hr after the last injection. After perfusion with 0.9% NaCl solution and 4% paraformaldehyde, the brains were post-fixed overnight and cryoprotected in 20% glycerol. Fluorescence immunohistochemistry was performed using standard protocols. Briefly, 40 μ m sections were incubated with rabbit anti β -galactosidase antibody at 1:1000 dilution (ICN Biochemicals, Costa Mesa, CA) and Alexa 594 goat anti-rabbit IgG at 3 μ g/ml (Molecular Probes, Eugene, OR). Sections were counterstained with 4',6-diamidino-2-phenylindole (DAPI) (see below). Images were captured on a Zeiss Axioskop fluorescence microscope.

Phosphorylation of CREB. After blocking with BSA and goat serum, free-floating sections were incubated overnight with anti-phosphorylated CREB (phospho-CREB) antibody from rabbit (New England Biolabs, Beverly, MA) at 1:250 dilution. A second overnight incubation was performed with 2.5 μ g/ml fluorescein-labeled anti-rabbit IgG antibody (Jackson ImmunoResearch, West Grove, PA), and a third was performed with 4 μ g/ml Alexa 488-anti-fluorescein antibody (Molecular Probes). This method involving three antibodies results in a considerable enhancement of the fluorescence signal. Between each incubation step with antibody, stringent washing steps with PBS containing NaF and Triton X-100 were performed.

Counterstaining. Vectashield Mounting Medium with DAPI (Vector Laboratories, Burlingame, CA) was used for counterstaining. The DAPI fluorophore binds to cellular DNA, allowing for fluorescence visualization of the cells. DAPI produces a blue fluorescence with excitation at 360 nm and emission at 460 nm when bound to DNA. There is no emission overlap with fluorescein, rhodamine, Texas Red, or the fluorophores used for β -galactosidase or phospho-CREB immunolabeling.

Data analysis. Images were evaluated using a four point score system (0 = minimal, 1 = weak, 2 = intermediate, 3 = strong immunoreactivity) by two independent investigators blinded to the treatment condition. The Kruskal–Wallis test was performed for statistical analysis. The level of significance was $p < 0.05$. A trend or tendency was assumed by $p < 0.10$. Similar results were obtained when quantified by densitometry of digitalized images. For this approach, the images were converted into gray-scale pictures. The appropriate areas were outlined, and staining intensity was determined using IPLab Spectrum–Scientific Image Processing, Version 3.1.2 (Scanalytics, Inc.).

RESULTS

Chronic antidepressant administration increases CRE-mediated gene expression

To measure CRE-mediated gene expression in the CRE–LacZ transgenic mice, levels of the LacZ gene product, β -galactosidase, were determined by fluorescence immunohistochemistry. Relatively low levels of β -galactosidase were observed in most brain regions in saline-treated mice, particularly the dentate gyrus of hippocampus and cerebral cortex (Figs. 1, 2). Relatively higher levels of immunoreactivity were observed in amygdala and hypothalamus (Figs. 1, 2).

The influence of chronic administration of several different classes of antidepressants, including a 5-HT selective reuptake inhibitor (fluoxetine), a norepinephrine selective reuptake inhibitor (desipramine), and a monoamine oxidase inhibitor (tranylcypromine), on levels of CRE-induced LacZ expression were tested. Mice were administered saline or antidepressants for 14 d, and the brains were processed for levels of β -galactosidase immunoreactivity 6 hr after the last treatment. The results demonstrate that chronic antidepressant administration significantly increases levels of CRE-mediated gene expression in several brain

regions, although differences were observed in the regional pattern of expression between the different classes of antidepressants. Chronic administration of each of the antidepressants significantly increased levels of β -galactosidase immunoreactivity in the amygdala. The subregions of amygdala that were influenced include the central, basolateral, and basomedial nuclei. Chronic administration of tranylcypromine or fluoxetine, but not desipramine, also induced CRE-mediated gene transcription in the cerebral cortex, including superficial and deep layers. Chronic administration of tranylcypromine additionally increased levels of β -galactosidase immunoreactivity in the hippocampus (CA3 pyramidal cell layer), and there was a tendency for an increase in the hypothalamus (dorsomedial and ventromedial nuclei). A tendency for an increase was also observed in the dentate gyrus granule cell layer, but this effect was small and did not achieve significance. Chronic fluoxetine administration also resulted in a tendency for induction of β -galactosidase immunoreactivity in the thalamus, and there were as well tendencies toward elevated levels in the CA3 pyramidal cell layer and hypothalamus. No alterations were observed in the CA1 pyramidal cell layer of the hippocampus. For most experiments, the immunohistochemical results were quantified using a subjective scoring scale (see Materials and Methods). However, similar results were obtained for chronic fluoxetine when the digitalized images were quantified by densitometry (Table 1).

In contrast to the chronic paradigms, acute antidepressant administration induced CRE-mediated gene expression to a much lower level. For the acute studies, mice received saline injections for 13 d, and on day 14 received either saline or an antidepressant at the same dose used for the chronic paradigm. The repeated saline injections were conducted to acclimate the mice to the stress associated with the handling and injections. The brains were then processed for levels of β -galactosidase immunohistochemistry 6 hr after saline or drug treatment. No significant effects were observed in any of the brain regions examined after acute administration of fluoxetine or desipramine (Table 2). Acute administration of tranylcypromine significantly increased β -galactosidase immunoreactivity in hypothalamus, and there was a similar trend in the thalamus and dentate gyrus. However, there was no significant effect of acute tranylcypromine administration in any of the other brain regions examined.

To determine whether the induction of CRE-mediated gene expression is specific to antidepressants, two other classes of psychotropic drugs were examined. This included an antipsychotic (haloperidol) and a psychostimulant (cocaine). However, chronic administration of these nonantidepressant drugs did not significantly influence levels of β -galactosidase immunoreactivity in any of the brain regions examined (Table 3).

Chronic antidepressant administration increases the phosphorylation of CREB

To study the possible mechanisms underlying the induction of CRE-mediated gene expression, the influence of antidepressants on the phosphorylation of CREB was examined. Levels of phospho-CREB were determined by fluorescence immunohistochemistry using an antibody specific for the phosphorylated form of CREB. Alternate sections from the same brains used for analysis of β -galactosidase immunohistochemistry were used for most experiments. Phospho-CREB immunoreactivity was observed in most brain regions of saline-treated mice, including the dentate gyrus granule cell layer and the cerebral cortex. Rela-

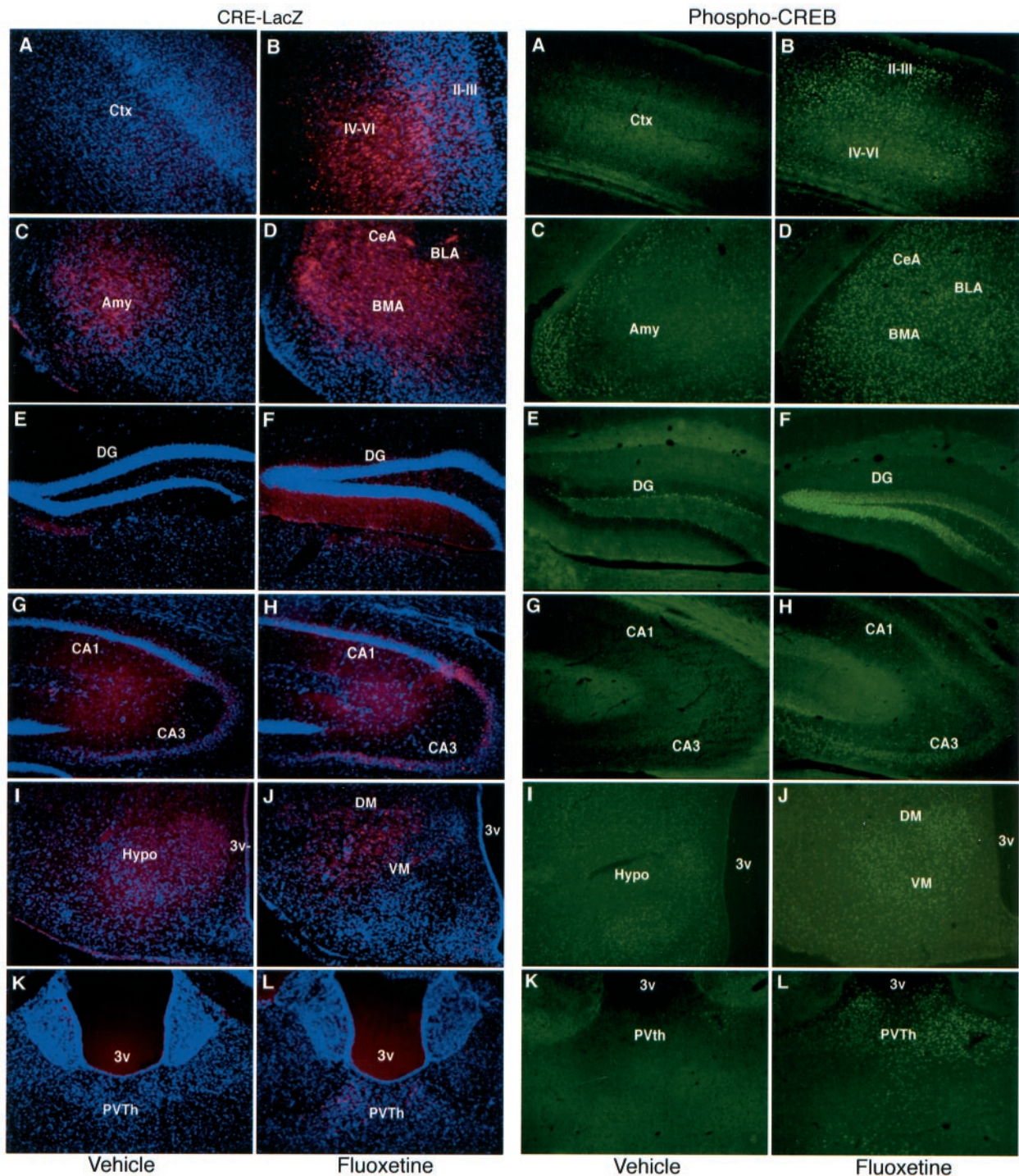


Figure 1. Chronic fluoxetine administration increases CRE-mediated gene expression and CREB phosphorylation in brain. Mice were administered saline or fluoxetine once daily for 14 d, and the brains were processed for β -galactosidase (CRE-LacZ) or phospho-CREB immunohistochemistry 6 hr after the last treatment. Representative images of β -galactosidase (left/red, A–L) or phospho-CREB (right/green, A–L) are shown. The brain regions that were examined include cerebral cortex (A, B), amygdala (C, D), dentate gyrus granule cell layer (E, F), CA3 pyramidal cell layer (G, H), hypothalamus (I, J), and thalamus (K, L). Images from saline-treated (A, C, E, G, I, K) and fluoxetine-treated (B, D, F, H, J, L) animals are shown. The subregions that are labeled include layers II–III and IV–VI of cerebral cortex; central (CeA), basolateral (BLA), and basomedial (BMA) nuclei of the amygdala; dentate gyrus (DG) granule cell layer, and CA3 and CA1 pyramidal cell layers of hippocampus; dorsomedial (DM) and ventromedial (VM) nuclei of the hypothalamus; paraventricular nucleus of the thalamus (PVTh); and third ventricle (3v).

tively high levels of immunoreactivity were found in amygdala and hypothalamus.

Chronic administration of fluoxetine significantly increased levels of phospho-CREB immunoreactivity in several brain re-

gions, including amygdala, cerebral cortex, dentate gyrus granule cell layer, thalamus, and hypothalamus (Figs. 1, 3). The subregions that were influenced within each of these areas were similar to those observed for β -galactosidase immunohistochemistry.

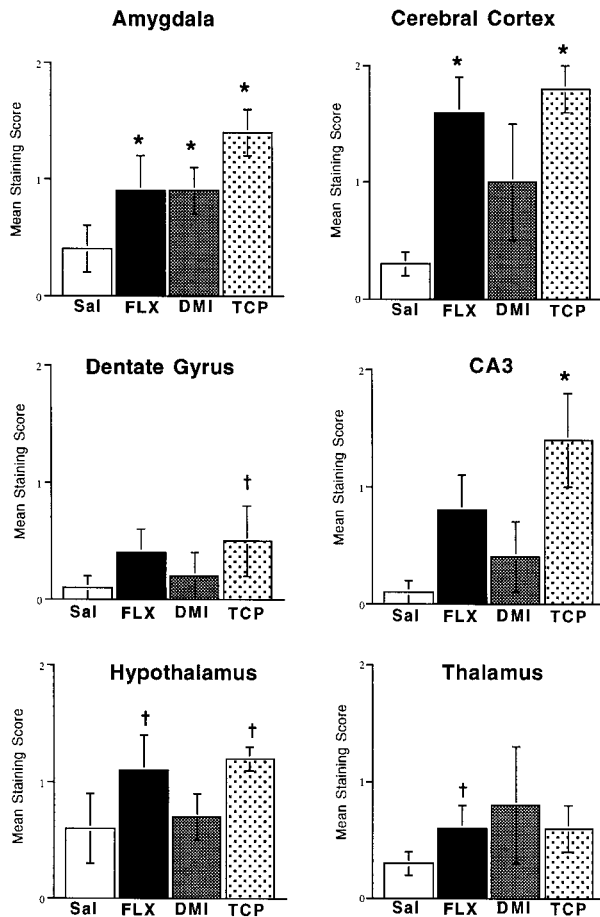


Figure 2. Chronic antidepressant administration increases CRE-mediated gene expression in brain. Mice were administered saline, fluoxetine, desipramine, or tranlycypromine once daily for 14 d, and 6 hr after the last treatment β -galactosidase immunohistochemistry was conducted as described in Materials and Methods. Levels of β -galactosidase were scored as described in Materials and Methods. The results are presented as the mean \pm SEM of four to six separate determinations. * $p < 0.05$, † $p < 0.10$ compared with saline (Kruskal–Wallis).

Table 1. Influence of chronic fluoxetine administration on CRE-mediated gene expression

	β -Galactosidase expression	
	SAL	FLU
Amygdala	173 \pm 15	217 \pm 11*
Cortex	196 \pm 17	241 \pm 9*
CA3	138 \pm 4	184 \pm 17*
Dentate gyrus	135 \pm 7	141 \pm 12
Hypothalamus	125 \pm 8	167 \pm 8*
Thalamus	120 \pm 4	152 \pm 14*

Mice received a chronic (14 d) administration of vehicle (SAL) or fluoxetine (FLU) and were killed 6 h after the last injection. Levels of β -galactosidase were determined by immunohistochemistry as described in Materials and Methods. Levels of β -galactosidase labeling were quantified by densitometry of digitalized images as described in Materials and Methods. The data are the mean \pm SEM; $n = 6$ separate determinations. * $p < 0.05$ compared with saline (Kruskal–Wallis).

Chronic administration of desipramine also increased phospho-CREB immunoreactivity in the dentate gyrus granule cell layer, although to a lower level than observed with fluoxetine. No significant effects were observed in response to chronic desipra-

Table 2. Influence of acute antidepressant administration on CRE-mediated gene expression

	β -Galactosidase expression			
	SAL	FLU	DMI	TCP
Amygdala	0.4 \pm 0.2	0.5 \pm 0.4	0.2 \pm 0.2	1.0 \pm 0.6
Cortex	0.3 \pm 0.1	0.2 \pm 0.2	0.3 \pm 0.2	1.8 \pm 0.7
CA1	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0	0.4 \pm 0.4
CA3	0.3 \pm 0.1	0.1 \pm 0.1	0.4 \pm 0.3	1.1 \pm 0.6
Dentate gyrus	0.1 \pm 0.1	0.1 \pm 0.1	0.1 \pm 0.1	0.8 \pm 0.4
Hypothalamus	0.6 \pm 0.3	0.3 \pm 0.3	0.3 \pm 0.2	1.6 \pm 0.6*
Thalamus	0.3 \pm 0.1	0.2 \pm 0.2	0.2 \pm 0.1	1.4 \pm 0.4*

Mice received a single dose of vehicle (SAL), fluoxetine (FLU), desipramine (DMI), or tranlycypromine (TCP) and were killed 6 h later. Levels of β -galactosidase were determined by immunohistochemistry as described in Materials and Methods. The data are the mean \pm SEM; $n = 4$ –5 separate determinations. * $p < 0.05$ compared with saline (Kruskal–Wallis).

Table 3. Influence of chronic cocaine or haloperidol administration on CRE-mediated gene expression

	Phospho-CREB		
	Saline	Cocaine	Haloperidol
Amygdala	0.0 \pm 0.0	0.0 \pm 0.0	0.4 \pm 0.4
Cortex	0.3 \pm 0.2	0.3 \pm 0.3	0.4 \pm 0.4
CA1	0.1 \pm 0.4	0.0 \pm 0.0	0.0 \pm 0.0
CA3	0.1 \pm 0.4	0.0 \pm 0.0	0.2 \pm 0.2
DG	0.2 \pm 0.3	0.0 \pm 0.0	0.2 \pm 0.2
Hypothalamus	0.4 \pm 0.2	0.3 \pm 0.3	0.4 \pm 0.4
Thalamus	0.3 \pm 0.2	0.3 \pm 0.3	0.8 \pm 0.6

Vehicle, cocaine, or haloperidol was administered chronically for 14 d, and mice were killed 6 h after the last injection. Levels of β -galactosidase were determined by immunohistochemistry as described in Materials and Methods. The data are the mean \pm SEM; $n = 7$ separate determinations.

mine administration in any of the other brain regions examined. The influence of tranlycypromine administration on levels of phospho-CREB immunoreactivity was not determined.

The influence of acute antidepressant administration on levels of phospho-CREB was also examined. Alternate sections from the mouse brains used for β -galactosidase immunohistochemistry were used for these experiments. Acute administration of fluoxetine or desipramine did not significantly influence levels of phospho-CREB immunostaining in any of the brain regions examined (Table 4). However, there was a tendency for acute administration of these antidepressants to increase levels of phospho-CREB in the dentate gyrus granule cell layer of hippocampus.

Several other classes of psychotropic drugs were also examined to determine whether the induction of phospho-CREB was specific to antidepressants. Chronic administration of haloperidol or cocaine did not significantly influence levels of phospho-CREB immunoreactivity in any of the brain regions examined (data not shown).

DISCUSSION

The results of this study demonstrate that chronic antidepressant administration increases CRE-mediated gene expression and CREB phosphorylation in a region- and drug-specific manner. The most consistent effects observed between the different antidepressants tested were in the amygdala and cerebral cortex. In addition, significant effects were observed in several other limbic brain regions, including the hippocampus, hypothalamus, and

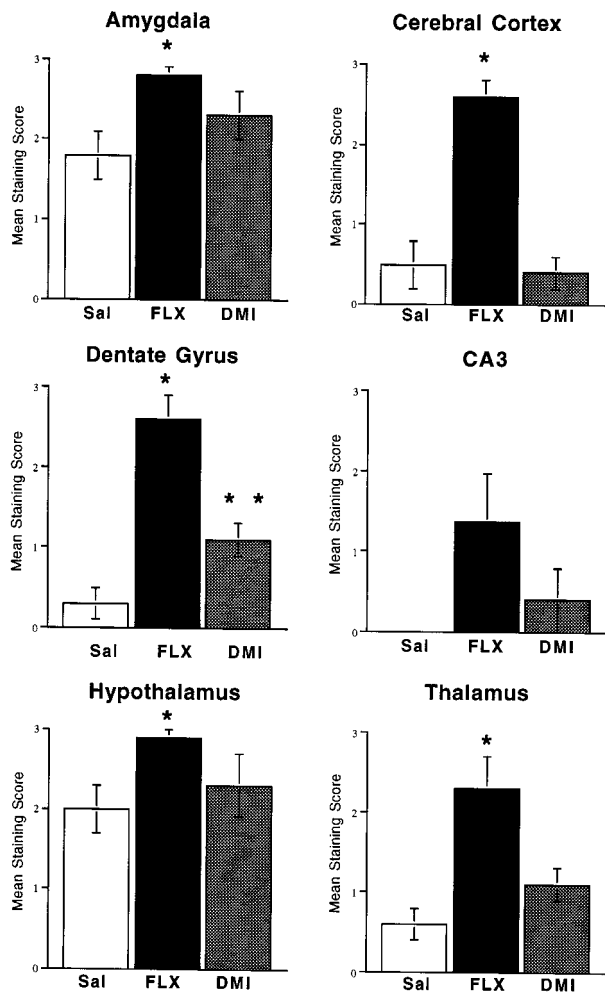


Figure 3. Chronic antidepressant administration increases CREB phosphorylation in brain. Mice were administered saline, fluoxetine, or desipramine once daily for 14 d, and 6 hr after the last treatment phospho-CREB immunohistochemistry was conducted as described in Materials and Methods. Levels of phospho-CREB staining were scored, and results are presented as the mean \pm SEM of four to six separate determinations. * $p < 0.05$ compared with saline (Kruskal–Wallis).

thalamus. Induction of CRE-mediated gene expression and CREB phosphorylation were observed in response to chronic, but not acute, antidepressant administration, consistent with the time course for the therapeutic action of these agents. Induction of CRE-mediated gene expression in these brain regions appears to be relatively specific to antidepressants in that administration of nonantidepressant psychotropic drugs does not result in similar effects. These results demonstrate that induction of CRE-mediated gene expression in specific limbic brain structures is an intracellular target of different classes of antidepressants.

The time lag for induction of CRE-mediated gene expression could result from the time required for upregulation of one or more of the upstream components of the cAMP cascade. Previous studies have demonstrated that upregulation of PKA enzyme activity and induction of CREB expression is dependent on chronic antidepressant treatment (Duman et al., 1999). There is also a corresponding delay in the upregulation of CRE binding in response to antidepressant administration (Nibuya et al., 1996; Frechilla et al., 1998). In addition, it is possible that adaptations of 5-HT and NE autoreceptors and monoamine neurotransmission

Table 4. Influence of acute antidepressant administration on levels of phospho-CREB immunohistochemistry

	Phospho-CREB		
	SAL	FLU	DMI
Amygdala	1.8 \pm 0.3	1.9 \pm 0.1	1.1 \pm 0.3
Cortex	0.5 \pm 0.3	0.3 \pm 0.2	0.0 \pm 0.0
CA1	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0
CA3	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0
Dentate gyrus	0.3 \pm 0.2	1.0 \pm 0.3	1.1 \pm 0.2
Hypothalamus	2.0 \pm 0.3	1.9 \pm 0.1	2.1 \pm 0.1
Thalamus	0.6 \pm 0.2	1.1 \pm 0.1	0.8 \pm 0.2

Mice received a single dose of vehicle (SAL), fluoxetine (FLU), or desipramine (DMI) and were killed 6 h later. Levels of phospho-CREB were determined by immunohistochemistry as described in Materials and Methods. The data are the mean \pm SEM; $n = 4$ –5 separate determinations. * $p < 0.05$ compared with saline (Kruskal–Wallis).

contribute to a stable upregulation of the cAMP cascade and consequently of CRE-mediated gene expression (Blier and de Montigny, 1994).

Both the NE and 5-HT selective reuptake inhibitors increase CRE-mediated gene expression and CREB phosphorylation in amygdala. In contrast, fluoxetine, but not desipramine, significantly increases β -galactosidase and CREB phosphorylation in cerebral cortex. Chronic fluoxetine administration also increases phospho-CREB in several other limbic structures, including the dentate gyrus, hypothalamus, and thalamus. The reasons for these regional differences between fluoxetine and desipramine are not clear because these structures receive diffuse projections from both the NE and 5-HT systems. However, there is evidence for a more rapid adaptation of autoreceptor inhibition of the 5-HT than the NE neurotransmitter system (Blier and de Montigny, 1994). Alternatively, it is possible that higher brain levels of fluoxetine are achieved because of its relatively long half-life. It is notable that there is a tendency for desipramine to increase phospho-CREB in brain regions in which fluoxetine produces a significant effect (i.e., dentate gyrus and thalamus). It is possible that a higher dose of desipramine would produce a significant response, although we were not able to test this possibility because of the side effects of this antidepressant. It will be interesting to examine more selective NE reuptake inhibitors with fewer side effects, such as reboxetine, to further study the influence of the NE system on CRE-mediated gene expression.

The corresponding induction of phospho-CREB and β -galactosidase in amygdala and cerebral cortex indicates that CREB phosphorylation could mediate the induction of CRE-mediated gene expression in these brain regions. However, it was somewhat surprising to find that there was not a significant induction of CRE-mediated gene expression in some brain regions where there was an induction of phospho-CREB (i.e., dentate gyrus, hypothalamus, and thalamus in response to fluoxetine). It is conceivable that transcription factors other than CREB are also required for induction of CRE-mediated gene expression and that these transcription factors are not regulated by antidepressants in all brain regions. This is consistent with previous reports that phosphorylation of CREB at Ser¹³³ is not always sufficient to induce CRE-mediated gene expression (Enslin et al., 1994; Thompson et al., 1995; Impey et al., 1998). Alternatively, the induction of β -galactosidase may not be as responsive as phospho-CREB. This possibility is supported by the observation that there was a trend for an induction of β -galactosidase in

some of the brain regions where an induction of phospho-CREB was observed (i.e., thalamus and hypothalamus).

Antidepressant induction of CREB phosphorylation and CRE-mediated gene expression is observed in brain structures that are thought to play a role in the regulation of emotion and responses to stress. Moreover, clinical studies have reported that there are alterations in blood flow, volume, and neurochemistry of many of these same brain regions in depressed patients. For example, previous studies have demonstrated that amygdala mediates some of the behavioral actions of antidepressants and that neurochemical adaptations to antidepressants are observed in this brain region (Ordway et al., 1991; Beck and Fibiger, 1995; Duncan et al., 1996; Dawes et al., 1998; Morelli et al., 1999). In addition, clinical brain imaging studies report alterations in blood flow and glucose metabolism in amygdala of depressed patients (Drevets et al., 1992). The amygdala plays a significant role in fear conditioning and conditioned avoidance behavior and is thought to encode the emotional component of aversive stimulus conditioning (Fanselow et al., 1999; Holland and Gallagher, 1999). The possibility that CREB influences the function of amygdala is supported by a recent report that overexpression of CREB in this brain regions alters long-term memory of fearful conditions (Joselyn et al., 2000). On the basis of these observations, it is also possible that neurochemical alterations in amygdala could contribute to the displaced emotion, as well as anxiety, that is often observed in depressed patients. The possibility that CREB influences amygdala function can be directly tested by studying the influence of the cAMP-CREB cascade on behavioral models of depression, as well as behaviors that are controlled by amygdala (i.e., fear conditioning and conditioned avoidance behavior).

Chronic antidepressant treatment also results in neurochemical and cellular adaptations in cerebral cortex and hippocampus. A role for CREB is supported by the results of our previous reports and the present study, which demonstrate that chronic antidepressant treatment increases the expression and function of CREB in these brain regions (Nibuya et al., 1995, 1996; Duman et al., 1997, 1999). In addition, we have found that antidepressant administration increases the expression of BDNF in hippocampus. The possibility that induction of BDNF is mediated by CREB is supported by recent reports that the promoter of the BDNF gene contains a CRE (Shieh et al., 1998; Tao et al., 1998). Upregulation of CREB and BDNF could act to oppose the damaging effects of stress on hippocampal neurons (Sapolsky, 1996; McEwen, 1999). Clinical studies also report a reduction in the volume and/or number of neurons in hippocampus and cerebral cortex of depressed patients (Sheline et al., 1996; Drevets et al., 1997; Ongur et al., 1998; Rajkowska et al., 1999). A role for CREB in the pathophysiology of depression is supported by a postmortem study demonstrating that CREB levels are decreased in the cerebral cortex of depressed patients and increased in patients receiving antidepressant medication at the time of death (Dowlatshahi et al., 1998).

Alterations in the function of hippocampus and cerebral cortex could also contribute to the cognitive deficits that are often observed in depressed patients. The cAMP cascade is reported to be integrally involved in the cellular adaptations underlying learning and memory in these brain regions (Abel et al., 1997; Taylor et al., 1999). On the basis of these findings and the results of the present study, it is possible that amelioration of the cognitive deficits in depressed patients could result, at least in part, from upregulation of CRE-mediated gene expression in response to antidepressant treatment.

The actions of antidepressants on the hypothalamus and thalamus are not as well characterized. The dorsomedial and ventromedial regions of the hypothalamus are reported to play a role in vegetative behaviors (e.g., eating and sexual drive) that are also abnormal in depression. It is possible that induction of CRE-mediated gene expression in these hypothalamic nuclei normalizes these abnormalities. There was no effect in the paraventricular nucleus of the hypothalamus, which controls the hypothalamic–pituitary adrenal (HPA) axis in response to stress. Interestingly, the paraventricular nucleus of the thalamus is another stress-responsive brain structure that is reported to exert negative control on the HPA axis (Bhatnagar and Dallman, 1998). Upregulation of CRE-mediated gene expression in this structure could contribute to normalization of HPA function in response to chronic antidepressant treatment.

The results of this study demonstrate that induction of CRE-mediated gene expression is a common action of antidepressant treatment. The challenge now is to directly test the role of CREB on the cellular, behavioral, and endocrine responses that are regulated by specific limbic brain structures. Studies are currently underway to address these questions using viral vectors and transgenic mice to determine the functional responses to overexpression of CREB in specific brain regions. Another challenge is to identify the target genes, in addition to BDNF, that are influenced by CREB and antidepressant treatment. One approach is to use DNA microarray technology to identify gene targets of the cAMP-CREB cascade and antidepressants. The results of the present study, in combination with these future approaches, should provide a more complete characterization of the role of CREB in the action of antidepressants, as well as the gene targets that mediate the therapeutic response to these agents.

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