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## Corticotropin-releasing factor and vasoactive intestinal polypeptide activate gene transcription through the cAMP signaling pathway in a catecholaminergic immortalized neuron

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### Abstract

Corticotropin-releasing factor (CRF) and vasoactive intestinal polypeptide (VIP) are neuropeptides displaying a variety of short-term effects in the nervous system. It is shown here in transfection experiments of an immortalized noradrenergic locus coeruleus-like cell line that both CRF and VIP also trigger a signaling cascade capable of activating gene transcription. To elucidate the signaling pathway leading to transcriptional induction, cells were transfected with an inhibitor for cAMP-dependent protein kinase, targeted to the nucleus via a nuclear-localization signal. Transcriptional induction of a reporter gene by CRF and VIP was blocked in these cells, indicating that the cAMP-dependent protein kinase is required for transducing CRF and VIP generated signals into the nucleus. Additionally, transfection experiments with a reporter gene containing cAMP response elements in its regulatory region demonstrate that CRF and VIP receptor activation induce transcription through this genetic regulatory element. We conclude that long-term effects of CRF and VIP in neurons are likely to be mediated by the transcriptional regulation of CRF and VIP-responsive genes via the cAMP signaling pathway.

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### 1. Introduction

Corticotropin-releasing factor (CRF), a neuropeptide of 41 amino acids that is highly conserved in evolution, acts as the major physiological regulator of the basal and stress-induced release of corticotropin (ACTH),  $\beta$ -endorphin and other proopiomelanocortin-

derived peptides from the anterior pituitary gland (for review, see Owens and Nemeroff, 1991). High CRF-immunoreactivity has been found in the paraventricular nucleus of the hypothalamus but also in many extrahypothalamic regions of the brain, indicating that CRF, in addition to its endocrine activity, displays extrahypophysiotropic effects, mainly as a regulator of stress responses. Overproduction of CRF has been implicated in neuropsychiatric disorders, such as major depression, anxiety-related disorders, and anorexia nervosa (for review, see Dunn and Berridge, 1990). Vasoactive intestinal polypeptide (VIP), a regulatory peptide of 28 amino acids, is widely expressed in the central and peripheral nervous system exhibiting neurotransmitter and hormonal roles (for review, see Gozes and Brenneman, 1989). Particularly, VIP plays

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*Abbreviations:* CRE, cAMP response element; CREB, cAMP response element binding protein; CRF, corticotropin-releasing factor; dbcAMP-N6, 2'-O-dibutyryladenosine 3'; 5'-monophosphate; GST, glutathione S-transferase; LC, locus coeruleus; NLS, nuclear localization signal; PACAP, pituitary adenylate cyclase-activating polypeptide; PKA, protein kinase A; VIP, vasoactive intestinal polypeptide.

an important role in systemic vasodilation, leading to a relaxation of cerebral blood vessels (Lee et al., 1984).

CRF and VIP mediate their actions through binding to receptors which belong to the family of seven transmembrane helix, Gs-coupled receptors (Ishihara et al., 1992; Chang et al., 1993; Chen et al., 1993; Lutz et al., 1993; Lovenberg et al., 1995). VIP additionally binds to the PAC<sub>1</sub> and VPAC<sub>1</sub> receptors (Sprengler et al., 1993; Rawlings et al., 1996; Harmar et al., 1998). Binding of CRF and VIP to their cognate receptors is associated with the activation of adenylyl cyclase (Chen et al., 1986, 1993; Chang et al., 1993; Ciccarelli et al., 1994; Lovenberg et al., 1995). The increased intracellular concentration of cAMP mediates the activation of cAMP-dependent protein kinase (PKA) by triggering the dissociation of catalytic and regulatory subunits of the holoenzyme. The activation of PKA is believed to transduce all known effects of cAMP in mammalian cells. PKA phosphorylates regulatory proteins in the cytoplasm such as tyrosine hydroxylase and synapsin I, and also proteins in the plasma membrane, such as ion channels and receptors (Girault et al., 1988; Moss et al., 1996; Hwang et al., 1997; Puri et al., 1997; Gray et al., 1998). In addition, the catalytic subunit of PKA translocates to the nucleus and activates gene transcription by phosphorylating transcription factors such as CREB (for review, see Montminy, 1993; Lalli and Sassone-Corsi, 1994; Vallejo and Habener, 1994). Unfortunately, in most of the studies dealing with cAMP-mediated gene transcription, membrane-penetrating cAMP analogues such as dbcAMP or unphysiological activators of adenylyl cyclase like forskolin were used, and only few studies were performed with physiological activators.

In this study, the role of CRF and VIP in activating gene transcription in CATH.a cells has been investigated. From the fact that both peptides initiate the cAMP signaling pathway we suggested that one of the functions of CRF and VIP in neurons is the activation of CREB, thus leading to transcription of CREB controlled genes. One CRF responsive target gene in the pituitary was identified as the proopiomelanocortin gene. This gene, however, lacks a canonical cAMP response element (CRE). This discrepancy has been explained by the fact that CRF receptor activation induces *c-fos*, that subsequently controls proopiomelanocortin gene expression (Boutillier et al., 1991). Alternatively, a CRF responsive element has been identified in the rat proopiomelanocortin gene, together with a DNA-binding protein termed PCRH-REB-1 (Jin et al., 1994). The CRF response element and the cognate DNA-binding protein have no similarities with the CRE and the CRE-binding proteins of the bZIP family, respectively. VIP potentiated *c-fos* expression in cultured cortical neurons. VIP, however, did not activate the *c-fos* gene directly but rather

potentiates the effect of glutamate, which activates transcription by stimulating *N*-Methyl-D-aspartate receptors (Martin et al., 1995).

To clearly elucidate the role of CRF and VIP upon CREB activation and gene transcription and other transcription factors, we used model promoters and activators to avoid interference of other transcriptional elements. In addition, we designed and used a high specific nuclear inhibitor of cAMP dependent protein kinase to verify the impact of the cAMP signaling cascade upon CRF and VIP regulated gene transcription. As a model system, to investigate the role of CRF and VIP upon gene transcription, an immortalized locus-coeruleus-like cell line, CATH.a, was used that contained both CRF, VIP and PAC<sub>1</sub> receptors (Duman et al., 1992; Iredale et al., 1996; Muller et al., 1997). The locus coeruleus (LC), a group of specialized neuronal cells in the brainstem, is the largest noradrenergic center in the brain, containing approximately 50% of brain noradrenergic neurons. The activation of noradrenergic LC neurons has been associated with stress due to local release of CRF (Valentino et al., 1991). In addition, the LC contains one of the highest densities of VIP binding sites in the brain (Martin et al., 1987).

## 2. Experimental procedures

### 2.1. Reporter constructs

Plasmid 5 × GAL4OVEC has been described (Cibelli et al., 1996a). Plasmid pCgBCRE<sup>4</sup>OVEC contains four copies of the CRE sequence derived from the murine chromogranin B gene and was constructed by inserting the synthetic annealed oligonucleotides 5'-TCGAGCTGCTCCGTGACGTCAGGGGCTGAG-3' and 5'-TCGACTCAGCCCTGACGTCACGGAGCAGC-3', comprising chromogranin B promoter sequences from -110 to -86, into the Sal I and Xho I sites of plasmid pHIVTATA-CAT (Thiel et al., 1996). This sequence was subsequently multimerized as described, excised with Sal I and Sac I and cloned into the plasmid OVEC (Westin et al., 1987).

### 2.2. Expression plasmids

Plasmid pGAL4-CREB, containing the human CREB cDNA fused to the DNA-binding domain of GAL4, has been described recently (Cibelli et al., 1996a). Plasmid pEBG-PKI encodes a fusion protein consisting of *Schistosoma japonicum* glutathione S-transferase (GST) and amino acids 1 to 31 of the PKA inhibitor PKI (Scott et al., 1985). This expression construct was generated by inserting a blunt ended BamHI/XbaI-fragment from plasmid pLXX-PKI(1-31) (Grove et al., 1987) into the filled in Not I site of

the eukaryotic GST-expression vector pEBGN. This vector encodes *S. japonicum* GST under the control of the EF-1 $\alpha$  promoter. To accomplish nuclear targeting of the GST fusion protein, two synthetic, annealed oligonucleotides 5'-GATCACCTCCTAAAAAGAA-GCGCAAGGTAGGCGGAG-3' and 5'-GATCCT-CCGCCTACCTTGCCTTCTTTTAGGAGGT-3', encoding the nuclear localization sequence of the SV40 large T antigen, were inserted into the BamHI site of pEBG (Sánchez et al., 1994), a kind gift of Bruce J. Mayer, Howard Hughes Medical Institute, Harvard University, Cambridge, MA, USA.

### 2.3. Cell culture and transfections

The locus coeruleus-like cell line CATH.a (Suri et al., 1993) was a generous gift of Donna Chikaraishi, Neuroscience Program, Tufts University School of Medicine, Boston, MA, USA. Cells were cultured in 88% RPMI medium supplemented with 4% fetal calf serum, 8% horse serum, 100 units/ml penicillin, and 100 mg/ml streptomycin. Cells were transfected by the calcium phosphate coprecipitation procedure (Thiel et al., 1991). The  $\beta$ -globin reporter plasmid (5  $\mu$ g), the GAL4-CREB expression plasmid (5  $\mu$ g) and the internal reference plasmid ICPOref (2.5–5  $\mu$ g) were transfected into cells on 100 mm plates. 24 h later, cells were treated with either 1 mM N6,2'-*O*-dibutyryl-adenosine 3',5'-monophosphate (dbcAMP, Sigma), 0.5  $\mu$ M ovine CRF (Serva or Calbiochem), 0.5  $\mu$ M  $\alpha$ -helical CRF9-41 (Sigma), or 1  $\mu$ M VIP (Serva), respectively. N6,2'-*O*-dibutyryl-adenosine 3',5'-monophosphate and VIP were dissolved in water, and CRF and  $\alpha$ -helical CRF9-41 in 0.05 M acetic acid, before addition to the culture medium. Total cytoplasmic RNA was extracted 48 h after transfection and the  $\beta$ -globin mRNA was detected by RNase protection mapping (Thiel et al., 1994). COS cells were transfected using the DEAE-dextran method (Thiel et al., 1998) and whole cell extracts were prepared as described (Cibelli et al., 1996b).

### 2.4. Miscellaneous techniques

For Western blot analysis, proteins from transfected COS cells were separated on a 14% SDS-PAGE and transferred to nitrocellulose membranes (pore size 0.2  $\mu$ m). Antibodies recognizing *S. japonicum* GST were raised in rabbits with purified recombinant GST protein. Blots were developed using an alkaline phosphatase-conjugated secondary antibody. Indirect immunofluorescence was performed as described (Cicchetti et al., 1995).

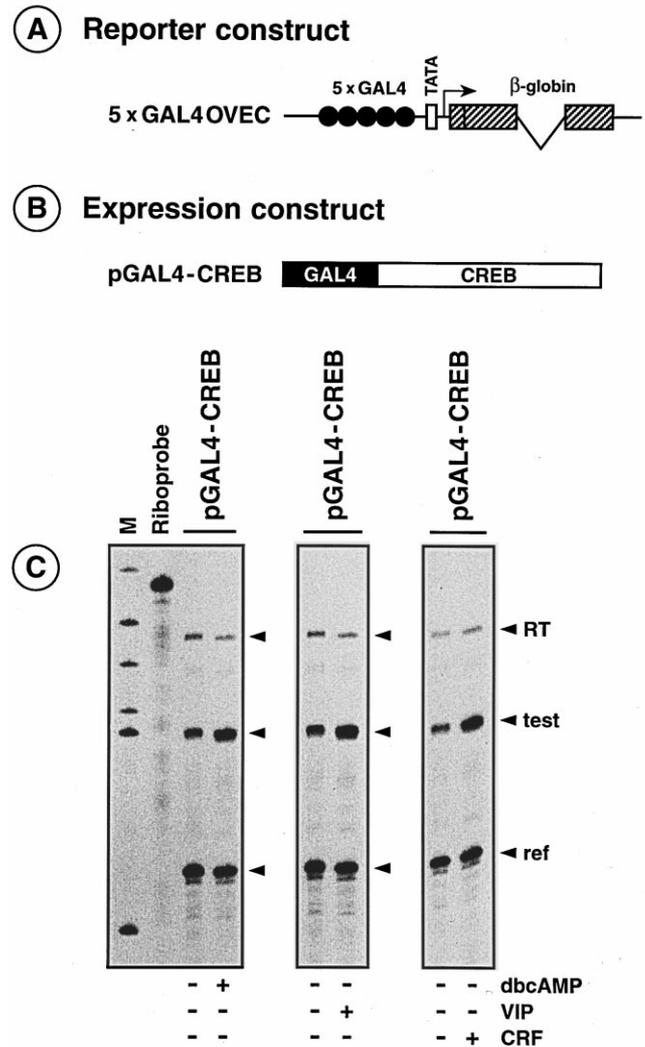


Fig. 1. VIP, CRF and dbcAMP activate gene transcription in CATH.a cells. (A) Reporter plasmid containing rabbit  $\beta$ -globin gene as a reporter gene, a TATA box and five copies of the GAL4 binding site upstream of the TATA box. (B) Expression vector encoding a fusion protein between the DNA-binding domain of the yeast transcription factor GAL4 and the cDNA of human CREB (plasmid pGAL4-CREB). (C) Reporter plasmid, expression plasmid and the ICPOref internal standard plasmid were introduced into CATH.a cells. Cells were incubated in the absence (-) or presence (+) of 1 mM dbcAMP, 1  $\mu$ M VIP or 0.5  $\mu$ M CRF for 24 h. Cytoplasmic RNA was prepared and analysed using RNase protection mapping. The bands labeled 'test' indicate correctly initiated  $\beta$ -globin transcripts, and the bands labeled 'ref' were generated by the internal standard plasmid ICPOref. RT: incorrectly initiated read-through transcripts of the test templates; riboprobe: aliquot of undigested cRNA; M: size marker, HaeIII digested pBR322.

## 3. Results

### 3.1. VIP and CRF induced gene transcription in a catecholaminergic LC-like cell line

The 5'-flanking regions from mammalian genes contain binding sites for many transcription regulating

proteins that exhibit effects upon transcriptional activation or repression. The tyrosine hydroxylase gene, for example, contains in its 5'-flanking region the *cis*-acting regulatory elements AP2, Ap1, E-box, POU, Sp1, and CRE showing positive and negative effects on tyrosine hydroxylase gene transcription (Lazaroff et al., 1998). To investigate whether VIP and CRF have an impact on gene transcription in CATH.a cells, model promoters and model activator proteins were used. The use of model promoters and activators allowed us to test the effect of CRF and VIP on gene transcription without influences from other promoter elements. As a model transactivator protein, a fusion protein consisting of the DNA binding domain of the yeast transcription factor GAL4 fused to the cDNA of human CREB was constructed (Fig. 1B). Since GAL4 does not bind to any known mammalian gene promoter element, interference by other competing CRE-binding proteins was avoided. As a reporter, five copies of GAL4 binding sites were inserted upstream of the  $\beta$ -globin reporter gene (Fig. 1A). These plasmids were introduced into the catecholaminergic cell line CATH.a, derived from a tumor developed in transgenic mice carrying the SV40 T antigen under the control of the rat tyrosine hydroxylase promoter (Suri et al., 1993). To control for variability in transfection efficiency, cells were additionally transfected with the plasmid ICPOref, containing a mutated  $\beta$ -globin gene under control of the herpes virus ICPO gene promoter. Two days post-transfection, cytoplasmic RNA of the transfected cells was harvested, hybridized to a  $\beta$ -globin-derived cRNA probe, and analyzed for the amount of  $\beta$ -globin mRNA by RNase protection assay. The correctly initiated  $\beta$ -globin message is indicated ('test'). The internal reference plasmid ICPOref protected a shorter fragment of the  $\beta$ -globin riboprobe (labeled 'ref'). Fig. 1C shows that the cAMP analogue dbcAMP (left panel) and the neuropeptides VIP and CRF (middle and right panel) were able to induce transcription of the  $\beta$ -globin reporter gene. This experiment revealed that VIP and CRF receptor activation induced gene transcription at levels similar to that resulting from treatment of cells with a cAMP analogue. In contrast, a truncated CRF peptide,  $\alpha$ -helical CRF9-41, that acts as a CRF antagonist, was unable to activate transcription from the reporter gene (data not shown). We conclude that receptor activation, mediated by CRF or VIP is capable of transducing a signaling cascade required for activating gene transcription.

### 3.2. Design and expression of a nuclear inhibitor for PKA

CRF, VIP and PAC<sub>1</sub> receptor activation stimulate adenylyl cyclase in CATH.a cells (Duman et al., 1992;

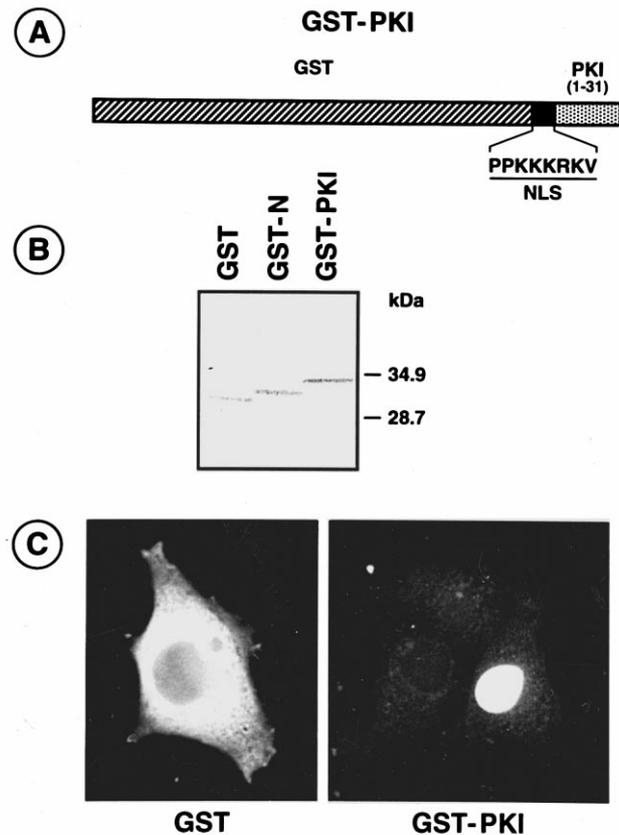


Fig. 2. Design and expression of GST-PKI, a nuclear PKA inhibitor. (A) Structure of the GST-PKI fusion protein. The GST and the PKI moieties are separated by the NLS derived from the large T antigen of SV40 virus. (B) Immunoblot analysis of GST and GST fusion proteins expressed in COS cells. Whole cell extracts of COS cells transfected with expression vectors encoding *S. japonicum* GST, a GST protein tagged with the SV40 large T antigen NLS (GST-N), and the GST-PKI fusion protein were analyzed by immunoblotting using a polyclonal antibody against GST followed by an alkaline phosphatase-labeled secondary antibody. Position of molecular markers are shown on the right. (C) Nuclear targeting of the GST-PKI fusion protein. COS cells were transfected with expression vectors encoding GST and GST-PKI and expression was monitored by indirect immunofluorescence using a polyclonal antibody against GST followed by an FITC-labeled secondary antibody.

Rawlings and Hezareh, 1996; Muller et al., 1997), leading to an increase in the intracellular concentration of cAMP. However, not all of the effects of CRF and VIP have been attributed to a stimulation of adenylyl cyclase. The inhibitory effect of VIP on N-type calcium channels of sympathetic neurons was shown to be independent of PKA (Zhu and Ikeda, 1994). In anterior pituitary corticotrophs, it was reported that CRF and VIP increased the intracellular free calcium levels (Luini et al., 1995). Calcium activates calcium/calmodulin-dependent protein kinases and one of these kinases, calcium/calmodulin-dependent protein kinase IV, translocates into the nucleus and activates transcription, as does PKA, via the CRE/CREB pathway

(Matthews et al., 1994; Sun et al., 1994). Moreover, we found recently that CRF increased MAP kinase activity in CATH.a cells<sup>1</sup>, suggesting that CREB could also be activated by the MAP kinase activated pp90 ribosomal S6 kinase RSK in CATH.a cells as shown for PC12 cells (Xing et al., 1998). To specify the signaling pathway leading from CRF and VIP/PACAP receptor activation to an increase in gene transcription, a nuclear inhibitor for PKA was constructed consisting of *S. japonicum* GST and amino acids 1 to 31 of the PKA inhibitor PKI. PKI, a 75 amino acid, heat-stable protein, binds with a high specificity to the catalytic subunit of PKA. The regulatory subunit of PKA and PKI bind and inhibit the catalytic subunit of PKA by analogous mechanisms. The exact physiological role of PKI is still unclear. It has been postulated that PKI fine-tunes the level of the catalytic activity of the catalytic subunit of PKA (Fantozzi et al., 1994; Seasholtz et al., 1995). The designed GST-PKI fusion protein additionally contained a nuclear localization signal to accomplish nuclear targeting (Fig. 2A). The expression of the GST-PKI protein and control proteins GST and an NLS-tagged GST, encoded by the expression vectors pEBG and pEBGN, respectively, was confirmed by transient transfection in COS cells followed by Western blot analysis using anti-GST antibodies (Fig. 2B). Indirect immunofluorescence was used to verify the intracellular distribution of the GST-PKI protein in COS cells transfected with the GST-PKI expression vector. We chose COS cells to test the expression vectors, because these cells can replicate the pEBG plasmids due to the expression of SV40 large T antigen in the cell and the presence of the SV40 replication origin in the plasmids. Thus, high expression levels of recombinant proteins can be obtained. The results show that the GST-PKI fusion protein was entirely expressed in the nuclei of the transfected cells (Fig. 2C). COS cells transfected with the expression vector for the GST protein alone, showed a clear cytoplasmic distribution (Fig. 2C).

### 3.3. The nuclear targeted GST-PKI fusion protein blocked transcriptional activation of a reporter gene by the CRF and VIP initiated signaling pathway

CATH.a cells were transfected with the 5×GAL4OVEC reporter plasmid, the GAL4-CREB expression plasmid and the internal reference plasmid ICPOref. In addition, cells were transfected with an expression vector encoding GST-PKI, or, as control, the GST expression vector pEBGN. Following transfection, CATH.a cells were treated with dbcAMP, VIP or

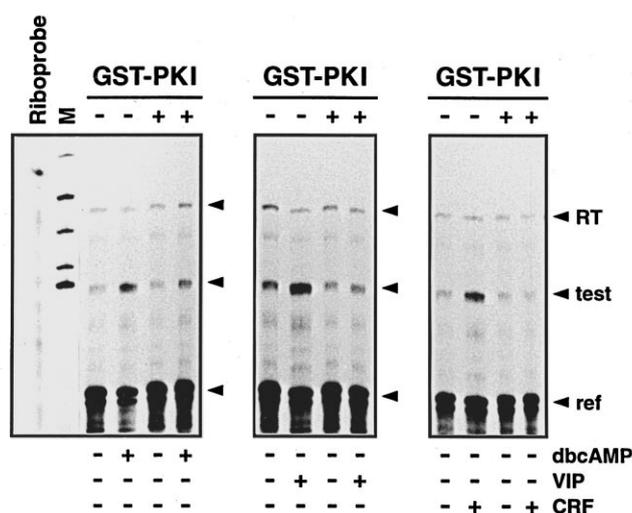


Fig. 3. The GST-PKI fusion protein inhibits VIP and CRF induced transcriptional activation in CATH.a cells. Reporter plasmid 5×GAL4OVEC, expression plasmid pGAL4-CREB and the ICPOref internal standard plasmid were introduced into CATH.a cells together with an expression vector encoding either a nuclear targeted GST (-) or the GST-PKI fusion protein. Cells were incubated in the absence (-) or presence (+) of 1 mM dbcAMP, 1 μM VIP or 0.5 μM CRF for 24 h. Cytoplasmic RNA was analyzed for β-globin mRNA by RNase protection mapping.

CRF for 24 h. Transcription was monitored by RNase protection mapping of the β-globin mRNA. The cAMP analogue dbcAMP and the neuropeptides VIP and CRF increased transcription of the reporter gene in the absence of the PKA inhibitor (Fig. 3). Expression of the GST-PKI fusion protein blocked the transcriptional induction of the reporter gene, clearly indicating the role of PKA in the signalling cascades initiated by CRF and VIP leading to transcription. Furthermore, basal transcription of the reporter gene was impaired by GST-PKI, most likely due to an inhibition of basal PKA activity in the nucleus.

### 3.4. CRF and VIP induced transcription of a reporter gene containing cAMP response elements in its regulatory region

The experiments described so far have shown that CRF and VIP receptor activation leads to changes in gene transcription and, additionally, that PKA is necessary for this activity. These data suggest that endogenous CREB or related bZIP proteins are activated by CRF and VIP via PKA to increase transcription of those genes containing CREB binding sites, such as the CRE sequence, in their regulatory regions. To demonstrate that VIP and CRF induced transcriptional activity can, in fact, be mediated by the CRE sequence, a reporter gene containing in its regulatory region four copies of the CRE derived from the chro-



LC, revealed that both CRF and VIP receptor activation set off a signaling cascade that results in gene transcription of the  $\beta$ -globin reporter gene. Moreover, the signaling pathway required PKA activity in the nucleus, as shown in an experiment using a nuclear PKA inhibitor.

Further clarification of the role of CRF and VIP in regulating gene transcription will be possible through an identification of those genes that are induced by these stimuli in different neuronal populations. Based on the data presented in this study, the most promising candidate genes should contain a CRE sequence in their regulatory regions. Recently, the macrophage migration-inhibitory factor (MIF) gene was identified as CRF-responsive in anterior pituitary cells. The CRF-mediated increase in transcription was executed by a CRE within the 5-regulatory sequence of the MIF gene (Waeber et al., 1998). Another likely CRF-responsive candidate gene is the gene encoding tyrosine hydroxylase. This enzyme catalyzes the reaction from L-tyrosine to 3,4-dihydroxy-L-phenylalanine which is the rate limiting reaction in catecholamine biosynthesis. Interestingly, repeated administration of CRF increased, as does chronic stress, the level of tyrosine hydroxylase in LC neurons (Melia and Duman, 1991). Moreover, it has been reported that transcription of a tyrosine hydroxylase promoter/chloramphenicol acetyltransferase gene could be induced in CATH.a cells by adding dbcAMP to the cells. The CRE was shown to be responsible for >80% of basal (uninduced) transcription of the reporter gene (Lazaroff et al., 1995). Furthermore, when this study was in progress, it was reported that VIP was able to induce transcription of a tyrosine hydroxylase promoter/chloramphenicol acetyltransferase gene in CATH.a cells (Muller et al., 1997). It would be of interest to know whether CRF exhibits a similar effect.

CRF receptor activation leads to an increase in proopiomelanocortin transcription in the pituitary. This effect was attributed to an activation of the transcription factor *c-fos* that subsequently induces proopiomelanocortin gene expression (Boutillier et al., 1991). However, a proopiomelanocortin promoter/chloramphenicol acetyltransferase gene lacking the major exonic *fos* binding site was still CRF responsive (Boutillier et al., 1995). Alternatively, a DNA-binding protein termed PCRH-REB-1 (Jin et al., 1994) might trigger transcriptional induction of the proopiomelanocortin gene via a CRF response element in its regulatory region. PCRH-REB-1 and the CRF response element exhibit no similarities with the CRE-binding proteins of the bZIP family or the CRE sequence, respectively. However, PCRH-REB-1 contains nine possible PKA phosphorylation sites suggesting that the CRF-induced activation of PKA might have an impact on PCRH-REB-1 activity via phosphorylation.

Although PKA induces gene transcription in a variety of genes by phosphorylating bZIP proteins such as CREB, PKA is also able to phosphorylate other transcription factors in the nucleus and change their activities (Rohlf, 1997; Zhong et al., 1998).

Further work will be needed to elucidate the connection between CRF receptor activation and PCRH-REB-1 activity. Moreover, a nonclassical CRE was recently identified in the human proopiomelanocortin gene that functioned as a binding site for CREB and conferred cAMP responsiveness to a reporter gene (Kraus and Hölt, 1995). Interestingly, the genes encoding the neuropeptides CRF and VIP contain CRE sequences in their regulatory regions and activation of the cAMP signaling pathway increased CRF and VIP gene transcription (Tsukada et al., 1987; Dorin et al., 1993; Guardiola-Diaz et al., 1994; Itoi et al., 1996). Therefore, CRF and VIP might control the expression of their own genes by inducing the cAMP signaling pathway.

In conclusion, using a catecholaminergic neuronal cell line, it is shown here that CRF and VIP induce gene transcription mediated by the cAMP signaling pathway. Future work to identify CRF and VIP responsive genes, quite likely mediated by CRE sequences in their regulatory regions, appears promising. The identification of such target genes will help to clarify the molecular basis for the long-term effects of these neuropeptides in the nervous system.

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