Corticotropin Releasing Factor Receptors and Their Ligand Family

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ABSTRACT: The CRF receptors belong to the VIP/GRF/PTH family of G-protein coupled receptors whose actions are mediated through activation of adenylate cyclase. Two CRF receptors, encoded by distinct genes, CRF-R1 and CRF-R2, and that can exist in two alternatively spliced forms, have been cloned. The type-1 receptor is expressed in many areas of the rodent brain, as well as in the pituitary, gonads, and skin. In the rodent, one splice variant of the type-2 receptor, CRF-R2 α , is expressed mainly in the brain, whereas the other variant, CRF-R2B, is found not only in the CNS, but also in cardiac and skeletal muscle, epididymis, and the gastrointestinal tract. The poor correlation between the sites of expression of CRF-R2 and CRF, as well as the relatively low affinity of CRF for CRF-R2, suggested the presence of another ligand, whose existence was confirmed in our cloning of urocortin. This CRFlike peptide is found not only in brain, but also in peripheral sites, such as lymphocytes. The broad tissue distribution of CRF receptors and their ligands underscores the important role of this system in maintenance of homeostasis. Functional studies of the two receptor types reveal differences in the specificity for CRF and related ligands. On the basis of its greater affinity for urocortin, in comparison with CRF, as well as its brain distribution, CRF-R2 may be the cognate receptor for urocortin. Mutagenesis studies of CRF receptors directed toward understanding the basis for their specificity, provide insight into the structural determinants for hormone-receptor recognition and signal transduction.

INTRODUCTION

It is a matter of life or death for an organism that is exposed to environmental perturbations (stressors) to be able to respond in a manner that maintains physiological homeostasis. A major regulator of homeostasis is the corticotropin-releasing-factor (CRF), which mediates the autonomic, behavioral, and neuroendocrine responses to stress. In addition, there are many data suggesting that CRF and its family members are involved in the function of the immune, reproductive, and cardiovascular systems,^{1–3} as well as in pathophysiologic states, such as depression and Alzheimer's disease.^{4–8} Recently, the observation that CRF affects the growth of mammary and lung cancer cells, and that it protects neuronal cells from hypoxia-induced cell death,^{9–11} suggests that CRF may play a role in cell growth and survival.

The family of CRF agonists includes (fish) urotensin, (frog) sauvagine, and the new mammalian CRF agonist, urocortin. The actions of these CRF agonists are initiated by binding and activating receptors that belong to the family of seven-

transmembrane domain, G-protein–coupled receptors that activate adenylate cyclase. In order to understand the mechanism of CRF action, a detailed description of the ligand-receptor interaction at a molecular level is required. This paper describes the cloned CRF receptors and the structural determinants for ligand recognition. Data on the tissue distribution and receptor binding of urocortin are included.

CRF RECEPTOR TYPES AND VARIANTS

To date, two types of CRF receptors, each encoded by a different gene, have been cloned. Using an expression cloning approach, the cDNA for CRF-R1 was obtained from a human Cushing's tumor¹² and from the mouse AtT-20 cell line.¹³ The homologous receptors in rat and human were also cloned and were found to have only minor amino acid differences.^{13–15} Type-1 CRF receptors have subsequently been cloned in sheep (Meyers, Trinh, Myers, Gene Bank #AF054582), chicken,¹⁶ and *Xenopus*.¹⁷ A splice variant of CRF-R1, in which 29 amino acids are inserted into the first intracellular loop, was cloned from the Cushing's tumor.¹² To date, the latter variant has not been found in any other tissue. Another variant, from which the third exon—encoding amino acids 40–80—is deleted, was cloned from human hypothalamus.¹⁸ This variant has been detected in human fetal and placental tissue,¹⁹ as well as in rat testis.²⁰

After CRF-R1 had been cloned, its expression sites in the rat brain were found to be widespread, but not completely overlapping with previously detected binding sites. For example, there are CRF binding sites in the hypothalamus,²¹ but there is very little binding to CRF-R1,²² or expression of its message.^{23,24} The existence of another CRF receptor was confirmed when rodent CRF-R2 was cloned.^{25–28} In the rodent, there are two splice variants, CRF-R2 α and CRF-R2 β , that differ in their N-terminal domains. Subsequently, the human homologs of CRF-R2 α and CRF-R2 β were cloned,^{29,30} as well as a third N-terminal splice variant, CRF-R2 γ .³¹ The sequences of cloned CRF receptors are compared in FIGURE 1.

The CRF receptors belong to the family that includes the growth hormone releasing factor, parathyroid hormone, glucagon, vasoactive intestinal peptide, calcitonin, and secretin receptors. The rodent CRF-R1 and CRF-R2 β are overall 68% similar, at the amino acid level, but they are 79% similar in the transmembrane domains, and 84% similar in the intracellular domains. The major sequence differences between the two types of receptors occur in the extracellular domains (FIG. 1).

All receptors have putative signal peptides, N-glycosylation sites in their N-termini, and eight conserved cysteine residues in their extracellular domains. Consistent with the signal transduction pathway of the native receptors, the cloned receptors transduce a CRF-stimulated accumulation of intracellular cAMP in transfected cells.²⁶ It is relevant, in this respect, that all receptors have the same sequence in the third intracellular loop, which is presumed to play an important role in coupling to the G-proteins (FIG.1).

The genomic structures of the rat and human receptors show that they contain 13 exons and 12 introns, and a general structure similar to that of the genes for parathyroid hormone and glucagon receptors.^{20,32} The splice variant encoding the insertion

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FIGURE 1. Sequence alignment of the cloned CRF receptors./Continued on next two pages.

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in the first intracellular loop, isolated from the human Cushing's tumor, was not detected in the rat gene, but was found in the human gene.³²

## **CRF RECEPTOR EXPRESSION**

After the receptors had been cloned, their expression sites were determined by means of *in situ* hybridization, RNAse protection, and RT-PCR. These results show that there is a wide distribution of CRF-R1 in the rat brain, with high levels in the forebrain, subcortical limbic structures in the septal region, amygdala, cerebellar cortex, and deep nuclei; and low or no expression in the hypothalamus. In keeping with the effects of CRF on the pituitary, CRF-R1 is also expressed in the anterior and intermediate lobes of the rat pituitary.²³ Additionally, CRF-R1 expression is found in mouse testis,³³ human ovary,³⁴ skin,^{35,36} human endometrial stromal cells,³⁷ human leukemic mast cells,³⁸ rat mammary carcinoma,¹⁰ and melanoma cells.³⁹

In the rodent, CRF-R2 $\alpha$  is found mainly in the brain, where it is confined to subcortical structures; with high expression in the lateral septal nucleus, the ventromedial hypothalamus, and in the choroid plexus; with very low expression in the pituitary.²⁴ A distinct site of expression for CRF-R2 $\beta$  is in cerebral arterioles throughout the rat brain.²⁴ Interestingly, in the rodent, CRF-R2 $\beta$  is expressed both in the brain and in the periphery—namely, the epididymis, skeletal muscle, gastrointestinal tract, and heart.^{26–28,40} Additionally, CRF-R2 $\beta$  is expressed in mouse AT-1 myocyte tumor cells.⁴¹ In contrast to the rodent, the human CRF-R2 $\alpha$  is found in the brain and periphery, whereas CRF-R2 $\beta$  and CRF-R2 $\gamma$  are found mainly in the brain.^{30,31}

Using pharmacological differences in affinities for ovine CRF, the binding sites that correspond to CRF-R1 and CRF-R2 in the rat brain were shown to be in distinct, nonoverlapping regions each associated with a specific message expression.²² Pharmacological characterization of CRF-like effects on relaxation of mesenteric small arteries of the rat suggests that the receptors that mediate these effects are CRF-R2;⁴² as are the mediators for rat neonatal cardiomyocytes and a murine cardiomyocyte tumor cell line.⁴¹

# A SECOND MAMMALIAN CRF AGONIST, UROCORTIN

The impetus behind the search for another mammalian, CRF-like ligand was based on four different observations. First, the peptides, urotensin and sauvagine, in fish and frogs, respectively, were thought to be the corresponding orthologs of CRF until peptides more closely related to CRF were cloned in these species.^{43,44} Second, urotensin and sauvagine are nearly equipotent with CRF on CRF-R1, but are more potent than CRF on CRF-R2.^{25–27} Third, there was specific urotensin-like immunoreactivity in rat brain and other tissues. Finally, the sites of expression of CRF-R1 and CRF-R2 in the rat brain are largely nonoverlapping,²⁴ showing a lack of correspondence, in some areas, with the expression of CRF. Using an antibody to urotensin, a peptide named urocortin was cloned from rat brain.⁴⁵ Subsequently, the human homolog was cloned.⁴⁶ Urocortin is 63% homologous to urotensin and 43%

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FIGURE 2. Sequences of CRF family members.

Compound		Inh	ibitory binding	constant, K _i (n]	q(W		cAMP, E	C ₅₀ (nM) ^c
	hCRF-R1 ^d	hCRF-R1 ^e	rCRF-R2α ^d	$rCRF-R2\alpha^{e}$	mCRF-R2β ^d	mCRF-R2β ^ε	rCRF-R1	mCRF-R2β
Astressin	2.0	1.6	1.5	2.8	1.0	0.87	n/a	n/a
	(1.8-2.3)	(0.95 - 2.7)	(0.81 - 2.8)	(1.7 - 4.7)	(0.39-2.8)	(0.56 - 1.4)		
α-helical CRF(9–41)	17	49	5.0	4.1	0.97	0.81	n/a	n/a
	(13–21)	(29–82)	(2.6-10)	(2.5–6.7)	(0.43 - 2.2)	(0.52 - 1.3)		
[DPhe ¹² ,Nle ^{21,38} ]r/	56	75	5.2	31	8.4	6.9	n/a	n/a
hCRF(12-41)	(41–75)	(64–88)	(2.6-10)	(19–51)	(4.4 - 16)	(5.0 - 9.5)		
r/hCRF	11	5.2	44	13	38	17	$1.9\pm0.8$	$1.7 \pm 0.4$
	(8.4–15)	(2.9 - 9.3)	(26–75)	(7.2–22)	(21–67)	(10-29)		
rUcn	13	0.79	1.5	0.58	0.97	0.41	$0.8 \pm 0.1$	$0.18 \pm 0.04$
	(0.71 - 2.6)	(0.43 - 1.4)	(0.74 - 3.2)	(0.42 - 0.82)	(0.46 - 2.0)	(0.26 - 0.66)		
sfUrotensin I	3.1	2.8	9.8	3.4	6.4	3.0	$3.1 \pm 1.7$	$0.74 \pm 0.1$
	(2.4 - 4.0)	(2.4 - 3.3)	(5.5–17)	(2.6 - 4.4)	(3.8-11)	(1.8-4.8)		
Sauvagine	9.4	11	9.9	1.4	3.8	2.0	$2.5 \pm 1.4$	$0.5 \pm 0.2$
	(7-13)	(8.8 - 13)	(4.4-22)	(1.1-1.8)	(2.3 - 6.2)	(1.1 - 3.6)		
^a The values for this tab	le are taken from	1 Refs. 45 and 60						

TABLE 1. Relative potencies of CRF ligands^a

b 95% confidence limits in parentheses.  $c \pm \text{SEM}$ .  $d ^{125}\text{I} - [\text{DTyr}^1] \text{Astressin as radioligand}$ .  $e ^{125}\text{I} - [\text{Tyr}^0]\text{rUcn as radioligand}$ .

PERRIN & VALE: CRF RECEPTORS AND THEIR LIGANDS

homologous to CRF. The sequences for urocortin and the other CRF family members are shown in FIGURE 2, from which it can be seen that urocortin is more closely related to urotensin than to CRF.

Urocortin is detected in many regions of the rat brain, including the Edinger-Westphal nucleus, lateral superior olive,⁴⁵ substantia nigra, ventral tegmental area, linear and dorsal raphe nuclei,^{47,48} and the hypothalamus.⁴⁹ In the human brain, urocortin-like immunoreactivity and message are observed in every region, with the highest concentrations found in the frontal cortex, temporal cortex, and hypothalamus.⁵⁰

Urocortin is also found in peripheral sites. For example, immunoreactive urocortin is detected in the rat digestive system and the pituitary.^{49,51} Human anterior pituitary was also found to express both the message and immunoreactivity for urocortin.⁵² Other sites of expression of urocortin mRNA and of the peptide are mucosal inflammatory cells in the human gastrointestinal tract,⁵³ human placenta, and fetal membranes.⁵⁴ Using RT-PCR, urocortin mRNA is detected in primary cultures of rat cardiomyocytes as well as in cardiac myocyte cell lines.⁵⁵

By combining RT-PCR and radioimmunoassay, it was found that normal human lymphocytes produce urocortin but not CRF.⁵⁶ The involvement of CRF-like ligands in immune function is further highlighted by the observation that both CRF and urocortin suppress experimental autoimmune encephalomyelitis in rats.⁵⁷ These neuropeptides may act directly on the components of the immune system, rather than indirectly through the activation of the HPA.

The regions of urocortin immunoreactivity overlap with the regions of expression of CRF-R2 in many areas of the brain⁴⁷ and some effects of urocortin appear to be mediated by CRF-R2. For example, urocortin was found to be approximately six times more potent than CRF as an inhibitor of heat-induced edema.⁵⁸ Furthermore, blockade of CRF-R1 does not affect the CRF- and urocortin-induced decrease in food intake in rats.⁵⁹ Also, urocortin protects myocytes, cells that express CRF-R2, from hypoxia-induced cell death.⁵⁵

Pharmacologic studies have shown that urocortin is more potent than CRF on both types of CRF receptors, but in general, the difference in potencies is greater for CRF-R2.^{45,60} The affinities of selected CRF ligands and the potencies in stimulating cAMP in cells expressing the two types of CRF receptors are given in TABLE 1. Astressin is a CRF peptide antagonist that has high affinity for both types of CRF receptors.⁶¹ From TABLE 1 it can be seen that the antagonist  $\alpha$ -helicalCRF(9–41) has a greater affinity for CRF-R2 than for CRF-R1. This observation may explain the fact that  $\alpha$ -helicalCRF(9–41) blocks heat-induced edema at doses that do not block CRF-stimulated ACTH release.⁵⁸

# STRUCTURAL REQUIREMENTS FOR CRF RECEPTOR BINDING—MUTAGENESIS STUDIES

One approach to assessing the structural determinants for receptor function uses mutagenesis to determine regions of the receptor that may be involved in binding and signaling. In one study, chimeric receptors were created in which domains of the GRF receptor were replaced by the corresponding domains of the CRF-R1. A chi-

meric receptor in which the ECD-1 of the CRF-R1 was replaced by the ECD-1 of the GRF-R, does not bind either labeled astressin or labeled urocortin. A chimera in which the ECD-1 of the GRF-R was replaced by the corresponding domain of the CRF-R, binds astressin and urocortin with a dissocuation constant  $K_d \sim 10$  nM. Additional chimeras were created to explore the role of the extracellular loops, and it was found that a chimera expressing all three extracellular loops, together with the N-terminus of the CRF-R, displays nearly the same affinity as the wild-type receptor.⁶²

As a further test of the role of the ECD-1 in binding CRF analogs, a chimera was created, in which the ECD of the activin type-2 receptor was replaced by the ECD-1 of CRF-R1 (CRF-R/Act-R). The activin receptor is a single transmembrane serine/ threonine kinase.⁶³ This chimera binds astressin with nanomolar affinity (see TABLE 2). In another study, the ECD-1 was expressed as a soluble protein and it was found to display only low binding affinity for a CRF agonist.⁶⁴ These data suggest that the N-terminal domain of the CRF-R1 contains major binding determinants for CRF agonists and antagonists, but that it must be anchored to the cell membrane.

In CRF-R1, there eight conserved cysteine residues in ECD-1, -2, and -3. The role of these residues was studied by mutating them, either singly or in pairs. It was found that there is potential pairing of the Cys residues, #44 with #102, #68 with #87; and the two residues in ECD-2 and -3, #188 with #258. Mutations of C30 and C54 appear to have little effect on receptor binding. Additionally, mutations of the cysteine residues within the transmembrane and intracellular domains have no effect on either binding or signaling.⁶⁵

Another question relates to the origin of CRF receptor selectivity as manifested by greater potency of CRF in signaling by CRF-R1 compared to CRF-R2.²⁵ The amino acids that differ between CRF-R1 and CRF-R2 were exchanged and it was found that substitution of Val266, Tyr267, and Thr 268 in CRF-R1 by the corresponding residues in CRF-R2 increases the EC-50 for CRF-stimulated cAMP accumulation by a factor of about ten. Other residues in ECD-2 (#175–#178, and #189) also appear to be involved in the CRF response of CRF-R1.^{66,67}

At the amino acid level, the *Xenopus* CRF-R1 is about 80% homologous to the human CRF-R1, but the two receptors differ in their affinities for sauvagine, r/h CRF, and ovine CRF.¹⁷ Whereas hCRF-R1 is not selective for these ligands, the xCRF-R1 has about one tenth the affinity for ovine CRF and sauvagine, as compared to CRF. Using mutagenesis, it was found that the five amino acids #76, #81 #83, #88, and #89 in the N-terminal domain are responsible for the observed differences.⁶⁸ These data are consistent with some published results,⁶² but they contrast with others.^{66,67}

The avian CRF-R1 is about 88% identical to that of the human, rat, and mouse CRF-R1, with many amino acid differences in the ECD-1, and the remainder of the differences scattered throughout the rest of the sequence.¹⁶ In the avian receptor the relative potencies are urotensin  $\cong$  sauvagine > CRF, for both binding and signaling. This pattern is more like that seen for the CRF-R2 than for CRF-R1. No mutagenesis study has yet been reported for the avian receptor, but the ECD-1 is again implicated in the ligand specificity because most of the amino acid differences appear in this domain.

Recently a series of nonpeptide CRF antagonists have been developed that bind, specifically and with high affinity, to type-1 receptors; and inhibit, both *in vitro* and

	Receptor	Ast/Ast* $K_i (nM)^b$	Ucn/Ucn* $K_i (nM)^b$
Ŵ	rCRF-R (wt)	1.8 (1.1–2.9)	3.0 (1.6-5.7)
- <del>M</del>	rGRF-R (wt)	c	c
Ŵ	E1 _g /CRF-R	c	c
	E1 _c /GRF-R	13 (7.7–22)	12 (1.9–74)
	E1 _c /E2 _c /GRF-R	10 (7.1–14)	10 (4.5–22)
Ŵ	E1 _c /E3 _c /GRF-R	7.1 (3.8–13)	c
	E1 _c /E4 _c /GRF-R	4.2 (3.0–5.8)	1.6 (0.58–4.4)
	E1 _c /E2 _c /E3 _c /GRF-R	5.4 (3.5–8.4)	<i>c</i>
	E1 _c /E2 _c /E4 _c /GRF-R	7.9 (5.3–12)	7.0 (3.5–14)
	E1 _c /E3 _c /E4 _c /GRF-R	11 (5.2–23)	7.6 (2.1–27)
	E1 _c /E2 _c /E3 _c /E4 _c /GRF-R	4.3 (1.9–9.6)	1.3 (0.71–2.4)
	E1 _c /ActIIB-R	3.5 (1.8–7.0)	c

TABLE 2. Inhibitory dissociation constants,  $K_i$  (nM), for astressin and urocortin bound to COSM6 cells transfected with various chimeric and mutant receptors a 

 $\overline{{}^{a}$ The  $K_i$  values were determined from homologous displacement assays, assuming that the  $K_d$  values for Ast* and Ucn* were not significantly different from those for astressin and urocortin, ^bValues in parentheses are 95% confidence limits.

^{*c*}The specific binding was too low to measure the  $K_i$  accurately.

*in vivo*, CRF actions mediated by them.^{69–72} One study found that the mutations H199V and M276I, two of the residues in transmembrane domains 3 and 5 that differ between CRF-R1 and CRF-R2, reduce the affinity of CRF-R1 for a nonpeptide antagonist but have no effect on the affinity for CRF.⁶⁷ In another study, a different nonpeptide antagonist, antalarmin,⁷² displaced the labeled antagonist, astressin, to a much smaller degree than it displaced labeled oCRF from both the cloned CRF-R1 and the native cerebellar receptor—a tissue that predominantly expresses CRF-R1 (Perrin, unpublished). These data suggest that the binding domains of peptide and nonpeptide antagonists are different.

## CONCLUSIONS

To date, four CRF receptors and one novel CRF ligand have been cloned. The type-1 receptor, widely expressed in the central nervous system and in the pituitary, is the major mediator for activation of the HPA axis. The fact that CRF-R1 is found in the reproductive and immune systems, as well as in spleen, adrenal, and skin, suggests that this receptor mediates many diverse actions. The type-2 receptor exists in different splice-variant forms that differ in their N-terminal sequences and in their tissue expression. In both the rodent and human, one variant is restricted to the central nervous system, whereas the other variants are found not only in the brain but also in peripheral sites.

A novel CRF agonist, urocortin, has been cloned and found to have characteristics that are similar, but not identical, to those of CRF. Urocortin is widely expressed in the brain, as well as in many of the peripheral tissues that express the type-2 receptor. Urocortin binds with high affinity to both types of receptors but is significantly more potent than CRF on the type-2 receptor.

Studies of mutant CRF receptors have shown that recognition of CRF peptides is governed, to a large extent, by residues in the extracellular domains, especially the N-terminus. Furthermore, the binding of nonpeptide CRF-R1–selective antagonists involves receptor domains that differ from those involved in binding of the peptide antagonists.

The future holds exciting prospects for cloning other CRF receptors and discovering novel sites of expression and action. The cloning of urocortin raises the possibility that other CRF-like molecules exist in mammals. Future studies of CRF receptor and/or urocortin knock-out mice will undoubtedly disclose expanded roles for the CRF family and increase our understanding of the roles of CRF in normal and pathological states.

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