

Immune Challenge and Immobilization Stress Induce Transcription of the Gene Encoding the CRF Receptor in Selective Nuclei of the Rat Hypothalamus

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The present study investigated the effect of intraperitoneal (i.p.) administration of endotoxin lipopolysaccharide (LPS) and immobilization stress on the genetic expression of corticotropin-releasing factor receptor (CRF-R) in the brains of conscious male Sprague–Dawley rats. One group of rats was killed at 1, 3, 6, 9, and 12 hr after a single intraperitoneal injection of either the LPS (250 μ g/100 gm of body weight) or the vehicle solution; the other group was killed before, immediately after, 1.5, 3, 6, and 12 hr after a 90 min acute session of immobilization stress. Rats were deeply anesthetized and rapidly perfused with a solution of 4% paraformaldehyde–borax. Frozen brains were mounted on a microtome and cut from the olfactory bulb to the medulla in 30 μ m coronal sections. mRNA encoding the rat CRF-R was assayed by *in situ* hybridization histochemistry using a 35 S-labeled riboprobe, and CRF-R localization within CRF-immunoreactive neurons in the PVN was determined using a combination of immunocytochemistry and *in situ* hybridization techniques. Strong basal levels of CRF-R transcripts were observed in several regions of the brain (piriform cortex, medial and basolateral nuclei of the amygdala, red nucleus, pontine gray, cerebellum, laterodorsal tegmental nucleus, caudal division of the zona incerta, nucleus incertus, spinal and principal sensory nuclei of the trigeminal nerve, and various layers of the cortex). A low to moderate signal was also detected in multiple sites (medial septal nucleus, nucleus of the diagonal band, supraoptic nucleus, arcuate nucleus of the hypothalamus, interpeduncular nucleus, and nucleus prepositus). Whereas vehicle-treated and control rats displayed hardly detectable signals of CRF-R mRNA in the paraventricular nucleus (PVN), CRF-R gene transcription was highly stimulated by LPS administration and immobilization stress in this hypothalamic structure. Indeed, the CRF-R mRNA signal was positive in the dorsomedial parvocellular PVN 3 hr after LPS injection, strong and maximum in both

parvo- and magno-PVN at 6 hr postinjection, and declined 9 and 12 hr after treatment. Similarly, 90 min and 3 hr after the immobilization session, mRNA encoding the CRF-R was highly expressed in the parvo-PVN and totally vanished 12 hr after the stress. A lower but significant increase in the CRF-R transcript signal was also observed in the supraoptic nucleus 6 hr after the LPS treatment. Systemic endotoxin and immobilization stress did not modulate the expression of CRF-R gene in other regions, which suggests that these types of challenges can induce a highly selective activation of CRF-R within hypothalamic nuclei directly involved in the regulation of neuroendocrine functions. CRF-immunoreactive neurons of the parvo-PVN expressed the CRF-R transcript after both the stress and the systemic LPS administration. Thus, CRF may play a role as neuromodulator, controlling directly the activity of neuroendocrine CRF motoneurons during immune challenge and other stressful circumstances for the organism.

[Key words: corticotropin-releasing factor, hypothalamic-pituitary-adrenal axis, *in situ* hybridization histochemistry, immunocytochemistry, immune response, male rats, neuroendocrinology, paraventricular nucleus of the hypothalamus, supraoptic nucleus of the hypothalamus]

Stress is associated with activation of the hypothalamic-pituitary-adrenal (HPA) axis, and this activation has indeed been proposed as a defining feature of the stress response. The primary mediator of this response is corticotropin-releasing factor (CRF), a 41-residue peptide acknowledged as the principal hypophysiotropic factor driving stress-induced adrenocorticotrophic hormone (ACTH) secretion (Vale et al., 1981; Rivier and Plotzky, 1986). Although CRF is widely distributed within the CNS, the paraventricular nucleus (PVN) of the hypothalamus is the principal site of the parvocellular neurosecretory neurons responsible for delivering CRF to the hypophyseal portal system, an event that initiates the activity of the pituitary–adrenal axis (for review, see Sawchenko and Swanson, 1990). Indeed, destruction of this hypothalamic nucleus abolishes the increase in plasma ACTH and corticosterone levels caused by various acute challenges (Bruhn et al., 1984; Makara et al., 1986; Rivest and Richard, 1990; Rivest and Rivier, 1991). Moreover, stressors such as single and repeated exercise sessions (Rivest et al., 1992), foot-shock stress (Imaki et al., 1991; Rivest and Rivier, 1994), immobilization (Imaki et al., 1992a), systemic administration of the bacterial lipopolysaccharide (LPS) (Laflamme and Rivest, 1994), and treatment with interleukin-1 β (Gong et al., 1991; Rivest et al., 1992; Brady et al., 1994; Ericsson et al.,

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1994; Rivest and Rivier, 1994) all stimulate the immediate *early* gene (IEG) *c-fos* and CRF gene expression in the parvocellular division of the rat PVN. These independent indices of neuronal activation provide evidence that neuroendocrine CRF motoneurons are highly stimulated during various stressful conditions, and this phenomenon appears crucial to the appropriate control of ACTH release from the adenohypophysis.

Stress-induced alteration of CRF neuronal activity is not, however, limited to the PVN; exposure to foot shocks, for example, causes a rapid upregulation of CRF mRNA in Barrington's nucleus, a pontine cell group involved in the autonomic control of micturition (Imaki et al., 1991; Sawchenko et al., 1993). Imaki et al. (1991) have also reported that foot shock can produce a downregulation of CRF gene expression in the olfactory bulb, a structure known as the main relay for a sensory modality governing most rodent behavior (Sawchenko et al., 1993). Interestingly, although levels of CRF transcripts in the PVN are negatively regulated by glucocorticoids, Makino et al. (1994) found that chronic treatment with corticosterone increases CRF mRNA expression in the central nucleus of the amygdala. Within the PVN itself, CRFergic systems seem to respond differently to glucocorticoids because in adrenalectomized rats treated with corticosterone, CRF mRNA increased in the autonomic component of this hypothalamic nucleus (dorsal cap) and decreased in the medial parvocellular neurons (Swanson and Simmons, 1989). Glucocorticoids may therefore restrain their inhibitory influence on neuroendocrine CRF motoneurons during stress. Salt-loading challenge, on the other hand, causes a decrease in CRF transcripts in Barrington's nucleus and a marked increase in the magnocellular neurosecretory cells, an effect accompanied by a paradoxical reduction in CRF mRNA levels in the parvocellular PVN (Young, 1986; Imaki et al., 1992b; Sawchenko et al., 1993). Exercise training on a treadmill can also modulate CRF gene expression in sites other than the parvocellular PVN, such as the medial part of the medial geniculate complex and the lateral subdivision of the medial mammillary nucleus (Rivest et al., 1992). Relative levels of CRF mRNA in the parvocellular PVN are also significantly increased following an acute immune challenge with systemic LPS injection, but endotoxin did not alter the genetic expression of this stress-related neuropeptide in other brain regions (Laflamme and Rivest, 1994). Although the stimulation of CRF neurons in various regions of the brain depends on the type and intensity of the stress, little is known about the sites of action of the peptide in the brain of stressed animals.

In addition to being a potential modulator of the stress responses in various regions of the brain, CRF can also positively modulate its own expression. Indeed, we recently reported that central administration of the neuropeptide causes a rapid induction of *c-fos* and NGFI-B in CRF-immunoreactive (ir) neurons of the rat PVN (Parkes et al., 1993). This IEG induction was also followed by an increase in the levels of CRF mRNA (Parkes et al., 1993), suggesting that central CRF per se can participate in the stimulation of neuroendocrine CRF motoneurons. Whether this effect is direct and whether the gene encoding the CRF receptor is expressed in the PVN during stress are open questions.

Vale's group (Chen et al., 1993; Perrin et al., 1993) recently cloned a cDNA coding for the CRF receptor that is widely distributed throughout the rat brain (Potter et al., 1994), but regulation of that gene in the brains of stressed animals has not been studied. It is in fact crucial to determine the effects of stressful

situations on the gene encoding the CRF-R in the brain, because although CRF appears to be a determinant modulator of many stress responses, the exact sites of action as well as the type of cells targeted by the peptide in the brain of challenged animals remain to be fully established. The purposes of the present study were therefore to examine the influence of a severe acute immune challenge and immobilization stress on the expression of the mRNA encoding the CRF receptor (CRF-R) in the brains of adult male rats; and to investigate whether the CRF-R was expressed in CRF-immunoreactive neurons in the PVN during normal or emergency circumstances for the organism. Immune challenge (systemic administration with the bacterial endotoxin LPS) and immobilization were used to verify the effects of a systemic and a neurogenic stress, respectively.

Materials and Methods

Animals. Adult male Sprague-Dawley (~230–260 gm) were acclimated to standard laboratory conditions (14 hr light, 10 hr dark cycle; lights on at 0600 and off at 2000) with free access to rat chow and water. Each rat was used for experimentation only once, and all protocols were approved by the Laval University Animals Welfare Committee.

Systemic LPS administration. In the morning (~0830), rats received either a single intraperitoneal injection of LPS (Sigma, L-2880, lot 122H4024, 250 µg/100 gm of body weight) diluted in 300 µl of sterile saline (0.9%) or the vehicle solution. This dose was selected on the basis of our previous findings that it elicits a robust activation of both IEG *c-fos* and NGFI-B in multiple regions of the brain and CRF gene expression in the PVN of adult male rats (Laflamme and Rivest, 1994). The purpose of that dose was also to investigate the influence of a severe immune activation on the expression of the gene encoding the CRF-R in the brain. The rats were conscious and freely moving at all times throughout the experimental procedure and no mortality was observed in LPS-treated animals. One, three, six, nine, and twelve hours after the intraperitoneal treatments with the bacterial endotoxin or the vehicle solution, the animals were deeply anesthetized via an intraperitoneal injection of 0.3 ml of a mixture of ketamine hydrochloride and xylazine, and then rapidly perfused transcardially with 0.9% saline, followed by 4% paraformaldehyde in 0.1 M borax buffer (pH 9.5 at 4°C). The time points were chosen on the basis of preliminary studies, which showed a strong signal for IEG mRNAs and proteins as well as CRF transcript 3 and 6 h after intraperitoneal LPS injection in the rat brain (Laflamme and Rivest, 1994). Brains were removed from the skull, postfixed for 4–8 d, and then placed in 10% sucrose in the solution of 4% paraformaldehyde–borax buffer overnight at 4°C. The frozen brains were mounted on a microtome (Reichert-Jung, Cambridge Instruments Company, Deerfield, IL) and cut into 30 µm coronal sections. The slices were collected in a cold cryoprotectant solution (0.05 M sodium phosphate buffer, 30% ethylene glycol, 20% glycerol) and stored at –20°C.

Acute immobilization session. Rats were either killed before (time 0: morning at ~0830) or placed in individual immobilizers for 90 min. Immobilization was performed using adjustable restraining cages (Centrap cages, Fisher Scientific 01-282-10). Cages were adjusted very tightly so that animals were unable to move once in the immobilizer, which explains our use of the term immobilization rather than restraint stress. After the stress session, rats were either killed or placed in their home cages until killed. Immediately, 1.5, 3, 6, and 12 hr after the end of the stress session, the animals were deeply anesthetized with an intraperitoneal injection of 0.3 ml of a mixture of ketamine hydrochloride and xylazine, and then rapidly perfused transcardially with saline, followed by 4% paraformaldehyde in 0.1 M borax buffer (pH 9.5 at 4°C). The brains were removed from the skull, postfixed for 4–8 d, and then placed in a solution of 10% sucrose and 4% paraformaldehyde–borax buffer overnight at 4°C. Frozen brains were mounted on a microtome (Reichert-Jung, Cambridge Instruments Company, Deerfield, IL) and cut in 30 µm coronal sections. The slices were collected and stored as described previously (see LPS treatment).

In situ hybridization histochemistry. Hybridization histochemical localization of CRF-R transcript was carried out in one in six series (every sixth section) of brain slices through the brain (from the olfactory bulb to the end of the medulla) using ³⁵S-labeled cRNA probes. Protocols for riboprobe synthesis, hybridization, and autoradiographic localization of mRNA signal were adapted from Simmons et al. (1989). All solu-

tions were treated with diethylpyrocarbonate (Depc) and sterilized to prevent RNA degradation. Tissue sections mounted onto poly-L-lysine-coated slides were desiccated under vacuum overnight, fixed in 4% paraformaldehyde for 30 min, and digested by proteinase K (10 µg/ml in 50 mM Tris HCl, pH 8.0, and 50 mM EDTA, at 37°C for 25 min). Thereafter, the brain sections were rinsed in sterile Depc water followed by a solution of 0.1 M triethanolamine (TEA, pH 8.0), acetylated in 0.25% acetic anhydride in 0.1 M TEA, and dehydrated through graded concentrations of alcohol (50, 70, 95, and 100%). After vacuum drying for a minimum of 2 hr, 90 µl of hybridization mixture (10⁷ cpm/ml) was spotted on each slide, sealed under a coverslip, and incubated at 60°C overnight (~15–20 hr) in a slide warmer. Coverslips were then removed and the slides were rinsed in 4× SSC at room temperature. Sections were digested by RNase A (20 µg/ml, 37°C, 30 min), rinsed in descending concentrations of SSC (2×, 1×, 0.5× SSC), washed in 0.1× SSC for 30 min at 60°C (1× SSC: 0.15 M NaCl, 15 mM trisodium citrate buffer, pH 7.0), and dehydrated through graded concentrations of alcohol. After being dried for 2 hr under the vacuum, the sections were exposed at 4°C to x-ray film (Kodak) for 12–20 hr, defatted in xylene, and dipped in NTB2 nuclear emulsion (Kodak; diluted 1:1 with distilled water). Slides were exposed for 7–10 d, developed in D19 developer (Kodak) for 3.5 min at 14–15°C, and fixed in rapid fixer (Kodak) for 5 min. Thereafter, tissues were rinsed in running distilled water for 1–2 hr, counterstained with thionin (0.25%), dehydrated through graded concentrations of alcohol, cleared in xylene, and coverslipped with DPX.

cRNA probe synthesis and preparation. The rat CRF-R probe (1.3 kb) was generated from the PstI-PstI fragment of the rat prCRF PP1.3-BS cDNA (Dr. W. Vale, Peptide Biology Laboratory, The Salk Institute) (Perrin et al., 1993), subcloned into pBluescript II SK (Stratagene, La Jolla, CA), and linearized with BamH I and HindIII (Pharmacia) for antisense and sense probes, respectively (Perrin et al., 1993). Radioactive cRNA copies were synthesized by incubation of 250 ng linearized plasmid in 6 mM MgCl₂, 40 mM Tris (pH 7.9), 2 mM spermidine, 10 mM NaCl, 10 mM dithiothreitol, 0.2 mM ATP/GTP/CTP, α-³⁵S-UTP, 40 U RNasin (Promega, Madison, WI), and 20 U T7 and T3 RNA polymerase for CRF-R antisense and sense, respectively, for 60 min at 37°C. Unincorporated nucleotides were removed using the ammonium-acetate method; 100 µl of DNase solution (1 µl DNase, 5 µl of 5 mg/ml tRNA, 94 µl of 10 mM Tris/10 mM MgCl₂) was added and 10 min later an extraction was accomplished using a phenol-chloroform solution. The cRNA was precipitated with 80 µl of 5 M ammonium acetate and 500 µl of 100% ethanol for 20 min on dry ice. The pellet was washed with 500 µl ethanol, dried, and resuspended in 100 µl of 10 mM Tris/1 mM EDTA. A concentration of 10⁷ cpm probe was mixed into 1 ml of hybridization solution [500 µl formamide, 60 µl 5 M NaCl, 10 µl 1 M Tris, pH 8.0, 2 µl 0.5 M EDTA, pH 8.0, 20 µl 50× Denhardt's solution, 200 µl 50% dextran sulfate, 50 µl 10 mg/ml tRNA, 10 µl 1 M DTT (118 µl Depc water – volume of probe used)]. This solution was mixed and heated for 5 min at 65°C before being spotted on slides.

Quantitative analysis. Semiquantitative analysis of hybridization signals for CRF-R mRNA was carried out in nuclear emulsion-dipped slides over the confines of cells within various structures expressing the CRF-R using an Olympus Optical System (BX-50, βMax) coupled to a Macintosh computer (PowerPC 7100/66) and IMAGE software (version 1.55 non-FPU, W. Rasband, NIH). The OD of the hybridization signal was measured under dark-field illumination at a magnification of 10× within a 10 mm diameter circular frame. Sections from the experimental and control animals were matched for rostrocaudal level. The size of the frame was chosen on the basis of the mean size range of neurons in these respective regions. The regions (PVN, SON, and so on) were digitized and subjected to densitometric analysis, yielding measurements of integrated OD (area of nucleus × average optical density). The OD of each specific region was then corrected for the average background signal, which was determined by sampling cells immediately outside the cell group of interest (McCabe and Pfaff, 1989).

Combination of immunocytochemistry with *in situ* hybridization. Immunocytochemistry (CRF-immunoreactive neurons) was combined to the *in situ* hybridization histochemistry protocol (CRF-R mRNA) to determine whether CRF neurons located particularly in the PVN express the gene encoding the receptor for this stress-related peptide. Every sixth tissue slice was processed by using the avidin-biotin bridge method with peroxidase as a substrate. Briefly, slices were washed in sterile Depc-treated 0.05 M potassium phosphate-buffered saline (KPBS) and incubated at 4°C with CRF antibody mixed in sterile KPBS, 0.4% Triton

X-100, 0.25% heparin sodium salt USP (ICN Biomedicals Inc., Aurora, OH), and 1% bovine serum albumin (fraction V, Sigma, St. Louis, MO). Rabbit antihuman/rat CRF serum (code PBL rc 70, 8/9/83 bleed), a generous gift of Dr. Wylie Vale (Peptide Biology Laboratory, The Salk Institute, La Jolla, CA), was used at a concentration of 1:2000. Approximately 48 hr after incubation with the primary antibody, the brain slices were rinsed in sterile KPBS and incubated with a mixture of KPBS + heparin + biotinylated goat anti-rabbit IgG (1:1500 dilution; Vector Laboratories, CA) for 60 min. Sections were then rinsed with KPBS and incubated at room temperature for 60 min with an avidin-biotin-peroxidase complex (Vectastain ABC elite kit, Vector Laboratories, CA). After several rinses in sterile KPBS, the brain slices were reacted in a mixture containing sterile KPBS, the chromogen 3,3'-diaminobenzidine tetrahydrochloride (DAB, 0.04%), and 1% hydrogen peroxide (H₂O₂).

Thereafter, tissues were rinsed in sterile KPBS, mounted onto poly-L-lysine-coated slides, desiccated under vacuum overnight, fixed in 4% paraformaldehyde for 30 min, and digested by proteinase K (10 µg/ml in 100 mM Tris HCl, pH 7.5, and 50 mM EDTA, at 37°C for 25 min). Prehybridization, hybridization, and posthybridization steps were performed as described above with the difference of dehydration (alcohol 50, 70, 95, 100%), which was shortened to avoid decoloration of CRF cells (brown staining). After being dried for 2 hr under the vacuum, sections were exposed at 4°C to x-ray film (Kodak) for 24–48 hr, defatted in xylene, and dipped in NTB2 nuclear emulsion (Kodak; diluted 1:1 with distilled water). Slides were exposed for 3 weeks, developed in D19 developer (Kodak) for 3.5 min at 15°C, and fixed in rapid fixer (Kodak) for 5 min. Thereafter, tissues were rinsed in running distilled water for 1–2 hr, rapidly dehydrated through graded concentrations of alcohol, cleared in xylene, and coverslipped with DPX. The presence of CRF-R transcripts was evident as silver grains in perikarya, and CRF immunoreactivity within the cell cytoplasm was stained in brown.

Statistical analysis. Data shown in Figures 6 and 7 are expressed as OD for CRF-R mRNA in the hypothalamic PVN and SON of control or challenged rats. The results of the LPS experiment were analyzed by a 2 × 5 analysis of variance (ANOVA), and post hoc comparisons were done using an adaptation of the Dunn-Sidak test (Kirk, 1982). Factors were identified as follows: *systemic treatment*, which was composed of two levels (intraperitoneal vehicle or intraperitoneal LPS), and *time postinjection*, which combined five levels (1, 3, 6, 9, and 12 hr postinjection). Data for the stress experiment were analyzed by a 1 × 6 ANOVA, and post hoc comparisons were done using an adaptation of the Dunn-Sidak test. Factor was identified as the *time related to immobilization*, including six levels: before (time 0), immediately after (time 1.5), 1.5 (time 3), 3 (time 4.5), 6 (time 7.5), and 12 (time 13.5) hr after the session.

Results

Distribution of CRF-R transcript throughout the rat brain

Figures 1 and 2 illustrate representative examples of the rostrocaudal distribution of the CRF gene expression in the brains of male Sprague-Dawley rats. Hybridized tissues with sense probe did not exhibit detectable signal in any of the regions that showed positive signal with antisense probe (results not shown). As recently reported by Potter et al. (1994), strong basal levels of CRF-R transcript were observed in several regions of the brain, such as the piriform cortex, the medial nucleus of the amygdala, the basolateral nucleus of the amygdala, the caudal division of the zona incerta, the red nucleus, the laterodorsal tegmental nucleus, the pontine gray, the cerebellum, the nucleus incertus, the spinal nucleus of the trigeminal nerve (oral part), the principal sensory nucleus of the trigeminal nerve, the supra-geniculate nucleus, the external cuneate nucleus, and various layers of the cerebral cortex. A low to moderate signal was also detected in multiple sites, including the medial septal nucleus, the nucleus of the diagonal band, the bed nucleus of the stria terminalis, the supraoptic nucleus (SON), the dorsomedial nucleus of the hypothalamus, the arcuate nucleus of the hypothalamus, the parafascicular nucleus, the interpeduncular nucleus (central subnucleus), the nucleus prepositus, the medial vestib-

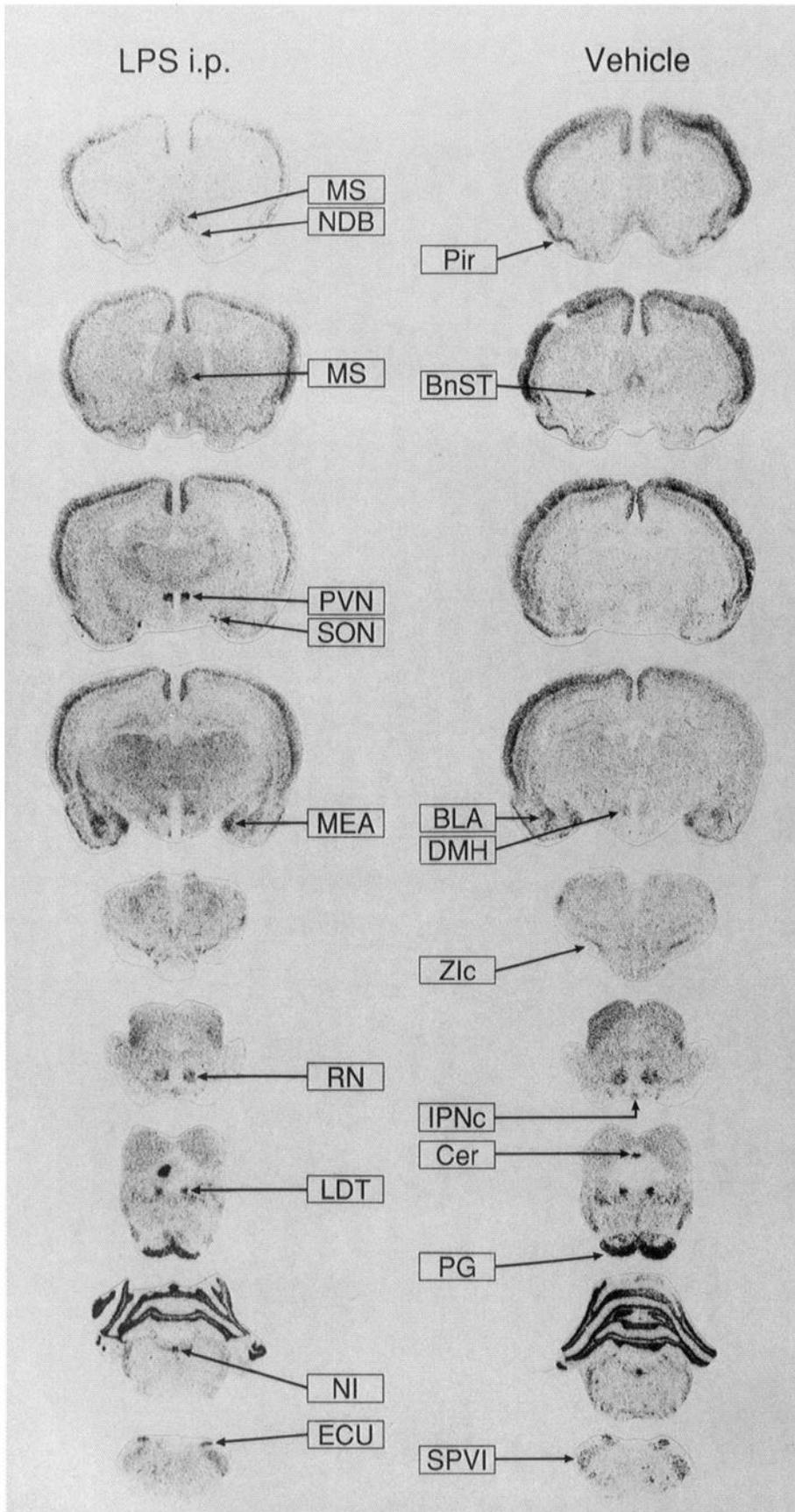


Figure 1. Representative example of the distribution of the mRNA encoding the receptor for corticotropin-releasing factor (CRF-R) in the rat brain after the intraperitoneal administration of endotoxin lipopolysaccharide (LPS) or vehicle. Animals were deeply anesthetized and rapidly perfused with 4% paraformaldehyde 6 hr after treatment with LPS (250 μ g/100 gm body weight) or the vehicle solution. These rostrocaudal coronal sections (30 μ m) of both LPS- and vehicle-treated rat brains exhibit positive signal on x-ray film (Kodak XAR 5) for the CRF-R transcript in the basolateral nucleus of the amygdala (BLA), the bed nucleus of the stria terminalis (BnST), the cerebellum (Cer), the dorsomedial nucleus of the hypothalamus (DMH), the external cuneate nucleus (ECU), the interpeduncular nucleus, central subnucleus (IPNc), the laterodorsal tegmental nucleus (LDT), the medial nucleus of the amygdala (MEA), the medial septal nucleus (MS), the nucleus incertus (NI), the nucleus of the diagonal band (NDB), the piriform cortex (Pir), the pontine gray (PG), the red nucleus (RN), the spinal nucleus of the trigeminal nerve, interpolar part (SPVI), the caudal division of the zona incerta (Zlc), and in various layers of the cerebral cortex. Note the selective induction of the CRF-R mRNA in the hypothalamic paraventricular nucleus (PVN) and the supraoptic nucleus (SON) of LPS-injected rat (left column).

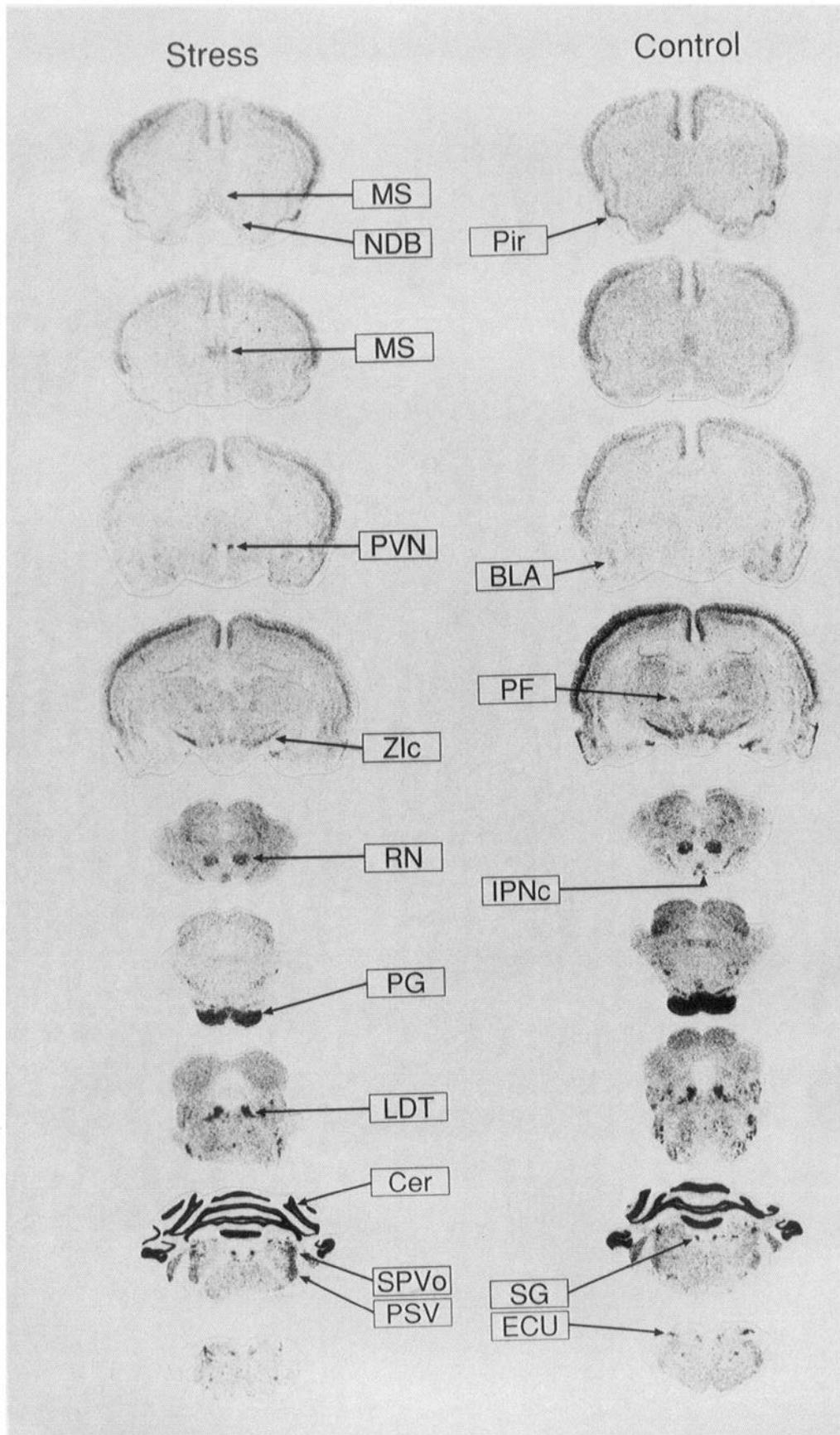


Figure 2. Representative example of the distribution of the gene encoding the corticotropin-releasing factor receptor (CRF-R) in the brain of stressed and control rats. Animals were deeply anesthetized and rapidly perfused with 4% paraformaldehyde before (*Control*) or 90 min after the end of the acute immobilization stress, which lasted 90 min (*Stress*, left column). These rostrocaudal coronal sections (30 μ m) of both stressed and

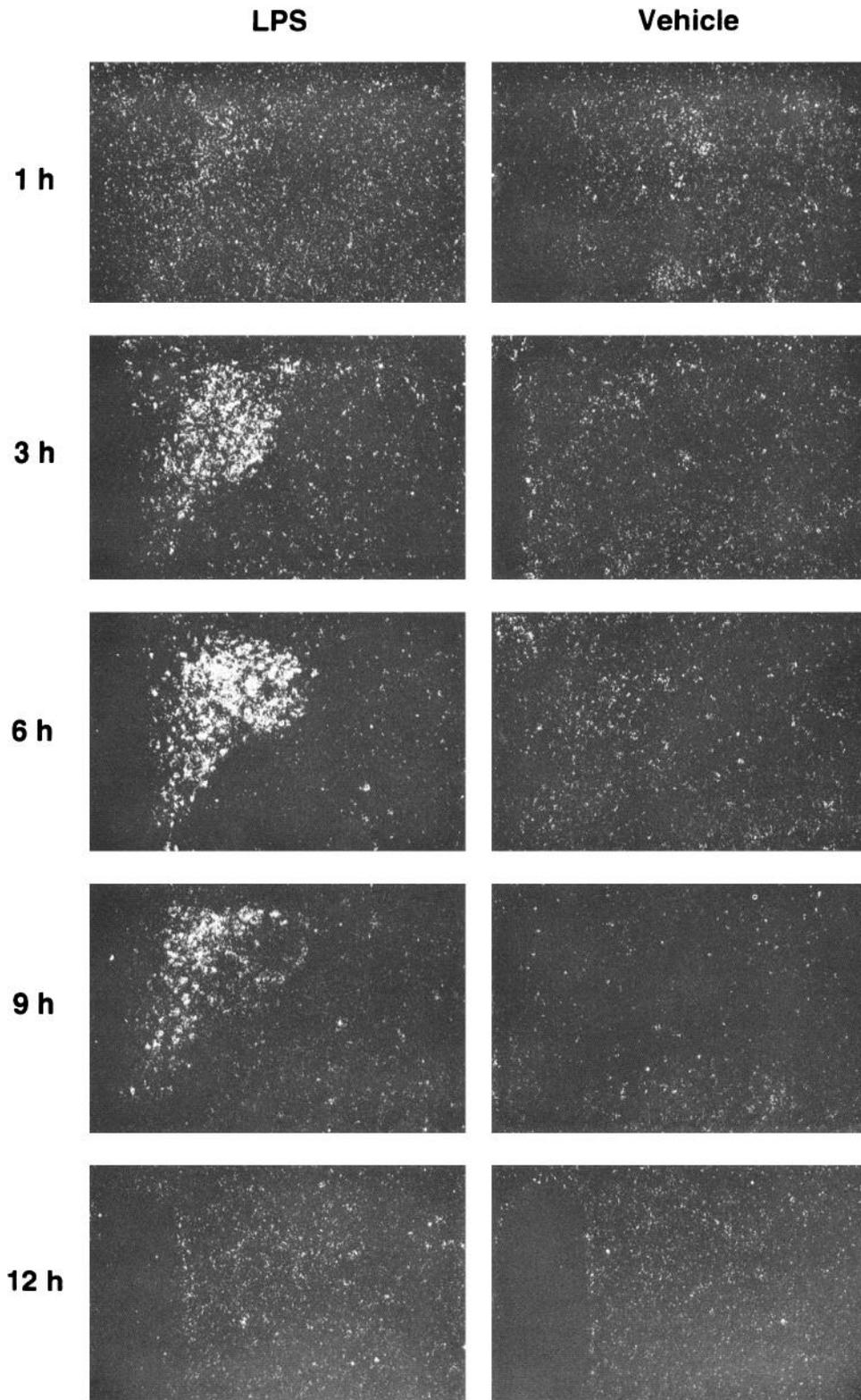


Figure 3. Influence of systemic (intrapertitoneal) administration of endotoxin lipopolysaccharide (*LPS*) or vehicle on the expression of CRF-R mRNA in the paraventricular nucleus (PVN) of the rat hypothalamus. These photos depict dark-field photomicrographs of dipped autoradiographs of hybridized 30 μ m sections with CRF-R riboprobe through identical areas of the right PVN. Animals were deeply anesthetized and rapidly perfused with 4% paraformaldehyde 1, 3, 6, 9, and 12 hr after the treatment with *LPS* (250 μ g/100 gm body weight, *left column*) or the vehicle solution (*right column*). Note the strong signal in both parvo- and magnocellular divisions of the PVN 6 hr following endotoxin injection, whereas the PVN of control rats displayed hardly detectable CRF-R transcript. Magnification 25 \times .

control rat brains display positive signal on x-ray film (Kodak XAR 5) for CRF-R mRNA in the basolateral nucleus of the amygdala (*BLA*), the bed nucleus of the stria terminalis (*BnST*), the cerebellum (*Cer*), the dorsomedial nucleus of the hypothalamus (*DMH*), the external cuneate nucleus (*ECU*), the interpeduncular nucleus, central subnucleus (*IPNc*), the laterodorsal tegmental nucleus (*LDT*), the medial nucleus of the amygdala (*MEA*), the medial septal nucleus (*MS*), the nucleus incertus (*NI*), the nucleus of the diagonal band (*NDB*), the parafascicular nucleus (*PF*), the piriform cortex (*Pir*), the pontine gray (*PG*), the principal sensory nucleus of the trigeminal nerve (*PSV*), the red nucleus (*RN*), the spinal nucleus of the trigeminal nerve, interpolar part (*SPVI*), the spinal nucleus of the trigeminal, oral part (*SPVo*), the supragenulate nucleus (*SG*), the caudal division of the zona incerta (*ZIc*), and various layers of the cerebral cortex. Note the selective induction of the mRNA encoding the CRF-R in the paraventricular nucleus of the hypothalamus (PVN) of stressed rat (*left column*).

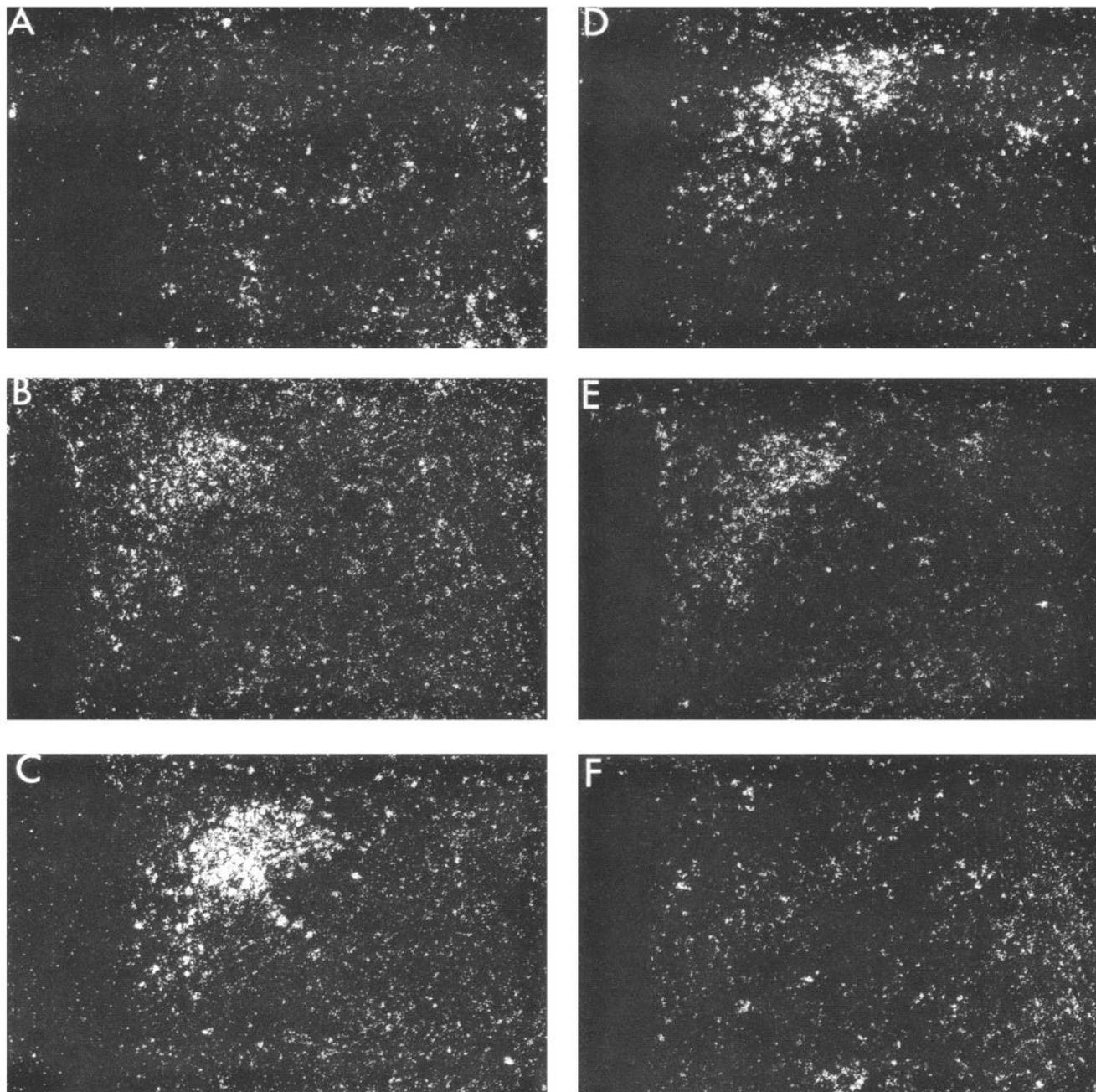


Figure 4. Acute immobilization stress induces expression of the mRNA encoding the CRF-R in the paraventricular nucleus (PVN) of the male rat hypothalamus. These photos depict dark-field photomicrographs of dipped autoradiographs of hybridized 30 μ m sections with CRF-R cRNA probe through identical areas of the right PVN. Animals were killed before (A), immediately (B), 1.5 (C), 3 (D), 6 (E), and 12 hr (F) after the 90 min stress session. Note that no visual CRF-R transcripts were observed before the stress session (A), whereas the parvocellular division of the PVN exhibits a robust and selective signal 1.5 (C) and 3 hr (D) following the acute stress session. Magnification 25 \times .

ular nucleus, and the spinal nucleus of the trigeminal nerve (interpolar part).

Site-specific induction of CRF-R transcription in the brain of challenged animals

Vehicle-treated (Fig. 1) and control (Fig. 2) rats displayed undetectable signals of CRF-R mRNA in the paraventricular nucleus (PVN), but CRF-R gene transcription was highly stimulated by LPS administration (Fig. 1) and immobilization stress (Fig. 2) in this hypothalamic structure. Indeed, the CRF-R

mRNA signal was positive in the dorsomedial parvocellular PVN 3 hr after LPS injection, strong and maximum in both parvo- and magnocellular divisions of the PVN at 6 hr postinjection, and declined 9 and 12 hr after treatment (Fig. 3). Interestingly, CRF-R was almost undetectable in the magno-PVN of animals treated with a lower dose of LPS intraperitoneal (25 μ g/100 gm of body weight) at 6 hr postinjection (results not shown). In these rats, the gene encoding CRF-R was still highly expressed in the parvocellular part of the PVN (results not shown). Similarly, the signal for CRF-R mRNA was detected in the par-

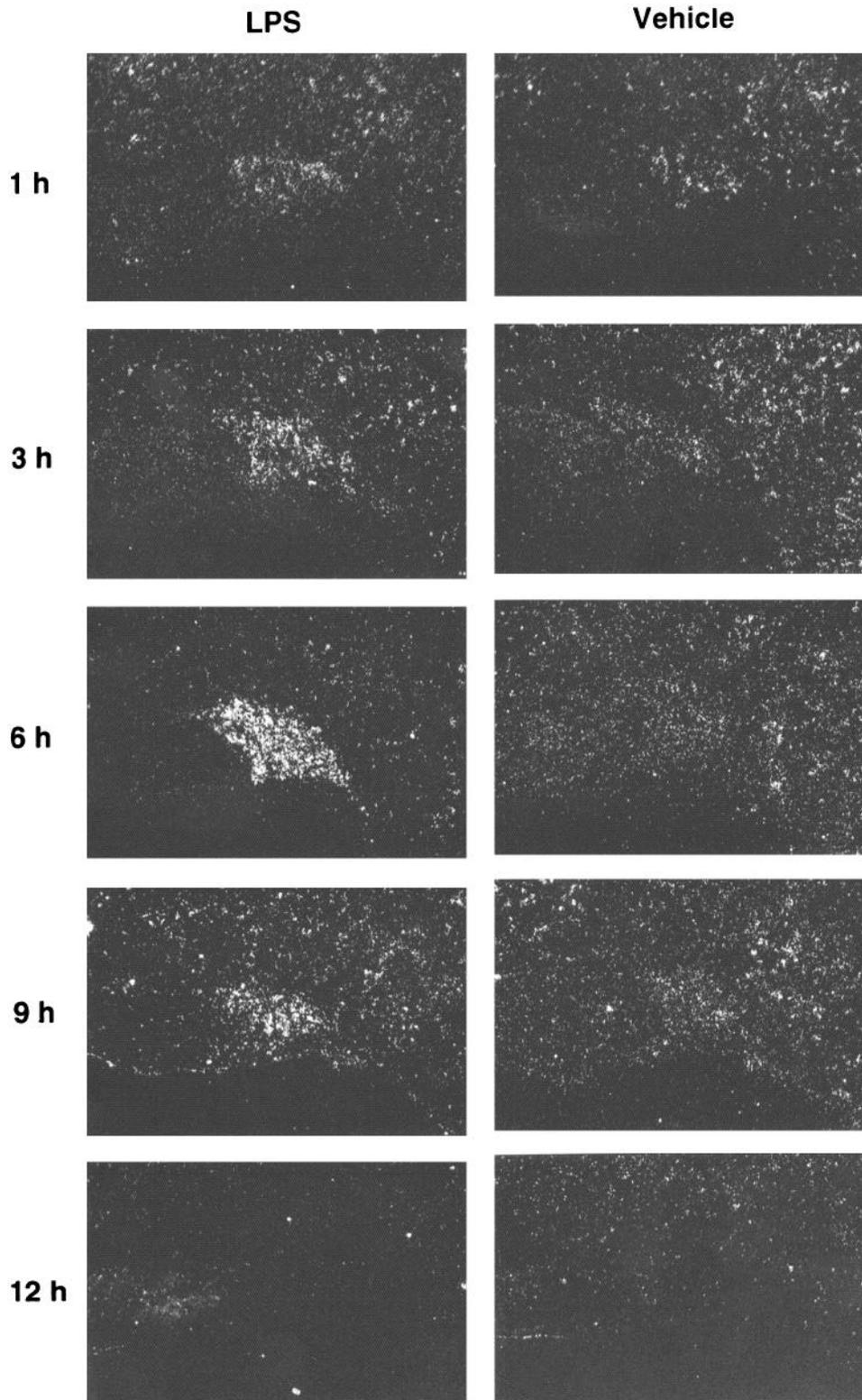


Figure 5. Effect of systemic (intra-peritoneal) administration of endotoxin lipopolysaccharide (LPS) or vehicle on the expression of CRF-R mRNA in the supraoptic nucleus (SON) of the rat hypothalamus. These photos depict dark-field photomicrographs of dipped autoradiographs of hybridized 30 μm sections with CRF-R riboprobe through identical areas of the right SON. Animals were deeply anesthetized and rapidly perfused with 4% paraformaldehyde 1, 3, 6, 9, and 12 hr after the treatment with LPS (250 $\mu\text{g}/100$ gm body weight, *left column*) or the vehicle solution (*right column*). Note the strong signal at 6 hr post-LPS injection (*left column*) compared to the small detectable expression of the CRF-R transcript in the SON of several vehicle-treated rats. In fact, the basal levels of CRF-R mRNA in the SON could be classified as low to barely detectable in some animals. Magnification 25 \times .

vo-PVN immediately after the 90 min stress session (Fig. 4B), highly expressed 90 min (Fig. 4C) and 3 hr (Fig. 4D) after the immobilization session, decreased at 6 hr (Fig. 4E), and vanished 12 hr after the stress session (Fig. 4F). Whereas CRF-R mRNA expression was located mainly in the parvocellular division of the PVN in immobilized rats (Fig. 4C), intraperitoneal

injection of a high dose of LPS induced transcription of this receptor in both parvocellular and magnocellular divisions of this hypothalamic nucleus (Fig. 3, 6 hr).

A lower but significant increase in the signal for CRF-R transcript was also observed in the SON after treatment with LPS (Fig. 5). Although a small endogenous expression of CRF-R was

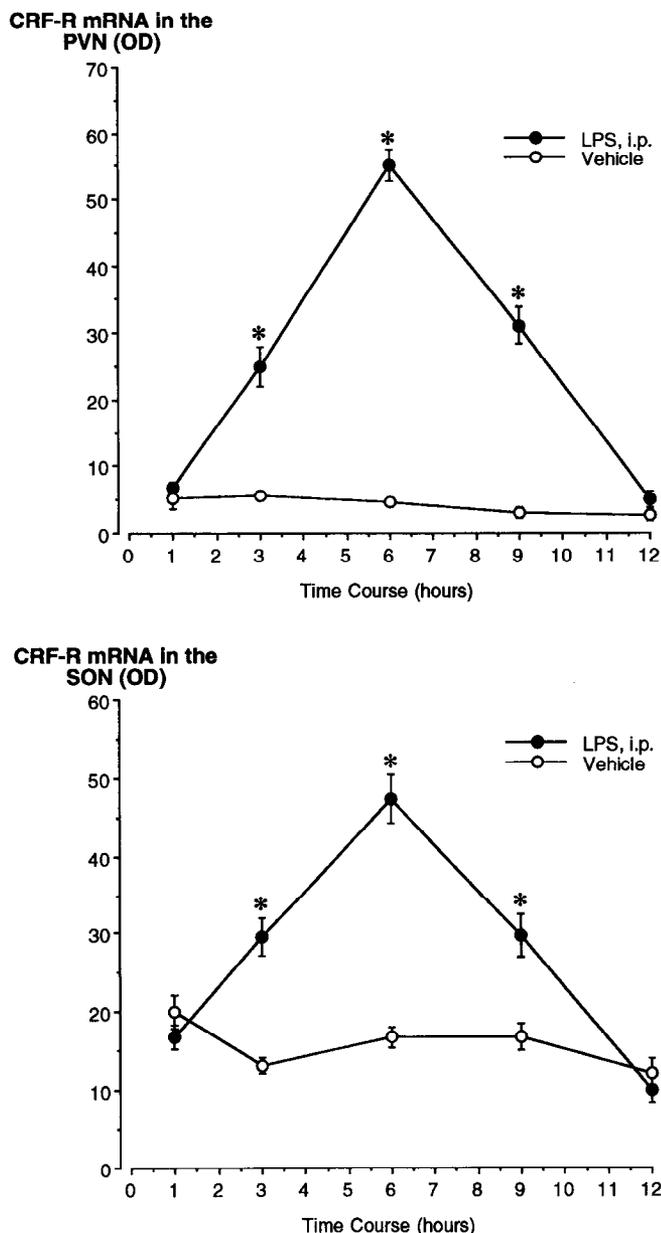


Figure 6. Influence of systemic (intraperitoneal) lipopolysaccharide (LPS; 250 $\mu\text{g}/100$ gm body weight) or vehicle administration on the average optical density (OD) for CRF-R mRNA signal in the paraventricular nucleus (PVN, top) and the supraoptic nucleus (SON, bottom) of the rat hypothalamus. The OD was quantified in both sides of the medial PVN and the SON using an Olympus Optical System (BX-50, βMax) coupled to a Macintosh computer (PowerPC 7100/66), and IMAGE software (version 1.55 non-FPU, W. Rasband, NIH). Results represent means \pm SEM of 6–8 rats. Statistical analysis was performed using a two-way analysis of variance (2×5) followed by a Dunn–Sidak post hoc test (STAVIEW 4.01). *, $P < 0.05$ from vehicle-treated rats. For more information on image analysis, see Materials and Methods.

detected in the SON of several vehicle-treated rats, a significant ($P < 0.05$) increase in the levels of CRF-R mRNA was obtained 3, 6, and 9 hr following the systemic treatment with the bacterial endotoxin. As for the PVN, a rapid downregulation of the gene encoding the CRF-R was also observed in this hypothalamic nucleus 12 hr after LPS injection. However, the transcription of the gene encoding the CRF-R was not notably increased in the

SON of rats treated with a lower dose of LPS (25 $\mu\text{g}/100$ gm of body weight, results not shown). As well, the immobilization session did not increase the relative levels of CRF-R transcript in the SON, although a small positive signal was detected in several control and stressed rats.

Systemic endotoxin and immobilization stress did not appear to modulate significantly the expression of CRF-R mRNA in other regions, which provides evidence that stress induced the transcription of the gene encoding the CRF-R in very selective hypothalamic nuclei directly involved in the regulation of neuroendocrine functions.

Time course of induction

Figures 6 and 7 depict the average optical density (OD) of the hybridization signal for CRF-R transcripts after immune challenge and stress, respectively. The OD was quantified on dark-field photomicrographs of dipped autoradiographs of hybridized 30 μm sections. The average OD for the CRF-R hybridized signal in both the PVN (Fig. 6, top) and the SON (Fig. 6, bottom) peaked 6 hr after the single intraperitoneal LPS administration and returned to the level of vehicle-treated rats at 12 hr postinjection. On the other hand, the influence of immobilization stress on the activation of CRF-R gene expression was significant in the PVN as soon as at the end of the 90 min session, but the highest average OD occurred 90 min after the stress period (3 hr after the beginning of the immobilization). Twelve hours after the stress session, the signal for CRF-R transcripts in the PVN returned to the control level. In contrast to the severe immune challenge, the OD for CRF-R mRNA in the SON was not significantly altered by the immobilization stress (Fig. 7).

Intensity of expression

Figure 8 shows bright-field photomicrographs of representative examples of the number of silver grains in LPS-induced CRF-R transcription in neurons of both the PVN and the SON. Although no positive cells were observed in the PVN of vehicle-treated rats, several neurons displayed a robust level of silver grains of this hypothalamic nucleus 6 hr after the immune challenge (Fig. 8, top). An increase in the number of positive neurons as well as in the number of silver grains per cell was also observed in the SON of LPS-treated rats (Fig. 8, bottom). Several positive neurons containing a high level of silver grains were also detected in the parvo-PVN of immobilized animals (Fig. 9).

Combination of immunocytochemistry and in situ hybridization

To determine whether CRF-ir neurons express the gene encoding CRF-R during specific challenges, immunocytochemistry was performed before the *in situ* hybridization histochemistry on the same brain sections. Figure 10 shows examples of CRF-ir neurons expressing the CRF-R mRNA (filled arrowheads) in the parvocellular PVN of LPS-treated (middle) and immobilized (bottom) rats, respectively. However, not all the CRF-ir neurons of the parvo-PVN expressed the gene encoding the CRF-R protein (open arrowheads), and CRF-R-expressing perikarya not colocalized in CRF-ir neurons were also noticed in the PVN of LPS and stressed rats. In vehicle-treated (Fig. 10, top) and control (not shown) rats, none of the CRF-ir neurons (open arrowheads) exhibited positive signal for CRF-R transcript.

Discussion

The present study provides solid evidence that the gene encoding the CRF-R can be finely regulated in selective hypothalamic

neurons, which could be directly related to the control of neuroendocrine functions such as the activity of the HPA axis. Although the CRF-R mRNA is widely distributed in the brain (Potter et al., 1994; present study), neither the immune challenge nor the immobilization stress seem to modulate the transcription of this receptor in most of these spontaneously expressing regions. In contrast, whereas the PVN displayed almost undetectable levels of CRF-R mRNA in control animals, a robust signal for this transcript was detected in this hypothalamic structure following both challenges. Interestingly, immobilization stress induced a specific expression of the gene encoding the CRF-R in the parvo-PVN, whereas systemic endotoxin produced a strong activation of CRF-R transcription in both the parvo- and the magnocellular divisions of the hypothalamic PVN. Similarly, systemic LPS caused a significant increase in the levels of CRF-R mRNA in the SON, suggesting that CRF can also modulate the activity of magnocellular neurons located in both the PVN and the SON during a severe immune activation. Animals treated with a lower dose of LPS (25 μ g/100 gm of body weight) still displayed strong expression of CRF-R in the parvo-PVN but not in the magnocellular division of this hypothalamic nucleus and the SON (results not shown). In the parvocellular division, CRF perikarya can express the gene encoding the CRF-R, but whether these neurons project to the infundibular system is not known. It nevertheless seems likely that CRF plays a direct role in controlling the activity of neuroendocrine CRF motoneurons during immune challenge as well as during other types of stressful conditions for the organism.

We recently reported that intracerebroventricular administration of CRF to conscious rats can activate the expression of CRF mRNA in the parvocellular neurons of the PVN (Parkes et al., 1993), the principal seat of the neurosecretory neurons responsible for delivering CRF to the hypophyseal portal vasculature (Sawchenko and Swanson, 1990; Swanson, 1991) and driving ACTH secretion (Rivier and Plotsky, 1986). This observation, together with the fact that CRF cell bodies of the parvo-PVN can express the CRF-R, indicates that CRF may be a potential modulator of its own biosynthesis selectively in the PVN of stressed rats. Moreover, central administration of CRF induces a rapid expression of various immediate *early* genes (*c-fos* and *NGFI-B*) in the CRF-ir neurons of the parvo-PVN (Parkes et al., 1993), a phenomenon also observed with systemic endotoxin injection and immobilization stress (Imaki et al., 1992a; Laflamme and Rivest, 1994). The temporal changes in both *c-fos* and *NGFI-B* gene expression are consistent with the hypothesis that the transcription factors encoded by these genes may be acting as early intracellular signals for direct or indirect activation of target genes such as CRF or CRF-R. In fact, intraperitoneal LPS treatment can stimulate both *c-fos* and *NGFI-B* genes in the parvocellular PVN as early as 1 hr after the administration (Laflamme and Rivest, 1994). The SON and the magnocellular PVN also exhibit a strong signal for these two transcripts 3 hr after endotoxin injection (Laflamme and Rivest, 1994). The signal for *c-fos* and *NGFI-B* mRNA in these brain nuclei was maximum 3 hr postinjection, declined at 6 hr, and vanished between 9 and 12 hr after LPS treatment. The induction of CRF-R was maximum at 6 hr postinjection, and relative levels of CRF mRNA in the parvocellular PVN significantly increased 6 and 9 hr after LPS treatment (Laflamme and Rivest, 1994). Whether AP-1 or NBRE can directly regulate the gene encoding the CRF-R has not been yet reported, but a relationship between IEGs, CRF-R,

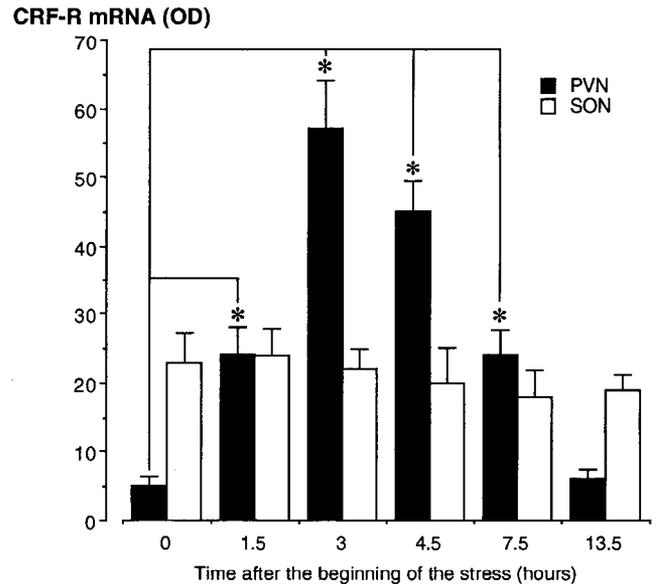
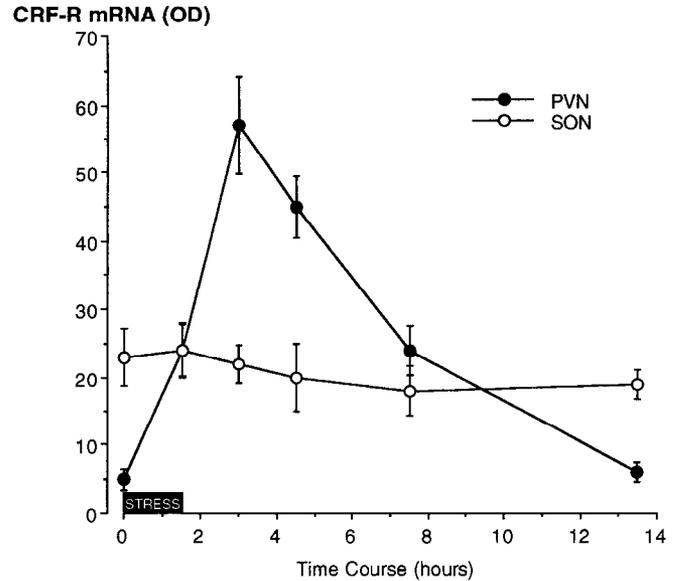


Figure 7. Effect of an acute 90 min session of immobilization stress on the average optical density (OD) for CRF-R mRNA signal in the paraventricular nucleus (PVN) and the supraoptic nucleus (SON) of the rat hypothalamus. The OD was quantified in both sides of the medial PVN and the SON using an Olympus Optical System (BX-50, β Max) coupled to a Macintosh computer (PowerPC 7100/66), and IMAGE software (version 1.55 non-FPU, W. Rasband, NIH). Results represent means \pm SEM, and statistical analysis was performed using a one-way analysis of variance (1×5) followed by a Dunn-Sidak post hoc test (STAVIEW 4.01). *Bottom*, The areas under the curve with the statistical analyses. *, $P < 0.05$ from control rats (time 0, *bottom*). For more information on image analysis, see Materials and Methods.

and CRF biosynthesis during immune activation and other types of stressful circumstances remains possible in the parvo-PVN.

The theory of autoregulation by endogenous neuroendocrine factors has been widely studied regarding negative suppression of protein release and biosynthesis (Dallman and Jones, 1973; Piva et al., 1979; Dallman et al., 1992). However, few neuroendocrine systems, apart from the positive effects of estradiol on

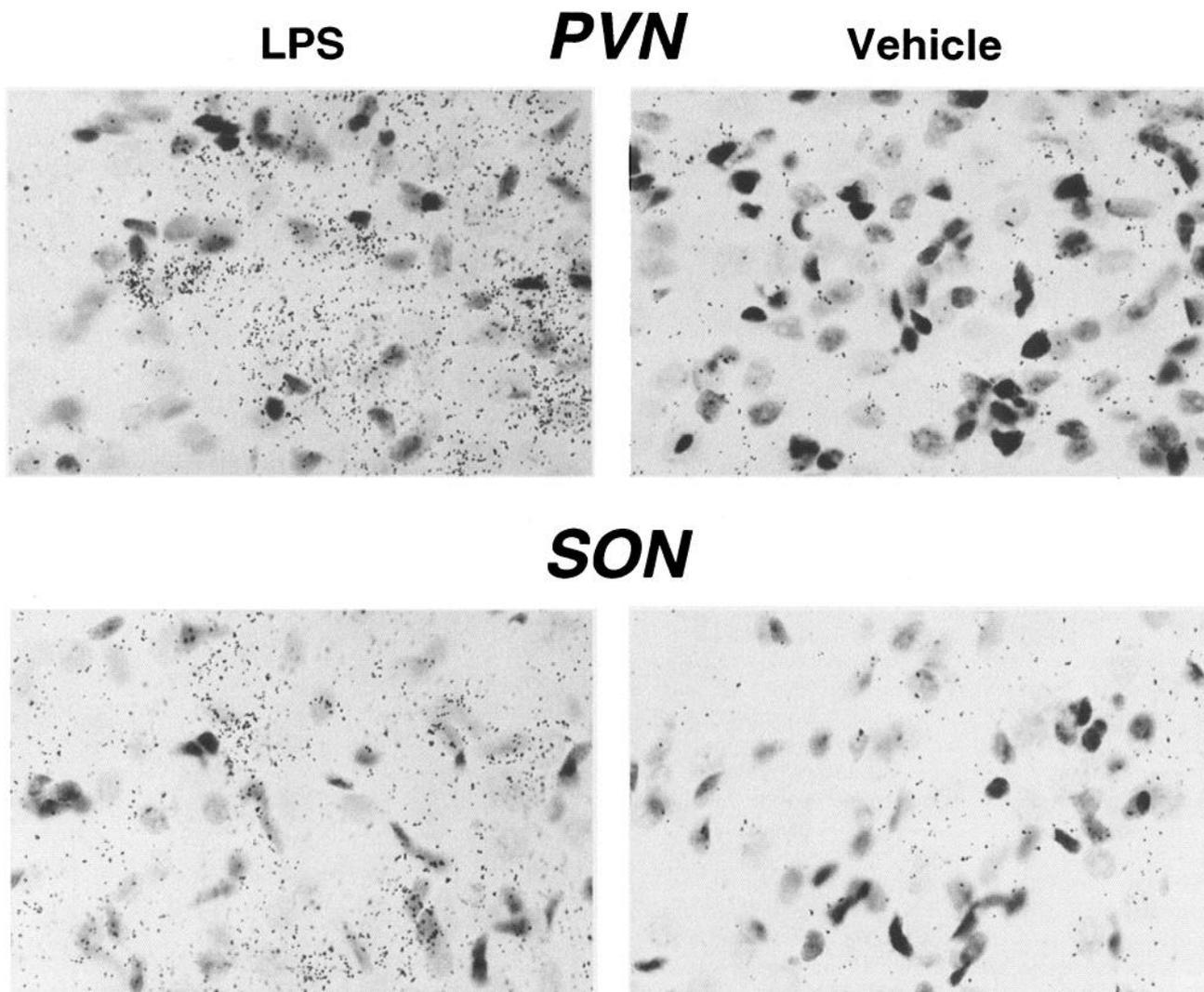


Figure 8. High-power bright-field photomicrographs of dipped autoradiographs of hybridized 30 μm sections with CRF-R riboprobe through similar areas of the paraventricular nucleus (PVN) and the supraoptic nucleus (SON). Animals were deeply anesthetized and rapidly perfused with 4% paraformaldehyde 6 hr after the intraperitoneal administration of lipopolysaccharide (LPS; 250 $\mu\text{g}/100$ gm body weight, *left panels*) or the vehicle solution (*right panels*). Note the number of silver grains delineating several neurons expressing the gene encoding the CRF-R in both the PVN and the SON of LPS-treated animals. Magnification 250 \times .

the activity of luteinizing hormone–releasing hormone (LHRH) neurons during the preovulatory LH surge, have exhibited evidence for positive feedback in either a direct or indirect manner. Negative autoregulatory mechanisms were demonstrated for oxytocin, somatostatin, and LHRH (Lumpkin et al., 1983, 1985; Richardson and Twente, 1986; Valenca et al., 1987). This decapeptide, which controls the secretion and biosynthesis of LH from the adenohypophysis, has been suggested as a candidate for short-term negative autoregulation (Sarkar, 1987; Valenca et al., 1987). In contrast, CRF per se could be involved in the stimulation of transduction pathways within PVN CRF cells to trigger the transcription of the peptide in challenged rats.

The existence of ultrashort-loop positive feedback control of CRF secretion during stress states was implied by Ono and co-workers (Ono et al., 1985). Detailed histochemical mapping of CRF neuronal fiber projections throughout the mammalian brain have shown that these fibers do indeed terminate and connect with parvocellular CRF perikarya in PVN (Silverman et al., 1989). In combining the retrograde transport of wheat-germ-con-

jugated gold particles and immunocytochemistry for CRF, Beaulieu and Drolet (1994) recently described the origin of CRF innervation to the PVN. CRF-ir neurons located in the perifornical hypothalamic nucleus, the dorsolateral hypothalamic area, bed nucleus of the lamina terminalis (dorsolateral and dorsomedial), medial preoptic region, and laterodorsal tegmental nucleus/Barrington's nucleus project to this hypothalamic nucleus. Besides these CRFergic projecting sites, the PVN itself could represent the origin of CRF fibers that terminate on neuroendocrine CRF perikarya. Immunohistochemical studies indicate that the PVN contains about 2000 CRF cells, which are found in all eight major parts of the nucleus; as mentioned, however, a large proportion is concentrated in the dorsomedial parvocellular part (Sawchenko et al., 1984, 1993; Swanson et al., 1986, 1983; Sawchenko and Swanson, 1990). Furthermore, CRF is expressed in a small group of neurons that project to the brainstem and the spinal cord, as well as in a subset of oxytocin cells (Sawchenko et al., 1984). In the dorsomedial parvocellular PVN, a large percentage of CRF motoneurons initially follow a laterally directed

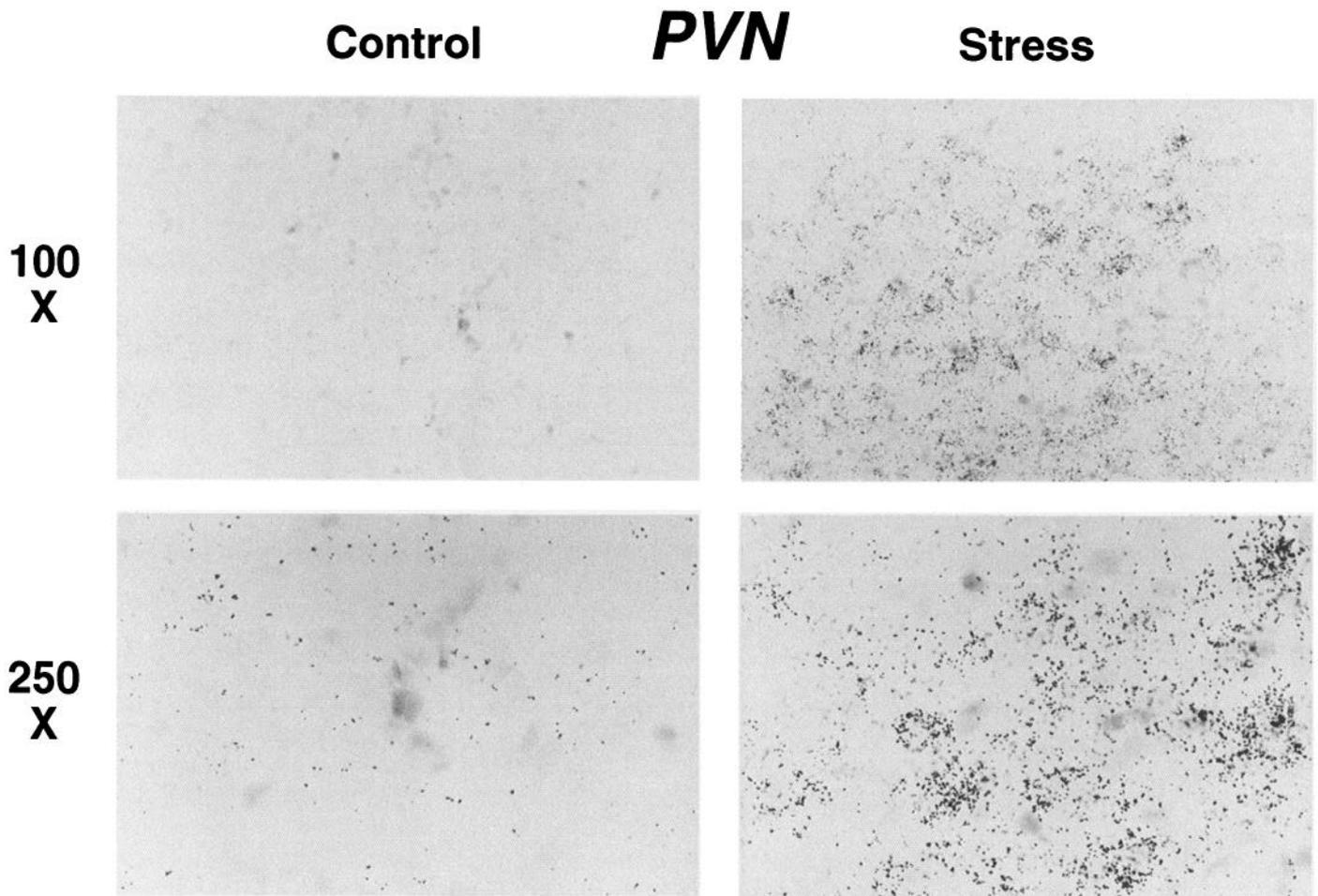


Figure 9. High-power bright-field photomicrographs of dipped autoradiographs of hybridized 30 μm sections with CRF-R riboprobe through similar areas of the hypothalamic paraventricular nucleus (PVN) in control and stressed rats. Rats were killed before (*Control*) or 90 min after the immobilization stress session (*Stress*), which lasted 90 min. Note the number of silver grains delineating several perikarya expressing the mRNA encoding the receptor for CRF in the parvo-PVN of stressed rats. Magnification 100 \times (*upper panels*), 250 \times (*lower panels*).

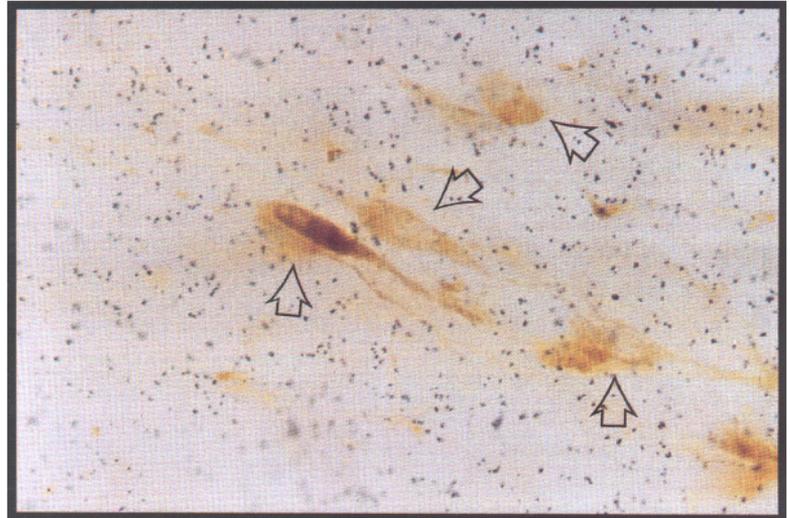
course through and around the fornix before arching medially to end in the median eminence (Makara et al., 1981; Swanson et al., 1983). However, an elegant tract-tracing study indicated that the axons of at least some CRF neurons in the dorsomedial PVN give rise to terminal boutons around the lateral margin of the PVN, and that these cells typically display 1–3 dendrites, which branch only once or twice, and bear a sparse complement of spines (Swanson et al., 1987). CRF–CRF synapses could therefore, in theory, represent either local connections within the PVN itself or CRF neuronal projections originating from other hypothalamic or extrahypothalamic areas.

The involvement of CNS CRF in regulating the activity of neuroendocrine CRF motoneurons during immune challenges and other types of neurogenic stresses could subservise as a potential physiological relevance of the present results. Indeed, whereas CRF-immunoreactive and -expressing neurons can be detected in several structures of the CNS, only the CRF neurons located in the PVN seem to express the CRF-R gene after stressful conditions. The selectivity in the induction of the CRF-R mRNA in the CRF neurons of the PVN might indicate that (1) neuroendocrine CRF is the only CRF that can be regulated by its own receptor during stress; (2) induction of the CRF-R in the parvo-PVN is a determinant mechanism involved in the restoration of the peptide depletion after the stress period; and (3)

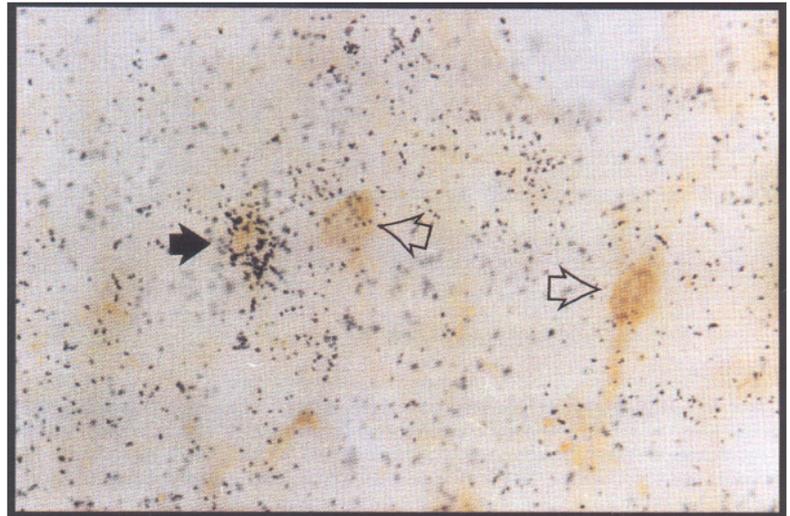
the basal expression of the gene encoding the CRF-R in most of the brain structures is not regulated in a stress-dependent manner. It is noteworthy that CRF is among the most widely distributed neuropeptides in the CNS, but little is known about the role of CRF located outside the PVN. Even in this hypothalamic nucleus, the exact function of CRF neurons that do not send their axons to the infundibular system still remains poorly described. The fact that stimulation of CRF-R gene transcription during immobilization and immune challenge is limited to the PVN and the SON (immune stress) might suggest that CRF selectively targets these regions during stress and modulates a confined group of neurons in the brains of challenged animals. We also observed by means of CRF intronic probe technology that both immobilization and systemic LPS administration generated production of CRF heteronuclear RNA only in the PVN and not in any other sites of the brain (S. Rivest, N. Laflamme, and B. Bonaz, unpublished observations).

Interestingly, immobilization stress induced a very selective expression of CRF-R in the parvocellular division of the PVN, whereas severe acute immune challenge caused a robust transcription of this receptor in both the parvocellular and magnocellular divisions of the PVN as well as in the SON. CRF therefore appears to selectively target parvocellular neurons during immobilization stress, although parvo-PVN, magno-PVN, and

Veh



LPS



Stress

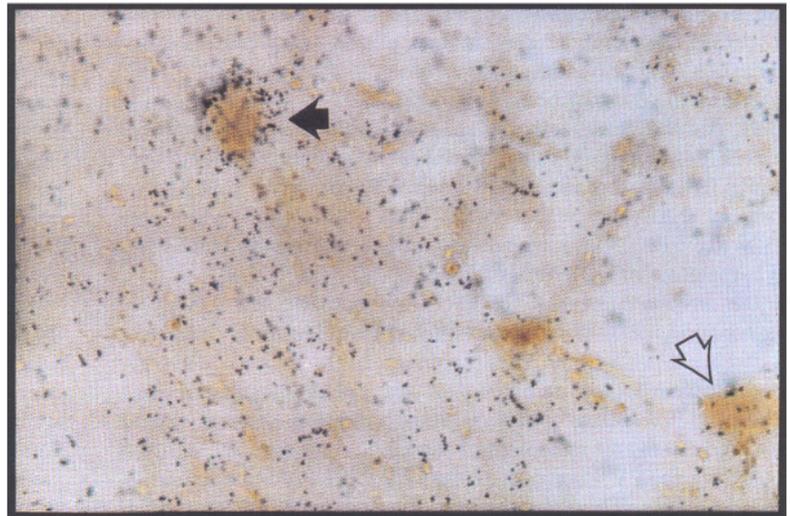


Figure 10. Expression of CRF-R mRNA in CRF perikarya located in the hypothalamic paraventricular nucleus (PVN) of LPS-treated and stressed rats. Animals were killed 6 hr after an intraperitoneal administration of vehicle (*Veh*) or endotoxin (*LPS*; 250 μ g/100 gm body weight), and 90 min after a 90 min immobilization stress session. Immunocytochemistry (CRF protein, *brown neurons*) was performed on the same brain sections (30 μ m) before *in situ* hybridization histochemistry (CRF-R mRNA, *silver grains*). For more details on the combination of both immunocytochemistry and *in situ* hybridization techniques, see Materials and Methods. *Filled arrowheads*, CRF neurons expressing the mRNA encoding the CRF-R (*middle, bottom*); *open arrowheads*, CRF perikarya alone.

SON can receive specific CRFergic stimulation during systemic LPS treatment. This effect seems dependent on the intensity of immune challenge, because at lower doses (25 $\mu\text{g}/100$ gm of body weight) systemic LPS activated the transcription of CRF-R selectively in the parvo-PVN. It is possible that the action of CRF during neurogenic stress and a middle immune activation is limited primarily to the parvo-PVN and its influence tightly connected to the regulation of neuroendocrine CRF motoneurons. In addition to this action, CRF could be able to modulate the activity of vasopressinergic (AVP) or oxytocinergic cells located in the magnocellular division of the PVN and in the SON during a severe immune challenge. The fact that systemic LPS treatment can cause a strong induction of various immediate *early* genes in the PVN (parvo- and magnocellular divisions) and the SON (Laflamme and Rivest, 1994) suggests that immune challenge can stimulate not only the parvo-PVN, but also the activity of neurons located in the magno-PVN and the SON. Although the large proportion of magnocellular neurons send their axons to the posterior pituitary, it is possible that these neurons, in particular AVP neurons, contribute to the control of ACTH from the adenohypophysis. AVP plays a supportive role for CRF to stimulate the release of ACTH (Rivier and Plotsky, 1986), and immune challenge is known to activate the release of AVP into the hypophyseal-portal circulation (Harbuz et al., 1992) and increase the levels of AVP heteronuclear RNA in the magnocellular PVN (Rivest and Laflamme, unpublished observations). Although the parvocellular division of the PVN contributes to synthesizing and releasing AVP into the infundibular system, most AVP released in portal plasma is probably derived from magnocellular neurosecretory neurons and thus participates in the control of the HPA axis during various stressors (Buma and Nieuwenhurs, 1987; Antoni et al., 1990; Sawchenko et al., 1993).

The development of models of immune activation is actively sought because these models would produce a better understanding of the mechanisms through which infectious diseases activate the HPA axis. Endotoxins are widely used to mimic some of the events that occur during sepsis (Kushner, 1982; Dinarello, 1984; Ertel et al., 1992). Although not a true model of infection, this model represents an accepted means of increasing the release of endogenous cytokines from the peripheral immune system (Staruch and Wood, 1985; Cybulsky et al., 1987; Shalaby et al., 1989; Bristow et al., 1991). Increased production of cytokines, proteins released by activated macrophages and lymphocytes upon presentation of an antigen (Dinarello, 1989; Rabin et al., 1990), represent an essential feature of the early events of immune activation, called the *acute-phase response*. A primary function of these proteins is directed toward expansion of the immunologic mass and activity by stimulating the production of growth factors. However, cytokines (such as interleukin-1) also reach the general circulation and thus can act on distant endocrine organs. The demonstration that these proteins can stimulate the neuropeptide CRF (Berkenbosch et al., 1987; Sapolsky et al., 1987; Tsagarakis et al., 1989; Barbanel et al., 1990; Suda et al., 1990; Navarra et al., 1991; Rivest et al., 1992) and the finding that subsequent activation of adrenal steroid release can in turn convey feedback signals to immune cells (Solomon, 1969) have paved the way for a better understanding of the bilateral communication pathways linking the immune and the HPA axis. Although long considered virtually independent of each other, the immune and neuroendocrine axes have thus special relevance for restoring physiological balance when chal-

lenged by antigens. In this concept, the PVN and the SON, which contain many of the peptides that modulate neuroendocrine functions, seem to play a determinant role during the acute phase of an immune challenge. As mentioned, systemic administration of bacterial endotoxin induces profound activation of PVN and SON neurons (Laflamme and Rivest, 1994). The present study also presents evidence that the CRF-R could be a potential modulator in both the PVN and the SON during severe immune challenge because transcription of the gene encoding this protein is turned on selectively in these hypothalamic nuclei.

In conclusion, the gene encoding the CRF-R is widely distributed in the rat brain, but immune challenge and immobilization stress activated the transcription of that gene in very selective structures of the hypothalamus. Indeed, immobilization stress and low doses of LPS (25 $\mu\text{g}/100$ gm of body weight) induced a specific expression of CRF-R mRNA in the parvo-PVN, whereas high doses of endotoxin injected intraperitoneally (250 $\mu\text{g}/100$ gm of body weight) produced a strong activation of CRF-R transcription in both the parvo- and the magnocellular divisions of this hypothalamic structure. This treatment also caused a significant increase in the levels of CRF-R mRNA in the SON, suggesting that CRF can modulate the activity of magnocellular neurons located in both the PVN and the SON of severely immune-challenged animals. In the parvocellular division of the PVN, CRF-ir neurons exhibit positive signal for the CRF-R transcript. Taken together, these results provide evidence that the gene encoding the CRF-R is finely regulated in selective regions of the brain and may play a crucial role in the control of neuroendocrine CRF motoneurons present in the dorsomedial parvo-PVN during immune challenge and other types of neurogenic stresses. Induction of CRF-R in the magnocellular PVN and the SON could also be a determinant step involved in the cascade of events modulating the activity of other neurons, such as vasopressinergic and oxytocinergic neurons, during a severe immune activation.

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