

# Region-Specific Regulation of Glutamic Acid Decarboxylase (GAD) mRNA Expression in Central Stress Circuits

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Neurocircuit inhibition of hypothalamic paraventricular nucleus (PVN) neurons controlling hypothalamo–pituitary–adrenocortical (HPA) activity prominently involves GABAergic cell groups of the hypothalamus and basal forebrain. In the present study, stress responsiveness of GABAergic regions implicated in HPA inhibition was assessed by *in situ* hybridization, using probes recognizing the GABA-synthesizing enzyme glutamic acid decarboxylase (GAD65 and GAD67 isoforms). Acute restraint preferentially increased GAD67 mRNA expression in several stress-relevant brain regions, including the arcuate nucleus, dorsomedial hypothalamic nucleus, medial preoptic area, bed nucleus of the stria terminalis (BST) and hippocampus (CA1 and dentate gyrus). In all cases GAD67 mRNA peaked at 1 hr after stress and returned to unstimulated levels by 2 hr. GAD65 mRNA upregulation was only observed in the BST and dentate gyrus. In contrast, chronic intermittent stress increased GAD65 mRNA in the anterior hypothalamic area, dorsomedial nucleus, medial preoptic area, suprachiasmatic nucleus, ante-

rior BST, perifornical nucleus, and periparaventricular nucleus region. GAD67 mRNA increases were only observed in the medial preoptic area, anterior BST, and hippocampus. Acute and chronic stress did not affect GAD65 or GAD67 mRNA expression in the caudate nucleus, reticular thalamus, or parietal cortex. Overall, the results indicate preferential upregulation of GAD in central circuitry responsible for direct (hypothalamus, BST) or multisynaptic (hippocampus) control of HPA activity. The distinct patterns of GAD65 and GAD67 by acute versus chronic stress suggest stimulus duration-dependent control of GAD biosynthesis. Chronic stress-induced increases in GAD65 mRNA expression predict enhanced availability of GAD65 apoenzyme after prolonged stimulation, whereas acute stress-specific GAD67 upregulation is consistent with *de novo* synthesis of active enzyme by discrete stressful stimuli.

**Key words:** acute stress; chronic stress; hypothalamus; hippocampus; preoptic area; hypothalamo–pituitary–adrenocortical axis

Physiological responses to stress typically demand changes in energy use, cardiovascular tone, endocrine status, and arousal level, which are driven in large part by adrenocortical glucocorticoid hormones. Release of glucocorticoids is initiated by a discrete set of executive neurons localized in the hypothalamic paraventricular nucleus (PVN) (Antoni, 1986; Whitnall, 1993). These neurons synthesize and secrete a cocktail of neuropeptides that promote release of adrenocorticotrophic hormone (ACTH) and, subsequently, glucocorticoids.

Efficient inhibition of glucocorticoid secretion is required to limit the magnitude and duration of stress responses at the level of the PVN. This is accomplished by both neuronal inhibitory circuitry and blood-borne glucocorticoid negative feedback. Central circuitry regulating neuronal inhibition of the PVN prominently involves the hippocampus. This region is known to play a role in the inhibition of basal ACTH secretagog expression and in limiting the duration of stress-induced glucocorticoid secretion (Herman et al., 1989; Jacobson and Sapolsky, 1991). Inhibitory effects of hippocampal action seem to be driven by the ventral

subiculum, because effects of total hippocampectomy on PVN corticotropin-releasing hormone (CRH) mRNA expression and stress duration can be mimicked by lesions confined to this region (Herman et al., 1995).

Anatomical data do not support a direct connection between limbic neurons and the medial parvocellular PVN. However, ventral subiculum projects to a number of forebrain regions that in turn innervate this region, including the bed nucleus of the stria terminalis (BST), medial preoptic area, dorsomedial hypothalamic nucleus, and anterior hypothalamic area (Cullinan et al., 1993). Combined anterograde–retrograde tracing studies indicate that ventral subiculum efferents contact BST, preoptic area, dorsomedial hypothalamic, and anterior hypothalamic area neurons that are retrogradely labeled by PVN injections of Fluorogold (Cullinan et al., 1993). Notably, the vast majority of these PVN-projecting neurons contain the inhibitory neurotransmitter GABA (Cullinan et al., 1993). Because projection neurons of ventral subiculum are likely to use excitatory amino acid transmitters (glutamate and aspartate) (Walaas and Fonnum, 1980), the anatomical data suggest the possibility of a bisynaptic subiculum–PVN connection, essentially switching excitatory hippocampal signals into inhibition at the PVN.

This hypothesis suggests that PVN-projecting GABAergic populations are critical components of hypothalamo–pituitary–adrenocortical (HPA) inhibition of stress responses and predicts that these cell groups should be activated by stressful stimuli. In accordance with this notion, recent studies indicate that neurons in the BST and hypothalamus increase cFOS expression after

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acute stress exposure (Cullinan et al., 1995; Sawchenko et al., 1996), consistent with stress activation. Furthermore, double-label studies indicate that a large proportion of stress-activated neurons in these PVN-projecting regions express the GABAergic marker glutamic acid decarboxylase (GAD) mRNA (Cullinan et al., 1996). Involvement of GABA in HPA inhibition is supported by microinjection studies, which document reduced PVN neuronal activity and attenuated corticosterone (CORT) secretion after local application of GABAergic drugs (Boudaba et al., 1996; W. E. Cullinan, unpublished observations). To address activation of GABAergic neurocircuits by stress directly, the present study assesses stress regulation of GAD isoform (GAD65/GAD67) mRNA expression in brain regions responsible for monosynaptic or disinynaptic control of HPA activation.

## MATERIALS AND METHODS

**Animals.** Male Sprague Dawley rats initially weighing between 240 and 320 gm were included in both studies. All rats were housed three per cage on a 12 hr:12 hr light/dark cycle with food and water available *ad libitum*.

**Acute stress protocol.** Animals were divided into three groups. Unstressed animals ( $n = 7$ ) were not exposed to stress before death and thus represented the control group. The 60 min group ( $n = 7$ ) was exposed to 1 hr of restraint stress in plastic restraint tubes and killed immediately after the stress. The 120 min group ( $n = 7$ ) was exposed to the same 60 min restraint stressor, returned to their home cages, and subsequently killed 1 hr later.

**Chronic stress protocol.** Animals were divided into two groups. The chronic intermittent stress group ( $n = 6$ ) was subjected to a 15 d variable intermittent stress paradigm using the following stressors:

- (1) Restraint: rats were placed in plastic restraint cages for 2 hr.
- (2) Isolation: rats were housed in individual cages until the next stress period.
- (3) Cold restraint: rats were placed in plastic restraint cages in a cold room (4°C) for 2 hr.
- (4) Crowding: rats were placed six per cage until the next stress period.
- (5) Rotation/crowding: rats were placed six per cage on an orbit shaker and rotated for 2 hr.
- (6) Swim: rats were placed in an aquarium filled with 28–32°C water for 40 min.
- (7) Cold swim: rats were placed in an aquarium filled with 15–18°C water for 10 min.

Stressed rats were exposed to two stressors per day, varied randomly among the above list. Control rats ( $n = 6$ ) were individually handled for 1 min each time the experimental group was stressed.

All rats in both the acute and chronic studies were killed by rapid decapitation between the hours of 9 and 11 A.M. All brains were removed and frozen in isopentane cooled on dry ice at  $-40$  to  $-50^{\circ}\text{C}$ . Core blood samples were collected in heparinized tubes and centrifuged at  $1500 \times g$ , and plasma samples were frozen at  $-20^{\circ}\text{C}$ . Brains were stored at  $-80^{\circ}\text{C}$  until processing. All brains were sectioned in series at  $15 \mu\text{m}$  using a Bright-Hacker cryostat, mounted on Superfrost Plus slides, and stored at  $-20^{\circ}\text{C}$ .

**In situ hybridization.** Series of tissue sections were taken from the  $-20^{\circ}\text{C}$  freezer and fixed in 4% phosphate-buffered paraformaldehyde for 10 min. Slides were rinsed twice in 5 mM potassium PBS (KPBS) for 5 min, twice in 5 mM KPBS with 0.2% glycine for 5 min, and twice in KPBS for 5 min. Slides were then acetylated in 0.1 M triethanolamine, pH 8.0, with 0.25% acetic anhydride for 10 min. Slides were rinsed twice in  $2 \times \text{SSC}$  for 5 min and dehydrated through graded alcohols.

Antisense rat GAD65 and GAD67 probes were produced by *in vitro* transcription using [ $^{33}\text{P}$ ]UTP. Plasmids containing the GAD65 (courtesy of A. Tobin, University of California, Los Angeles) insert were linearized with *Stu*I and transcribed with T3 RNA polymerase. Plasmids containing the GAD67 insert (courtesy of A. Tobin, University of California, Los Angeles) were linearized with *Hinc*II and transcribed with T3 RNA polymerase. The transcription reaction consisted of  $10 \times$  transcription buffer that contained 125  $\mu\text{Ci}$  of [ $^{33}\text{P}$ ]UTP, 200  $\mu\text{M}$  ATP, CTP, and GTP, 10  $\mu\text{M}$  cold UTP, 100 mM dithiothreitol, 40 U/ $\mu\text{l}$  placental RNase inhibitor, and 20 U/ $\mu\text{l}$  T3 RNA polymerase. The mixture was incubated for 90 min at  $37^{\circ}\text{C}$ , and probe was separated from free nucleotides by ammonium acetate precipitation.

**Table 1. Plasma ACTH and corticosterone responses to acute stress**

Group	ACTH (pg/ml)	Corticosterone (ng/ml)
Unstressed	73.5 $\pm$ 25.1	4.3 $\pm$ 2.5
60 min	179.9 $\pm$ 29.5*	266.8 $\pm$ 61.1*
120 min	50.9 $\pm$ 9.1	9.7 $\pm$ 2.1

\*Significant differences.

Probes were diluted in hybridization buffer to yield  $\sim 1,000,000$  cpm/50  $\mu\text{l}$  of buffer. Diluted aliquots of 50  $\mu\text{l}$  were applied to each slide, with slides coverslipped and incubated overnight at  $55^{\circ}\text{C}$  in chambers containing filter paper moistened with 50% formamide. Coverslips were then removed in  $2 \times \text{SSC}$ , and slides were incubated in RNase A (100  $\mu\text{g}/\text{ml}$ ) for 30 min at  $37^{\circ}\text{C}$ . Slides were briefly rinsed in  $2 \times \text{SSC}$  and washed three times in  $0.2 \times \text{SSC}$  for 10 min, followed by a 1 hr bath in  $0.2 \times \text{SSC}$  at  $65^{\circ}\text{C}$ . Slides were dehydrated through graded alcohols, exposed to Kodak Biomax MR-2 film for 3–6 d, and subsequently dipped in Kodak (NTB2) emulsion. Dipped slides were stored at  $4^{\circ}\text{C}$  in light-tight boxes for 25 d, developed in Kodak D-19 developer, and coverslipped with DPX mountant.

**Image analysis.** Semiquantitative analyses of *in situ* hybridization films were performed with Macintosh-based NIH Image 1.59 software. Sections from control and experimental animals were matched for rostro-caudal level, and regions of interest were captured. Anatomical areas of interest were determined from the Paxinos and Watson (1986) atlas and sampled manually. The rostrocaudal levels for BST and hypothalamic regions selected for analysis were based on distinctions outlined in the Paxinos and Watson atlas as follows: anteromedial BST (encompassing atlas divisions BSTMA) and anterodorsal BST (encompassing atlas divisions BSTLD, BSTLJ, and BSTI), plates 19–20; posterior intermediate BST, posterolateral BST, and medial preoptic area, plates 21–22; suprachiasmatic nucleus, plates 23–24; anterior hypothalamic area, plates 24–25; arcuate nucleus, plates 25–29; and dorsomedial hypothalamus, plates 29–30 (Paxinos and Watson, 1986). Background signal was determined by sampling nonhybridized regions of each section (white matter). Background signal was subtracted from raw gray level measures, and the resulting regional corrected gray level measures were averaged for each animal.

Assessment of grain density over neurons of the peri-PVN zone and perifornical region was performed by a semiautomated computerized grain-counting protocol. Images of lightly counterstained neurons of the peri-PVN region and perifornical nucleus were captured at  $63 \times$ , through a blue Wratten filter (no. 47). This filter reduced the intensity of the counterstain and allowed grains to be clearly distinguished from Nissl-counterstained cellular profiles. Cellular profiles were manually sampled, and area determinations were made within Image 1.59. Images were then thresholded to visualize grains only, and area determinations were repeated. The sampling template was then moved to an unhybridized region of tissue to establish background grain area. Results were expressed as the percent of area occupied by grains, calculated as:

$$\% \text{ area} = \frac{(\text{sampled area occupied by grains}) - (\text{background area occupied by grains})}{\text{total area of sampling template}} \times 100.$$

**Image processing.** Images were obtained from x-ray film autoradiographs or negatives using a Polaroid SprintScan 35 slide scanner and Adobe Photoshop 4.0 software. Images imported into Photoshop were contrast and brightness adjusted and assembled into composite images (Figs. 1, 2).

**Plasma hormone assays.** Plasma from trunk blood samples was processed for radioimmunoassay (RIA) for CORT and ACTH. Plasma CORT and ACTH levels were obtained via RIA kits using  $^{125}\text{I}$  tracers from ICN Biomedicals (Cleveland, OH) and IncSTAR, respectively.

**Data analysis.** Acute stress data were analyzed by one-way ANOVA, with time differences then evaluated by Duncan's multiple range test. Chronic stress data were analyzed by unpaired Student's *t* test.

## RESULTS

### Physiological impact of acute and chronic intermittent stress

Effects of restraint on HPA activation are summarized in Table 1.

**Table 2. Effects of chronic stress on adrenal weight, thymus weight, plasma ACTH, and plasma corticosterone**

Group	ACTH (pg/ml)	CORT (ng/ml)	Adrenal wt. (mg)	Adrenal wt. (mg/100gm)	Thymus wt (mg)	Thymus wt (mg/100gm)
Handled	50.4 ± 2.0	52.4 ± 10.3	37.0 ± 1.5	12.7 ± 4.0	436 ± 25	149.6 ± 4.7
Stress	79.6 ± 17.9	49.6 ± 23.5	42.4 ± 1.0*	16.5 ± 4.0*	336 ± 22*	130.4 ± 8.4

\*Significant differences. wt, Weight.

As expected, there was a significant effect of restraint on both ACTH [ $F_{(2,19)} = 6.76$ ;  $p < 0.05$ ] and CORT [ $F_{(2,19)} = 19.46$ ;  $p < 0.05$ ] secretion. Effects were carried by significant elevations in both ACTH and CORT at the 60 min time point after stress (Duncan's multiple range test).

Chronic intermittent stress data are summarized in Table 2. There was no overall effect of stress on basal ACTH or CORT secretion, likely reflecting the fact that animals were killed 16 hr after the last stressor. A significant long-term impact of chronic stress is verified by adrenal hypertrophy [raw adrenal weight,  $t(10) = 5.50$  and  $p < 0.05$ ; adrenal weight/100 gm of body weight,  $t(10) = 7.26$  and  $p < 0.05$ ] and decreased thymus weight [raw thymus weight,  $t(10) = 2.96$  and  $p < 0.05$ ]. Effects of stress on thymus weight/100 gm of body weight approached statistical reliability [ $t(10) = 1.99$ ;  $p = 0.07$ ].

### Localization of GAD65 and GAD67 mRNA

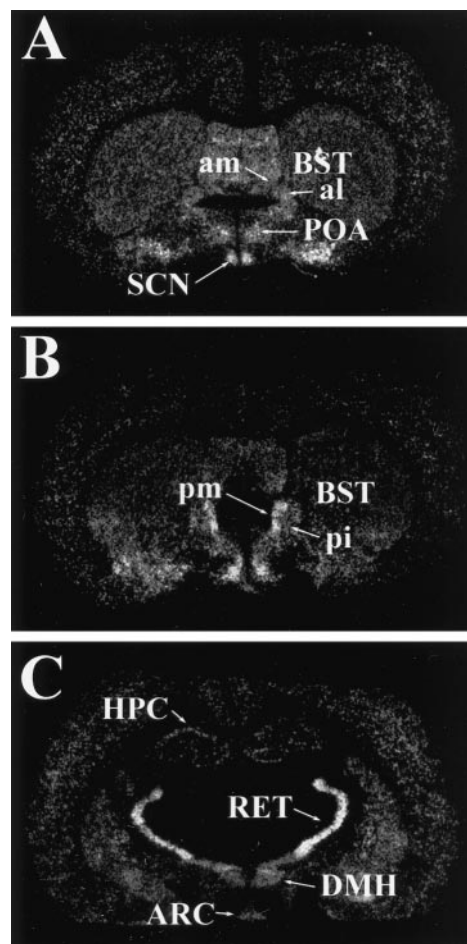
*In situ* hybridization analysis indicated that GAD65 and GAD67 mRNAs were highly abundant in numerous CNS loci (Fig. 1). Of particular relevance to the present study, both GAD65 and GAD67 mRNAs were present in numerous PVN-projecting nuclei, including the medial preoptic area, anterior hypothalamic area, dorsomedial hypothalamic nucleus, arcuate nucleus, and the anterodorsal, anteromedial, posteromedial, and posterointermediate divisions of the BST. In addition, significant hybridization was also seen in the lateral septum, posterolateral BST, and suprachiasmatic nuclei, regions projecting to the immediate surround of the PVN. Neurons in close proximity to the PVN (peri-PVN zone and perifornical nucleus) also express high levels of both GAD mRNAs (Fig. 2).

GAD65 and GAD67 mRNAs were highly expressed outside the hypothalamic-basal forebrain continuum. In the hippocampus, a region that has been repeatedly implicated in HPA regulation, GAD65- and 67-positive neurons were scattered throughout the CA1 and CA3 subfields and in the dentate gyrus (Fig. 1C), consistent with localization to interneurons. Positive hybridization was also observed in neurons throughout the cerebral cortex and striatum. Robust expression was observed in the thalamic reticular nucleus and anteroventral thalamic nucleus. In all cases, overlap of regions hybridized for GAD65 and GAD67 was extensive, suggesting a high degree of colocalization (Escalapez et al., 1993; Feldblum et al., 1993).

### Acute stress

Semiquantitative *in situ* hybridization was used to assess changes in GAD65 mRNA expression after acute restraint stress. Results of densitometric analysis are summarized in Figure 3A. Exposure to acute restraint elevated GAD65 mRNA expression in the anteromedial subnucleus of the bed nucleus of the stria terminalis. However, GAD65 mRNA expression was not affected in any other subdivisions of the bed nucleus of the stria terminalis or in any region of the hypothalamus. No effects of stress were observed in reticular thalamus or caudate nucleus.

Expression of GAD65 mRNAs in the peri-PVN region and

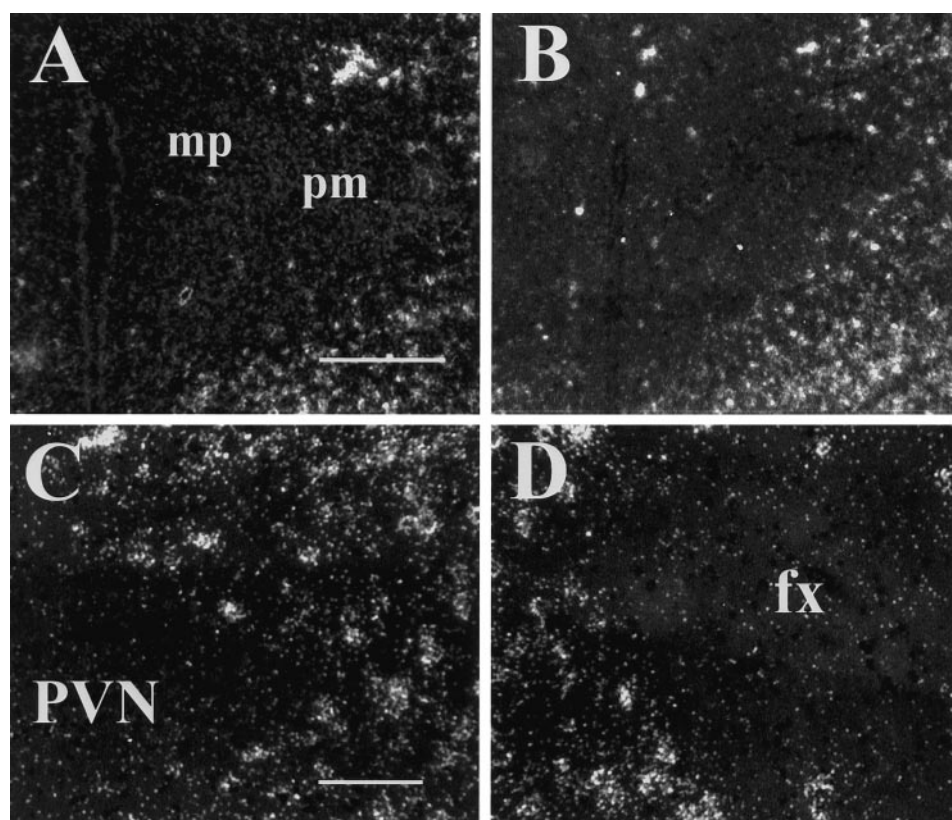


**Figure 1.** Localization of GAD65 mRNA in central stress circuits. GAD65 mRNA is expressed in numerous forebrain stress-relevant nuclei, including the anteromedial (am), anterolateral (al), posteromedial (pm), and posterointermediate (pi) subdivisions of the bed nucleus of the stria terminalis; the medial and lateral preoptic area (POA); the suprachiasmatic nucleus (SCN); the dorsomedial hypothalamic nucleus (DMH); the arcuate nucleus (ARC); and the hippocampal formation (HPC). GAD65 mRNA is also localized to additional regions not implicated in stress regulation, such as the reticular thalamic nucleus (RET). Scale bar, 1 mm.

perifornical nucleus were assessed by semiautomated grain density measures (Fig. 3B). There was a significant effect of stress on GAD65 mRNA expression in the immediate surround of the PVN [ $F_{(2,19)} = 4.04$ ;  $p < 0.05$ ]; this effect was carried by a decrease in the 120 min group relative to unstressed controls ( $p < 0.05$ ). There was no effect of stress on the perifornical cell group.

Acute stress did not affect expression of GAD65 mRNA in hippocampal pyramidal cell layers or in parietal cortex (Fig. 3C). However, significant effects of acute stress on GAD65 expression were observed in the dentate gyrus [ $F_{(2,18)} = 4.95$ ;  $p < 0.05$ ]. Post





**Figure 2.** Localization of GAD65 (*A*) and GAD67 (*B*) mRNAs in the immediate surround of the PVN. Note that very few GAD mRNA-expressing neurons are present in the medial parvocellular (*mp*) or posterior magnocellular (*pm*) PVN; however, aggregates of GAD-positive neurons can be observed to cluster just outside the PVN proper (*A, B*) and in the neighboring perifornical region. Higher power photomicrographs indicate high levels of GAD65 (*C, D*) and GAD67 (data not shown) mRNA in scattered neurons in both regions. *fx*, Foramen. Scale bars: *A, B*, 200  $\mu$ m; *C, D*, 100  $\mu$ m.

*hoc* analysis (Duncan's multiple range test) revealed significant increases in the 60 min stress group relative to unstressed rats.

In contrast with GAD65, significant effects of stress on GAD67 mRNA were observed in several regions of the hypothalamus, including the medial preoptic area [ $F_{(2,19)} = 5.17$ ;  $p < 0.05$ ], arcuate nucleus [ $F_{(2,19)} = 12.56$ ;  $p < 0.05$ ], and dorsomedial hypothalamic nucleus [ $F_{(2,19)} = 6.11$ ;  $p < 0.05$ ] (Fig. 4*A*). In all cases, expression was increased only at the 60 min time point after stress. No changes in GAD67 expression were observed in the anterior hypothalamic area or suprachiasmatic nucleus. Within the BST, GAD67 mRNA was increased in the anteromedial [ $F_{(2,18)} = 6.36$ ;  $p < 0.05$ ] and anterodorsal [ $F_{(2,18)} = 4.65$ ;  $p < 0.05$ ] subnuclei. No changes were observed in the posteromedial and posterointermediate or posterolateral subdivisions of the BST. Similarly, no changes in cellular GAD67 mRNA expression were observed in the peri-PVN region or perifornical nucleus (Fig. 4*B*).

Increased GAD67 mRNA was also observed in the hippocampus (Fig. 4*C*). Significant effects of stress on GAD67 expression were observed in CA1 [ $F_{(2,18)} = 5.82$ ;  $p < 0.05$ ] and dentate gyrus [ $F_{(2,18)} = 6.61$ ;  $p < 0.05$ ]. In both cases, 60 min groups were distinguished from unstressed animals ( $p < 0.05$ ). No stress effects were observed in the CA3, parietal cortex, reticular thalamic nucleus, or caudate nucleus.

#### Chronic stress

In contrast to the results of the acute stress experiments, chronic stress primarily affected GAD65 mRNA expression (Figs. 5, 6). GAD65 mRNA was increased in several hypothalamic nuclei, including the medial preoptic area [ $t(9) = 3.92$ ;  $p < 0.05$ ], anterior hypothalamic area [ $t(9) = 2.35$ ;  $p < 0.05$ ], and dorsomedial hypothalamic nucleus [ $t(9) = 2.51$ ;  $p < 0.05$ ] (Fig. 5*A*). GAD65 mRNA expression was also increased in the anterodorsal

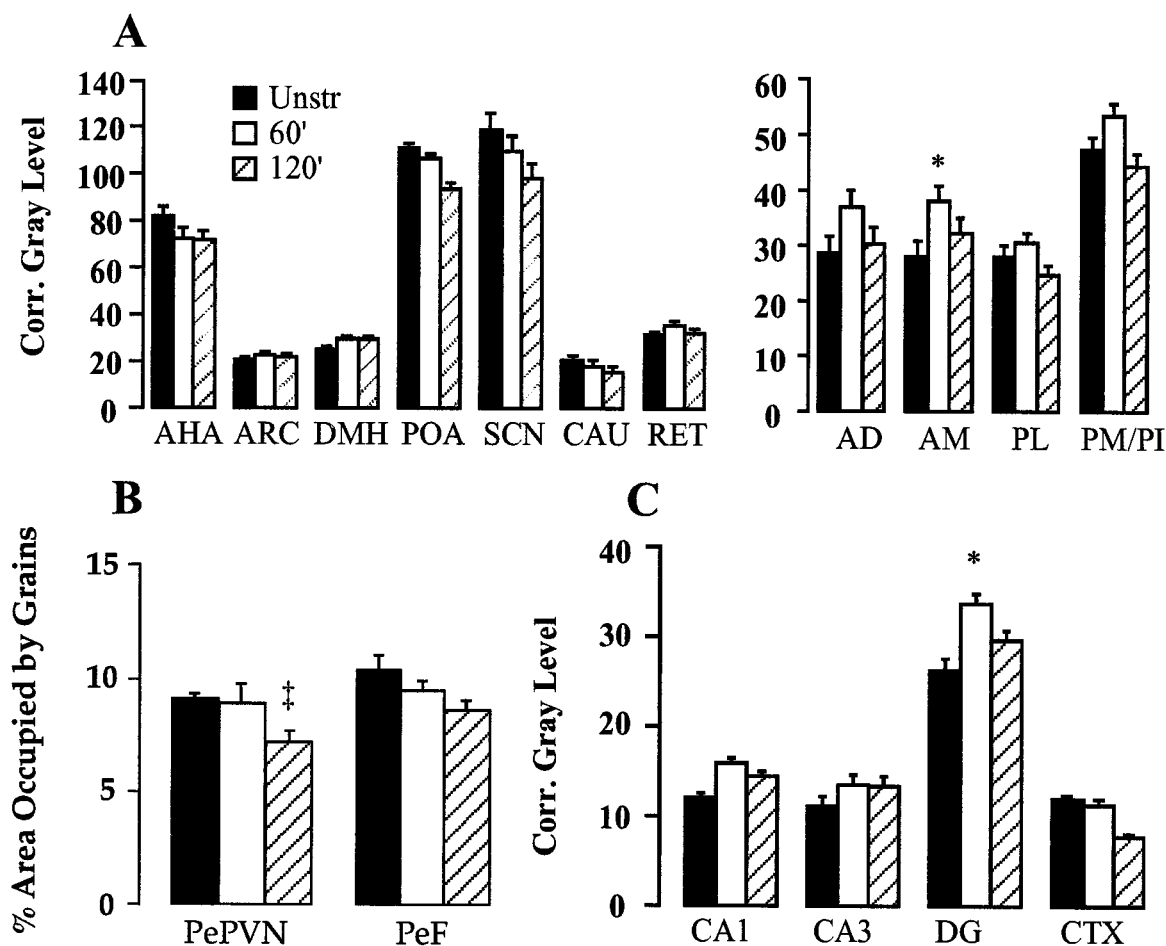
[ $t(10) = 5.07$ ;  $p < 0.05$ ] and anteromedial [ $t(10) = 4.83$ ;  $p < 0.05$ ] BST; no changes were observed in the posterolateral or posteromedial and posterointermediate subnuclei. GAD65 hybridization density was increased over neurons in the peri-PVN region [ $t(8) = 2.28$ ;  $p < 0.05$ ] and in the perifornical nucleus [ $t(8) = 2.71$ ;  $p < 0.05$ ] (Fig. 5*B*). No changes in GAD65 mRNA expression were observed in the hippocampus (Fig. 5*C*), caudate, parietal cortex, or reticular thalamic nucleus.

Chronic stress-induced changes in GAD67 mRNA were considerably more limited than that in GAD65 mRNA (Fig. 6). Increased levels of GAD67 mRNA were observed in the medial preoptic area [ $t(9) = 2.21$ ;  $p = 0.05$ ] and in the anteromedial [ $t(9) = 2.48$ ;  $p < 0.05$ ] and anterodorsal [ $t(9) = 2.37$ ;  $p < 0.05$ ] subnuclei of the BST (Fig. 6*A*). In contrast, no GAD67 changes were seen in any other region of the hypothalamus proper or in individual neurons of the peri-PVN region and perifornical nucleus (Fig. 6*B*). GAD67 mRNA was upregulated by stress in the hippocampal subfield CA3 [ $t(9) = 4.64$ ;  $p < 0.05$ ] and in the dentate gyrus [ $t(9) = 2.22$ ;  $p = 0.05$ ] (Fig. 6*C*). No changes in GAD67 mRNA were observed in CA1, cortex, caudate nucleus, or reticular thalamus.

## DISCUSSION

### GAD65 and GAD67 mRNA regulation in PVN-projecting nuclei

The present study is consistent with the hypothesis that PVN-projecting GABAergic neurons are instrumental in central stress regulation. Expression of GAD65 and GAD67 mRNA was markedly increased by stress in numerous hypothalamic and BST cell groups known to have efferents to the medial parvocellular PVN. These include the medial preoptic area, dorsomedial hypothalamic nucleus, arcuate nucleus, peri-PVN region, perifornical

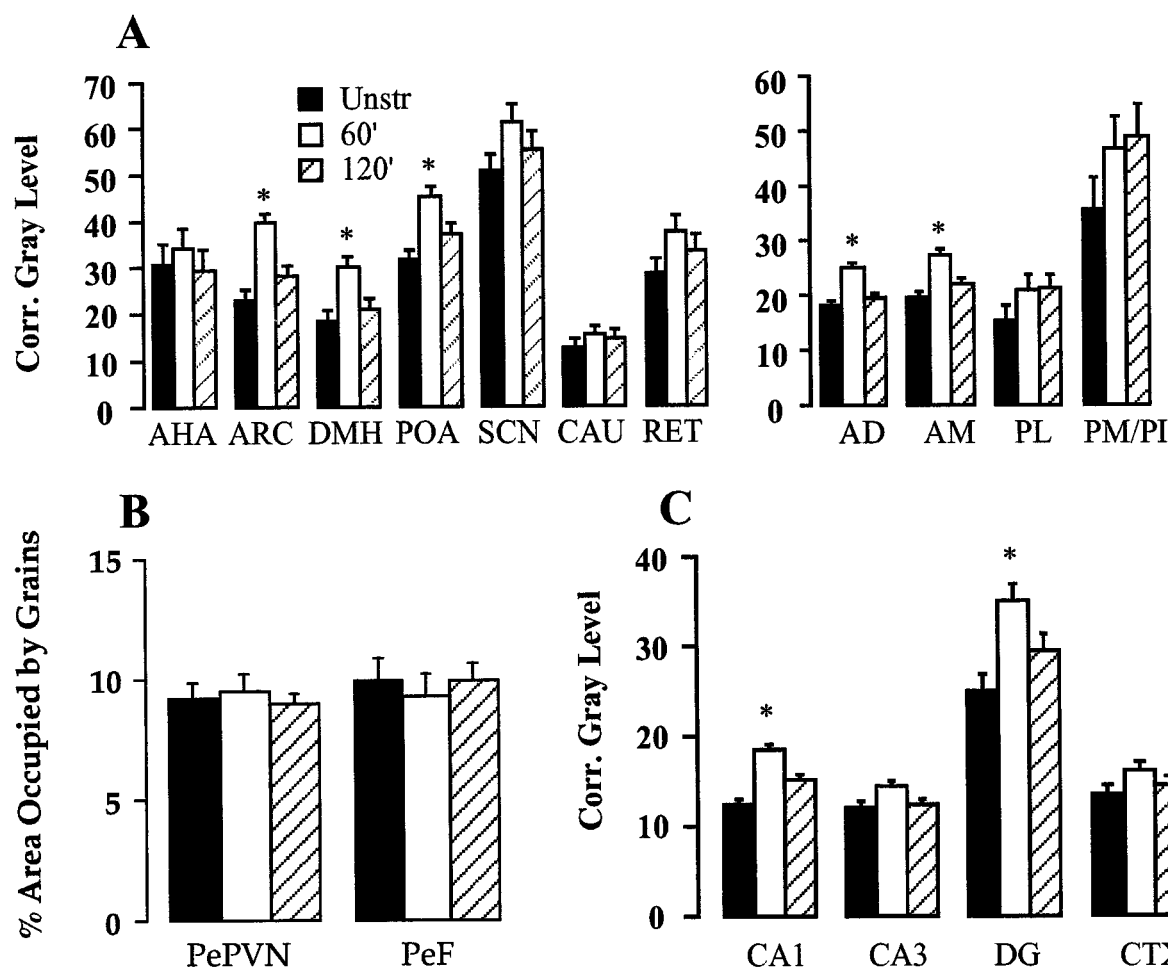


**Figure 3.** Semiquantitative assessment of the effect of acute restraint on GAD65 mRNA expression in forebrain stress-relevant nuclei. *A*, No changes in GAD65 mRNA expression were observed in the anterior hypothalamic nucleus (AHA), arcuate nucleus (ARC), dorsomedial hypothalamic nucleus (DMH), medial preoptic area (POA), suprachiasmatic nucleus (SCN), caudate nucleus (CAU), and reticular thalamic nucleus (RET). GAD65 mRNA is elevated 60 min after stress in the anteromedial subnucleus (AM) of the bed nucleus of the stria terminalis but is not altered in the anterodorsal (AD), posterolateral (PL), or posteromedial and posterointermediate subnuclei (PM/PI). *B*, Grain density analysis was used to assess GAD65 mRNA expression in the peri-PVN region (PePVN) and the perifornical nucleus (PeF). Reduced grain density was observed over neurons in the peri-PVN region 120 min after stress exposure. *C*, Densitometric analysis of GAD65 mRNA in the hippocampus revealed significant elevation in dentate gyrus (DG) 60 min after stress exposure. CTX, Parietal cortex; Unstr, Unstressed.

nucleus, and anterior subnuclei of the BST. Importantly, these regions have all been implicated in central control of the HPA axis. The medial preoptic area is known to inhibit ACTH and corticosterone secretion and is a potential site of glucocorticoid negative feedback inhibition of the HPA axis (Viau and Meaney, 1996). The dorsomedial nucleus maintains a strong GABAergic projection to the PVN that is directly activated by stressful stimuli (Cullinan et al., 1995, 1996). Lesions of the arcuate nucleus increase stress-induced corticosterone secretion (Magarinos et al., 1988; Larsen et al., 1994), suggestive of an inhibitory role of this region in PVN regulation. Electrophysiological studies note that neurons in the peri-PVN region and perifornical nucleus inhibit neuronal activity of medial parvocellular PVN neurons (Boudaba et al., 1996). Finally, neurons in the anterior subdivisions of the BST are involved in increased CRH mRNA expression seen after anterior BST lesions (Herman et al., 1994). Thus, all PVN-projecting regions showing increased GAD expression after stress have been associated with HPA regulation, strengthening the hypothesis that GABAergic neurons contained in these areas play an important role in inhibition of the stress axis.

Exposure to stress results in distinct patterns of GAD65 and GAD67 mRNA upregulation. In the case of acute stress, GAD67 appears to be preferentially induced. The number of regions showing upregulation of GAD67 mRNA is substantially more widespread than that of GAD65. Induction of GAD67 is rapid and transient, occurring at 1 hr of stimulation and returning to levels indistinguishable from baseline within 2 hr of stress induction. The limited effects of acute stress on GAD65 mRNA levels (BST and dentate gyrus only) suggest that this gene is less responsive than GAD67; however, it is also possible that induction of the GAD65 gene may occur more slowly and thus not be visible in the present study. The rapid activation and inactivation of GAD67 mRNA expression suggests transcriptional regulation by immediate early genes; indeed, it is important to note that all regions showing GAD67 upregulation express cFOS after restraint (Cullinan et al., 1995), consistent with transcriptional activation by AP-1.

In contrast to GAD67 mRNA, GAD65 mRNA is preferentially activated by prolonged stress exposure. After a 2 week exposure to chronic intermittent stress, rats show pronounced increases in



**Figure 4.** Semiquantitative assessment of the effect of acute restraint on GAD67 mRNA expression in forebrain stress-relevant nuclei. *A*, In contrast with GAD65 mRNA, GAD67 mRNA was elevated in the *ARC*, *DMH*, medial *POA*, and anterodorsal and anteromedial BST 60 min after initiation of stress. In all cases, GAD67 mRNA levels returned to baseline by 120 min. *B*, No changes in GAD67 grain density were observed over neurons in the peri-PVN region or *PeF* after acute stress. *C*, GAD67 mRNA was elevated in the hippocampal subfield *CA1* and *DG* 60 min after stress initiation. Abbreviations are given in the Figure 3 legend.

GAD65 mRNA expression in most of the same cell groups manifesting increased GAD67 expression with acute stress (i.e., medial preoptic area, dorsomedial nucleus, and anterodorsal and anteromedial BST). Increased GAD65 mRNA was also seen in PVN-projecting regions of the anterior hypothalamic area, peri-PVN zone, and perifornical nucleus. GAD67 upregulation was substantially more limited, being confined to the anterodorsal and anteromedial BST and medial preoptic area.

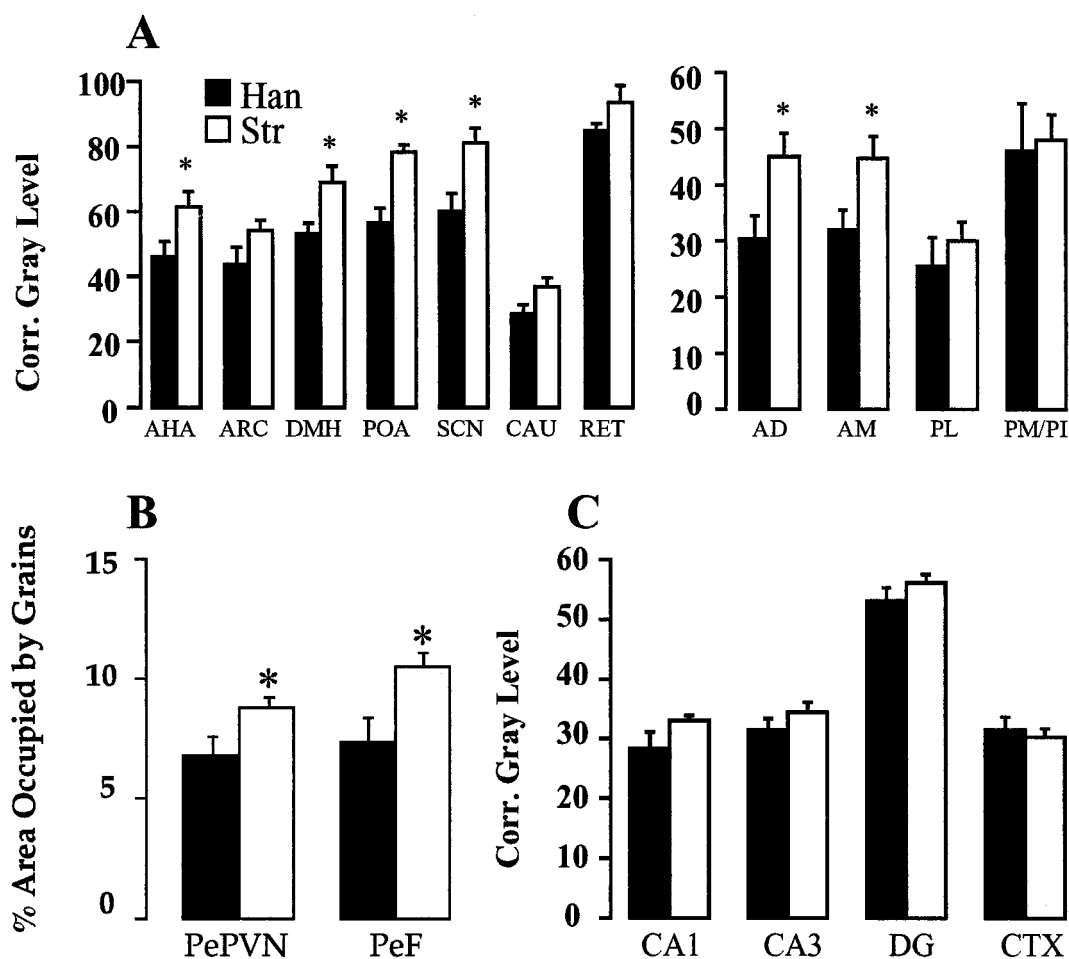
Expression of GAD65 mRNA is slightly decreased in the immediate surround of the PVN 2 hr after induction of acute restraint. Interestingly, this area is the only brain region to show decreased GAD expression after stress. This decrease may be related to heavy innervation of this region by GABAergic brain regions activated during stress, such as the lateral septum (Cullinan et al., 1995; Risold and Swanson, 1996). Evidence of chronic stress-induced increases in GAD65 mRNA expression in this region suggests that such inhibition may be overcome by prolonged stimulation.

#### GAD65 and GAD67 mRNA regulation in non-PVN projecting loci

Other stress-relevant regions, such as the suprachiasmatic nucleus and hippocampus, showed distinctive patterns of GAD65 and

GAD67 mRNA expression after acute and chronic stimulation. The suprachiasmatic nucleus is known to play a role in circadian regulation of corticosterone secretion (Cascio et al., 1987) and is implicated in inhibition of PVN activation by acute stress (Kalsbeck et al., 1992; Buijs et al., 1993a). The suprachiasmatic–PVN connection appears to be mediated by way of interneurons, perhaps in the subparaventricular zone (Watts et al., 1987; Buijs et al., 1993b), indicating a trans-synaptic influence. Upregulation of suprachiasmatic nucleus GAD mRNA expression was only observed after chronic stress and was specific for GAD65. It remains to be determined whether altered GABAergic neurotransmission may be involved in stress-induced disruption of circadian rhythms.

The hippocampus is known to inhibit the HPA axis (Jacobson and Sapolsky, 1991; Herman and Cullinan, 1997). Hippocampal actions appear to be trans-synaptic and may involve connections between hippocampal outflow neurons in the ventral subiculum and subcortical GABAergic pathways, notably including the medial preoptic area, BST, anterior hypothalamus, dorsomedial hypothalamic nucleus, and peri-PVN region (Cullinan et al., 1993; Herman et al., 1995). As such, it was of interest to determine whether hippocampal GAD was regulated in parallel with PVN-



**Figure 5.** Semiquantitative assessment of GAD65 mRNA expression in forebrain stress-relevant nuclei after chronic stress or handling. *A*, GAD65 mRNA expression was significantly increased in the *AHA*, *DMH*, *POA*, *SCN*, and the *AD* and *AM* divisions of the BST of rats exposed to chronic stress. *B*, A significant increase in GAD65 mRNA expression/cell was seen in both the peri-PVN region and the perifornical nucleus by grain density analysis. *C*, GAD65 mRNA expression in the hippocampus was unaffected by chronic stress. *Han*, Handled; *Str*, stressed. Other abbreviations are given in the Figure 3 legend.

projecting hypothalamic and BST cell groups. In agreement with this notion, GAD67 mRNA is increased by acute stress in the CA1 and dentate gyrus. However, dentate gyrus GAD65 mRNA was also increased with acute stress, and chronic stress increased GAD67 mRNA in both CA3 and dentate gyrus. Thus, GAD regulation does not obey the same pattern in hippocampus as in the hypothalamic–BST continuum, indicating differential regulation among potential stress-regulatory pathways.

Notably, no changes in GAD65 or GAD67 mRNA were observed in the parietal cortex, caudate nucleus, or reticular thalamus. These regions have not been directly implicated in stress regulation, suggesting that stress-induced changes are not generalized throughout the nervous system. These data also verify that positive findings were not attributable to random variance in hybridization efficiency across sections. Thus, changes in GAD expression seem to be relegated to CNS pathways implicated in control of stress responsiveness.

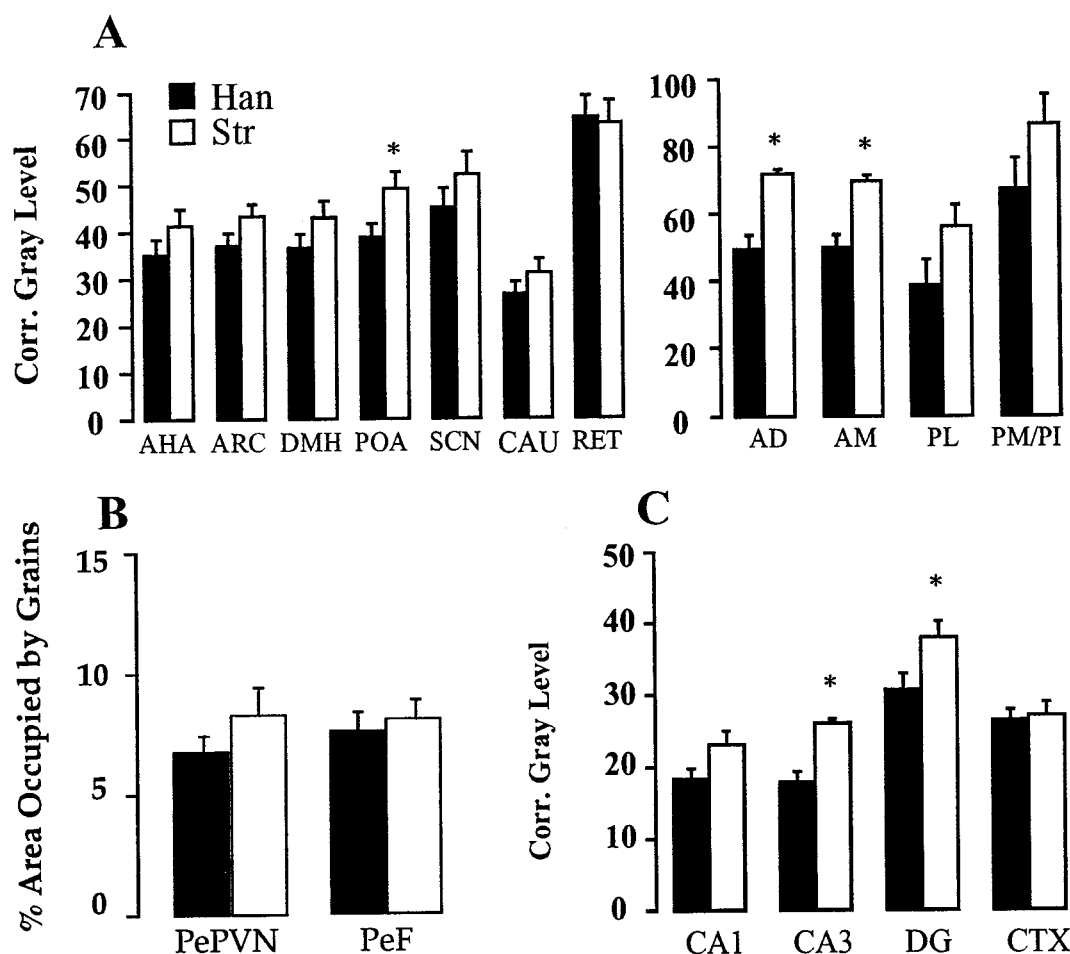
#### Differential regulation of GAD65 and GAD67 mRNA by acute and chronic stress: functional implications

The distinctive induction of GAD65 and GAD67 mRNAs by chronic or acute stress, respectively, seems in keeping with the

perceived role of the two isoforms in neuronal function. Approximately 50% of GAD65 appears as inactive apoenzyme (Kaufman et al., 1991). Much of GAD65 immunoreactivity is localized to nerve terminals, suggesting that GAD65 is stored in inactive form in presynaptic endings (Esclapez et al., 1994). This suggests that chronic stress-induced upregulation of GAD65 may increase apo-GAD stores, presumably to compensate for greater rates of stimulation. Conversely, GAD67 does not appear to be sequestered as an inactive apoenzyme and is enriched in neuronal cell bodies and dendrites (Kaufman et al., 1991; Esclapez et al., 1994). These data suggest the rate of GAD67 usage is higher than that of GAD65, given diminished localization in sites of storage (e.g., terminals) and the relative lack of detectable apo-GAD67. Upregulation of GAD67 mRNA by acute stress may thus reflect biosynthesis keyed to neuronal activity.

Precedent for physiological modulation of GAD65 and GAD67 mRNA expression has been noted in the literature. For example, striatal GAD67 mRNA and protein expression are upregulated after cortical ischemia (Salin and Chesselet, 1993). Striatal GAD activity and GAD67 mRNA levels are also increased after 6-hydroxydopamine lesion of the nigrostriatal pathway (Lindfors et al., 1989; Segovia et al., 1990). Interestingly, GAD65 and





**Figure 6.** Semiquantitative assessment of GAD67 mRNA expression in forebrain stress-relevant nuclei after chronic stress or handling. *A*, Chronic stress induction of GAD67 mRNA expression was observed only in the medial *POA* and the *AD* and *AM* divisions of the BST of rats. *B*, No changes in GAD67 mRNA were seen in the peri-PVN region or perifornical nucleus by grain density analysis. *C*, Chronic stress exposure increased GAD67 mRNA expression in subfield *CA3* of the hippocampus and in the *DG*. Abbreviations are given in the legends of Figures 3 and 5.

GAD67 mRNAs are differentially regulated in this model system; in general, changes in GAD65 mRNA are more circumscribed than are those in GAD67 mRNA, showing no induction in the striatum (Soghomonian et al., 1992) and minimal induction (relative to GAD67) in the thalamic reticular nucleus, a downstream target of striatal efferents (Delfs et al., 1996). Pharmacological analyses indicate that long-term treatment with neuroleptic drugs increases GAD67 mRNA expression in the entopeduncular nucleus and globus pallidus (Mercugliano et al., 1992), further consistent with an integral role for GABA in extrapyramidal regulation. In the cerebellum, neurotoxic lesions of the climbing-fiber pathway increased GAD67 mRNA expression and GAD activity in Purkinje cell populations (Litwak et al., 1990), indicating induction of GAD67 gene transcription by cellular activity. Nonetheless, changes in GAD67 mRNA occur after chronic stimulation, indicating that GAD expression is capable of responding dynamically to changes in activity in multiple neuronal systems.

In the present study, long-term stress exposure does not affect expression of GAD67 mRNA in the majority of the stress-related regions examined; rather, chronic stress seems to differentially increase expression of GAD65 mRNA. These results suggest that, unlike the extrapyramidal system and cerebellum, activity-

dependent upregulation of GAD in stress pathways occurs via increased expression of GAD65 mRNA. Alternatively, this difference may also reflect a greater sensitivity of the GAD65 gene to induction by stress; GAD67 mRNA induction may require the more prolonged and consistent stimulation afforded by lesion or pharmacological stimulation.

Induction of GAD by stress has important implications for HPA regulation. First, regions showing GAD induction correspond with those showing cFOS expression after stress (Cullinan et al., 1996). These neural populations also contain inhibitory neuropeptides, including CRH (Champagne et al., 1998), further supporting inhibitory actions on HPA activity. Together, these data suggest that BST and hypothalamic GABAergic cell populations are activated by stress and are likely to convey inhibition to the PVN. Second, hypothalamic and BST neurons are in a position to interconnect regions such as the hippocampus and amygdala with the PVN (Price et al., 1987; Swanson, 1987; Swanson et al., 1987). Thus, this collection of GABAergic neurons may translate limbic output into appropriate integration of stress responses. Connections between PVN-projecting neuronal populations and extrahypothalamic glucocorticoid-receptive sites (e.g., the hippocampus/ventral subiculum) (Herman, 1993) also raise the possibility that GABA may play a role in translating glucocorticoid



feedback signals into PVN inhibition. Third, induction of GAD65 mRNA is consistent with a role for GABA in attenuating activation of the HPA axis in the face of chronic drive. Increased GAD65 mRNA availability predicts increased GAD65 levels, which may stand to enhance inhibition at the PVN. Finally, induction of GAD in the hippocampus suggests that local GABA may modulate cognitive function after stress. Stress is known to have deleterious effects on learning and memory (Luine et al., 1993; Diamond and Rose, 1994; Diamond et al., 1994) and induces dendritic atrophy in subfield CA3 (Magarinos and McEwen, 1995). The observed increases in GAD expression in the hippocampus raises the possibility that altered inhibition may contribute to stress-induced behavioral changes.

In summary, GABA mRNA synthesis is specifically increased in PVN-projecting brain regions after acute and chronic stress exposure. Acute increases in GAD67 are likely keyed toward replenishing GABA released after stimulation, whereas chronic increases in GAD65 may serve to attenuate the effects of repetitive stimulation on central stress circuitry. The results point toward a prominent role for GABAergic pathways in central stress integration.

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