

Forum Original Research Communication

Redox Modulation of Gi Protein Expression and Adenylyl Cyclase Signaling: Role of Nitric Oxide

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ABSTRACT

Nitric oxide (NO) has been shown to regulate a variety of physiological functions, including vascular tone. The inhibition of NO synthase by *N*^ω-nitro-L-arginine methyl ester (L-NAME) has been reported to increase arterial blood pressure. The present studies were undertaken to investigate if the increased blood pressure by L-NAME is associated with enhanced expression of Gi proteins, implicated in the pathogenesis of hypertension. L-NAME was administered orally into Sprague–Dawley rats for a period of 4 weeks. Control rats were given plain tap water only. The systolic blood pressure was enhanced in L-NAME-treated rats as compared with control rats; however, the heart-to-body weight ratio was not different in the two groups. The levels of Gi α -2 and Gi α -3 proteins and their mRNA as determined by western and northern blotting, respectively, were significantly augmented in hearts from L-NAME-treated rats, whereas the levels of Gs α and G β were unaltered. In addition, the effect of low concentrations of GTP γ S on forskolin-stimulated adenylyl cyclase activity (receptor-independent functions of Gi α) was significantly enhanced, whereas the receptor-dependent inhibitions of adenylyl cyclase were completely attenuated in L-NAME-treated rats. Whereas cholera toxin-mediated stimulation of adenylyl cyclase was unaltered in both group of rats, the stimulatory effects of some agonists on adenylyl cyclase activity were diminished in L-NAME-treated rats. These results suggest the implication of NO in the modulation of Gi protein expression and associated adenylyl cyclase signaling. *Antioxid. Redox Signal.* 6, 385–392.

INTRODUCTION

GUANINE NUCLEOTIDE REGULATORY PROTEINS (G proteins) are a family of GTP-binding proteins that play an important role in the regulation of a variety of signal transduction systems, including the adenylyl cyclase/cyclic AMP (cAMP) system. The adenylyl cyclase system is composed of three components: receptor, catalytic subunit, and stimulatory (Gs) and inhibitory (Gi) guanine nucleotide regulatory proteins (16, 38). The stimulation and inhibition of adenylyl cyclase by hormones are mediated by two distinct G proteins, Gs and Gi, respectively, that couple the receptor to the catalytic subunit. The G proteins are heterotrimeric and are composed of α , β , and γ subunits. The α subunits bind and hydrolyze GTP and confer specificity in receptor and effector interactions. Molecular cloning has revealed four different forms of Gs α result-

ing from the differential splicing of one gene (10, 36, 44) and three distinct forms of Gi α , Gi α -1, Gi α -2, and Gi α -3, encoded by three distinct genes (23). All three forms of Gi α (Gi α 1–3) are implicated in adenylyl cyclase inhibition (23) and activation of atrial K⁺ channels (53).

The adenylyl cyclase/cAMP system has been implicated in both the control of heart contractility and vascular smooth muscle tone (25, 49). The levels of cAMP are regulated by Gs and Gi proteins. Gi protein and associated adenylyl cyclase signaling have been shown to be implicated in a variety of cellular functions, including vascular permeability (20, 39), salt and water transport (15, 26), and catecholamine release (50), all of which play a key role in the regulation of blood pressure. Alterations in the levels of Gi proteins and cAMP that result in impaired cellular functions lead to various pathological states, including hypertension. Several abnormalities in

G-protein expression, adenylyl cyclase activity, and cAMP levels have been reported in cardiovascular tissues from genetic [spontaneously hypertensive rats (SHR)] and different models of experimentally induced hypertensive rats (1, 2, 6, 7, 9, 48). An increased expression of Gi protein and Gi-protein mRNA in heart and aorta from SHR and hearts from DOCA-salt (deoxycorticosterone acetate) hypertensive rats with established hypertension has been reported (2, 7, 48). On the other hand, the levels of G_{α} were shown to be unaltered in SHR, but were decreased in DOCA-salt hypertensive rats (2, 6, 7). We have recently shown that the increased expression of G_{α} proteins occurs before the onset of hypertension in SHR (34) and DOCA-salt hypertensive rats (35), suggesting that the enhanced levels of G_{α} proteins may be one of the contributing factors for the pathogenesis of hypertension. This was supported by our recent studies showing that inactivation of enhanced expression of Gi proteins by pertussis toxin treatment attenuates the development of hypertension in SHR (31).

The implication of free radicals in the pathogenesis of hypertension has been well established (27). Increased levels of superoxide anion ($O_2^{\cdot-}$) and hydrogen peroxide (H_2O_2) have been reported in hypertensive patients (28). The endothelium plays an important role in the regulation of cardiovascular functions by its ability to produce a variety of vasoactive agents, including nitric oxide (NO) (51). NO is synthesized from L-arginine by the action of NO synthase and regulates a variety of physiological processes, including vascular tone, platelet aggregation, neurotransmission, cell differentiation, cell migration, hormone release, and apoptosis (30). Endothelial dysfunction has been shown to be an important contributor in the pathogenesis of hypertension due to the impaired generation of NO, which regulates vasodilation. Vitamin C, an antioxidant, has been reported to recover endothelial functions by restoring NO-mediated vasodilation of the endothelium in hypertensive patients (47). NO reacts with superoxide anion $O_2^{\cdot-}$ to form peroxynitrite ($ONOO^-$), which is a powerful oxidant and has been implicated in the vasoconstriction of vascular smooth muscle cells and atherosclerosis (37, 52). NO synthase has been shown to be inhibited by several arginine analogues, including N^{ω} -nitro-L-arginine methyl ester (L-NAME) (42). Acute *in vivo* administration of L-NAME elicits marked arterial hypertension without hypertrophy (8), presumably as a consequence of an abrupt inhibition of NO synthesis (42) that results in the inhibition of vasorelaxation.

The present studies were undertaken to investigate if the increased blood pressure by L-NAME is associated with enhanced expression of Gi proteins and adenylyl cyclase signaling. We have shown that L-NAME treatment enhances the Gi protein expression and adenylyl cyclase signaling in rat heart, suggesting a role of NO in the modulation of Gi protein expression and thereby blood pressure.

MATERIALS AND METHODS

Animals

Male Sprague-Dawley rats (200–250 g) purchased from Charles River Canada (St. Constant, Quebec, Canada) were used in the present studies. L-NAME (100 mg/kg/day) was administered orally into rats in their drinking tap water for a period

of 4 weeks, whereas control rats received plain tap water only. The arterial blood pressure (mm Hg) was measured after 4 weeks of treatment by the tail-cuff method without anesthesia. After 4 weeks of treatment, the rats were killed by decapitation, and the hearts were removed for adenylyl cyclase activity determination and mRNA and protein quantification. All the animal procedures used in the present studies were reviewed and approved by the Comité de déontologie de l'expérimentation sur les animaux of the University of Montreal (no. 99050).

Preparation of heart particulate fraction

The heart particulate fraction was prepared as described previously (34, 35). Frozen hearts were quickly pulverized to a fine powder using mortar and pestle cooled in liquid N_2 . The powder was stored at $-80^{\circ}C$ until assayed. The powder was homogenized (12 strokes) in a Teflon glass homogenizer, in a buffer containing 10 mM Tris-HCl, 1 mM EDTA, pH 7.5. The homogenate was centrifuged at 1,000 g for 10 min. The supernatant fraction was discarded, and the pellet was finally resuspended and homogenized in a buffer containing 10 mM Tris-HCl and 1 mM EDTA, pH 7.5, and used for adenylyl cyclase activity determination and immunoblotting studies.

Cholera toxin (CT) treatment of heart particulate fraction

The heart particulate fraction was treated with CT as described earlier (34, 35). CT (500 μ g/ml) was preactivated for 20 min at $37^{\circ}C$ in a mixture containing 20 mmol/L dithiothreitol, 1 μ g/ml bovine serum albumin, and 25 mmol/L KH_2PO_4 , pH 8.0. To study the effect of CT on adenylyl cyclase activity, the heart particulate fraction was pretreated with and without CT for 30 min at $30^{\circ}C$ in a reaction mixture containing 250 mM KH_2PO_4 , pH 6.8, 1 mM $MgCl_2$, 0.5 mM EDTA, pH 8.0, 5 mM ATP, 15 mM thymidine, 0.15 mM GTP, 20 mM dithiothreitol, and 1 mM NAD. The particulate fractions were washed twice with buffer containing 10 mM Tris and 1 mM EDTA, pH 7.5, and finally suspended in the same buffer for adenylyl cyclase activity determination.

Adenylyl cyclase activity determination

Adenylyl cyclase activity was determined by measuring [^{32}P]cAMP formation from [α - ^{32}P]ATP, as described previously (34, 35). The assay medium contained 50 mM glycylglycine, pH 7.5, 0.5 mM MgATP, [α - ^{32}P]ATP ($1-1.5 \times 10^6$ cpm), 5 mM $MgCl_2$, 100 mM NaCl, 0.5 mM cAMP, 1 mM 3-isobutyl-1-methylxanthine, 0.1 mM EGTA, 10 μ M guanosine 5'-(γ -thio)triphosphate (GTP γ S) (or otherwise as indicated) and an ATP regenerating system consisting of 2 mM phosphocreatine, 0.1 mg of creatine kinase/ml, and 0.1 mg of myokinase/ml in a final volume of 200 μ L. Incubations were initiated by addition of the heart particulate fraction (50–100 μ g) to the reaction mixture, which had been thermally pre-equilibrated for 2 min at $37^{\circ}C$. The reactions, conducted in triplicate for 10 min at $37^{\circ}C$, were terminated by the addition of 0.6 ml of 120 mM zinc acetate. cAMP was purified by coprecipitation of other nucleotides with $ZnCO_3$, addition of 0.5 ml of 144 mM Na_2CO_3 , and subsequent chromatography by the double column system, as described by Salomon *et al.* (45). Under

the assay conditions used, adenylyl cyclase activity was linear with respect to protein concentration and time of incubation.

Protein was determined essentially as described by Lowry *et al.* (33), with crystalline bovine serum albumin as standard.

Immunoblotting

Immunoblotting of G proteins was performed as described previously (34, 35). After sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE), the separated proteins were transferred electrophoretically to a nitrocellulose sheet (Schleicher and Schuell) with a semidry transblot apparatus (Bio-Rad) at 15 V for 45 min. After transfer, the membranes were washed twice in phosphate-buffered saline (PBS) and incubated in PBS containing 3% skim milk at room temperature for 2 h. The blots were then incubated with antibodies against G proteins in PBS containing 1.5% skim milk and 0.1% Tween 20 at room temperature overnight. The antigen–antibody complexes were detected by incubating the blots with goat anti-rabbit IgG (Bio-Rad) conjugated with horseradish peroxidase for 2 h at room temperature. The blots were then washed three times with PBS before reaction with enhanced chemiluminescence western blotting detection reagents (Amersham).

Northern analysis

Total RNA was isolated by the guanidinium thiocyanate–phenol–chloroform method described by Chomczynski and Sacchi (12). cDNA inserts encoding for $G\alpha_2$, $G\alpha_3$, $G\alpha$, and catalytic subunit V of adenylyl cyclase were radiolabeled with [α - 32 P]dCTP by random priming essentially as described by Feinberg and Vogelstein (13). Dimethyl sulfoxide/glyoxal-treated total RNA was resolved on 1% agarose gels and transferred to nylon membrane as described by Sambrook *et al.* (46). Filters, after prehybridization at 65°C for 6 h in hybridization solution (600 mM NaCl, 8 mM EDTA, 120 mM Tris, pH 7.4) containing 0.1% sodium pyrophosphate, 0.2% SDS, and heparin (500 U/ml), were hybridized overnight in hybridization solution containing 10% (wt/vol) dextran sulfate and cDNA probe ($1\text{--}3 \times 10^6$ cpm/ml). Filters were then rinsed at 65°C for 2×30 min in 300 mM NaCl, 4 mM EDTA, 60 mM Tris, pH 7.4, and 0.1% SDS. Autoradiography was performed with x-ray films at -70°C . To assess the possibility of any variations in the amounts of total RNA in individual samples applied to the gel, each filter was hybridized with the ^{32}P end-labeled oligonucleotide, which recognizes a highly conserved region of 28S ribosomal RNA. The blots that had been probed with the G-protein cDNA were dehybridized by washing for 1 h at 65°C in 50% formamide, 300 mM NaCl, 4 mM EDTA, and 60 mM Tris, pH 7.4, and rehybridized overnight at room temperature with the oligonucleotide. Quantitative analysis of the hybridization of probes bound was performed by densitometric scanning of the autoradiographic film using the enhanced laser densitometer, LKB Utroskan XL, and quantified using the gel scan XL evaluation software (v. 2.1) from Amersham Pharmacia Biotech (Baie d'Urfe, PQ, Canada).

Data analysis

Results are expressed as means \pm SEM. Comparisons between groups (control and L-NAME-treated rats) were made with Student's *t* test for unpaired samples, whereas ANOVA

followed by Dunnett's multiple comparison tests or Newman–Keuls multiple comparison tests were performed for the dose-response curve of GTP γ S on forskolin (FSK)-stimulated adenylyl cyclase activity. $p < 0.05$ was considered statistically significant.

RESULTS

Treatment of rats with L-NAME for 4 weeks significantly increased the arterial systolic blood pressure (190 ± 9.2 mm Hg versus 121 ± 6.3 mm Hg); however, it did not affect the heart-to-body weight ratio (2.8 ± 0.36 versus 2.7 ± 0.26 mg/g).

Effect of L-NAME treatment on Gi protein levels

To investigate if the increased blood pressure by L-NAME treatment is associated with increased expression of $G\alpha$ proteins, the levels of Gi proteins were determined in hearts from control and L-NAME-treated rats by immunoblotting techniques using specific antibodies AS/7 against $G\alpha_1$ and $G\alpha_2$ and EC/2 against $G\alpha_3$. The results depicted in Fig. 1 show that the AS/7 recognized a single protein of M_r 40 kDa referred to as $G\alpha_2$ [$G\alpha_1$ has been shown to be absent in heart (24)], whereas EC/2 antibodies recognized a single protein of 41 kDa referred to as $G\alpha_3$ in hearts from control and L-NAME-treated hearts; however, the amounts of immunodetectable $G\alpha_2$ and $G\alpha_3$ protein were increased in L-NAME-treated rats by $38.6 \pm 5.9\%$ and $87.9 \pm 9.7\%$ ($n = 4$), respectively, as determined by densitometric scanning. In addition, the levels of $G\beta$ and $G\gamma$ were also determined by using antibodies RM/1 and SW/1 against $G\beta$ and $G\gamma$ (common), respectively; however, no differences in the amounts of immunodetectable $G\beta$ or $G\gamma$ protein were detected between the two groups (data not shown).

We extended our studies further to investigate if mRNA levels of G proteins change concomitantly with protein levels and determined the mRNA content of G proteins from hearts from control and L-NAME-treated rats using cDNA probes encoding for $G\alpha_2$, $G\alpha_3$, and $G\alpha$. The results depicted in Fig. 2 show that the $G\alpha_2$ and $G\alpha_3$ probes detected a message of 2.3 and 3.5 kb, respectively, in both control and L-NAME-treated heart; however, the $G\alpha_2$ and $G\alpha_3$ mRNA levels were increased by $40 \pm 3.6\%$ ($n = 4$) and $66 \pm 4.2\%$ ($n = 4$), respectively, in hearts from L-NAME-treated rats as compared with their control rats as determined by densitometric scanning. On the other hand, the cDNA probe encoding $G\alpha$ detected a message of 1.8 kb in both control and L-NAME-treated rats; however, no difference in $G\alpha$ mRNA levels was detected between the groups (data not shown). The alterations in $G\alpha_2$ and $G\alpha_3$ mRNA in L-NAME-treated hearts as compared with control rats may not be attributed to the variation in the amounts of total RNA applied to the gels, because hybridization with 32-mer oligonucleotide, which recognizes a highly conserved region of 28S rRNA, showed a similar amount of 28S rRNA loaded from control and L-NAME-treated rats onto the gels (data not shown).

Effect of L-NAME treatment on Gi protein functions

To investigate if the enhanced expression of Gi proteins was reflected in an increased function of Gi proteins, the ef-

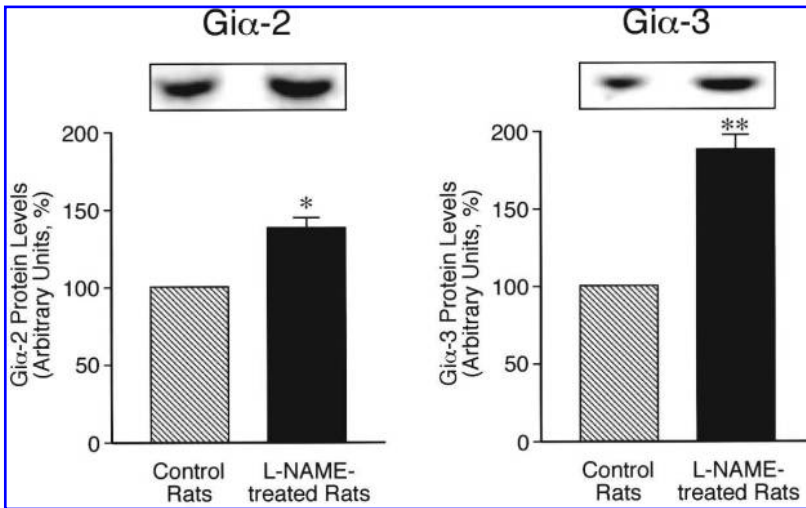


FIG. 1. Quantification of G proteins by immunoblotting in hearts from control and L-NAME-treated rats. (Upper panel) The membrane proteins (50 μ g) from control and L-NAME-treated rats were separated by SDS-PAGE electrophoresis and transferred to nitrocellulose, which was then immunoblotted using antibody AS/7 for Gi α -1 and Gi α -2 and antibody EC/2 for Gi α -3 as described in Materials and Methods. The autoradiogram is representative of four separate experiments. The detection of different G proteins was performed by using enhanced chemiluminescence western blotting detection reagents. **(Lower panel)** Densitometric scanning of G proteins from control and L-NAME-treated rats. The results are expressed as arbitrary units and values from control rats were taken as 100%. Values are means \pm SEM of four separate experiments. * p < 0.05.

fect of low concentrations of GTP γ S on FSK-stimulated adenylyl cyclase activity was examined, and the results are shown in Fig. 3. GTP γ S inhibited FSK-stimulated adenylyl cyclase activity in a concentration-dependent manner in both groups; however, the inhibition of FSK-stimulated adenylyl cyclase by GTP γ S was significantly enhanced in L-NAME-treated rats as compared with their control rats. For example, GTP γ S at 1 nM inhibited FSK-stimulated enzyme activity by ~20% in control rats and ~50% in L-NAME-treated rats.

As Gi proteins couple the inhibitory hormone receptors to adenylyl cyclase and mediate the inhibitory responses of hormones on adenylyl cyclase activity, it was interesting to determine if enhanced expression and functions of Gi α were also associated with enhanced inhibition of adenylyl cyclase by inhibitory hormones. To explore this, the effects of inhibitory hormones on adenylyl cyclase activity were examined in hearts from control and L-NAME-treated rats. Figure 4 shows that oxotremorine, C-ANP₄₋₂₃ (a ring-deleted analogue of atrial natriuretic peptide), and angiotensin II (Ang II) inhibited ade-

nylyl cyclase activity to various degrees in control rats, but the inhibition was completely attenuated in L-NAME-treated rats.

Effect of L-NAME treatment on Gs protein functions

To corroborate our results of Gs α levels with Gs α function, we analyzed the effect of CT treatment on GTP or GTP γ S-sensitive adenylyl cyclase in heart membranes from both control and L-NAME-treated rats. CT-treatment augmented GTP or GTP γ S-sensitive adenylyl cyclase activity in both groups, however, the percentage stimulation was not significantly different in L-NAME-treated rats as compared with control rats (data not shown), indicating that Gs α functions were also not altered in L-NAME-treated rats. As Gi α has been shown to modulate Gs α functions (11, 14), it was of interest to examine if increased Gi α levels in L-NAME-treated rats could affect Gs α -mediated hormonal stimulations of adenylyl cyclase. As shown in Table 1, isoproterenol, *N*-ethylcarboxamido adenosine (NECA), and glucagon stimulated adenylyl cyclase ac-

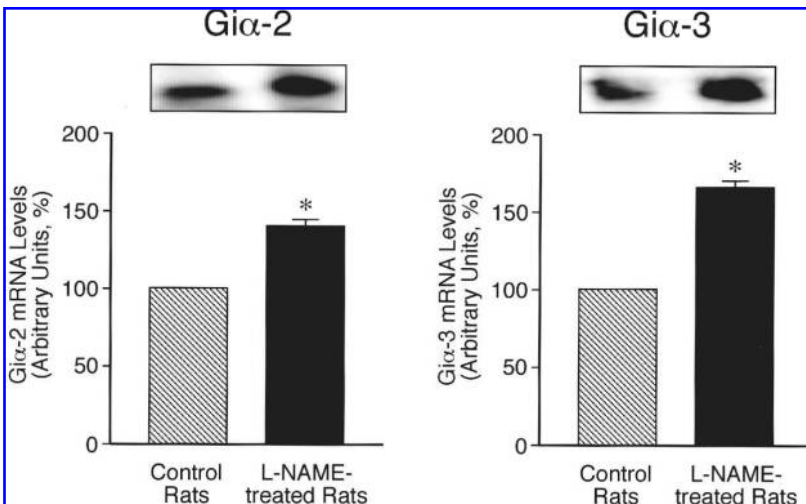


FIG. 2. mRNA expression of G proteins by northern blot in hearts from control and L-NAME hypertensive rats. (Upper panel) Total RNA (10 μ g) extracted from heart from control and L-NAME hypertensive rats was separated on 1% agarose and transferred to nylon membranes, which was then hybridized with full-length cDNA probe encoding Gi α -2 and Gi α -3 as described in Materials and Methods. The autoradiogram is representative of four separate experiments. **(Lower panel)** Densitometric scanning of G-protein mRNA from control and L-NAME-treated rats. The results are expressed as arbitrary units, and values from control rats were taken as 100%. Values are means \pm SEM of four separate experiments. * p < 0.05.

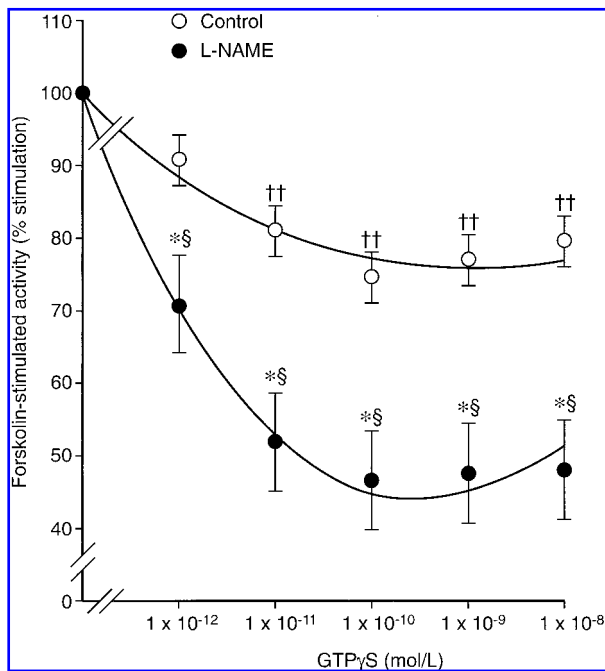


FIG. 3. Effect of GTP γ S on FSK-stimulated adenylyl cyclase activity in heart membranes from control and L-NAME-treated rats. Adenylyl cyclase activity in the presence of 100 μ M FSK was determined in the absence (basal) or presence of increasing concentrations of GTP γ S in control (○) and L-NAME hypertensive rats (●) as described in Materials and Methods. The values are means \pm SEM of three separate experiments. Adenylyl cyclase activities in the presence of 100 μ M FSK in control and L-NAME-treated rats in the absence of GTP γ S were $2,934.5 \pm 84.8$ and $3,072.5 \pm 37.7$ pmol of cAMP/mg of protein/10 min, respectively. Statistical analysis was performed by ANOVA followed by Dunnett's test for comparison between basal and all other doses of GTP γ S ($\$p < 0.01$, $\dagger\dagger p < 0.05$), whereas Newman-Keuls test was used for comparison between control and L-NAME-treated rats ($*p < 0.001$).

tivity to various degrees in heart membranes from both groups; however, the extent of stimulation of adenylyl cyclase was significantly diminished in L-NAME-treated rats as compared with control rats. In addition, FSK- and NaF-stimulated

adenylyl cyclase activities were also significantly diminished in L-NAME-treated rats as compared with control rats. To investigate if the decreased stimulation of adenylyl cyclase by FSK in L-NAME-treated rats is attributed to the decreased levels of catalytic subunit, the mRNA levels of type V enzyme were determined in hearts using a cDNA probe encoding type V enzyme. However, no difference in the amount of mRNA type V was observed between the two groups as quantified by densitometric scanning (data not shown).

DISCUSSION

In the present studies, we have shown that inhibition of NO levels by L-NAME treatment of rats for 4 weeks increases the levels of Gi proteins, as well as blood pressure.

The hearts from L-NAME-treated rats exhibited an increased expression of Gi proteins as compared with control rats as demonstrated by immunoblotting studies. These data suggest that the reduction of NO levels by L-NAME that results in the augmentation of Gi proteins may be responsible for the enhanced blood pressure in L-NAME-treated rats. This notion is supported by our earlier studies showing the implication of enhanced expression of Gi proteins and associated functions in the pathogenesis of hypertension (31, 34, 35). However, L-NAME treatment of rats did not affect the levels of Gs proteins, as well as Gs mRNA. In addition, the functions of Gs were not altered by L-NAME treatment, suggesting that NO may not play a role in the modulation of Gs protein expression and Gs-mediated functions.

The increased expression of Gi α -2 and Gi α -3 in L-NAME-treated rats was also reflected in Gi α functions as was shown by the inhibition of FSK-stimulated adenylyl cyclase activity by GTP γ S, which was greater in L-NAME-treated rats as compared with control rats. These results are in agreement with the previous studies reported in SHR and other models of hypertension showing a relationship between enhanced expression of Gi proteins and Gi functions (2, 17, 34, 35). However, a complete attenuation of adenylyl cyclase inhibition by inhibitory hormones such as oxotremorine, C-ANP₄₋₂₃, and Ang II in hearts from L-NAME-treated rats is in contrast with other studies conducted in genetic and experimental hypertensive rats, where a correlation between the enhanced levels of Gi

FIG. 4. Effect of inhibitory hormones on adenylyl cyclase activity in heart membranes from control and L-NAME-treated rats. Adenylyl cyclase was determined in the presence of GTP γ S alone (basal) or in combination with 10 μ M Ang II, 50 μ M oxotremorine (OXO), or 0.1 μ M C-ANP₄₋₂₃ as described in Materials and Methods. The values are means \pm SEM of three separate experiments. The basal adenylyl cyclase activities in control and L-NAME-treated rats in the presence of 10 μ M GTP γ S were 481.2 ± 10.4 and 460.5 ± 11.2 pmol of cAMP/mg of protein/10 min, respectively. $*p < 0.05$, $**p < 0.01$.

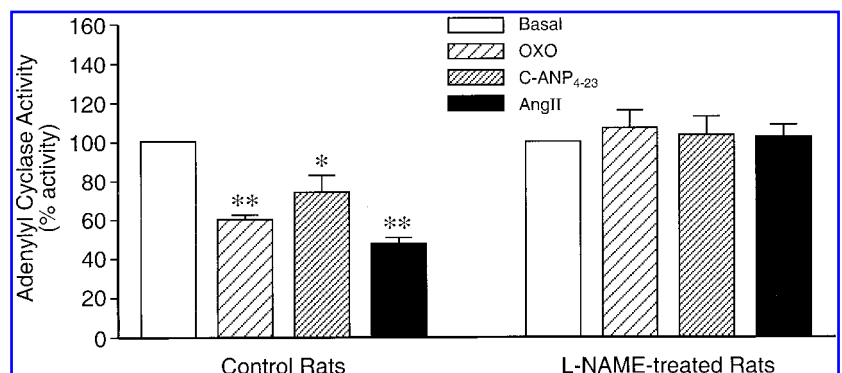


TABLE 1. EFFECT OF SOME AGONISTS ON ADENYLYL CYCLASE ACTIVITY IN HEART MEMBRANES FROM CONTROL AND L-NAME-TREATED RATS

Addition	Adenylyl cyclase activity (pmol of cAMP/mg of protein/10 min)	
	Control rats (% stimulation)	L-NAME-treated rats (% stimulation)
None	32.4 ± 2.4	34.2 ± 2.8
GTP	66.6 ± 1.4	68.3 ± 2.8
GTP + ISO	373.0 ± 33.4 (460)	293.7 ± 22.5* (330)
GTP + GLU	266.4 ± 26.4 (300)	218.6 ± 6.83* (220)
GTP + NECA	326.3 ± 26.6 (390)	266.4 ± 17.1* (290)
FSK	2,332.8 ± 194.4 (7,200)	1,949.4 ± 239* (5,800)
NaF	421.2 ± 36.5 (1,300)	324.9 ± 30.8† (950)

Adenylyl cyclase activity in heart membranes from control and L-NAME-treated rats was determined in the absence or presence of 50 μ M FSK, 10 mM NaF, or 10 μ M GTP alone or in combination with 50 μ M isoproterenol (ISO), 1 μ M glucagon (GLU), or 10 μ M NECA as described in Materials and Methods. The values are means \pm SEM of three separate experiments.

* $p < 0.05$.

† $p < 0.01$.

protein and functions has been reported (17, 34, 35). However, the reason for the lack of correlation between the enhanced levels of Gi proteins and receptor-dependent Gi functions in the present studies is not clear. It may be possible that the hormone receptors are down-regulated by increased activity of the renin-angiotensin system induced by L-NAME treatment (43). In this regard, we have previously shown an attenuation of C-ANP₄₋₂₃ and Ang II-mediated inhibition of adenylyl cyclase by Ang II treatment in A10 smooth muscle cells (40).

We have also shown that L-NAME treatment of rats resulted in decreased Gs α -mediated stimulations of adenylyl cyclase by NECA, glucagon, and isoproterenol as compared with control rats, which may be attributed to down-regulation of hormone receptors (32) or decreased levels of Gs α . However, L-NAME treatment did not alter the levels of Gs α protein and functions, suggesting that Gs α may not be responsible for the observed attenuated responsiveness of adenylyl cyclase to stimulatory hormones. On the other hand, the levels of plasma catecholamines have been shown to be enhanced by L-NAME treatment (54), which may be responsible for the desensitization of β -adrenergic receptors in L-NAME-treated rats. However, Laflamme *et al.* (29) did not observe any changes in density and affinity of cardiac β -adrenergic receptors in L-NAME-induced hypertensive rats as compared with sham-operated control rats. The modulation of Gs α functions by Gi α has been reported by several investigators (11, 14). An increased expression of Gi α -2 and Gi α -3 has been shown to be associated with attenuated responsiveness of adenylyl cyclase to stimulatory hormones (2, 34, 35), whereas a decreased expression of Gi α -2 resulted in the augmentation of stimulatory hormones on adenylyl cyclase (3, 4). Taken together, it may be possible that the enhanced levels of Gi α -2 and Gi α -3 in hearts by L-NAME treatment may be responsible for the diminished sensitivity of adenylyl cyclase to isoproterenol, NECA, and glucagon stimulation.

The decreased stimulation of adenylyl cyclase by FSK in L-NAME-treated rats may be due to the impaired catalytic subunit of adenylyl cyclase or to the overexpression of Gi α or

both. However, as the mRNA expression of type V enzyme was not decreased in hearts by L-NAME treatment and the levels of type VI enzyme were not determined, the overexpression of Gi α -2 and Gi α -3 proteins induced by L-NAME and type VI enzyme that may be impaired by L-NAME treatment may be responsible for the decreased sensitivity of adenylyl cyclase to FSK stimulation. In this regard, the role of Gi α in FSK-mediated stimulation for adenylyl cyclase has been reported (5). In addition, the requirement of Gs α and guanine nucleotides for FSK activation of adenylyl cyclase has also been shown (21). As the levels and functions of Gs α were not altered in hearts by L-NAME treatment, the role of Gs α in eliciting decreased FSK stimulation of adenylyl cyclase cannot be attributed to the impaired Gs α .

In conclusion, we have demonstrated that L-NAME treatment of rats that results in decreased levels of NO increased the expression of Gi α -2 and Gi α -3 genes and translated proteins, whereas the levels and functions of Gs α were not altered by such treatment.

NO has also been shown to play a crucial role as mediator of ischemic preconditioning (18). In this regard, a natural antioxidant, resveratrol, found in grapes and wine has been reported to protect ischemic myocardium through NO, because L-NAME, which inhibits the production of NO, abolished resveratrol-mediated cardioprotective effects (19). Furthermore, resveratrol was also unable to precondition hearts from inducible NO synthase knockout mice (22). A role of Gi proteins in ischemic preconditioning has also been reported (41). Taken together, it may be suggested that NO, by modulating the expression of Gi α proteins, may be responsible for ischemia preconditioning.

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ABBREVIATIONS

Ang II, angiotensin II; cAMP, cyclic AMP; C-ANP₄₋₂₃, a ring-deleted analogue of atrial natriuretic peptide, ANP (des-[Gln¹⁸,Ser¹⁹,Gln²⁰,Leu²¹,Gly²²]ANP₄₋₂₃-NH₂); CT, cholera toxin; FSK, forskolin; G protein, guanine nucleotide regulatory protein; Gi, inhibitory guanine nucleotide regulatory protein; Gs, stimulatory guanine nucleotide regulatory protein; GTPγS, guanosine 5'-(γ-thio)triphosphate; L-NAME, N^ω-nitro-L-arginine methyl ester, NECA, N-ethylcarboxamido adenosine; NO, nitric oxide; PBS, phosphate-buffered saline; SDS-PAGE, sodium dodecyl sulfate–polyacrylamide gel electrophoresis; SHR, spontaneously hypertensive rats.

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