# REGULATION OF INOSITOL 1,4,5-TRISPHOSPHATE METABOLISM BY GUANINE NUCLEOTIDES IN MEMBRANES OF CULTURED NEWBORN RAT CARDIOMYOCYTES

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Abstract—Membranes of cultured newborn rat cardiomyocytes contain enzymatic activities that regulate the formation and the breakdown of inositol 1,4,5-trisphosphate (1,4,5-IP<sub>3</sub>). GTP $\gamma$ S increased the rate of exogenous [³H]phosphatidyl 4,5-bisphosphate ([³H]PIP<sub>2</sub>) hydrolysis (EC<sub>50</sub>: 40  $\mu$ M). This effect was dependent on the presence of deoxycholate and maximal at 2 mM deoxycholate. GTP $\gamma$ S increased the efficacy of phospholipase C (PLC) (by 2.3-fold), but did not alter the apparent affinity of the enzyme for PIP<sub>2</sub>. Other nucleotides, GDP $\beta$ S and ATP $\gamma$ S, and pyrophosphate also stimulated PIP<sub>2</sub> hydrolysis, while AIF<sub>4</sub> was ineffective. The effect of GTP $\gamma$ S was not inhibited by GDP $\beta$ S. The agonists nor-epinephrine and thrombin, which by themselves had no effect, did not potentiate the response to GTP $\gamma$ S. In contrast, 1,4,5-IP<sub>3</sub> hydrolysis was decreased by GTP $\gamma$ S (EC<sub>50</sub>: 100  $\mu$ M) as well as by other nucleotides and by pyrophosphate, but not by AIF<sub>4</sub>. GDP $\beta$ S did not antagonize the GTP $\gamma$ S-induced inhibition of IP<sub>3</sub> hydrolysis. These results suggest that GTP can stimulate the hydrolysis of exogenous PIP<sub>2</sub> by an action on membrane-bound PLC at a site beyond the G protein activating PLC and inhibit the hydrolysis of 1,4,5-IP<sub>3</sub> by a mechanism common to all nucleotides. Thus, GTP can regulate 1,4,5-IP<sub>3</sub> metabolism by stimulating its formation and inhibiting its breakdown.

Phosphoinositides are membrane-bound phospholipids that are rapidly hydrolyzed by phospholipase C (PLC||) to inositol phosphates and diacylglycerol in response to a variety of hormones and transmitters [1]. Inositol 1,4,5-trisphosphate (1,4,5-IP<sub>3</sub>), the major physiologically active inositol phosphate, mediates the release of calcium from nonmitochondrial stores. The phosphoinositides and the inositol phosphates are metabolically closely related. They coexist in a dynamic equilibrium which is maintained by a complex system of specific kinases and phosphatases [2]. Kinetic studies in many intact tissues and isolated cells have provided indirect evidence that stimulation of phosphatidyl 4,5bisphosphate (PIP2) hydrolysis by PLC is the preferred pathway for agonist-induced formation of 1,4,5-IP<sub>3</sub>[1]. The agonist-induced activation of PLC, like adenylate cyclase activation, has been shown in a number of tissues to be dependent upon guanine nucleotides, implying that a guanine nucleotide binding protein (G protein) is involved in the transduction of the hormonal signal from the receptor binding site to PLC [3, 4]. This has been confirmed recently by Smrcka et al. [5] who demonstrated that a G protein  $\alpha$  subunit purified from bovine brain selectively stimulates PIP<sub>2</sub> hydrolysis by a partially purified PLC.

The recent studies of Steinberg et al. [6] show that, in intact cardiomyocytes, norepinephrine-induced stimulation of phosphoinositide hydrolysis results in the rapid formation of 1,4,5-IP<sub>3</sub>. In addition, in myocyte membranes, the  $\alpha_1$ -adrenergic stimulation of inositol phosphate formation from endogenous phosphoinositides is dependent upon the presence of a GTP analogue, an indication that the receptor is coupled to PLC by a G protein [6]. Those studies, however, did not directly demonstrate a role for a guanine nucleotide in the hydrolysis of PIP<sub>2</sub> to 1,4,5-IP<sub>3</sub>. Furthermore, Coleman and Bilezikian [7] have shown recently that, in renal cortical membranes, guanine nucleotides also increase the level of 1,4,5-IP<sub>3</sub> by inhibiting its breakdown.

In the present study, we have used exogenous radiolabeled [3H]PIP<sub>2</sub> and [3H]1,4,5-IP<sub>3</sub> to explore whether guanine nucleotides have a direct regulatory role on PIP<sub>2</sub> and 1,4,5-IP<sub>3</sub> hydrolysis. Cardiomyocyte membranes from newborn rats served as the source of PLC and phosphatase. The results of our study show that guanine nucleotides can modulate 1,4,5-IP<sub>3</sub> levels by a dual action, stimulation of PIP<sub>2</sub> hydrolysis and inhibition of 1,4,5-IP<sub>3</sub> breakdown.

### **METHODS**

Chemicals and drugs. (-)Norepinephrine, GTP $\gamma$ S, GDP $\beta$ S, Gpp(NH)p, ATP, ATP $\gamma$ S, 2,3-diphosphoglycerate, hypoxanthine, and fibronectin were obtained from the Sigma Chemical Co. (St. Louis,

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<sup>||</sup> Abbreviations: PLC, phospholipase C; 1,4,5-IP<sub>3</sub>(IP<sub>3</sub>), inositol 1,4,5-trisphosphate; PIP<sub>2</sub>, phosphatidyl 4,5-bisphosphate; and DOC, deoxycholate.

MO); phosphatidylinositol 4,5-bisphosphate and 1,4,5-inositol trisphosphate from Boehringer Mannheim (Indianapolis, IN); pertussis toxin from the List Biological Co. (Campbell, CA); and  $[^3H]$ -phosphatidylinositol 4,5-bisphosphate and  $[^3H]$ -inositol 1,4,5-trisphosphate from Dupont-NEN (Boston, MA).  $\alpha$ -Thrombin was a gift from Dr. David M. Stern of the Department of Medicine at this institution. All other chemicals were reagent grade and were obtained from standard chemical suppliers.

Cardiomyocyte cultures and membrane preparation. Hearts from 2-day-old Wistar rats from Camm Research (Wayne, NJ) were removed under sterile conditions and placed in sterile salt solution. The atria were trimmed away and the ventricular cells were isolated according to a trypsin dispersion protocol described previously [8]. Briefly, isolated cells were pooled, centrifuged at 200 g for 5 min, and resuspended in Dulbecco's Minimum Essential Medium (MEM) supplemented with 10% horse serum, 5 µM hypoxanthine and 12 mM NaHCO<sub>3</sub>. The cells were preplated for 60 min at 37° to decrease fibroblast contamination. The muscle cells were then resuspended in MEM supplemented with 10% horse serum and grown in 100 mm plates, which had been coated previously with fibronectin (25 ng/mm<sup>2</sup>) for 45 min. For some experiments, myocytes were pretreated with pertussis toxin (100 ng/mL) for the last 24 hr in culture. This treatment has been shown previously to be associated with complete ADPribosylation of the pertussis toxin sensitive G protein in cardiomyocytes [9].

After 3-4 days of culture, the cells were detached and suspended in buffer containing 250 mM sucrose, 0.1 mM phenylmethylsulfonyl fluoride, 20 mM Tris-HCl, pH 7.5, homogenized and centrifuged at 43,000 g at 4° for 30 min. The pellet was taken up in homogenization buffer, and recentrifuged at the same speed. The final pellet was resuspended in homogenization buffer at a protein concentration of ~2 mg/mL, and stored in aliquots at -70°. Protein concentration was determined by the modified method of Lowry et al. [10].

Since PLC and phosphatase activities are present in both the membrane and the cytosol [11, 12], the activity of lactate dehydrogenase, a cytosolic marker, was measured in four membrane suspensions, with the Yellow Spring Instrument WSY-model 27. Negligible amounts of lactate dehydrogenase were found, indicating that the activities measured in our experiments were from membrane-associated PLC and phosphatases.

 $PIP_2$  hydrolysis assay. The reaction mixture consisted, unless otherwise indicated, of [ $^3H$ ]PIP $_2$ ,  $8\,\mu$ M (0.01 mCi); MgCl $_2$ , 0.5 mM; KCl, 25 mM; LiCl, 25 mM; 2,3-diphosphoglycerate, 1 mM; sodium deoxycholate, 2 mM; sodium cholate, 2 mM; the agents under study; the membrane suspension containing  $2\,\mu$ g of protein; sodium phosphate buffer, 100 mM, pH 7.0, in a total volume of  $50\,\mu$ L. After the addition of the membrane suspension, the reaction mixture was kept at 4° for 10 min. After this preincubation, the reaction proceeded at 30° for 5 min unless otherwise stated, and was terminated by the addition of 1.5 mL of ice-cold acidified

chloroform-methanol (1:2, v/v), followed by 0.5 mL chloroform and 1 mL distilled water to separate the aqueous and organic phase. Before use, PIP2 was dried under a stream of  $N_2$ , resuspended by sonication (2 × 30 sec) in 2 mM sodium deoxycholate and 2 mM sodium cholate in 100 mM phosphate buffer, pH 7.0. Lithium and 2,3-diphosphoglycerate were present in the reaction mixture to prevent the breakdown of IP3 by phosphatases [11, 13]. IP3, the product of PIP2 hydrolysis, was used as an index of PIP2 hydrolysis. An assay was performed in the absence of membrane (blank) to estimate the nonenzymatic hydrolysis of the substrate (PIP2) during the time of incubation. These blank values, <0.5% of total [3H]PIP2, were subtracted from the values obtained in the presence of membrane.

IP<sub>3</sub> hydrolysis assay. The reaction mixture was the same as for PIP<sub>2</sub> hydrolysis, except that 20 nM [<sup>3</sup>H]1,4,5-IP<sub>3</sub> (about 10,000 cpm) was included as substrate instead of [<sup>3</sup>H]PIP<sub>2</sub>; lithium and 2,3-diphosphoglycerate were omitted; 20 μg protein was routinely used per assay. The reaction proceeded at 30° for 20 min unless otherwise stated. Loss of IP<sub>3</sub> was used as index of IP<sub>3</sub> hydrolysis. An assay was also performed in the absence of membrane. In these conditions there was no significant loss of [<sup>3</sup>H]-IP<sub>3</sub>, indicating the lack of non-enzymatic hydrolysis during the time of incubation. This value was used as 100% to calculate percent hydrolyzed.

Identification of inositol phosphates. Inositol phosphates in the aqueous phase were separated and identified by anion exchange chromatography, as described by Berridge et al. [14]. The radioactivity of the eluate fractions, corresponding to [<sup>3</sup>H]IP<sub>1</sub>, -IP<sub>2</sub>, -IP<sub>3</sub> and -IP<sub>4</sub>, was determined by liquid scintillation spectrometry, using Aquasol-II as scintillant.

Data analysis. Triplicate determinations were performed with each membrane preparation in a single experiment. The membranes for repeated experiments were obtained from different cultures. Statistical differences were determined using ANOVA for repeated measures and, when the F value indicated significance, Student's paired t-tests with Bonferroni correction for multiple comparisons; P values <0.05 were considered significant.  $K_D$  and  $V_{\text{max}}$  values for each experiment were determined by linear regression [15].

## RESULTS

Effect of GTP analogues on PIP<sub>2</sub> hydrolysis. Membrane preparations of cultured cardiomyocytes from newborn rats contained an enzymatic activity that hydrolyzed [<sup>3</sup>H]PIP<sub>2</sub> (Fig. 1). There was no significant accumulation of IP<sub>2</sub>, IP<sub>1</sub>, or IP<sub>4</sub> over the 5-min time course of the assay. Thus, IP<sub>3</sub> formation reflected the hydrolysis of PIP<sub>2</sub> by PLC. The rate of IP<sub>3</sub> formation was linear with time up to 8 min (data not shown). The detergent deoxycholate (DOC) increased the rate of hydrolysis of PIP<sub>2</sub> with a maximum effect at 2 mM. At higher concentrations of DOC the rate of IP<sub>3</sub> formation declined and DOC became inhibitory (Fig. 1). Comparable biphasic effects of DOC on PIP<sub>2</sub> hydrolysis have been observed in other heart preparations [16, 17].

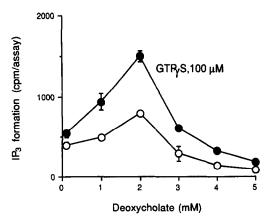


Fig. 1. Effect of sodium deoxycholate on basal- and GTPγS-stimulated PIP<sub>2</sub> hydrolysis. Cardiomyocyte membranes were incubated in the presence of increasing concentrations of sodium deoxycholate without (○) and with (●) GTPγS (100 μM) as described in Methods, using exogenous [³H]-PIP<sub>2</sub> as substrate. The results are expressed in cpm [³H]-IP3 formed per assay. Values are means ± SD of triplicate determinations from one of four experiments with membrane preparations from different cultures.

GTP $\gamma$ S, a phosphatase-resistant GTP analogue, did not increase significantly IP<sub>3</sub> formation in the presence of 0.2 mM DOC (+13 ± 14.3%, N = 4), but produced a gradual increase in IP<sub>3</sub> formation over the concentration range of DOC associated with stimulation, reaching a maximum at 2 mM (+57.9 ± 17.7%, N = 4,: P < 0.05) (Fig. 1). At inhibitory concentrations of DOC, the GTP $\gamma$ S response also declined. In all the subsequent experiments, DOC was used at 2 mM.

GTP $\gamma$ S produced a concentration-dependent increase in IP<sub>3</sub> formation; half-maximum activation occurred at 40  $\mu$ M and the maximum at 100  $\mu$ M (Fig. 2). Figure 3 shows the effect of 100  $\mu$ M GTP $\gamma$ S on PLC activity at increasing PIP<sub>2</sub> concentrations. Analysis of PIP<sub>2</sub> hydrolysis, according to Lineweaver-Burk, indicated that 100  $\mu$ M GTP $\gamma$ S significantly increased the apparent  $V_{\rm max}$  of PLC (2.3  $\pm$  0.25 fold, N = 4, P < 0.05), but did not alter the apparent affinity of the enzyme for PIP<sub>2</sub> ( $K_D$  11.1  $\pm$  2.4 vs 16.0  $\pm$  4.0  $\mu$ M in the absence and the presence of GTP $\gamma$ S, respectively).

Several other compounds were tested to evaluate the selectivity of the action of GTP $\gamma$ S for activation of the G protein coupled to PLC (Fig. 4). The analogue of GDP, GDP $\beta$ S, which competes with GTP and prevents G protein activation, by itself stimulated IP<sub>3</sub> formation and did not antagonize the increase induced by GTP $\gamma$ S, even when the membrane was preincubated with GDP $\beta$ S for 30 min before addition of GTP $\gamma$ S. ATP $\gamma$ S was about as effective as GTP $\gamma$ S. AIF $_4$ , which activates G proteins, did not increase IP<sub>3</sub> formation. Sodium pyrophosphate (1000  $\mu$ M) had a weak stimulatory effect (+37%).

Since GTP is an absolute requirement for agonist activation of G protein-mediated receptor-effector pathways, we next studied the effect of agonists on

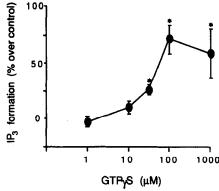


Fig. 2. Concentration-dependent effect of GTPγS on PIP<sub>2</sub> hydrolysis. Cardiomyocyte membranes were incubated in the absence and in the presence of increasing concentrations of GTPγS under standard assay conditions as described in Methods, using exogenous [<sup>3</sup>H]PIP<sub>2</sub> as substrate. Values, expressed in percent over control, i.e. IP<sub>3</sub> formed in the absence of GTPγS, are means ± SEM from eight experiments. Control value: 940 ± 110 cpm/assay. Key: (\*) significantly different from control (P < 0.05).

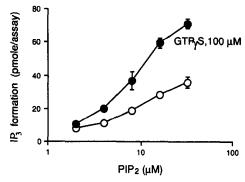


Fig. 3. PIP<sub>2</sub> hydrolysis as a function of substrate concentration and the effect of GTPγS. Cardiomyocyte membranes were incubated with increasing concentrations of exogenous PIP<sub>2</sub>, labeled with a fixed amount of [<sup>3</sup>H]-PIP<sub>2</sub>, in the absence (○) or in the presence (●) of GTPγS. Values are means ± SD of triplicate determinations from one of four experiments and are expressed in pmol IP<sub>3</sub> formed per assay, calculated from the specific activity of exogenous PIP<sub>2</sub>.

PIP<sub>2</sub> hydrolysis. Norepinephrine and thrombin, at concentrations which stimulate inositol phosphate accumulation in intact myocardial tissue [9, 18], did not increase IP<sub>3</sub> formation. GTPγS was not able to promote a response to either norepinephrine or thrombin (Fig. 4). The agonists were also ineffective in the presence of ATP, which was necessary in some preparations to demonstrate G protein activation and agonist potentiation of the guanine nucleotide response [19, 20]. Similarly, elimination of the 10-min preincubation period of membranes with detergent at 4°, which has been reported to decrease GTPγS and carbachol stimulation of PIP<sub>2</sub>

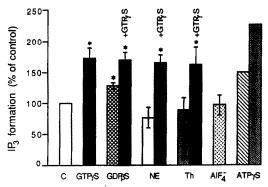
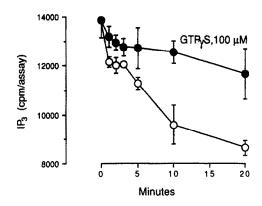


Fig. 4. Influence of nucleotides, agonists and fluoroaluminate on PIP2 hydrolysis. Cardiomyocyte membranes were incubated in the absence (C) or in the presence of the indicated test agents under standard assay conditions as described in Methods, using exogenous [3H]PIP2 as substrate. The concentration of GTPyS, alone and in combination, and of GDP $\beta$ S was 100  $\mu$ M; that of ATP $\gamma$ S was 100 µM (lighter shade) and 1000 µM (darker shade). AlF<sub>4</sub> consisted of a mixture of 10 mM sodium fluoride and 20 μM aluminum chloride. Abbreviations: NE, norepinephrine, 50 μM; Th, α-thrombin, one unit. Values, expressed as percent of corresponding control (C), are means ± SEM from three to six experiments, except for ATPyS where only two experiments were performed. The SD of the triplicates in each experiment with ATPyS was less than 12% at 100  $\mu$ M and 3% at 1000  $\mu$ M. The control values, expressed in IP<sub>3</sub> formed per assay, were: 1100 ± 190 cpm/assay for the GTPγS and GDPβS groups;  $1260 \pm 140$  for the norepinephrine and thrombin groups;  $1090 \pm 180$  for AlF<sub>4</sub>; and 930 for ATP $\gamma$ S. Key: (\*) significantly different from corresponding control (C) (P < 0.05).

hydrolysis [21], was also without effect (data not shown).

Several studies have suggested that PLC, like adenylate cyclase, is positively and negatively coupled to the agonist receptor by G proteins. Litosch [22] observed in rat cerebral cortical membranes that GTP analogues have a dual effect on exogenous PIP<sub>2</sub> hydrolysis, a stimulatory effect and a pertussis toxin-sensitive inhibitory effect. Since  $\alpha_1$ -adrenergic agonists produce a positive and a pertussis toxin-sensitive negative chronotropic response in cardiomyocytes [23], cells were pretreated with pertussis toxin to test whether GTPyS might also regulate cardiomyocyte PLC by a pertussis toxin-sensitive inhibitory mechanism. Pretreatment of cardiomyocytes with 100 ng/mL of pertussis toxin, a concentration sufficient to ADP-ribosylate virtually all accessible sites [9], decreased, though not significantly, both basal and GTPyS-stimulated PIP<sub>2</sub> hydrolysis. However, the response to GTP yS  $(1-100 \,\mu\text{M})$ , as a function of control activity, was similar in treated and untreated membranes (+54.3 and 51.7% at  $100 \,\mu\text{M}$  GTP $\gamma$ S, respectively). These results are in agreement with those obtained using membranes with endogenous phosphoinositide substrates [6] and indicate that the effect of GTPyS on PIP<sub>2</sub> hydrolysis does not involve a pertussis toxinsensitive inhibitory component.



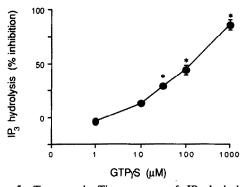


Fig. 5. Top panel: Time course of IP3 hydrolysis. Cardiomyocyte membranes were incubated in the absence (○) and in the presence of GTP<sub>V</sub>S (●) for the times indicated under standard assay conditions as described in Methods, using exogenous [3H]1,4,5-IP3 as substrate. The results are expressed in cpm [3H]IP3 remaining at the end of the incubation period. Values are means ± SD of triplicate determinations from one of two experiments. Bottom panel: Concentration-dependent effect of GTPyS on IP3 hydrolysis. Cardiomyocytes were incubated in the absence (control) and in the presence of increasing concentrations of GTPyS under standard assay conditions as described in Methods, using exogenous [3H]1,4,5-IP<sub>3</sub> as substrate. The results are expressed in percent inhibition of IP3 hydrolysis under control conditions, calculated from the equation  $(C-D)/C \times 100$ , where  $C = \text{cpm } [^3H]IP_3$ hydrolyzed under control conditions; and  $D = \text{cpm} [{}^{3}\text{H}]$ -IP<sub>3</sub> hydrolyzed in the presence of GTP $\gamma$ S. Control: 5190 ± 730 cpm [ $^3$ H]1,4,5-IP<sub>3</sub> hydrolyzed in 20 min, corresponding to 47% of [3H]1,4,5-IP3 incubated in the absence of membranes. Values are means ± SEM from five experiments. Key: (\*) significantly different from control (P < 0.05).

Effect of GTP analogues on  $IP_3$  hydrolysis. In view of the results of a recent study suggesting that guanine nucleotides may also increase  $IP_3$  levels by inhibiting the rate of  $IP_3$  hydrolysis [7], the next set of experiments was aimed at defining a role for GTP in  $IP_3$  hydrolysis. The rate of 1,4,5- $IP_3$  hydrolysis in myocyte membranes was very slow. At the protein concentration (2  $\mu$ g) used in the  $PIP_2$  assay, there was no detectable activity. This observation is consistent with the lack of accumulation of  $IP_2$  and  $IP_1$  in the  $PIP_2$  assay. At a higher protein concentration (20  $\mu$ g), 18%  $IP_3$  was hydrolyzed in 5 min and 34% after 20 min (P < 0.05) (Fig. 5, top

Table 1. Effect of nucleotide analogues, aluminum fluorate and sodium pyrophosphate on 1,4,5-IP<sub>3</sub> hydrolysis

Test agent	% Inhibition
GTPγS, 100 μM	49.6 ± 3.1*
GTP $\gamma$ S, 1000 $\mu$ M	93.7
$Gpp(NH)p$ , $100 \mu M$	$23.4 \pm 1.8*$
$Gpp(NH)p$ , $1000 \mu M$	$72.0 \pm 4.2*$
GDPβS, 100 μM	$31.7 \pm 3.2*$
GDPβS, 1000 μM	59.5
GDP $\beta$ S, 100 $\mu$ M + GTP $\gamma$ S, 100 $\mu$ M	$50.1 \pm 3.2*$
GDP $\beta$ S, 1000 $\mu$ M + GTP $\gamma$ S, 100 $\mu$ M	51.4
ATPγS, 100 μM	42.9
ATP $\gamma$ S, 1000 $\mu$ M	$92.6 \pm 13.2*$
AlF <sub>4</sub> †	$0.3 \pm 13.2$
Sodium pyrophosphate, 100 µM	$15.7 \pm 4.1^*$

Cardiomyocyte membranes were incubated in the absence (control) or in the presence of a test agent under the standard assay conditions as described in Methods, using exogenous  $[^3H]1,4,5\text{-}IP_3$  as substrate. The results are expressed in percent inhibition of  $IP_3$  hydrolysis under control conditions, calculated from the equation:  $(C-D)/C \times 100$ , where  $C = \text{cpm} [^3H]IP_3$  hydrolyzed under control conditions; and  $D = \text{cpm} [^3H]IP_3$  hydrolyzed in the presence of a test agent. The average control value was  $2340 \pm 460$  (SEM) cpm  $[^3H]1,4,5\text{-}IP_3$  hydrolyzed in 20 min, corresponding to 40% of  $[^3H]1,4,5\text{-}IP_3$  incubated in the absence of membranes. Values are from 2 to 4 experiments; when 3-4 experiments were performed, values are means  $\pm$  SEM; when only 2 experiments were done, no SEM is shown (the SD of the triplicates in each experiment ranged from 0.2 to 5%).

\* Significantly different from corresponding control (P < 0.05).

† AlF $_4$  consisted of a mixture of 10 mM sodium fluoride and 20  $\mu$ M aluminum chloride.

panel). In the following experiments, the assay contained  $20 \mu g$  of protein and proceeded for 20 min. There was no significant accumulation of  $IP_4$ ,  $IP_1$  or inositol during the incubation. After correction for recovery, the combined amounts of  $[^3H]IP_3$  and  $[^3H]IP_2$  at the end of incubation were similar to the total amount of  $[^3H]IP_3$  (data not shown). This indicated that the decrease in  $IP_3$  reflects mainly phosphatase activity

GTP \( \gamma \) reduced the rate of 1,4,5-IP3 hydrolysis (Fig. 5, top panel). The effect was concentration dependent between 10 and 1000 µM, with halfmaximum inhibition at  $100 \,\mu\text{M}$  (Fig. 5, bottom panel). Other nucleotides, AlF4 and sodium pyrophosphate were tested for their effect on IP<sub>3</sub> hydrolysis to evaluate the selectivity of the GTP effect (Table 1). Gpp(NH)p, another GTP analogue, as well as GDP $\beta$ S, inhibited IP<sub>3</sub> hydrolysis, though less effectively than GTP<sub>2</sub>S. When the membranes were preincubated with GDP $\beta$ S before the addition of GTP<sub>2</sub>S, the rate of IP<sub>3</sub> hydrolysis was similar to that of GTPyS alone. The adenosine derivative, ATPyS, was about equieffective as GTPyS as an inhibitor of IP3 hydrolysis. AlF4 had no effect, while sodium pyrophosphate, at the same concentration as GTPyS, produced a small, though significant, inhibitory effect on IP3 hydrolysis. Norepinephrine, 50 and 500  $\mu$ M, alone and in the presence of 100  $\mu$ M GTP $\gamma$ S, had no effect on IP<sub>3</sub> hydrolysis (data not shown).

#### DISCUSSION

The results of this study demonstrate the presence in cardiomyocyte membranes from newborn rats of GTP-sensitive enzymatic activities that regulate the formation and the hydrolysis of 1,4,5-IP<sub>3</sub>. The stimulatory effect of the GTP analogue GTP \( \gamma \) on the hydrolysis of exogenous PIP<sub>2</sub> confirms the results of earlier studies with intact cells and with prelabeled membranes using endogenous PIP<sub>2</sub> [6]. However, the mechanism of GTP stimulation may be different when an exogenous substrate is used. stimulation of IP<sub>3</sub> formation from endogenous substrate has been attributed to activation of the G protein regulating phosphoinositide hydrolysis. In our preparation GTPyS appears to have a broader role. The GTPyS response was not specific, as GDP and ATP analogues were also stimulatory. The concentrations of GTPyS stimulating PLC far exceeded those for activation of G protein-dependent processes. Furthermore, GDP analogues, which compete with GTP and prevent the activation of the G protein, did not antagonize the stimulation of GTP $\gamma$ S. AlF $_4$ , a potent activator of G proteins, was ineffective. Moreover, norepinephrine and thrombin in the presence of GTPyS, which is an absolute requirement for agonist stimulation of G proteinmediated receptor pathways, had no effect on PIP<sub>2</sub> hydrolysis.

The reason for the difference in the mechanism of GTP activation of endogenous and exogenous PIP<sub>2</sub> hydrolysis is not clear. The need to treat the membranes with a detergent to observe a GTP effect suggests that the accessibility and/or the presentation of the substrate PIP<sub>2</sub> to its binding sites is critical to demonstrate R-G protein-activation of PLC. Even though the detergent deoxycholate facilitates the stimulation of PLC by GTP 7S, it did not promote agonist stimulation, as was the case in brain membranes [21]. It is possible that disruption of the membrane structure by the detergent, while facilitating access of the substrate to PLC, dissociates PLC from the R-G protein complex. In a detailed study comparing the effects of several detergents on the activation of PLC by exogenous PIP2, Carter et al. [21] showed, in rabbit cortical membranes, that stimulation by GTPyS and carbachol required deoxycholate and was maximal at 1 mM. From their results, and those obtained in other similar studies, they concluded that the concentration of detergents that maintain a functionally active R-G protein-PLC complex and that which uncouples the complex varies among tissues and is critical to demonstrate R-G protein-activation of PLC. Future studies will explore whether with other detergents agonist activation of exogenous PIP<sub>2</sub> hydrolysis can be observed in cardiomyocytes.

The lack of nucleotide selectivity observed in our preparation suggests additional mechanisms to G protein activation for GTP stimulation of PIP<sub>2</sub> hydrolysis. Non-specific stimulation of exogenous PIP<sub>2</sub> hydrolysis by nucleotides has also been observed in other preparations and appears to be related to

activation of PLC or an associated protein downstream of the G protein [24, 25]. Ryu et al. [26] have shown that ATP and GTP activate a purified brain PLC. The stimulatory effect of sodium pyrophosphate suggests that the nucleotide effect may be due to a phosphorylation reaction. In rabbit thymocyte membranes [24], hydrolysis of endogenous PIP<sub>2</sub> was selectively stimulated by GTP<sub>2</sub>S, while that of exogenous PIP2 was stimulated by GTP- as well as GDP- and ATP-analogues. In contrast, in permeabilized 3T3 cells, which have a functionally activatable G protein, ATPyS stimulated the hydrolysis of endogenous PIP<sub>2</sub> [25]. These observations suggest that the GTP sensitivity of the processes involved in PLC activation may depend, in part, upon the structural properties of the membrane components of the pathway, as well as their orientation with the plane of the membrane.

The results of our study demonstrate that GTP analogues can also regulate 1,4,5-IP<sub>3</sub> hydrolysis. The rate of IP<sub>3</sub> hydrolysis, however, was much slower than its formation by PLC. This apparently low phosphatase activity probably reflects a small amount of membrane-bound enzyme, since the amount of protein present in the PIP<sub>2</sub> assay had to be increased in order to observe IP3 hydrolyzing activity. It has been reported that in canine renal cortical membranes or in coronary artery membranes, 80 and 20% [3H]-IP<sub>3</sub>, respectively, was hydrolyzed in 1 min [7, 12]. Even though it is difficult to compare these preparations because of differences in the assay conditions and the unknown endogenous IP<sub>3</sub> concentration, it appears that the membrane bound phosphatase activity which hydrolyzes 1,4,5-IP<sub>3</sub> varies greatly from tissue to tissue.

GTPyS had an inhibitory effect on IP<sub>3</sub> hydrolysis. The effect is not selective for  $GTP\gamma S$ , since other nucleotides were also inhibitory. The specificity of the effect appears to be in the phosphate group of the nucleotide, suggesting a phosphorylating mechanism. IP3 hydrolysis does not appear to be regulated by norepinephrine, since the agonist had no significant effect in the absence or in the presence of GTP\(gamma S\). This could indicate that the phosphatase hydrolyzing 1,4,5-IP<sub>3</sub> is not linked to the  $\alpha_1$ adrenergic receptor. However, this possibility cannot be ruled out completely since in the same membrane preparations the  $\alpha_1$ -adrenergic receptor uncoupled from PLC. Our results are comparable to those obtained in renal cortical membranes [7]. In this preparation, nucleotides inhibited 1,4,5-IP<sub>3</sub> formation and showed specificity best associated with the phosphate group of the nucleotide, GTPyS and ATPyS being most potent. In addition, parathyroid hormone, which stimulates phosphoinositide hydrolysis, has no effect on 1,4,5-IP<sub>3</sub> hydrolysis in the absence or in the presence of GTPyS [7]. Furthermore, in cardiomyocyte membranes the characteristics of the effect of GTPyS on the phosphatase and PLC are similar, suggesting that the nucleotide regulates the activity of both enzymes by a similar mechanism.

The results of our study indicate that, besides its action on the G protein mediating PLC activation observed in preparations using endogenous PIP<sub>2</sub>, GTP can modulate phosphoinositide metabolism by

actions at multiple sites along this pathway. Our data suggest that GTP can regulate 1,4,5-IP<sub>3</sub> at the level of membrane-bound PLC and 5'-phosphatase, stimulating its formation from PIP<sub>2</sub> and inhibiting its breakdown. Other possible sites of control have been described recently. GTP \( \gamma \) has been shown to stimulate the formation of the phosphoinositides in rat liver and brain plasma membranes [27, 28]. The relative influence of these various regulatory sites on the ultimate IP<sub>3</sub> level will depend upon many factors such as the amount of enzyme activities per se, the availability of substrates, as well as their accessibility to the components of the sytem involved.

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