

Molecular Characterization of ADP-ribosyl Cyclase Coupled with Receptors

メタデータ	言語: jpn 出版者: 公開日: 2017-12-22 キーワード (Ja): キーワード (En): 作成者: Higashida, Haruhiro メールアドレス: 所属:
URL	https://doi.org/10.24517/00049435

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KAKEN

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受容体とカップルする ADP リボシールシクラーゼ

分子の個体レベルでの解析

(課題番号 12480228)

平成 12 年度～平成 14 年度科学研究費補助金

基盤研究 B(2)研究成果報告書

平成 15 年 5 月

金沢大学附属図書館



0300-02120-8

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1. はしがき

私達は 10 数年来、受容体からイオンチャンネルに至るシグナル伝達について研究している。ことにムスカリン性アセチルコリンやブラジキニン両受容体刺激により、イオンコンダクタンスを減少させ膜を興奮させる M タイプのカリウム (K^+) イオンチャンネルに興味を持っている。M と K^+ チャンネルの遺伝子が KCNQ チャンネルであることがわかり、サイクリック ADP リボース (cADPR) とプロテインキナーゼ C が M KCNQ チャンネル抑制のセカンドメッセンジャーではないかという仮定のもとに研究を進めてきた。

後者の成果として KCNQ を抑制するプロテインキナーゼは AKAP (A Kinase anchoring protein) と結合する C-キナーゼであることを見出し、この 6 月に Nautre Neuroscience に発表する。また、cADPR 産生酵素 (ADP-ribosyl cyclase) 活性を持つリンパ球細胞膜表面抗原であるヒト CD38 cDNA の大量発現細胞を作り、cADPR の細胞内作用を研究しているうちに、思いがけなく、内在性の ADP-ribosyl cyclase がムスカリン受容体により、G タンパク質を介してその活性を上下させることを知った。この情報交換機構は、非骨格筋細胞内でリアノジン受容体の関与する。即ち、 Ca^{2+} induced Ca^{2+} -release 機構の関与する Ca^{2+} バイオシグナル分野で長らく疑問とされていた宿題に対する一つの回答であった。

そこで我々は、(非骨格筋細胞に存在する II 型の) リアノジン受容体に作用する cADPR の産生機構の分子機構解明を目指し、平成 9 ~ 11 年度に「脳型 ADP リボシルシクラーゼの精製と cDNA クローニング」という課題の研究補助を受け、研究をスタートさせた。また、12 年度から 3 年間にわたり、「受容体とカップルする ADP リボシルシクラーゼ分子の個体レベルでの解析」という課題で研究を深化させた。

その結果①ラット心臓心室筋細胞での ADP リボシルシクラーゼが β 受容体により活性化されること、②培養脳アストロ細胞で ADP リボシルシクラーゼが β 受容体により活性化されること、③神経腫瘍細胞 NG108-15 細胞に注入した cADPR により、L 型 Ca チャンネルが活性化されること、④生後 1 週間以内の心筋では、アンギオテンシン受容体と ADP リボシルシクラーゼがカップルしていること、⑤代謝型グルタミン酸受容体はサブタイプ特異的にカップルすること、⑥上頸神経節細胞のムスカリン受容体が ADP リボシルシクラーゼとカップルし、cADPR が M チャンネル抑制の感受性」を規定していることなどを CD38 ノックアウトマウスを使い証明した。

以上、ADP リボシルシクラーゼの遺伝子から細胞や個体レベルでの解析を行い成果をあげた。

2. 研究組織

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3. 研究経費

交付決定額（配分額）

	直接経費	間接経費	合計
平成 12 年度	5,300,000	0	5,300,000
平成 13 年度	3,600,000	0	3,600,000
平成 14 年度	4,300,000	0	4,300,000
総計	13,200,000	0	13,200,000

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5. 研究成果（日本語版）

パート I

ADP リボシルシクラーゼとカップルする代謝型グルタミン酸受容体サブタイプ

サイクリック ADP リボース (cADPR) はセカンドメッセンジャーであると考えられているが、代謝型グルタミン酸受容体 (mGluR) の下流での役割はわかっていない。ラットやマウスの上頸神経節 (SCG) や網膜から取った粗膜分画を酵素源として、グルタミン酸を投与すると NGD から cGDPR や [3H] cADPR の産成が増加した。薬理的検討から SCG はグループ I、網膜はグループ III の mGluR の関与が考えられた。mGluR 6 ノックアウトマウスの網膜ではグルタミン酸に対する反応は消失した。mgluR 1 が mGluR 7 の cDNA を NG108-15 細胞に発現し、クローン受容体レベルで ADPR シクラーゼ反応を確認した。mGluR 1, 3, 5, 6 が ADPR シクラーゼを上昇させ mGluR 2 は減少させた。グループ II, III の中にカップリング様式の異なるサブタイプがあることから、表に示すような新しいサブタイプ分類を提唱した。

Subclassification of mGluRs according to coupling preference and localization in synaptic regions.

Group	Subtype	Coupling to			Synaptic localization*	
		NC**	AC**	PLC**	Presynaptic	Postsynaptic
I a	mGluR1	↑	↑	↑	-	Peri, Extra
I a	mGluR5	↑	↑	↑	-	Peri, Extra
II a	mGluR3 ↑	↓			Extra	PSD, Peri, Extra
II b	mGluR2 ↓	↓			Extra	Extra
III a	mGluR6 ↑	↓		-		BCE
III c	mGluR4 →	↓			Active zone	-
III c	mGluR7 →	↓			Active zone	-

*Based upon observations described in the text. Peri, perisynaptic site; Extra, extrasynaptic site; PSD, postsynaptic density; BCE, base of central element; -, not detected.

**NC, ADP-ribosyl cyclase; AC, adenylyl cyclase; PLC, phospholipase C.

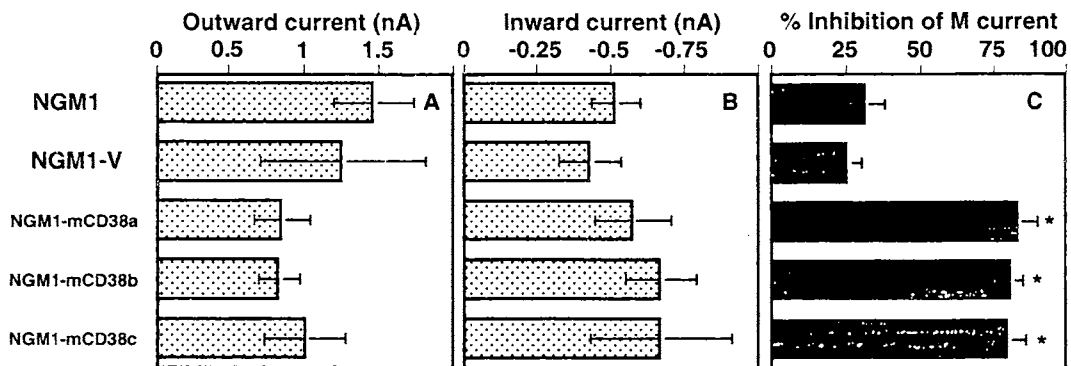
↑, activation; ↓, inhibition; →, no effect.

パート II

アセチルコリンによる M 電流抑制を CD38/ADP リボシルシクラーゼの遺伝子的変異により変化させた

ムスカリン性アセチルコリン受容体 (mAChR) による M 電流抑制における ADP リボシルシクラーゼ/CD38 の役割を検討した。上頸神経節 (SCG) の膜分画にアセチルコリンを投与すると ADP リボシルシクラーゼが濃度依存的に上昇した。ADPR が産生されない CD38 ノックアウトマウスノ SSSCG ではアセチルコリン投与では上昇しなかった。また、M 電流も正常ネズミの SCG 神経細胞と同じ電位量であったが、アセチルコリン投与により M 電流は 25% しか減少しなかった (正常ネズミでは 75%)。一方 CD38 を NG108-15 細胞に大量発現した NG108-15 細胞では、逆に M 電流減少が 3 倍以上増大した (下図 C 参照)。

以上の結果より cADPR より M 電流チャネルへの信号伝達を変化させることがわかった。



6. 研究成果 (英語版)

Part I

Subtype-specific coupling with ADP-ribosyl cyclase of metabotropic glutamate receptors in retina, cervical superior ganglion and NG108-15 cells

Running title: Selective coupling of mGluRs with ADP-ribosyl cyclase

Abstract

Cyclic ADP-ribose (cADP-ribose) is a putative second messenger or modulator. However, the role of cADP-ribose in the downstream signals of the metabotropic glutamate receptors (mGluRs) is unclear. Here, we show that glutamate stimulates ADP-ribosyl cyclase activity in rat or mouse crude membranes of retina via group III mGluRs or in superior cervical ganglion via group I mGluRs. The retina of mGluR6 deficient mice showed no increase in the ADP-ribosyl cyclase level in response to glutamate. GTP enhanced the initial rate of basal and glutamate-stimulated cyclase activity. GTP- γ -S also stimulated basal activity. To determine whether the coupling mode of mGluRs to ADP-ribosyl cyclase is a feature common to individual cloned mGluRs, we expressed each mGluR subtype in NG108-15 neuroblastoma x glioma hybrid cells. The glutamate-induced stimulation of the cyclase occurs preferentially in NG108-15 cells over-expressing mGluRs1, 3, 5, and 6. Cells expressing mGluR2 or mGluRs4 and 7 exhibit inhibition or no coupling, respectively. Glutamate-induced activation or inhibition of the cyclase activity was eliminated after pretreatment with cholera or pertussis toxin, respectively. Thus, the subtype-specific coupling of mGluRs to ADP-ribosyl cyclase via G

proteins suggests that some glutamate-evoked neuronal functions are mediated by cADP-ribose.

Keywords: cyclic ADP-ribose, second messengers, glutamate, signal transduction

Introduction

Metabotropic glutamate receptors (mGluRs) are coupled to G proteins and evoke a variety of neuronal functions, such as modulation of membrane currents, neurotransmitter release, plasticity, cortical differentiation, and addiction, by mediating intracellular signal transduction (Nakanishi, 1992 and 1994; Conn and Pin, 1997; Nakanishi et al., 1998; Chiamulera et al., 2001; Hannan et al., 2001; Miura et al., 2002). There are eight genetically-different subtypes designated mGluR1 to mGluR8 (Nakanishi, 1992 and 1994; Duvoisin et al., 1995). On the basis of their sequence similarity and pharmacology, mGluRs have been defined to belong to three distinct groups (groups I, II and III) (Nakanishi, 1992; Conn and Pin, 1997). However, from the view point of signal transduction, the mGluR subtypes can be largely divided into two classes: those that couple to phospholipase C and/or D (group I, mGluRs1 and 5) (Nakanishi, 1994; Nakanishi et al., 1998; Servitja et al., 1999), and those that negatively regulate adenylyl cyclase (group II, mGluRs2 and 3, and group III, mGluRs4, 6-8) (Nakanishi, 1992; Tanabe et al., 1992; Nakajima et al., 1993; Okamoto et al., 1994; Duvoisin et al., 1995). In contrast, recent *in situ* hybridization and immunohistological studies have revealed that the mGluR subtypes display diverse but distinct distribution in brain regions and more complex pre- and post-synaptic localizations (Masu et al., 1991; Tanabe et al., 1992; Abe et al., 1992; Nomura et al., 1994; Ottersen and Landsend, 1997; Shigemoto et al., 1997; Cartmell and Schoepp, 2000; Neto et al., 2000; Tamaru et al., 2001). Collectively, these results indicate that multiple intracellular signal pathways may exist for different mGluR subtypes to play numerous roles in neuronal function.

In mGluR-mediated Ca^{2+} signaling, group I mGluRs can increase intracellular Ca^{2+} concentrations in brain neurons by Ca^{2+} mobilization from inositol-1,4,5-trisphosphate-sensitive Ca^{2+} stores, interacting with Homer proteins between them (Nakanishi, 1992 and 1994; Crawford et al., 2000; Fagni et al., 2000; Ango et al., 2001; Morikawa et al., 2003). In addition, activation of group I mGluRs mediates increased Ca^{2+} entry through L- or N-type Ca^{2+} channels (Anwyl, 1999), especially in a ryanodine receptor-dependent manner (Chavis

et al., 1996). Although mGluRs sometimes cause inhibition of voltage-dependent Ca^{2+} channels (Ikeda et al., 1995; Anwyl, 1999), these mGluR-mediated Ca^{2+} elevations can often activate Ca^{2+} -dependent K^+ or non-selective cationic channels (Pollock et al., 1999; Fagni et al., 2000; Morikawa et al., 2003). In retina, depolarization of 'ON' bipolar cells is thought to be mediated by the inactivation of mGluR6 with light (Nawy, 2000), and increases conductance of cyclic GMP-dependent cation channels accompanied by subsequent Ca^{2+} concentration elevations (Shiells and Falk, 1999). However, little is known about how mGluRs interact with the Ca^{2+} amplification machinery, namely functional triads consisting of ryanodine receptors, Ca^{2+} channels, and Ca^{2+} -activated K^+ channels (Akita and Kuba, 2000).

One possibility is the involvement of cyclic ADP-ribose (cADP-ribose), a putative second messenger (Lee, 2001; Higashida et al., 2001). cADP-ribose potentiates Ca^{2+} -induced Ca^{2+} -release at neuronal ryanodine receptors triggered by Ca^{2+} influx, designated an 'orthograde signal' (Empson and Galione, 1997; Pollock et al., 1999; Hashii et al., 2000). Subsequently, increased Ca^{2+} affects Ca^{2+} channels indirectly through ryanodine receptors activated by cADP-ribose, referred to as a 'retrograde signal' (Chavis et al., 1996; Empson and Galione, 1997; Hashii et al., 2000). Therefore, our favored hypothesis is that cADP-ribose plays a role as a soluble factor for mGluR-mediated Ca^{2+} signaling.

We previously reported that stimulation of β -adrenergic and angiotensin II receptors modulates activity of ADP-ribosyl cyclase, a synthetic enzyme of cADP-ribose, and showed that cADP-ribose is a critical molecule in sympathetic potentiation of heart contraction, post-natal heart growth, and neuron-glia interaction (Higashida et al., 1999 and 2001). In the present study, we used the same strategy to explore the idea that cADP-ribose is a second messenger downstream of the mGluRs. We first measured ADP-ribosyl cyclase activity in crude membranes from various rat and mouse nervous tissues in the presence or absence of glutamate and/or GTP. Second, the coupling preference of mGluRs to ADP-ribosyl cyclase was examined in NG108-15 neuroblastoma x glioma hybrid cells over-expressing each

subtype.

Materials and methods

Membrane preparation

Wistar rats (1 to 12 weeks old) and adult wild-type or mGluR6^{-/-} mice (Sugihara et al., 1997) were anesthetized using diethylether and decapitated. Various brain regions, retina and superior cervical ganglion (SCG) were isolated and minced, and NG108-15 cells frozen at -80 °C were thawed. Nervous tissues or cells were then suspended in a hypotonic solution (10 mM Tris-HCl solution, pH 7.3, with 5 mM MgCl₂) at 4°C for 30 min. The cell suspensions were homogenized in a glass homogenizer, and the resultant homogenate was centrifuged at 4°C for 5 min at 1000 x g to remove unbroken cells and nuclei. Crude membrane fractions were prepared by centrifugation (twice) of homogenates at 105000 x g for 15 min. The supernatant was removed and the pellet was washed twice, the final pellet was dispersed in 10 mM Tris-HCl solution, pH 6.9, and used immediately for enzymatic reactions. In some experiments, NG108-15 cells were treated with chorela toxin (CTx) or pertussis toxin (PTx) (100 ng/ml) for 8-16 h in culture dishes. Rats were intraperitoneally injected with CTx (100 ng per g of body weight) 16 h before sacrifice.

ADP-ribosyl cyclase and cADP-ribose hydrolase assay

Cell membranes (1.2-6 µg), prepared as above, were placed in 20 µl of reaction mixture containing Tris-HCl (50 mM, pH 7.1), 100 mM KCl, 5 mM MgCl₂, 0.1 mM EDTA, 2 µMNAD⁺, [³H] β-NAD⁺ (0.11 µM, 0.06 µCi) with or without glutamate and GTP slightly modified to what was reported previously (Higashida et al., 1997 and 1999). Reaction mixtures were incubated for 2 min at 37 °C and stopped by the addition of 2 µl trichloroacetic acid (15%). Aliquots were centrifuged (14000 x g, 2.5 min) and 2 µl of the

supernatant was spotted on silica gel plastic thin layer chromatography sheets (20 x 10 cm). The layer was developed in the ascending direction for 40-60 min at room temperature in a mixture consisting of 30% water, 70% ethanol and 0.2 M ammonium bicarbonate (Higashida et al., 1997 and 2002). The positions of authentic cADP-ribose, ADPR and β -NAD⁺ after UV detection were detected and confirmed by autoradiogram of TLC with [³H] β -NAD⁺ obtained after exposure (24 - 36 hours) on a [³H] imaging plate (Fuji BAS 1000, Tokyo, Japan) in each run. Corresponding positions on the experimental runs were cut out of the gel (about 1 cm by 0.7 cm) and the radioactivity counted in a liquid scintillation counter. cADP-ribose hydrolase activities were also assayed in 20- μ l reaction mixtures contained 2 μ M [³H]cADP-ribose (0.015 μ Ci), according to a previously described formula (Higashida et al., 1997).

Fluorometrical measurement of ADP-ribosyl cyclase

ADP-ribosyl cyclase activity was determined fluorometrically using a technique based on the measurement of the conversion of β -nicotinamide guanine dinucleotide⁺ (β -NGD⁺) into the fluorescent product cyclic GDP-ribose (cGDP-ribose), as described previously (Graeff et al., 1994; Higashida et al., 1999). Briefly, 2.4 ml of reaction mixtures containing 60 μ M β -NGD⁺, 50 mM Tris-HCl, pH 6.9, 100 mM KCl, 10 μ M CaCl₂ and membranes (1.5-55.4 μ g of protein) were maintained at 37°C with constant stirring. The samples were then excited at 300 nm and fluorescence emission was continuously monitored at 410 nm in a Shimadzu RF-5300PC spectrofluorophotometer (Kyoto, Japan). Enzyme activity was determined as the slope of curves computed from data points obtained every 60 ms, and confirmed by curve fitting of smoothing-processed data.

Electrophysiological measurements

Collagenased rat SCG cells were cultured and membrane potentials of ganglion neurons were recorded with a sharp intracellular microelectrode, as previously described (Mochida et al., 1994).

Construction of expression plasmids

Expression plasmids carrying the entire coding sequences for the mGluRs1-7 were constructed as follows. The *EcoRV/NotI* fragments from pmGR1 (Masu et al., 1991) were ligated to *NotI/EcoRV*-cleaved pcDNA3 (Invitrogen) to obtain pcDmGR1a. The *EcoRI* fragments from pmGRs2, 3 and 4 (Tanabe et al., 1992), and pmGR6 (Nakajima et al., 1993) were cloned into the *EcoRI* sites of pcDNA3 to obtain pcDmGR2, pcDmGR3, pcDmGR4a, and pcDmGR6, respectively. The *EcoRI/NotI* fragments from pmGR5 (Abe et al., 1992) and pmGR7 (Okamoto et al., 1994) were ligated to *NotI/EcoRI*-cleaved pcDNA3 to yield pcDmGR5a and pcDmGR7a.

Transient or stable transfection of mGluR cDNAs into NG108-15 cells

Mouse neuroblastoma x rat glioma hybrid NG108-15 cells (5×10^5) were plated in 60-mm dishes (Higashida et al., 1997). Twenty-four h later, the cells were transfected with 10 μ g each of pcDmGR1a-7a or pcDNA3 mixed with 25 μ l Lipofectamine plus (Gibco-BRL). The transfection efficiency of pcDNA3 was 15-35% (n=5) in NG108-15 cells in dishes transfected with pcDNA3 inserted with neuronal calcium sensor 1 fused with green fluorescence protein, according to results from the fluorescence-positive cells. One day after transfection, harvested cells were further cultured in new 60-mm dishes for 4 days for the ADP-ribosyl cyclase assay. Some cells were characterized by Western blotting using six monoclonal or polyclonal antibodies (A52, mG2Na-5, Y33, K44, G53, and G74) which recognize mGluRs1a, 2, 3, 4a, 5, and 7a and an antiserum against mGluR6 (Shigemoto et al., 1997; Tamaru et al., 2001).

Stable NG108-15 clones over-expressing each mGluR were generated by transfecting the cells with pcDmGRs1-7 and selecting in 800 μ g/ml geneticin. At least two clones of transfectants expressing each mGluR subtype were selected, and characterized by Western blotting and measurement of adenylyl cyclase. Adenylyl cyclase activity was measured in a

100 μ l reaction mixture containing 0.32 mM [32 P]ATP (0.6 μ Ci), and [32 P]cAMP was separated with thin-layer chromatography by using the same solvent for separating cADP-ribose as described previously (Higashida et al., 2002).

Statistical analysis

All values are represented as mean \pm S.E.M. Homogeneity of variance was tested with Fischer's *F* test followed by Student' *t* test using two-way analysis of homogeneous variance.

Results

ADP-ribosyl cyclase activity was measured using radioisotopes (Higashida et al., 1997) or by the formation rate of nonhydrolyzable cGDP-ribose from β -NGD $^{+}$ (Graeff et al., 1994; Higashida et al., 1999) with crude membranes isolated from various regions of the nervous system. ADP-ribosyl cyclase activity was detected in the olfactory bulb, cerebral cortex, hippocampus, cerebellum, retina or SCG of 1- to 12-week-old rats. [3 H]cADP-ribose formation and cGDP-ribose fluorescence increased upon exposure to glutamate (100 nM and 1 μ M) in crude membrane fractions of both retina and SCG in the presence of endogenous GTP (Figs. 1 and 2), but was not much influenced in other brain regions, including in the olfactory bulb, cerebral cortex, hippocampus, cerebellum. The mean ADP-ribosyl cyclase activity was 49.3 ± 23.6 and 495.5 ± 226.9 pmol/min/mg of protein (n=4) for retina and SCG of 4-week-old rats, respectively. Total activities of ADP-ribosyl cyclase were about 0.86 pmol/min/SCG and 0.38 pmol/min/retina in 4-week-old rats, respectively. The increase in [3 H]cADP-ribose formation induced by 10 μ M glutamate was $188.3 \pm 46.1\%$ in retina and $195.0 \pm 36.4\%$ in SCG (n=4; vs. no glutamate, $p < 0.05$; Fig. 1), respectively. Differing from the responses in the rat heart to angiotensin II and isoproterenol (Higashida et al., 2001), coupling efficiency and preference of ADP-ribosyl cyclase to glutamate was not much changed developmentally in retina and SCG of 2-4-weeks old to young adult rats.

In contrast, cADP-ribose hydrolase activity was barely detectable in both retina and

SCG membranes, similar to NG108-15 cells (Higashida et al., 1997). The conversion ratio to [³H]ADP-ribose from [³H]cADP-ribose was unchanged in the absence or presence of 10 μM glutamate. This implies that the glutamate-induced accumulation of cADP-ribose in the membranes of both retina and SCG is due to accelerated production, rather than breakdown inhibition. These findings were further studied by measuring ADP-ribosyl cyclase fluorometrically.

Glutamate-induced activation of ADP-ribosyl cyclase in retina

The increase in ADP-ribosyl cyclase was dose-dependent (Fig. 2). The increase to 205.5 ± 16.6 and 220.0 ± 20.8% (n=6 and 4) of the pre-exposure level was obtained by 1 and 10 μM glutamate in retina, respectively (*p* < 0.001). This glutamate-induced increase was mimicked by an mGluR group III-selective agonist, L(+)-2-amino-4-phosphonobutyric acid (L-AP4), but not by (S)-3,5-dihydroxyphenylglycine (DHPG) or (2S,2'R,3'R)-2-(2',3'-dicarboxycyclopropyl)glycine (DCG-IV), group I- and II-selective agonists (Fig. 3A). Exposure to 10 μM L-AP4 increased the ADP-ribosyl cyclase level to 196.1 ± 8.8% (n=6) of the control level (*p* < 0.001). An increase obtained by the glutamate (10 nM) exposure subsequent to the initial application of 10 nM L-AP4 was small, but much higher by DHPG and DCG-IV (Table 1).

In the presence of 10 μM (RS)-α-cyclopropyl-4-phosphonophenylglycine (CPPG), a group III-selective antagonist, an application of glutamate had no effect (Table 2). In contrast, application of group I- and II-selective antagonists, (RS)-1-aminoindan-1,5-dicarboxylic acid (AIDA) and (2S)-α-ethylglutamic acid (EGLU), did not lead to an increase in the ADP-ribosyl cyclase level by themselves, but subsequent glutamate (10 nM - 1 μM) did, suggesting that this glutamate effect in retina is mediated via the group III mGluRs.

The most abundant subtype in retina is mGluR6 (Nakajima et al., 1993), though other mGluR subtypes are also expressed (Duvoisin et al., 1995; Hartveit et al., 1995). To define whether mGluR6 mediates ADP-ribosyl cyclase activation, we used mGluR6 deficient mice

(Sugihara et al., 1997). The mGluR6^{-/-} retina showed no increase in the ADP-ribosyl cyclase level in response to 1 μ M glutamate ($101.9 \pm 5.7\%$, $n=7$, vs. no glutamate, Fig. 4A), while the retina from wild-type mice showed an identical increase to the rat retina ($201.0 \pm 20.4\%$, $n=6$; $P<0.001$ vs. mGluR6^{-/-}). In the control experiment, glutamate increased ADP-ribosyl cyclase activity in the SCG of both mGluR6^{+/+} and mGluR6^{-/-} mice equally (Fig. 4B). These results indicate that retinal activation is primarily due to mGluR6, while that of SCG is due to other mGluR subtypes.

In separate experiments, optimal conditions for the assay and effects of glutamate were examined in more detail with retinal preparations. Production of cGDP-ribose was optimal at about pH 6.5 (at 37 °C), similarly shown in ADP-ribosyl cyclase activity of cardiac myocytes (Higashida et al., 1999). However, the glutamate-induced activation was obtained only at the more alkaline range than 6.8 (data not shown). Thus, it is confirmed that the identical activation (up to 2- to 3-fold) by glutamate was observed at the physiological pH range, though the basal activity was reduced.

Glutamate-induced activation of ADP-ribosyl cyclase in SCG

The maximum increase in ADP-ribosyl cyclase to $201.3 \pm 17.9\%$ ($n=7$) and $194.7 \pm 10.8\%$ ($n=3$) of the pre-exposure level was obtained by 1 or 10 μ M glutamate in the SCG (Fig 2; $P<0.001$). Unlike retina, this glutamate-induced increase was mimicked by DHPG (1 μ M, $198.7 \pm 15.5\%$, $n=3$; $P<0.01$), but not by DCG-IV or L-AP4 (Fig. 3B). Glutamate induced a further activation even after the treatment with DCG-IV or L-AP4, but not with DHPG (Table 1). In the presence of 10 μ M AIDA, but not EGLU or CPPG, glutamate (10 nM) failed to induce an increase in the ADP-ribosyl cyclase level (Table 2). These results suggest that, pharmacologically, the effect of glutamate in the SCG is mediated via the group I mGluRs.

Despite a lack of mGluR responses in adult rat SCG (Ikeda et al., 1995), mobilization of intracellular Ca²⁺ by mGluR5 has been shown in neonatal rat dorsal root ganglion neurons in culture (Crawford et al., 2000). Therefore, we dissected SCG neurons from 10-day-old rats

and cultured them for 5-6 weeks (Mochida et al., 1994), to evaluate functional mGluRs. Application of 300 μM glutamate to cultured SCG neurons resulted in a rapid depolarization (time to peak, ~ 15 s) with a mean amplitude of 3.1 ± 0.84 mV ($n=5$), followed by a small depolarization of 0.97 ± 0.09 mV ($n=5$) which lasted for up to 30 min (Fig. 5). This slow depolarization was associated with an increased membrane resistance and action potential firings (data not shown), similar to group I agonist-induced depolarizing responses in enteric neurons (Liu and Kirchgessner, 2000). Exposure of neurons to glutamate also elicited a frequency increase in spontaneous excitatory postsynaptic potentials in our records (data not shown), due to an enhanced release of acetylcholine from paired SCG neurons (Mochida et al., 1994). The depolarization was mimicked by 1.5 μM DHPG (1.40 ± 0.34 mV, $n=8$) and significantly antagonized by 100 μM AIDA (0.13 ± 0.06 mV, $n=5$; $p<0.001$), indicating that cultured SCG neurons possess group I mGluRs.

Effects of GTP or GTP analogs on ADP-ribosyl cyclase activity

The above stimulation induced by glutamate seems to be due to endogenous GTP and GTP-binding proteins in membranes. Therefore, we determined whether or not mGluR-mediated activation of ADP-ribosyl cyclase in retina and SCG is reproduced by GTP and its analogue. Addition of GTP resulted in an increase in enzyme activity in retina (Fig. 6A) and SCG (Fig. 6B). A relatively symmetrical biphasic (activating and inhibitory) relationship is apparent between the concentration (0.1 nM-100 μM) of GTP, as shown in the fat cell adenylyl cyclase system by Rodbell and his colleagues (Cooper et al., 1979). GTP was more potent than glutamate in terms of its effective concentration (less than micromolar) range. GTP (100 or 10 nM) stimulated basal activity by $84.2 \pm 11.4\%$ ($n=6$) in retina and $109.6 \pm 18.3\%$ ($n=6$) in SCG ($p<0.001$).

Next, we examined the effect of glutamate in the presence of various concentrations of GTP. GTP caused a 1.5- to 2.5-fold stimulation of the response of the enzyme system to 10 nM glutamate (Fig. 6C). The maximum increase in the presence of glutamate was $271.2 \pm$

22.9% (n=6) at 1 μ M GTP in retina or $341.8 \pm 41.4\%$ (n=4) at 10 nM GTP in SCG, respectively ($p < 0.001$).

The effect of a non-hydrolyzable analogue of GTP, GTP- γ -S, that persistently activates GTP-binding proteins, was examined. GTP- γ -S produced only an increase in cGDP-ribose forming activity in both retina and SCG. The dependency on the concentration is not bell shaped (data not shown). Increases to $207.7 \pm 3.3\%$ (n=6) in retina and $199.3 \pm 6.0\%$ (n=3) in SCG of the control were produced at 1 μ M or 100 nM GTP- γ -S ($p < 0.001$), respectively. Addition of 1 μ M GDP- β -S increased the cyclase activity slightly (about 10%), but blocked further increases by 1 μ M glutamate in retina and SCG (n=4).

Effect of glutamate and bacterial toxin on [3 H]cADP-ribose formation

Glutamate-induced activation of the cyclase activity in the crude membrane of both retina and SCG in rats was eliminated after pretreatment with 100 ng/ml CTx for 8-16 h ($p < 0.01$; Fig. 7A). This suggests the likelihood that the membrane delimited effect of glutamate is mediated via the CTx-sensitive subset of G proteins. The same line of data has been obtained in m1 and m3 muscarinic receptors in NG108-15 cells (Higashida et al., 1997) and β -adrenoceptors in cardiomyocytes and cortical astrocytes (Higashida et al., 1999 and 2001).

Cyclase activity in NG108-15 cells transiently overexpressing each mGluR subtype

To determine whether the above coupling mode of mGluRs to ADP-ribosyl cyclase is a feature common to individual cloned mGluRs, we expressed each mGluR subtype in NG108-15 neuroblastoma x glioma hybrid cells. ADP-ribosyl cyclase activity was detected in membranes prepared from NG108-15 cells transiently transfected with cDNAs of each mGluR subtype, as previously characterized (Higashida et al., 1997).

Upon application of 10 nM glutamate, ADP-ribosyl cyclase activity significantly increased in membranes prepared from NG108-15 cells expressing mGluRs1 and 5, with an average activation of $177.4 \pm 18.9\%$ (n=5) and $151.0 \pm 10.9\%$ (n=3) of the pre-application

level (Fig. 8, $p < 0.01$ and 0.05), respectively. As expected, 10 nM glutamate significantly increased ADP-ribosyl cyclase activity to $214.2 \pm 41.5\%$ ($n=5$) of the pre-exposure level in mGluR6 expressing cells ($p < 0.01$). Surprisingly, activation of ADP-ribosyl cyclase activity to $184.6 \pm 10.7\%$ ($n=7$) by glutamate was also obtained in mGluR3-transfected cells ($p < 0.01$). In the control experiments, non-transfected or mock-transfected cells did not exhibit glutamate-induced activation (101.3 to 108.1% ($n=4$ and 8)) of the pre-application level (Fig. 8). Thus, the values obtained in NG108-15 cells over-expressing mGluRs1, 3, 5, and 6 are significantly higher than those in wild-type and mock-transfected control cells ($p < 0.001$ or 0.05).

A decrease in ADP-ribosyl cyclase activity by 10 nM glutamate to $74.4 \pm 4.3\%$ ($n=5$) of the pre-application level was observed in mGluR2-transfected NG108-15 cells ($p < 0.001$; Figs. 7C and 8). No or little change in cyclase activity was obtained with glutamate in mGluR4- and mGluR7-expressing NG108-15 cells (Fig. 8). Moreover, group-selective agonists faithfully replicated the glutamate responses for different subtypes (data not shown).

To exclude dysfunctional receptor expression in NG108-15 cells, we measured adenylyl cyclase activity. Glutamate (10 μM) inhibited to $83.2 \pm 6.7\%$ ($n=11$, $p < 0.05$) of the non-treated level in 10 μM forskolin-activated adenylyl cyclase activity in cells over-expressing mGluR4. 10 μM prostaglandin E_1 -activated adenylyl cyclase activity was inhibited to $65.6 \pm 6.7\%$ ($n=11$, $p < 0.001$) of the non-application level in mGluR7-expressing cells. Likewise, as reported, stimulation of mGluRs2, 3 and 6 by 10 μM glutamate inhibited basal adenylyl cyclase activity to $79.4 \pm 14.4\%$ ($n=5$), $83.3 \pm 8.1\%$ ($n=3$), and $81.4 \pm 12.3\%$ ($n=7$) of the control, respectively ($p < 0.05$).

Signal transduction from mGluRs to ADP-ribosyl cyclase in stable clones

Finally, the coupling preferences of the mGluRs was examined in stably expressing NG108-15 cell lines. The results of stable clones matched those of transiently expressed cells, and exhibited nearly identical magnitudes of responses by 10 nM to 10 μM glutamate (Fig. 7B).

The glutamate-induced activation was blocked in mGluRs1-, 3-, and 6-expressing NG108-15 cells treated with CTx, but not with PTx. Inversely, the inhibition by glutamate in mGluR2-expressing cells was nullified with PTx (Fig. 7C).

Discussion

These results show that glutamate stimulates activity of the membrane-bound form of ADP-ribosyl cyclase in rat and mouse retina or rat SCG, and that GTP enhanced the initial rate of glutamate-stimulated activity in rat retina and SCG. This is the first biochemical demonstration of the mGluR/cADP-ribose signal transduction pathway via G proteins in the nervous system. Pharmacological effectiveness and the result obtained in mGluR6-gene-targeted mice revealed that group III (mGluR6) and group I (mGluRs1 and/or 5) mGluRs are responsible for glutamate-induced activation of ADP-ribosyl cyclase in retina and SCG, respectively. Our results of over-expression of cloned mGluRs in NG108-15 cells demonstrate subtype-specific coupling preferences of mGluRs to ADP-ribosyl cyclase, as summarized in Table 3. Cyclase activation was not only restricted to mGluRs1 and 5, but also mGluRs3 and 6. The reconstitution of mGluRs1, 5 and 6 in NG108-15 cells may represent those found in SCG and retina.

It will be interesting to show an increase of cADP-ribose concentrations by glutamate in vivo cells expressing mGluRs 1, 3, 5 and 6. However, we could not perform a kind of radioimmunoassay, because an ADPR-binding protein (antibody; Takahashi et al., 1995) was no more available. Since ryanodine receptors with a major band of 400 kd (Kuwajima et al., 1992) were detected with an anti-dog cardiac ryanodine receptor antibody in retina, SCG and NG108-15 cells in our preliminary results as shown by Ronde et al. (2000), the stimulatory mGluRs may modulate intracellular Ca^{2+} concentrations through interaction of ryanodine receptors with increased cADP-ribose. The similar evidence has recently been shown by Morikawa et al. (2003) that mGluRs can release Ca^{2+} in dopamine neurons in the

rat ventral midbrain through the activation of both inositol trisphosphate and ryanodine receptors

In the mGluRs belonging to group II, we found that inhibitory mGluR2 differs from excitatory mGluR3 with respect to glutamate-induced ADP-ribosyl cyclase activity, while both receptors equally inhibit adenylyl cyclase, as reported (Nakanishi et al., 1992 and 1994). Among group III mGluRs, which also exhibit the same effect on adenylyl cyclase, stimulatory mGluR6 is totally different from mGluRs4 and 7, which have no apparent coupling. The reason for such discrepancy in coupling modes to ADP-ribosyl cyclase and adenylyl cyclase between mGluRs within the same group is not clear. One possible explanation will be different G-protein selectivity of mGluRs belonging to the same group. Although this is not a direct evidence for mGluRs in group II and III, it has been recently shown that mGluR1 and mGluR5 of the same group-I mGluRs, which primarily couple to phospholipase C via the $G_{q/11}$ -type G protein, activate the extracellular signal-regulated kinase via distinct signaling pathways, in that mGluR1 utilizes a PTx-sensitive pathway but mGluR5 does not (Thandi et al., 2002).

Taking our findings on coupling to ADP-ribosyl cyclase into consideration, new subclasses in group II and III of mGluRs should be defined (Table 3). Both mGluRs1 and 5 remain in group I, given their identical signal transduction preference. If 'a' is used for mGluRs that activate, 'b' is given to mGluRs that negatively regulate, and 'c' stands for non-coupling receptors, mGluRs1 and 5 belong to group Ia. mGluR3 could be subdivided to group IIa, and mGluR2 into group IIb. The group IIIa subclass includes mGluR6, whereas group IIIc includes mGluRs4 and 7.

Is such sub-classification valid? We believe that new subdivision from functional aspects (i.e., coupling preference) may well accord to the difference in histological localization (Ottersen and Landsend, 1997; Shigemoto et al., 1997; Cartmell and Schoepp, 2000), as listed in Table 3. For instance, mGluRs3 and 2 are located at the same presynaptic regions, but not exactly the same postsynaptic site (Tamaru et al., 2001). In the hippocampal

dentate gyrus, the postsynaptic localization of mGluR3 is rather similar to that of group I mGluRs than mGluR2 (Tamaru et al., 2001). Similarly, while mGluRs4 and 7 are exclusively located presynaptically, mGluR6 shows postsynaptic localization (Nakajima et al., 1993; Nomura et al., 1994; Shigemoto et al., 1997; Neto et al., 2000). Interestingly, closer inspection of Table 3 reveals that coupling to ADP-ribosyl cyclase appears to be restricted to a subset of mGluRs whose postsynaptic existence has been proven. Therefore, such coincidence may support our new classification resulted from more multiple signal transduction pathways than previously appreciated (Nakanishi, 1992 and 1994; Duvoisin et al., 1995; Conn and Pin, 1997; Nakanishi et al., 1998).

The present study is one of the few to describe the possible role of peripheral mGluRs, given that activation of mGluR5 modulates nociception on unmyelinated sensory afferents (Bhave et al., 2001). Judging from our electrophysiological studies on cultured SCG cells, group I mGluRs increase cADP-ribose formation in ganglion neurons. However, glial cells also possess group I mGluRs (Biber et al., 1999), hence a glial contribution to our measurement of ADP-ribosyl cyclase can not be excluded. Preganglionic neurons in the SCG do not release glutamate (Conn and Pin, 1997; Anwyl, 1999), and thus one possibility is that glutamate released from surrounding glial cells may act on mGluR1 or mGluR5 in postganglionic neurons, *in vivo*. It will be interesting to test if release of acetylcholine from ganglion cells could be modulated by activation of ryanodine receptors by increased cADP-ribose.

mGluR6 has been shown to be located not on the tip of the central element but on its base at the mouth of the invagination, 400-800 nm from the release site in 'ON' bipolar cells (Vardi et al., 2000), and function to transfer on-signaling by depolarizing bipolar cells via opening of cGMP-gated cation channels by light stimulation (de la Villa et al., 1995). These channels could be partially phosphorylated with light by Ca^{2+} (Shiells and Falk, 1999; Nawy, 2000) and CaM-kinase II (Walters et al., 1998; Shiells and Falk, 2000), and should be dephosphorylated by calcineurin in the dark-adapted retina (Shiells and Falk, 2001). If this

explanation is true, a plausible mechanism is that cADP-ribose interacts with calcineurin (Shiells and Falk, 2001), which removes the rectification properties of cGMP-dependent channels (Shiells and Falk, 2001). Alternatively, since Ca^{2+} concentration is increased by caffeine in goldfish or carp retinal bipolar cells (Kobayashi et al., 1995; Wu and Zhu, 1999), cADP-ribose may potentiate this reaction through ryanodine receptors. The physiological relevance of mGluR6-mediated cADP-ribose formation remains to be determined.

The coupling of mGluRs1 and 5 to ADP-ribosyl cyclase via CTx-sensitive G proteins is not surprising, because mGluRs1 and 5 activate phospholipase C (Nakanishi et al., 1992 and 1994; Miyashita and Kubo, 2000; Miura et al., 2002), similarly observed in muscarinic receptor subtypes 1 and 3 (Higashida et al., 1997). However, the involvement of a CTx-sensitive subset of G proteins in mGluR6- and mGluR3-coupling is unexpected, since mGluRs6 and 3 are negatively coupled to adenylyl cyclase via PTx-sensitive G_i/G_o in this and other experiments (Nakanishi et al., 1994). Although the main G protein coupled with mGluR6 is G_o (Vardi, 1998), it has been reported that CTx-sensitive- G_i also weakly but significantly couples to mGluR6 (Weng et al., 1997). However, CTx on this mGluR-mediated activity is perplexing and CTx may not have any direct effects on any G proteins that bridge over mGluRs and ADP-ribosyl cyclase. Further validation is necessary of G protein species-linkage between mGluRs and ADP-ribosyl cyclase. In support of our hypothesis of G protein involvement, recently, cADP-ribose and CD38 were shown to regulate Ca^{2+} responses and chemotaxis of a specific subset of G protein-coupled receptors in human neutrophils (Partida-Sanchez et al., 2001) and endothelin was found to stimulate ADP-ribosyl cyclase via activation of both endothelin A and B receptors (Barone et al., 2002).

In conclusion, we have demonstrated that efficient coupling of mGluRs to ADP-ribosyl cyclase in neuronal tissues occurs in a subtype-specific fashion. The challenge for the future is to define in greater detail the role of cADP-ribose in neurons (Higashida et al., 2001), which may provide greater insight into the mechanisms of mGluR-mediated neuronal function.

Acknowledgements - We thank Masanobu Kano for comments on the manuscript. This work was supported grants from the Ministry of Education, Science, Sports, Culture and Technology of Japan.

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Table 1

Effects of glutamate and group-specific agonists on cGDP-ribose formation in rat retina and SCG

Group-specific agonist	ADP-ribosyl cyclase activity (% of pre-addition level)		
	Initial application of agonist	Subsequent application of glutamate	Δ Increase (%)
<u>Retina</u>			
None	-	167.5 \pm 16.9 (9)*	-
I (DHPG)	102.3 \pm 2.3 (4)	173.0 \pm 6.6 (4)**	70.7***
II (DCG-IV)	100.3 \pm 0.4 (3)	158.5 \pm 10.8 (3)*	58.2***
III (L-AP4)	172.3 \pm 8.6 (5) **	194.0 \pm 8.2 (5)**	21.7
<u>SCG</u>			
None	-	164.8 \pm 5.2 (5)**	-
I (DHPG)	173.3 \pm 4.3 (4)**	178.8 \pm 5.7 (4)**	5.5
II (DCG-IV)	111.3 \pm 6.6 (3)	171.0 \pm 4.7 (3)**	59.7***
III (L-AP4)	107.3 \pm 1.3 (3)	174.7 \pm 9.4 (3)**	67.4***

Cell membranes were incubated with 60 μ M β -NAD⁺, and the fluorescence of the resulting cGDP-ribose continuously monitored. Activity after application of 10 nM DHPG, DCG-IV, and L-AP4 was measured and calculated as percentage of the pre-application level, respectively. Subsequently, 10 nM glutamate was added. It is noted that only little increase was obtained by glutamate in the presence of L-AP4 or DHPG in retina and SCG. * and **, significantly different from control values (without agonists) at $p < 0.05$ and 0.001. ***, significantly different between values of initial application with agonists and subsequent application with glutamate at $p < 0.01$.

Table 2

Effect of glutamate on cGDP-ribose formation in the presence of each group-specific antagonist in rat retina and SCG.

Group-specific antagonist	Glutamate-induced activation (%)	
	Retina	SCG
I (AIDA)	159.5 ± 20.2 (4)*	103.0 ± 4.3 (4)
II (EGLU)	147.0 ± 5.0 (3)**	185.7 ± 1.5 (3)**
III (CPPG)	102.3 ± 1.5 (3)	185.3 ± 8.6 (3)**

Glutamate (10 nM) was applied in the presence of 10 µM antagonists. * and **, significantly different from the control value at $p < 0.05$ and $p < 0.001$.

Table 3

Subclassification of mGluRs according to coupling preference and localization in synaptic regions.

Group	Subtype	Coupling to			Synaptic localization*	
		NC**	AC**	PLC**	Presynaptic	Postsynaptic
I a	mGluR1	↑	↑	↑	-	Peri, Extra
I a	mGluR5	↑	↑	↑	-	Peri, Extra
II a	mGluR3	↑	↓		Extra	PSD, Peri, Extra
II b	mGluR2	↓	↓		Extra	Extra
III a	mGluR6	↑	↓		-	BCE
III c	mGluR4	→	↓		Active zone	-
III c	mGluR7	→	↓		Active zone	-

*Based upon observations described in the text. Peri, perisynaptic site; Extra, extrasynaptic site; PSD, postsynaptic density; BCE, base of central element; -, not detected.

**NC, ADP-ribosyl cyclase; AC, adenylyl cyclase; PLC, phospholipase C.

↑, activation; ↓, inhibition; →, no effect.

Figure legends

Fig. 1. Activation of ADP-ribosyl cyclase by glutamate in rat retina and SCG. ADP-ribosyl cyclase was measured from the initial velocity during 2 min with membranes prepared from rat retina (A) and SCG (B). Values were calculated by conversion of [³H]cADPR from [³H]NAD⁺ in the presence or absence of 10 μM glutamate in 4 experiments. Values indicate mean ± S.E.M. * indicates significantly different from the control at $p < 0.01$.

Fig. 2. Effect of glutamate on cGDP-ribose-producing activity in retina and SCG. Plot shows relationship between concentrations of glutamate and cGDP-ribose formation in retina (square) and SCG (circle), olfactory bulb (downward triangle), cerebral cortex (right-ward triangle), hippocampus (upward triangle), cerebellum (diamond). Data represent mean ± S.E.M. for 3-9 measurements. Values at 1 nM or higher concentrations of glutamate are significantly different ($p < 0.02$) from control activity (100% without agonists) in both retina and SCG. Inset, cell membranes isolated from retina (upper, 55.4 μg) and SCG (lower, 35.7 μg) of 5-week-old rats were incubated with 60 μM β-NGD⁺, and the fluorescence of the resulting cGDP-ribose continuously monitored. At the time indicated by arrow, 10 μM glutamate at the final concentration was added.

Fig. 3. Effects of various concentrations of group-specific agonists on cGDP-ribose-producing activity in rat retina (A) and SCG (B). Relationship between concentrations of L-AP4 (diamond, III), DCG-IV (circle, II), or DHPG (triangle, I) and cGDP-ribose formation. The fluorescence of the resulting cGDP-ribose was continuously monitored, and the velocity of cGDP-ribose formation calculated as shown in Fig. 2. Changes in fluorescence induced by application of indicated concentrations of the 3 agonists and glutamate (square, G), as in Fig. 2, were

plotted. Data represent mean \pm S.E.M. for 3-9 measurements. Values at 1 nM of L-AP4 in A and DHPG in B are significantly different ($p < 0.02$) from the control activity (100% without agonists) and all other values of L-AP4 in A and DHPG in B are at $p < 0.001$.

Fig. 4. Effect of glutamate on ADP-ribosyl cyclase activity in retina and SCG of wild-type and mGluR6 deficient mice. 1 μ M glutamate induced activation of ADP-ribosyl cyclase activity in membranes of retina (A) and SCG (B) of wild-type (+/+) or mGluR6-knockout mice (-/-). Data represent 6-8 measurements from three 12-month-old mice of each type. Bars= S.E.M. *, $p < 0.001$.

Fig. 5. Chart-recordings of membrane potential changes in cultured rat SCG neurons induced by glutamate and a group I agonist. (A, B) Three hundred μ M glutamate was added to the bath (arrows), in the absence (A) and the presence of 100 μ M AIDA (B). (C) DHPG was added to the bath producing 1.5 μ M. The resting membrane potential of the SCG neurons was -51 mV (A), -53 mV (B) and -52 mV (C), respectively. (Inset) Peak amplitude of slow depolarizations was measured and averaged from five experiments with 300 μ M glutamate (A), five experiments with glutamate in the presence of 100 μ M AIDA (B), and eight experiments with 1.5 μ M (S)-5,5-DHPG (C). Values represent the mean \pm SEM was plotted.

Fig. 6. Effect of GTP on cGDP-ribose-producing activities in retina and SCG. Changes in fluorescence induced by application of indicated concentrations of GTP were plotted in retina (A) and SCG (B) of 5-week-old rats. Each data point represents the mean \pm S.E.M of 3-11 determinations. Values at 0.1 nM of GTP in A and B are significantly different at $p < 0.05$ from the control activity. Values for 10 and 100 μ M are at $p < 0.01$, and all other values at $p < 0.001$. (C) Changes in

fluorescence induced by application of 10 nM glutamate in the presence of indicated concentrations of GTP were plotted in retina (square) and SCG (circle). Each data point represents the mean \pm S.E.M. of 6 or 4 determinations in retina and SCG, respectively. All values are significantly different at $p < 0.01$ from the control activity in both retina and SCG.

Fig. 7. Effects of pretreatment of rats or NG108-15 cells with CTx or PTx on glutamate-induced changes in ADP-ribosyl cyclase activity. Rats (A) or cells stably over-expressing mGluRs1, 3 and 6 (B) or mGluR2 (C) were intraperitoneally injected or incubated with CTx or PTx (100 ng/body weight or /ml) 8-16 h before being tested. Membranes of retina and SCG (A) or NG108-15 cells (B and C) treated (+) or untreated (-) were incubated and ADP-ribosyl cyclase activity determined before and after 1 μ M glutamate application. Data represent the mean (\pm S.E.M.) of 3-6 determinations. * and #, significantly different from non-treated control values at $p < 0.01$ or $p < 0.05$, respectively.

Fig. 8. Glutamate-induced activation or inhibition of ADP-ribosyl cyclase activity in various NG108-15 cells over-expressing mGluRs. ADP-ribosyl cyclase activity was measured with membranes prepared from NG108-15 cells, mock-transfected cells, or NG108-15 cells transiently transfected to express mGluRs1-7. The level of activity was compared in the presence or absence of 10 nM glutamate in the reaction mixture. Data represent means of 3-10 determinations. Bars represent S.E.M. * and #, significantly different from non-transfected (101.3%) and mock-transfected (108.1%) values, at $p < 0.001$ or 0.005, respectively.

Figure 1

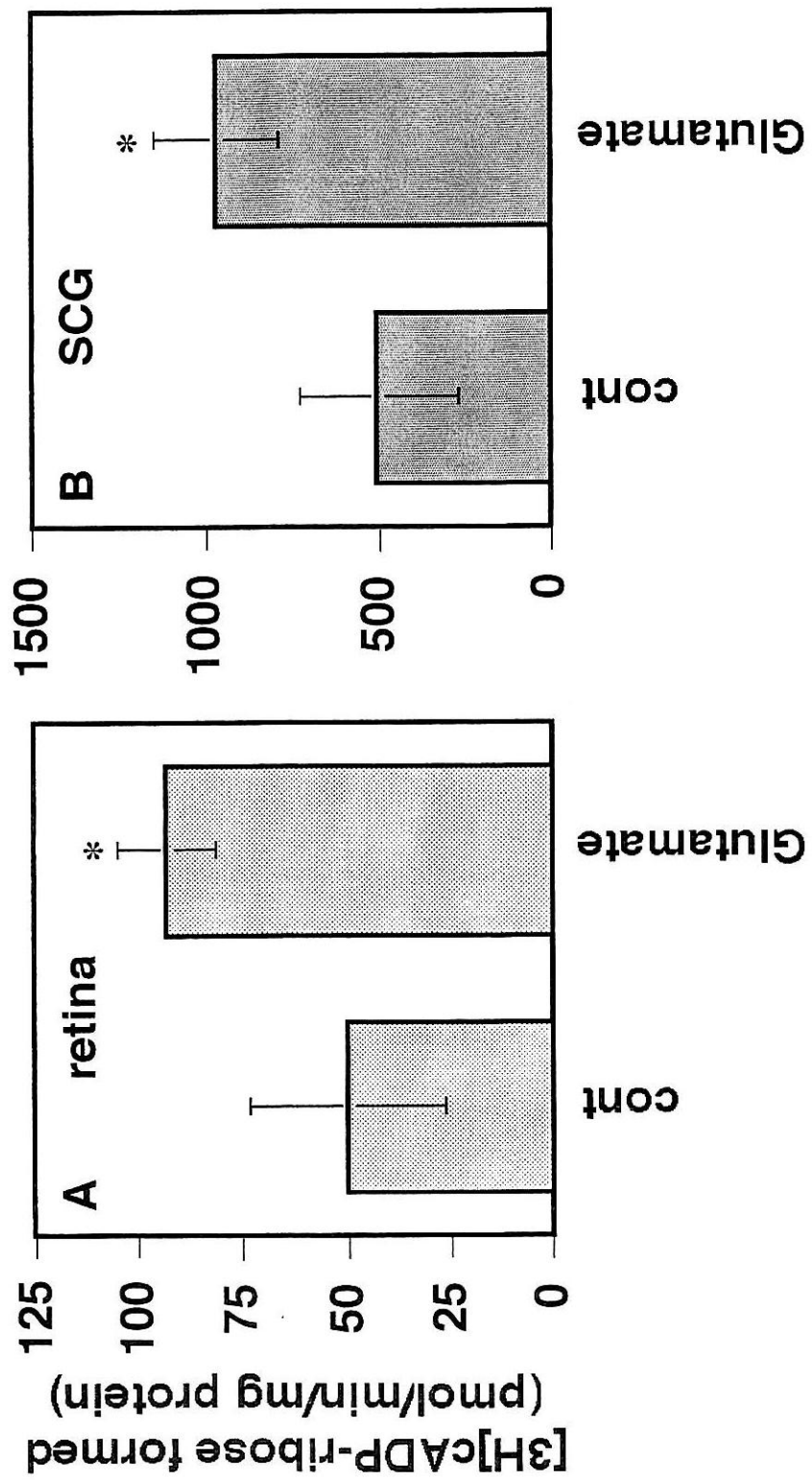


Figure 2

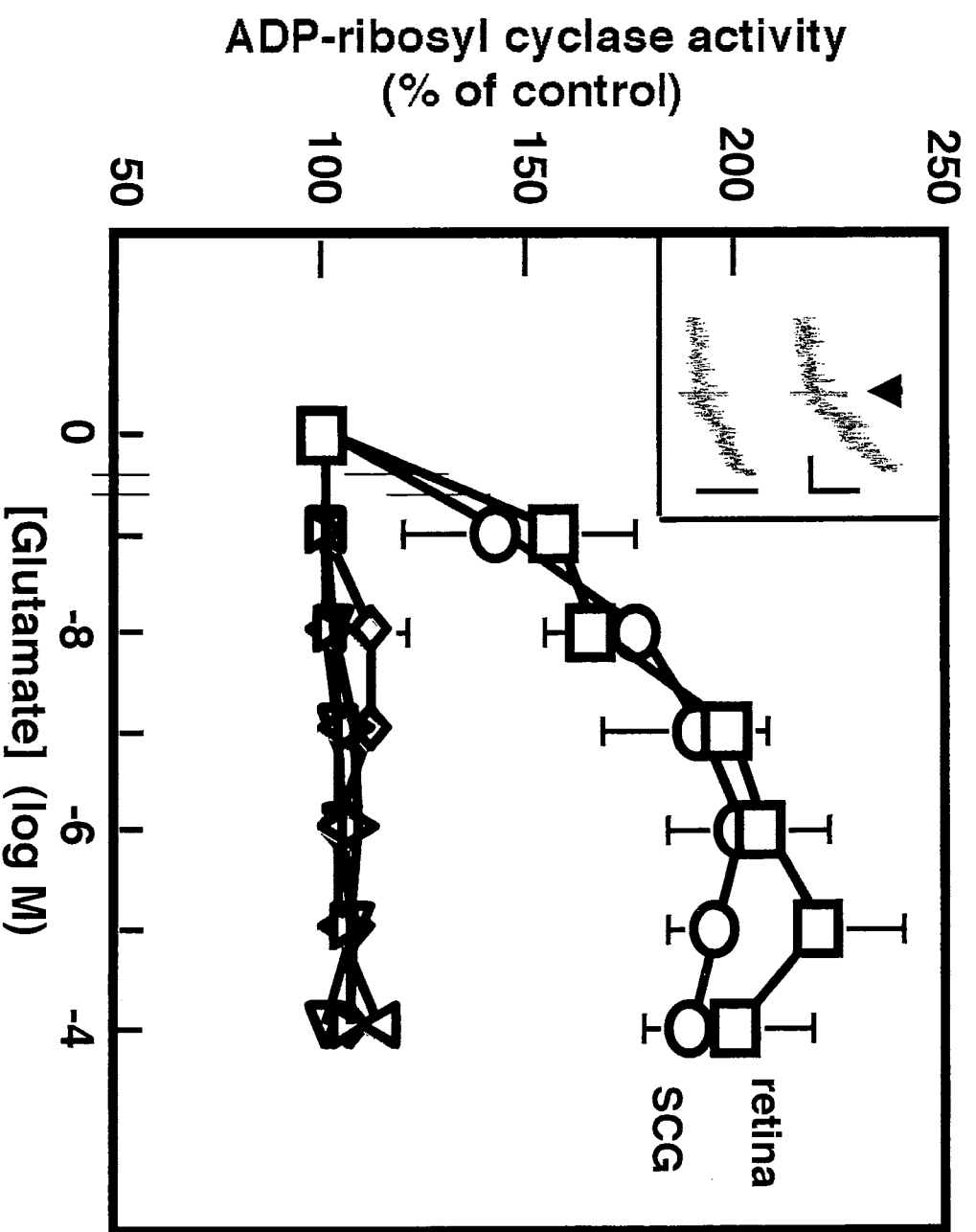
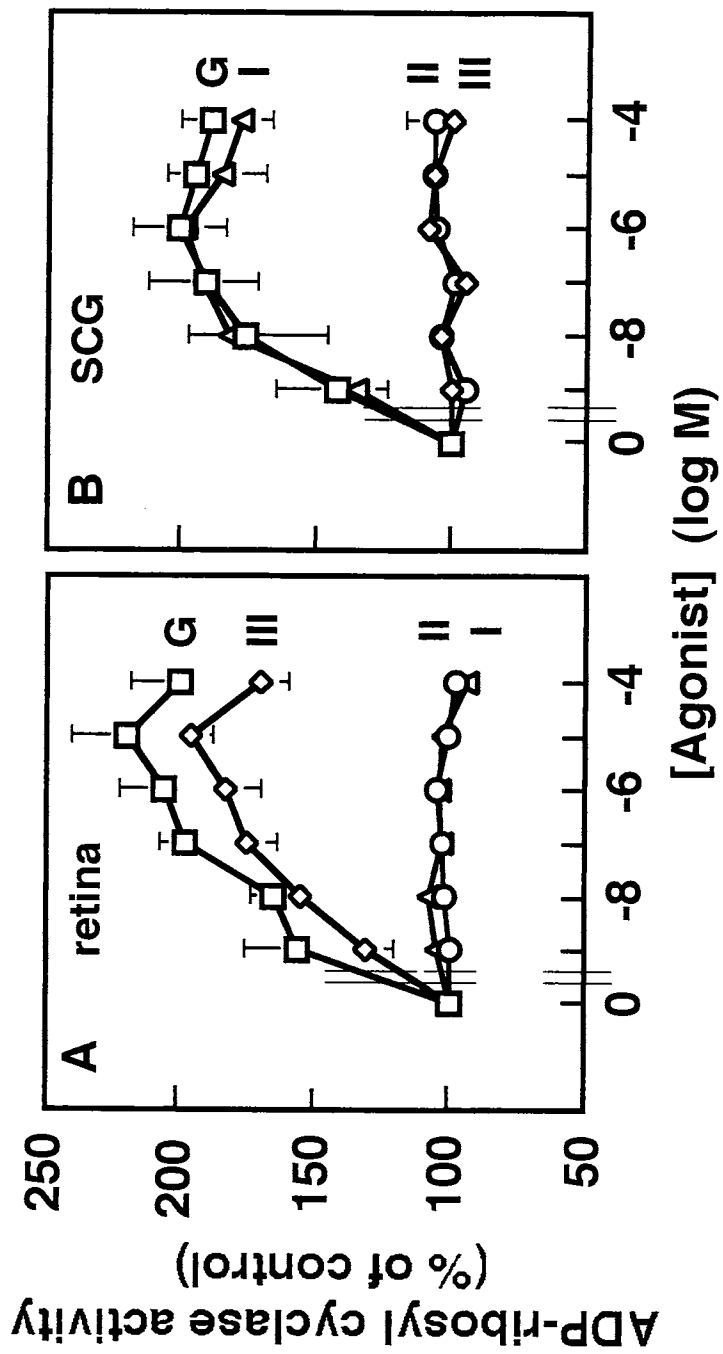


Figure 3



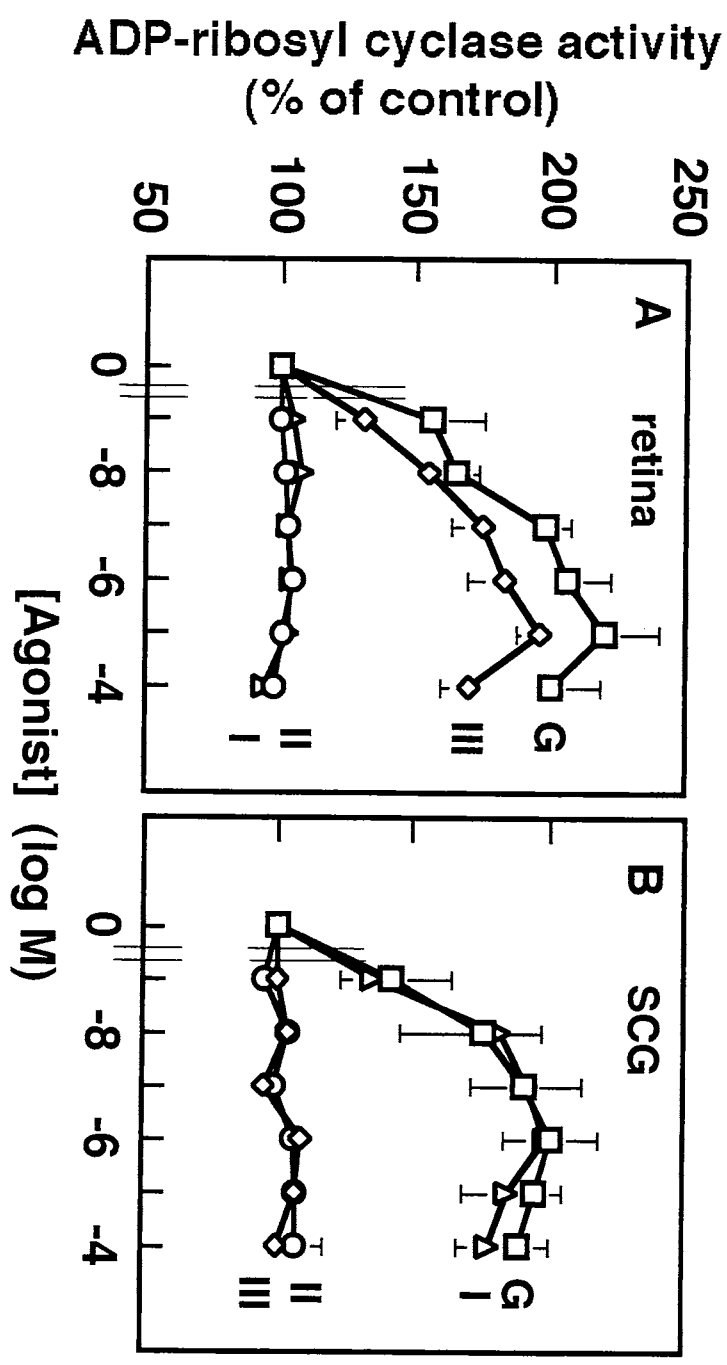
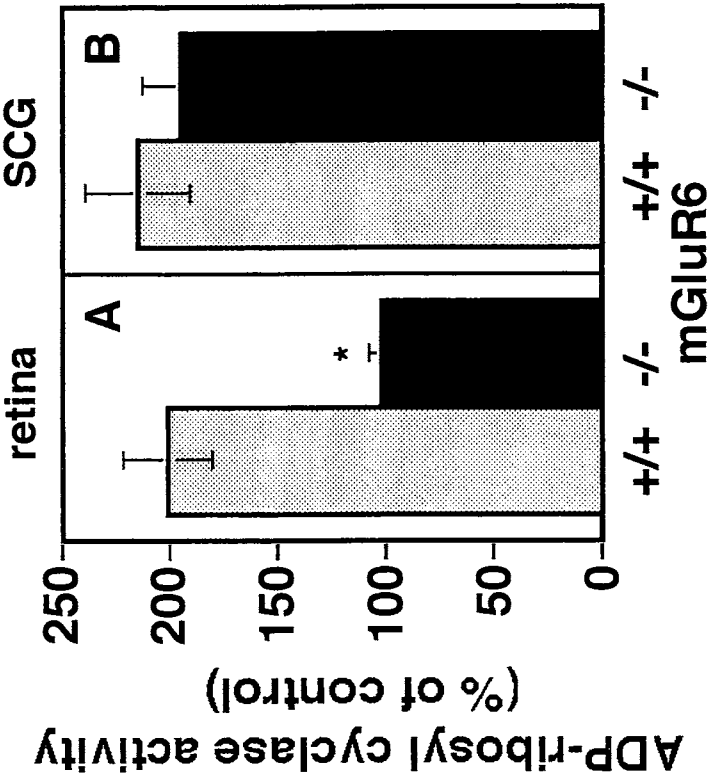


Figure 4

Figure 5



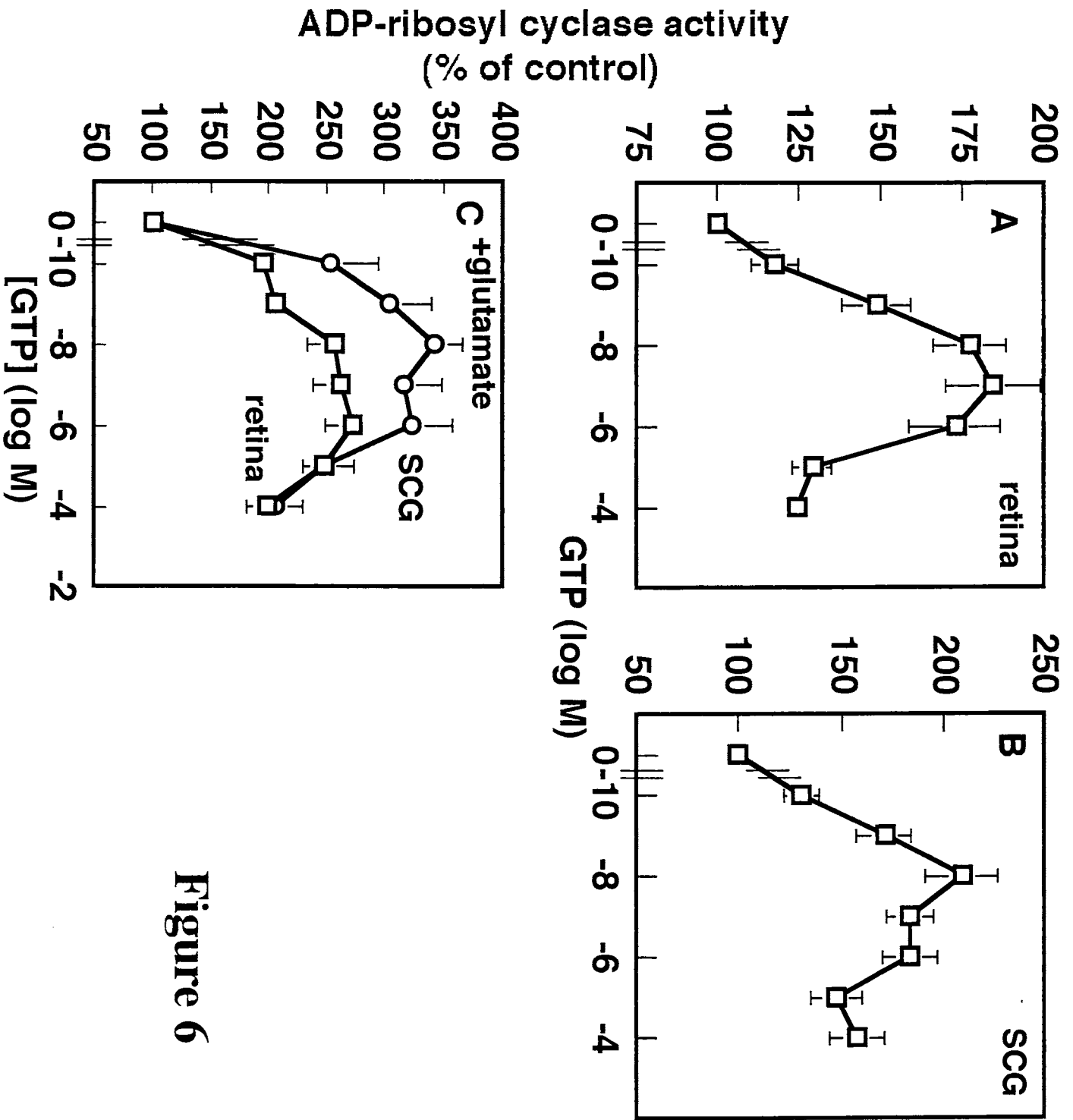


Figure 6

Figure 7

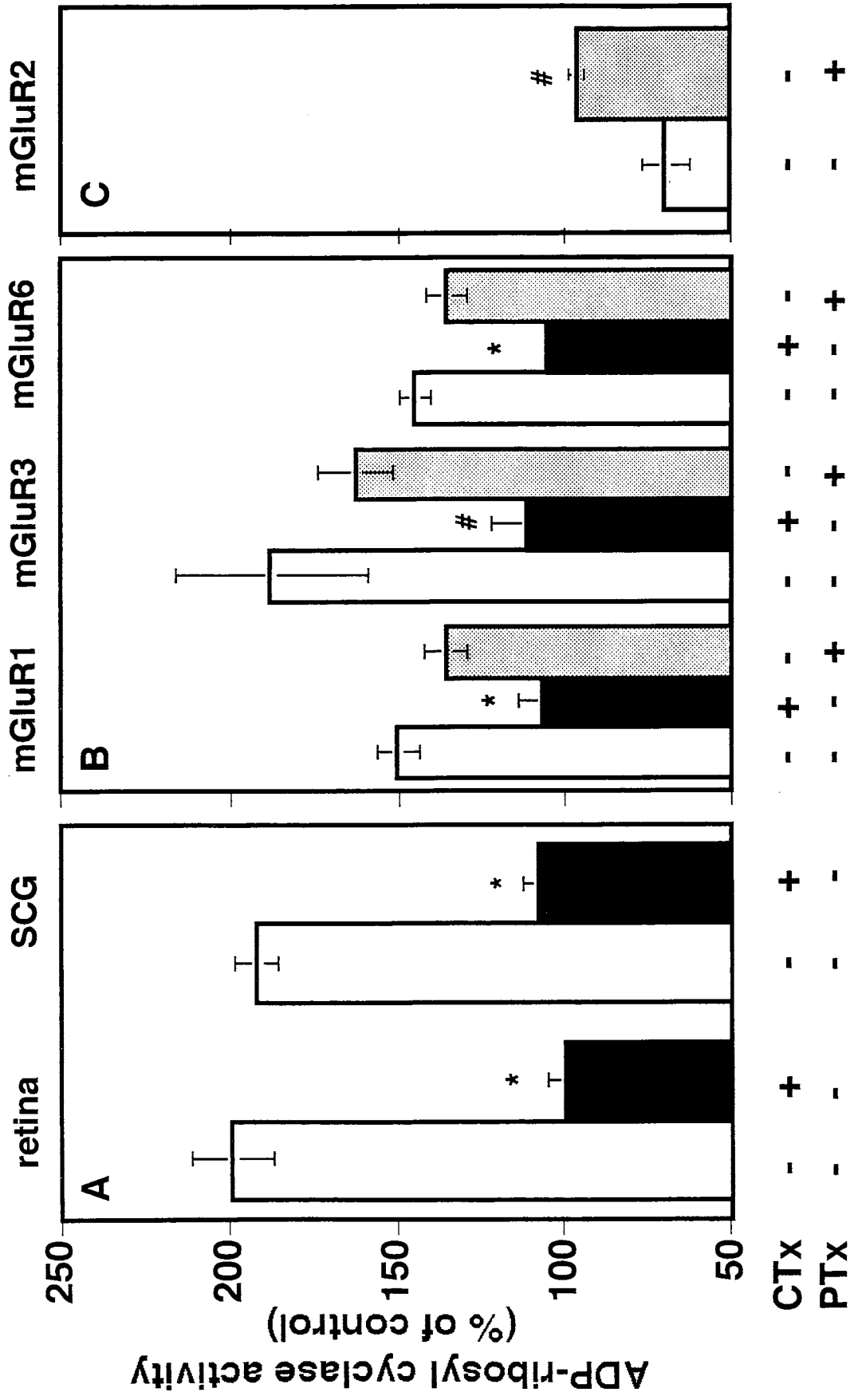
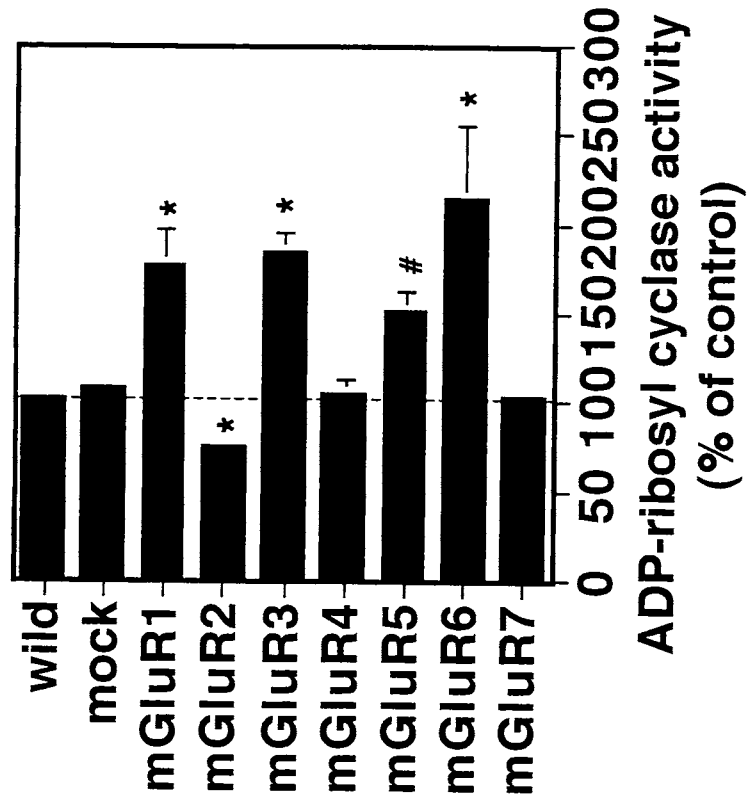


Figure 8



Part II

Genetic Manipulations of CD38/ADP-Ribosyl Cyclase Alter Acetylcholine-Induced M-Current Inhibition in Superior Cervical Ganglion Neurons and Neuroblastoma Cells

Key Words: cyclic ADP-ribose, CD38, M-current, ADP-ribosyl cyclase, potassium current, muscarinic receptor.

Abstract

The role of ADP-ribosyl cyclase/CD38 in muscarinic acetylcholine receptor (mAChR)-activated inhibition of voltage-gated M potassium currents was examined in normal rats and mice, CD38 knockout mice, and CD38-transformed NG108-15 cells. Application of ACh or other mAChR agonists increased ADP-ribosyl cyclase activity by 200-350% in crude membrane fractions from rat superior cervical ganglia (SCG). This increase was inhibited by atropine or by an M1-mAChR antagonist, pirenzepine, and was mimicked by GTP. ADP-ribosyl cyclase activity in membrane fraction from SCG of CD38 knockout (CD38^{-/-}) mice was negligible. M current inhibition by ACh in cultured SCG neurons from CD38^{-/-} mice was greatly reduced compared with that in neurons from wild-type (CD38^{+/+}) mice. Cloned M1-mAChR-transformed NG108-15 neuroblastoma x glioma hybrid cells over-expressing human CD38 which has been modified to only display ADP-ribosyl cyclase activity were developed. In these cells, ACh-induced inhibition of the fast (KCNQ) component of the M-like current ($I_{K(M,ng)}$) was significantly greater than in control cells. Inhibition of the slow (ERG) component of $I_{K(M,ng)}$ was largely unchanged, and the voltage dependent gating parameters of $I_{K(M,ng)}$ were not significantly different between the cell types tested. Thus, the extent of ACh-induced KCNQ/M-current inhibition positively correlated with ADP-ribosyl cyclase activity and/or CD38 expression level. These results suggest that the cADPR/ADP-ribosyl cyclase signalling system plays a role in KCNQ/M current inhibition following mAChR stimulation.

Introduction

Cyclic ADP-ribose (cADPR) is an endogenous modulator of ryanodine receptor Ca^{2+} releasing channels (Lee, 1997, 2001). In neurons, cADPR facilitates the release of Ca^{2+} from ryanodine-sensitive stores when Ca^{2+} entry is evoked through voltage-dependent Ca^{2+} channels (an 'orthograde' signal: Hua et al., 1994; Empson and Galione, 1997). There is also a 'retrograde' signal, since Ca^{2+} influx through N- or L-type Ca^{2+} channels can be modulated by activating type 2 ryanodine receptors (Davies et al., 1996; Chavis et al., 1996). Hashii et al. (2000) demonstrated that cADPR is involved in this bi-directional coupling between ryanodine receptors and neuronal Ca^{2+} channels in NG108-15 cells (Higashida et al., 2001a, b).

The key enzyme responsible for synthesis of cADPR is ADP-ribosyl cyclase (Kim et al., 1993; Lee, 1997, 2001; Higashida et al., 1997, 2001a, 2001b). ADP-ribosyl cyclase is a multi-functional ectoenzyme, catalysing both the synthesis of cADPR from $\beta\text{-NAD}^+$ and the hydrolysis of cADPR to ADP-ribose (Summerhill et al., 1993; Takasawa et al., 1993b). This enzyme is homologous to the mammalian T and/or B cell surface marker CD38 (Jackson and Bell, 1990; States et al., 1992; Harada et al., 1993; Koguma et al., 1994; Prasad et al., 1996). CD38 null mutant mice ($\text{CD38}^{-/-}$) show impaired ADP-ribosyl cyclase activity; In such mice, blood sugar level is significantly higher and ryanodine-sensitive Ca^{2+} signalling is attenuated in pancreas, compared to wild type animals (Kato et al., 1999; Fukushi et al., 2001). It has been shown that cysteine residues at 119th and 201st positions found in the human (h)CD38 sequence are involved in the cADPR hydrolase reaction (Takasawa et al., 1993b; Tohgo et al., 1994). Replacing these cysteines (C119K/C201E) converts hCD38 to a monofunctional enzyme with only cADPR synthesizing activity (Tohgo et al., 1994).

In previous experiments (Higashida et al., 1995; Higashida et al., 1996; Bowden et al., 1999; Higashida et al., 2000a), we have obtained information to suggest that cADPR may modulate the M-like K^+ current ($I_{\text{K(M.ng)}}$) in NG108-15 mouse neuroblastoma x rat glioma hybrid cells, and may play a part in the inhibition of this current by muscarinic acetylcholine-receptor (mAChR) stimulation. Thus, stimulation of those muscarinic receptors that couple to $I_{\text{K(M.ng)}}$ (M1 and M3 receptors: Fukuda et al., 1988; Robbins et al., 1991 and 1993), but not M2 or M4 receptors, stimulated ADP-ribosyl cyclase activity in membrane fractions from these cells (Higashida et al., 1997); elevation of the precursor to cADPR (NAD^+) reduced M1-mAChR-induced $I_{\text{K(M.ng)}}$ inhibition (Higashida et al., 1995); intracellular application of cADPR via the patch-pipette accelerated 'run-down' of $I_{\text{K(M.ng)}}$ (Higashida et al., 1995); and intracellular application of the cADPR antagonists, 8-amino-cADPR and 8-bromo-cADPR reduced M1-mAChR-induced inhibition of $I_{\text{K(M.ng)}}$ (Bowden et al., 1999).

The M-like current in NG108-15 cells is a composite current, carried in part by currents

through KCNQ2/3 channels (the molecular correlates of the M channel in sympathetic neurons: Wang et al., 1998), and in part by currents through mERG channels (Meves et al., 1999; Selyanko et al., 1999). Although both components appear to be affected by cADPR, and although both are inhibited by M1-mAChR stimulation in a manner sensitive to cADPR antagonists (Higashida et al., 2000a), the presence of the two components adds complexity to the interpretation of the above experiments.

Accordingly, in the present experiments, we have sought further information regarding the role of cADPR in M current regulation. First, we have tested whether M1-mAChR stimulation increases ADP-ribosyl cyclase activity in membrane fractions from rat superior cervical sympathetic ganglia (SCGs), the neurons of which only appear to express KCNQ currents, not ERG currents (Selyanko et al., 2002). Second, we have compared M1-mAChR-induced inhibition in sympathetic neurons from wild-type (CD38^{+/+}) and CD38^{-/-} null mutant mice. Third, we have tested the effect of over-expressing the cADPR hydrolase-deficient CD38 mutant on M1-mAChR-induced inhibition of $I_{K(M,ng)}$ in NG108-15 cells.

Methods and Materials

Materials

β -[2,8-adenine-³H]NAD⁺ (30.5 Ci mmol⁻¹) was purchased from New England Nuclear (Boston, MA, U.S.A.). cADPR was obtained from either Yamasa Shoyu (Choshi, Japan) or Sigma (St Louis, MO, U.S.A.). Oxidized-nicotinamide guanine dinucleotide (β -NGD⁺) was obtained from Sigma. Silica Gel 60 F254 plastic TLC sheets were obtained from Merck (Darmstadt, Germany).

Mice

Mice lacking CD38 were generated by homologous recombination. The generation and genotyping of the mice have been described in detail previously (Kato et al., 1999; Fukushi et al., 2001). The mice used for each experiment were derived from ICR background and were from the same litter or the same family. Sympathetic neurons were isolated from SCG of humanely killed 14- to 21-d-old mice and cultured using standard procedures as described previously (Selyanko et al., 2002).

Cell culture of NG108-15 cells and CD38 transfection

Neuroblastoma x glioma NG108-15 cells over-expressing the M1 muscarinic receptor (NGM1; Fukuda et al., 1988 and 1989) were cultured as described previously (Higashida et al., 1986). Briefly cells were maintained at 37 °C and 10% CO₂ in Dulbecco's modified Eagle's

medium supplemented with 5% foetal calf serum 100 μ M hypoxanthine, 0.1 μ M aminopterin and 16 μ M thymidine. The expression plasmids pZHCD38 and pZHCMCD38 were constructed as follows. The 0.55 Kb *Hind*III (vector)/*Pst*I (606) fragment and 0.36-kb *Pst*I (606)/*Bgl*II (969) fragment from pSV2-HCD38 (Takasawa et al., 1993b) were ligated with 3.0-kb *Bam*HI/*Hind*III fragment from pBSIIKS(+) to yield pBSHCD38. Restriction endonuclease sites are identified by numbers in parentheses indicating the 5'-terminal nucleotide generated by cleavage. The 0.91-kb *Hind*III (vector)/*Spe*I (vector) fragment containing the entire protein-coding sequence from pBSHCD38 was ligated with 3.4-kb *Spe*I/*Hind*III fragment from the expression vector pZeoSV to yield pZHCD38 and pZHCMCD38. The plasmids were multiplied in *E. coli* and purified with a plasmid DNA purification kit (Qiagen, Germany).

NGM1, m1 muscarinic receptor transformed NG108-15 cells were plated in six 35-mm dishes at a density 7.5×10^4 cells per dish. After 24 hours the cells were transfected with 10 μ g of pZHCD38 or pZHCMCD38 mixed with 25 μ g lipofectamine (Bethesda Research laboratories Life Technologies Inc., USA). One day after transfection, the cells were harvested and plated in twelve 60-mm dishes. One week later the cells were selected with 200, 250 and 500 μ g ml⁻¹ Zeocin. Independent colonies were obtained from cells treated with 200 μ g ml⁻¹ Zeocin after 4 weeks.

Membrane preparation

Both sexes of Wistar rats (1 to 4 weeks old) and wild-type or CD38^{-/-} mice (3-4 weeks old) were anaesthetized using diethylether and decapitated. SCGs were isolated and minced. NG108-15 cells frozen at -80 °C were thawed. Nervous tissues or cells were then suspended in a hypotonic solution (10 mM Tris-HCl, pH 7.3, with 5 mM MgCl₂) at 4°C for 30 min. The cell suspensions were homogenized and the resultant homogenate centrifuged at 4°C for 5 min at 1000 x g to remove unbroken cells and nuclei. Crude membrane fractions were prepared by centrifugation (twice) of homogenates at 105,000 x g for 15 min. The supernatant was removed and the pellet was washed twice, the final pellet was dispersed in 10 mM Tris-HCl solution, pH 6.9, and used immediately for enzymatic reactions. Protein was measured by the BioRad assay dye reagent.

ADP-ribosyl cyclase and cADP-ribose hydrolase assay

Cell membranes (1.2-6 μ g), prepared as above, were placed in 20 μ l of reaction mixture containing Tris-HCl (50 mM, pH 6.6), 100 mM KCl, 5 mM MgCl₂, 0.1 mM EDTA, 2 μ MNAD⁺, [³H] β -NAD⁺ (0.11 μ M, 0.06 μ Ci) slightly modified to what was reported previously (Higashida et al., 1997). Reaction mixtures were incubated for 0.5 to 16 min at 37

°C and stopped by the addition of 2 μ l trichloroacetic acid (10-48%). Aliquots were centrifuged (14,000 x g, 2.5 min) and 2 μ l of the supernatant was spotted on silica gel plastic thin layer chromatography (TLC) sheets (20 x 10 cm). The layer was developed in the ascending direction for 40-60 min at 24 °C in a mixture consisting of 30% water, 70% ethanol and 0.2 M ammonium bicarbonate (Higashida et al., 1999). The positions of authentic cADPR, ADPR and β -NAD⁺ after UV detection were confirmed in each run. Corresponding positions on the experimental runs were cut out of the gel (about 1 cm by 0.7 cm) and the radioactivity counted in a liquid scintillation counter. Autoradiography of TLC with [³H] β -NAD⁺ was obtained after exposure (24 - 36 hours) on a [³H] imaging plate (Fuji BAS1000, Tokyo, Japan). cADPR hydrolase activities were also assayed in 20- μ l reaction mixtures contained 2 μ M [³H]cADPR (0.015 μ Ci), according to a previously described formula (Higashida et al., 1997).

Fluorometrical measurement of ADP-ribosyl cyclase

ADP-ribosyl cyclase activity was determined fluorometrically using a technique based on the measurement of the conversion of β -NGD⁺ into the fluorescent product cGDPR, as described previously (Graeff et al., 1994; Higashida et al., 1999). Briefly, 2.4 ml of reaction mixtures containing 60 μ M β -NGD⁺, 50 mM Tris-HCl, pH 6.9, 100 mM KCl, 10 μ M CaCl₂ and membranes (1.5-50 μ g of protein) were maintained at 37°C with constant stirring. The samples were then excited at 300 nm and fluorescence emission was continuously monitored at 410 nm in a Shimadzu RF-5300PC spectrofluorophotometer (Kyoto, Japan). Enzyme activity was determined as the slope of curves computed from data points obtained every 60 ms, and confirmed by curve fitting of smoothing-processed data.

Electrophysiology

The whole cell variant of the patch-clamp technique was employed to record membrane currents from SCG neurons after 2 days in culture or from NG108-15 cells at 4 to 14 days after differentiation, as described in Robbins et al. (1993). Cells were superfused, at 5-10 ml/min, with HEPES-buffered MEM or DMEM or a solution containing 120 mM NaCl, 3 mM KCl, 11 mM glucose, 22.6 mM NaHCO₃, 1.2 mM MgCl₂, 5 mM HEPES, 2.5 mM CaCl₂ and 500 nM tetrodotoxin. The pH was 7.36 when gassed with 95%O₂ and 5% CO₂ at 35 °C. Glass microelectrodes of 2-4 MOhm resistance when filled with potassium acetate (90 mM), KCl (20 mM), MgCl₂ (3 mM), HEPES (40 mM), EGTA (3 mM), CaCl₂ (1 mM), ATP (2 mM), GTP (0.5 mM), pH adjusted to 7.4 with KOH (1M). Calculated free calcium for this solution was 66 nM (Chelator for Windows 1.1) and series resistance were typically 5 MOhm. Cells were routinely clamped at -20 to -30 mV and hyperpolarised by -30 or -40 mV to generate deactivating tail

currents. Voltage pulses were generated by an amplifier (Axoclamp 2B) in discontinuous mode, while currents were recorded via a signal conditioner (Cyberamp 320), digitised on an ADC (Digidata 1200) and displayed on a PC running commercial software (pClamp 6). All data was stored on optical disk (Panasonic, UK). Data is expressed as mean \pm S.E.M. and statistical analysis was performed using an ANOVA and a post hoc Dunnetts test if $p > 0.05$ on the ANOVA.

Membrane currents were recorded from SCG neurons using amphotericin-perforated patch electrodes, as described previously (Selyanko et al., 2002). The composition of the electrode solutions were (mM) for SCG neurons: K acetate 80; KCl 30, HEPES 40, and MgCl₂ 3. Solutions were adjusted to pH 7.3-7.4 with KOH and to 280 mOsmol/l with K acetate. Electrode resistances were 2-4 MOhm; access resistances after amphotericin perforation were 6-8 MOhm.

Western blotting

Control and transformed cell membranes were subjected to Western blot analysis to detect the presence of CD38 protein as described previously (Zhong et al., 1997). The membranes were incubated for 2 h with mouse anti-human or -mouse CD38 monoclonal antibody (Immunotech, France and Chemicon Int.), washed 3 times for 5-10 min in a blocking buffer. The membranes were then incubated for 1 h at room temperature in the blocking buffer containing horseradish peroxidase-cojugated goat anti-mouse or anti-rabbit antibodies. The membranes were then washed 3 times for 5-10 min with blocking buffer and reactive proteins visualised with a Western blotting analysis system (Amersham, UK).

Northern blotting

CD38 mRNA was detected using Northern blot analysis. Total cellular RNA was extracted from control NGM1 cells, mock-, and wild-type- or mutant CD38-transfected cells as described previously (Yokoyama et al., 1994). Briefly, 15 μ g of total RNA was separated on a 1.1% agarose/2.2 M formaldehyde gel and transferred onto a Zetaprobe nylon membrane. Hybridisation was performed at 42 °C in a solution whose recipe was reported previously (Yokoyama et al., 1994). The probe used was the 0.91-kb *HindIII/SpeI* fragment from pZHCD38 labelled by the random primer method. The filter was washed at 50 °C in 0.1 x SSG. Autoradiography was carried out using Fuji BAS1000.

Immunocytochemistry

Control and transfected NG108-15 cells grown on coverslips were washed twice for 3

min with PBS, pH 7.4, fixed for 20 min with freshly prepared 4% formaldehyde in PBS, pH 7.4, at room temperature, and washed 3 times with PBS. The fixed cells were further incubated with 0.1% Triton X-100 in PBS for 5 min; after washing for 3 min (3 times) with PBS, the cells were quenched for 20 min in PBS containing 0.1 M glycine and washed for 5 min with PBS. Immunoreactivity was then demonstrated by incubating the cells with mouse anti-hCD38 mAbs clone T16 (Immunotech, France), which were diluted 1:20 in PBS supplemented with 10% whole goat serum, (12 hr; 4°C), further rinsing 4 times for 3 min in PBS with 10% whole goat serum, followed by incubation with FITC-conjugated goat anti-mouse IgG antibodies diluted 1:200 (1 h; at room temperature). The cells were washed 5 times with PBS and mounted onto glass slides for fluorescence microscope examination.

Calcium measurements

Microspectrofluometric determination of intracellular calcium concentrations ($[Ca^{2+}]_i$) was performed with fura-2 acetoxymethylester (fura-2 AM) as described previously (Hashii et al., 2000). Briefly, the cells were loaded with 5 μ M fura-2 AM for 60 min in the dark at 37 °C after an initial DMEM wash. Emission light at 540 nm was monitored via an aperture (10-20 μ m in diameter) in response to alternating excitation wavelengths of 340 nm and 380 nm using a Ca^{2+} microspectrofluometrics system (OSP-3, Olympus, Japan).

The inward spread of an increase in $[Ca^{2+}]_i$ by the activation of mAChRs was measured by a confocal microscope as described previously (Akita and Kuba, 2000). Wild or CD38-transfected cells were loaded with OGB-1 (10 μ M) for 2 hours at 37°C. The cells were line-scanned at the rate of 8 msec/sec with Argon laser (488 nm) before, during and after the application of muscarine. The resultant changes in fluorescence were expanded into a X-t plane.

Results

ACh-induced activation of ADP-ribosyl cyclase in SCG

First, we tested whether or not application of ACh induces any changes in ADP-ribosyl cyclase and cADPR hydrolase activities in SCG membranes when measured by [³H]cADPR production (see Methods). In ganglion membranes from 4-week old rats, the mean rate of [³H]cADPR production from [³H]NAD⁺ was 495.5 ± 226.9 pmol min⁻¹ mg⁻¹ of protein (n=4). ACh (100 nM) increased [³H]cADPR formation to 1555.9 ± 336.7 pmol min⁻¹ mg⁻¹ of protein (n=4), by $314.0 \pm 36.4\%$ ($p < 0.01$), as shown in Fig. 1A. In contrast, cADPR hydrolase activity was negligible in SCG membranes (as with NG108-15 cells: Higashida et al., 1997). This implies that the ACh-induced accumulation of cADPR in the membranes of SCG is due predominantly to accelerated production.

The effect of ACh was confirmed in crude membrane preparations of SCG from 10-35 day old rats using fluorimetric measurement of cyclic GDP-ribose (cGDPR) production as an assay for cADP-ribosyl enzymatic activity (see Methods; Fig. 1B, insert). cGDPR fluorescence increased in a concentration-dependent manner upon exposure to ACh (Fig. 1B). The maximum increase in ADP-ribosyl cyclase was $272.5 \pm 28.1\%$ (n=4) of the pre-exposure level was obtained by 10 nM ACh ($p < 0.01$). This increase was prevented by simultaneous application of 0.1 – 10 μ M atropine (Fig. 1B open diamond). The action of ACh was replicated by 10 μ M carbamylcholine (CCh) ($213.7 \pm 7.8\%$, n=3; $p < 0.001$) or 1 μ M oxotremorine M (Oxo-M) ($256.3 \pm 15.5\%$, n=4; $p < 0.001$).

{Fig. 1 near here}

ADP-ribosyl cyclase stimulation was probably mediated by M1 muscarinic receptors since it was prevented by 100 nM - 10 μ M pirenzepine (an M1-selective antagonist; Fig. 2A) but was unaffected by 1 μ M AF-DX 116, 4-DAMP or tropicamide (M2, M3 and M4-prefering antagonists, respectively; Fig. 2B).

{Fig. 2 near here}

In order to test whether ADP-ribosyl cyclase stimulation by ACh required receptor coupling to an endogenous G protein, we examined the effect of exogenous GTP in well-washed membranes. Addition of GTP alone (10 - 100 nM) resulted in an increase in enzymatic activity in SCG membrane preparations, increasing basal ADP-ribosyl cyclase activity by $84.2 \pm 11.4\%$ (n=6; $P < 0.01$). GTP (10 nM) also enhanced the ACh-induced stimulation (1.5- to 2.5-

fold) of ADP-ribosyl cyclase activity to $341.8 \pm 41.4\%$ ($n=4$; $p < 0.001$). The non-hydrolysable GTP analog, GTP- γ -S, produced a larger increase in basal activity, by $199.3 \pm 6.0\%$ ($n=3$); under these conditions, however, no further activation by ACh could be observed (as previously noted in neuroblastoma hybrid cells: Higashida et al., 1997).

These results suggest that ADP-ribosyl cyclase activation and cADPR formation, like M current inhibition, is a downstream response of sympathetic neurons to stimulating G protein-coupled M1 muscarinic receptors. We therefore tested whether M current inhibition might be affected in ganglion cells from mice that genetically lack CD38/ADP-ribosyl cyclase (Kato et al., 1999).

Decrease in ACh-induced M-current inhibition in SCG of CD38 knockout mice

First, we compared ADP-ribosyl cyclase activity in wild type (CD38^{+/+}) and knockout (CD38^{-/-}) mice. As reported previously (Kato et al., 1999; Fukushi et al., 2001), ADP-ribosyl cyclase was greatly reduced in whole homogenate of cells isolated from knockout mouse SCG. Though some remaining activity was detected in the supernatant fraction, activity in the 10,000 x g membrane fraction was barely detectable ($< 13\%$, $n=5$) (0.18 arbitrary units in CD38^{-/-} vs 1.35 units in CD38^{+/+} mice; Fig. 3A). Further, ACh did not increase fluorescence intensity in membrane fractions prepared from SCG from CD38^{-/-} mice, but did increase fluorescence in membranes from CD38^{+/+} mice (Fig. 3B), as in membranes from rat ganglia (cf. Fig.1.)

{Fig. 3 near here}

Armed with this information, we cultured SCG cells from both types of mice for two days, and then measured M currents in individual neurons (Fig. 4). M-currents in ganglion cells from CD38^{-/-} mice were indistinguishable from those in cells from CD38^{+/+} mice. Thus, the mean amplitude of M-current deactivation tails evoked by step hyperpolarization from -20 to -50 mV were 14.1 ± 1.3 ($n=17$) and 15.9 ± 1.5 ($n=18$) pA pF⁻¹ for CD38^{+/+} and CD38^{-/-} mice respectively; see Fig. 4A and B. I-V curves were also comparable (not illustrated). However, the inhibition of the M current induced by 10 μ M Oxo-M was dramatically reduced in cells from CD38^{-/-} mice, as shown in Fig. 4B-D, from $74.9 \pm 2.1\%$ ($n=7$) in CD38^{+/+} cells to $28.0 \pm 2.2\%$ ($n=4$) ($p < 0.001$) in CD38^{-/-} (Fig. 4D). The time course of development and recovery of M current inhibition was similar in both sets of cells (Fig. 4C).

{Fig. 4 near here}

NGM1 cells over expressing dual cysteine mutant CD38

We next wanted to test the effect of over-expressing the C119K/C201E mutated human CD38/ADP-ribosyl cyclase cDNA, which is devoid of cADPR hydrolase activity and hence functions only as a cADPR synthesizing enzyme (Tohgo et al., 1994; see Introduction). For these experiments we used NGM1 neuroblastoma hybrid cells that has been stably transformed to express the M1 muscarinic acetylcholine receptor (Fukuda et al., 1988). These cells were transfected with a cDNA plasmid encoding the mutated CD38 gene and selected in the presence of 200 $\mu\text{g ml}^{-1}$ zeocin for 2-6 weeks. The presence of the hCD38 mRNA was verified by Northern blot analysis in zeocin-resistant clones. Expression of the human CD38 protein in the CD38 RNA-positive clones was examined by Western blot analysis using a mouse anti-hCD38 monoclonal antibody (data not shown). Among cells showing a protein band at ~47 kDa for the mutant CD38, three clones designated NGM1-mCD38a, NGM1-mCD38b, and NGM1-mCD38c were selected. Mock- or vector-transformed cells were designated NGM1-V and those transformed with the wild type human CD38, NGM1-CD38.

The efficiency with which the dual cysteine mutant CD38 had been transfected was also confirmed by immunocytochemical analysis (Fig. 5). hCD38-specific immunoreactivity was detected in NGM1-mCD38a, b and c cells (Fig. 5F, H, and J), as in NGM1-CD38 cells (data not shown). Wild type NGM1 or mock-transfected NGM1-V cells showed no or little specific staining under the same conditions (Fig. 5B and D). Our fluorescence microscopic observation did not identify the exact location of the hCD38-immunoreactivity –i.e., whether it was in the cell surface and intracellular membranes or in the perinuclear regions (as shown in MC3T3.E1 CD38-transfected fibroblast cells: Adebanjo et al., 1999). Since thiol-sensitivity of CD38, and its ability to oligomerize have been reported (Grimaldi et al., 1995), we exposed pre-fixed intact cells to 0.5 or 1 mM HgCl_2 for 2 min. This resulted in the enhancement of hCD38-specific staining at the plasma membrane area (Fig. 5K). All control (NGM1) or mock-transfected (NGM1-V) cells, as well as hCD38-transfected (NGM1-CD38) cells, possessed endogenous mouse CD38 (Fig. 5L), which was detected with an anti-mouse CD38 monoclonal antibody in Western blot analysis (data not shown).

{Fig. 5 near here}

ADP-ribosyl cyclase activities were studied by [^3H]cADPR formation from [^3H]NAD $^+$ as the substrate in all clones tested. The basal cyclase activity in NGM1-mCD38a, b, and c cells was 5.7-, 12.7-, and 12.1-fold higher than that ($116 \pm 14 \text{ pmol min}^{-1} \text{ mg}^{-1}$ of protein, n=24) of NGM1 cells and 0.74-, 1.64-, and 1.56-fold of that ($899 \pm 330 \text{ pmol min}^{-1} \text{ mg}^{-1}$ of protein, n=7)

of NG cells, most of which were increased in a clone-specific manner.

The first striking feature in hCD38 transfected cells was the higher rate of [³H]cADPR formation over [³H]ADPR, as shown in their time courses (Fig. 6). In NG control neuroblastoma hybrid cells, NGM1 M1-transformed NG cells and NGM1-V vector-transfected cells, [³H]ADPR formation was much greater than [³H]cADPR production (Fig. 6 A-C), probably due to the endogenous activity of cADPR hydrolase or NAD glycohydrolase (of CD38). Conversely, in C119K/C201E-CD38 transfected NGM1-mCD38a, b, and c cells, [³H]cADPR generation was higher, compared with the conversion rate to [³H]ADPR (Fig. 6D-F). As a cross reference, NGM1-CD38 cells transformed to express wild-type hCD38 with both cyclase and hydrolase activities exhibited the higher [³H]cADPR generation in two clones, but not in other three clones (data not shown). Thus, this clone-specific variation is the reason why we did not use NGM1-CD38 clones in M-current analysis in subsequent experiments.

{Fig. 6 near here}

CCh application to mutant CD38 expressing cells did not produce any activation of ADP-ribosyl cyclase activity, but instead a slight inhibition, to 87-93% of the control level. This was the case in NGM1-CD38 cells with over-expressed wild-type CD38 (data not shown). As shown previously (Higashida et al., 1997), endogenous ADP-ribosyl cyclase activity was increased in M1 mAChR over-expressing NGM1 cells and was inhibited by stimulating endogenous M4 mAChRs in NG cells by 10 μ M CCh. The absence of equivalent effects in CD38-transformed cells was not due to loss of M1 mAChRs, because the cells gave clear Ca²⁺ signals on mAChR stimulation (see below). One possibility is that ADP-ribosyl cyclase activity in mutant CD38-transfected cells is mostly derived from the exogenous CD38, which may not be the type of cyclase controlled by mAChRs.

Effect of muscarinic agonists on intracellular calcium in mutant CD38-expressing NGM1 cells

It has previously been shown that ACh produces large transient rises in [Ca²⁺]_i in M1 mAChR-transfected NG108-15 cells, as a result of phospholipase C-stimulated InsP₃ formation (Fukuda et al., 1988 and 1989). This provides a convenient measure of mAChR stimulation and downstream signaling. All three types of NG cells used in this study possessed comparable levels of resting [Ca²⁺]_i, between 69 and 77 nM. Application of 10 μ M CCh elicited an increase to 171 \pm 30% (n=9) of the pre-exposure level in NGM1 cells. The same or a greater increase to 286 \pm 45% (n=4), 232 \pm 16% (n=7) and 251 \pm 40% (n=5) was observed in NGM1-

mCD38a, b and c cells, respectively. Hence, receptor activation and downstream signaling were maintained in cells expressing the constitutively-active form of hCD38.

To address the question whether the intracellular distribution of the ACh-induced Ca^{2+} rise in C119K/C201E-CD38 over-expressing cells differed from that in control cells, we applied digital imaging analysis by confocal microscopy. Line scans were made in mutant CD38 over-expressing cells or wild-type NG cells loaded with the Ca^{2+} indicator, OGB-1, after local application of 100 μM muscarine. Although the magnitude of the rise was greater in NGM1-mCD38b cells, the elevation of $[\text{Ca}^{2+}]_i$ started from the submembrane compartment beneath the point of application and then spread at the same rate into the deeper regions in both NGM1 and NGM1-mCD38b cells. Thus, the topographical distribution of the sites of Ca^{2+} release was unchanged by the over-expression of mutated CD38.

Carbachol-induced biphasic membrane current response

CCh, when applied to parental NGM1 cells, produces an outward current due to the opening of Ca^{2+} -dependent K^+ channels, followed by an inward current caused principally by suppression of M-current (Fukuda et al., 1988; Robbins et al, 1992; Higashida et al., 1995, 1996). We therefore examined how these two CCh-induced current changes were modified in cells over-expressing mutant CD38.

Focal application of 3 μl of 10 μM CCh to the external surface of NGM1-mCD38a, b, and c cells voltage-clamped at -20 mV, produced either a similar biphasic current response to that in wild-type cells (Fig.7A), or a large inward current with a negligible initial outward current (Fig.7B). The mean outward current elicited by CCh in NGM1-mCD38a, b, and c cells appeared somewhat smaller than that in NGM1 and NGM1-V cells (Fig. 8A), whereas the mean inward current was similar or somewhat larger (Fig. 8B).

{Figs. 7 and 8 near here}

Inhibition of the M-like current $I_{\text{K(M,ng)}}$ (Robbins et al., 1992) at the peak of the second-phase inward current following local application of 3 μl of 10 μM acetylcholine (to give an estimated equilibrium concentration of 15 nM) was measured from the reduction in the amplitude of the deactivation relaxations that accompanied 20 mV hyperpolarizing steps from the holding potential of -20 mV (Fig.7A, B, lower records). The average inhibition in NGM1 and NGM1-V cells was $32.1 \pm 6.5\%$ (n=18) and $26.3 \pm 4.7\%$ (n=19), similar to that previously reported (Higashida et al, 1996). In contrast, the same treatment in NGM1-mCD38a, b and c cells produced significantly larger inhibitions ($p < 0.001$) up to $83.2 \pm 12.9\%$ (n=5), $80.3 \pm$

3.6% (n=23), and $80.8 \pm 6.2\%$ (n=8), respectively (Fig. 8C). Judging from the dose-response relationship obtained by perfusion of various concentrations of ACh (Fig. 9), the IC_{50} of ACh was calculated to be about 1.55 and 0.88 μ M ACh for NGM1-mCD38b and c cells, compared with 13 μ M for wild-type NGM1 cells.

{Fig. 9 near here}

Finally, we examined whether the voltage-dependent characteristics of M currents, *per se*, were altered in mutant CD38 expressing cells. Therefore control (NGM1), mock-transfected (NGM1-V) and the dual cysteine mutant CD38 transfected cells (NGM1-mCD38b) were recorded under identical conditions to measure $I_{K(M,ng)}$ in the whole-cell voltage-clamp mode. The total amplitude of the deactivation relaxation measured at -50 mV after a step from -20 mV was found to be not significantly different between the three cell types. Construction of an activation curve followed by a fit to a Boltzman function (see Robbins et al., 1992) demonstrated no significant difference between the half activation potential (V_0) or the slope value (s) among all three groups of cells (Table 1). Construction of a time constant- V_m curve (Robbins et al., 1992) also indicated no significant difference between the opening (α_0) or closing (β_0) rate constants or the slope factor (c). Thus, over-expression of the mutated CD38 had little effect on the gating, voltage dependence or kinetics of the channels responsible for $I_{K(M,ng)}$. [Note: for these analyses, a composite time-constant embracing both components of $I_{K(M,ng)}$ current relaxation (see Meves et al., 1999; Selyanko et al., 1999) were applied (see Robbins et al., 1992). The fact that this composite time-constant was unchanged implies that the two components of current carried by KCNQ and ERG channels were affected equally.]

{Table 1 near here}

Discussion

The results of previous experiments (see Introduction) have indicated that cADPR may play a role in inhibition of the M-like current in NG108-15 neuroblastoma hybrid cells that results from M1 mAChR stimulation. In this report we have investigated this possibility further by the unique approach of genetically manipulating the enzyme that may be responsible for generating cADPR, human CD38. This has led to two significant advances.

First, over-expression of a mutated form of the cADPR synthetic enzyme, CD38 /ADP-ribosyl cyclase, that was devoid of hydrolase activity (and therefore acts as a constitutively-active ADP-ribosyl cyclase, leading to higher levels of cADPR) strongly augmented the

inhibition of the M-like current in NG108-15 cells. This was specific to this component of M1-mAChR action since there was no corresponding increase in the initial elevation of intracellular Ca^{2+} or in the consequential outward current mediated by activation of Ca^{2+} -dependent K^+ channels (see Higashida & Brown, 1986; Brown & Higashida, 1988; Robbins et al., 1991 and 1993). Interestingly (and in contrast to control cells: Higashida et al., 1997), activation of the M1 receptors did not appear to enhance the activity of this mutated CD38 enzyme. Hence, the enhanced inhibition of the M-like current appears to result from the higher overall basal rate of ADP-ribosyl cyclase activity.

Second – and perhaps more significantly – we have extended the putative role for cADPR in M current regulation to include primary sympathetic neurons. Thus, we have shown that ACh accelerates the synthetic activity of ADP-ribosyl cyclase in membrane preparations from sympathetic ganglia, just as it does in NG108-15 cell membranes (Higashida et al., 1997). Tests with antagonists showed that this was due to stimulation of endogenous M1 mAChRs, and tests with GTP and GTP- γ -S indicated that it resulted from M1 mAChR-G protein coupling.

Finally, using CD38 knock-out mice, in which negligible ADP-ribosyl cyclase activity could be detected in the membrane fractions prepared from sympathetic ganglia, we found a very substantial reduction in the mAChR-induced inhibition of the native M currents in dissociated sympathetic neurons in CD38^{-/-} mice when compared with that in neurons from wild-type CD38^{+/+} mice, from ~75% to ~25%.

Some of the experimental manipulations previously used to investigate the role of cADPR in neuroblastoma cells – for example, intracellular dialysis of cADPR and its antagonists – are difficult to apply to sympathetic neurons because of the rapid M current ‘rundown’ usually experienced with whole-cell recording (see, for example, Brown et al., 1995). This has hitherto precluded extension of information drawn from NG108-15 cells to primary neurons. The present experiments are therefore particularly important in suggesting that such an extension can reasonably be made.

Nevertheless, when viewing cADPR as a potential ‘second messenger’ for M current inhibition (see further below), a number of questions remain. First, CD38 is a multifunctional ectoenzyme (Shubinsky and Schlesinger, 1997), which generates cADPR from β -NAD⁺ in the extracellular compartment. This conflicts with the established dogma that second messengers are generated and remain intracellularly (with the exception perhaps of nitric oxide). This apparent anomaly might be explained in a number of ways. First, perhaps it is extracellular cADPR that modulates the M-current. This seems unlikely, because cADPR has been shown to inhibit the neuroblastoma current when applied intracellularly via the patch-pipette (Higashida et al., 1995) and its antagonists are also effective from the inside (Bowden et al., 1999). Second,

it has been suggested that some of the CD38 protein may exist on intracellular membranes and therefore generate cADPR within the cell (Kato et al., 1995; Chini et al., 2002; Khoo and Chang, 2002). Thirdly it has been proposed by Zocchi et al. (1999) that plasma membrane-located CD38 internalises β -NAD⁺ into vesicles and, following conversion of β -NAD⁺ to cADPR, pumps cADPR into the cytosol. Similar considerations apply to the proposed involvement of cADPR in other cellular responses such as fertilization, insulin secretion and cardiac contraction (Galione et al., 1993; Lee et al., 1993; Takasawa et al., 1993a; Higashida et al., 1999, 2000b) and clearly need resolution.

Second, the *modus operandi* of cADPR as a potential M current inhibitor is not yet clear. The most widely-accepted view is that it acts by facilitating ryanodine receptor-induced intracellular Ca²⁺ release (Meszaros et al., 1993; Lee et al., 1994; Thomas et al., 2001). However, in previous experiments, no evidence could be adduced to support a role for either Ca²⁺ (Higashida et al., 1995) or ryanodine receptors (Bowden et al., 1999) in the regulation of the M-like currents in neuroblastoma hybrid cells. Equally, an alternative possibility – that of a direct effect on the M channels themselves – can be excluded because cADPR did not affect M channel activity when applied to the inside face of excised membrane patches from sympathetic neurons (Bowden et al., 1999).

Notwithstanding this paucity of mechanistic information, the experimental evidence for cADPR as a putative ‘second messenger’ for G protein-coupled receptor-induced M current inhibition now appears quite substantial (see Introduction), and comparable to that for some other suggested messengers such as Ca²⁺ and protein kinase C (see Marrion, 1997). Recent work has added phosphatidylinositol bisphosphate (PIP₂) depletion to that list of putative second messenger systems (Suh & Hille, 2002). One possibility that emerges is that changes in membrane phosphatidylinositide levels provide the unifying mechanism underlying the regulation of M channels by Gq/11-coupled receptors in general, but that other modulators generated by receptor stimulation enhance the sensitivity of the channels to changes in PIP₂ levels in a receptor-specific manner. Thus, for bradykinin B2 receptors (Cruzblanca et al., 1998) and nucleotide P2Y receptors (Bofill-Cardona et al., 2000), closure is facilitated by the InsP₃-induced rise in Ca²⁺ (perhaps resulting from a close proximity of the transmitter receptors to the InsP₃-receptors: Delmas et al., 2002) and the consequent effect of Ca²⁺ on channel activity (Selyanko & Brown, 1996). On the other hand, formation of diacylglycerol and consequent activation of protein kinase C bound to the A-kinase anchoring protein (AKAP) in a complex with the M channel acts as a sensitizer for mAChR-mediated inhibition (Hoshi et al., 2003). The present results strongly suggest that ADP-ribosyl cyclase activation, and consequential formation of cADPR, also falls into this category of ‘sensitizing mechanisms’.

An essential requirement for future research is to determine more precisely the interactions between the various regulatory mechanisms for these channels.

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Table 1. Amplitude and kinetic properties of $I_{K(M,ng)}$ in control and cysteine mutated CD38-transfected NGM1 cells.

Parameter	NGM1- $I_{K(M,ng)}$		
	NGM1 (n=9)	NGM1-V (n=8)	NGM1-mCD38b (n=9)
Amplitude (pA)	524 ± 85	1272 ± 408	700 ± 127
V_o (mV)	-38.9 ± 1.61	-36.3 ± 1.5	-38.5 ± 0.5
s	-7.7 ± 1.6	-6.5 ± 0.6	-6.8 ± 0.5
α_o (s ⁻¹)	1.9 ± 0.4	1.1 ± 0.2	1.0 ± 0.2
β_o (s ⁻¹)	1.4 ± 0.1	1.2 ± 0.1	1.2 ± 0.2
C (mV)	15.0 ± 0.8	15.9 ± 0.6	15.5 ± 0.6

Biophysical properties of $I_{K(M,ng)}$ in control (NGM1), vector transfected (NGM1-V) cells and dual cysteine mutant CD38 (NGM1-mCD38b) cells. Data are given as the mean ± s.e.m. for the deactivation tail amplitude at -50 mV, half activation potential (V_o), slope (s) obtained from activation curves fit by Boltzman equation $G/G_{max} = 1/\{1+\exp[V_{1/2}[(V_o-V_c)/s]]\}$. The opening (α_o) and closing (β_o) rate constants were calculated from tau- V_c curve fit by the equations $\alpha = \alpha_o \exp((V_c-V_o)/C)$ and $\beta = \beta_o((V_c-V_o)/-C)$. Number of cells is given in brackets.

Figure legends

Fig. 1. Activation of ADP-ribosyl cyclase by ACh in rat SCG. A. ADP-ribosyl cyclase was measured from the initial velocity during 2 min with membranes prepared from rat SCG. Values were calculated by conversion of [³H]cADPR from [³H]NAD⁺ in the presence or absence of 100 nM ACh in 4 experiments. Values indicate mean ± S.E.M. * indicates significantly different from the control at $p < 0.01$. B. Relationship between ACh concentration and ADP-ribosyl cyclase activity with (open diamonds) or without 1 μM atropine (filled circles) measured fluorometrically. The fluorescence of the resulting cGDPR was continuously monitored before and after application of 100 nM ACh (arrow), as shown in insert. Changes in velocity were plotted. Data represent the mean (± S.E.M.) in 3-5 measurements. Some S.E.M. values are within the size of symbols. * and ** significantly different from the value with atropine at the indicated ACh concentrations at $p < 0.001$ and 0.01, respectively. *Insert*, Bars indicate 30 s and 0.1 arbitrary unit.

Fig. 2. Effects of subtype-specific antagonists on ACh-induced activation of ADP-ribosyl cyclase activity in rat SCG. A. Relationship between ACh concentration and ADP-ribosyl cyclase activity with (open squares) or without 1 μM pirenzepine (filled circles) measured fluorometrically. Values indicate mean ± S.E.M. in 3-5 measurements. * significantly different from the value with pirenzepine at the indicated ACh concentrations at $p < 0.01$. B. Effects of 4 subtype-specific antagonists at 1 μM on ACh (100 nM)-induced activation of ADP-ribosyl cyclase. Values indicate mean ± S.E.M. in 3-5 measurements. * significantly different from control value at $p < 0.001$.

Fig. 3. Synthesis of cGMP-ribose and dose-dependent effect of ACh in SCG membranes of CD38^{+/+} wild type and CD38^{-/-} knockout mice. cGDPR synthesis was measured fluorometrically. A. Membrane preparations of SCG of CD38^{+/+} and CD38^{-/-} mice were incubated for 700s and basal activities (arbitrary units min⁻¹ mg protein⁻¹) were calculated. * significantly different from the value of wild type mice at $p < 0.001$ (n=7). B. Membranes were incubated with different concentrations of ACh and the velocity was calculated. Results are expressed as percentage of control rate of cGDP-ribose in 4 independent experiments. * and ** significantly different from values at the indicated ACh concentrations in CD38^{-/-} mice at $p < 0.002$ and 0.001 respectively.

Fig. 4. Comparison of M-current inhibition in cultured SCG neurons of wild-type CD38^{+/+} or CD38^{-/-} mutant mice. Typical current responses to a voltage step to -50 mV from holding

potential of -30 mV for M currents in CD38^{+/+} (A) or CD38^{-/-} (B) mice before (trace 1) during (2) and after (3) application of 10 μ M Oxo-M. C and D. Statistical evaluation of the effect of Oxo-M on M-current in both types of mice. Depicted are the amplitudes of M-current deactivation relaxations in response to the voltage steps in A and B and inhibition of M-current. Means \pm S.E.M. from between 4 and 7 experiments in CD38^{+/+} and CD38^{-/-} mice. *significantly different at $p < 0.005$.

Fig. 5. Immunofluorescence of human or mouse CD38 in various types of cells derived from NGM1 cells. Using a fluorescence microscopy, FITC for NGM1, (A, B), NGM1-V (C, D), NGM1-mCD38a (E, F), NGM1-mCD38b (G, H), and NGM1-mCD38c (I, J and K) cells. Images obtained with (B, D, F, H, J, and K) or without (A, C, E, G and I) anti-human CD38 mouse monoclonal antibody. (magnification, $\times 756$). In K, the cells were treated with 1 mM HgCl₂ for 2 min in order to facilitate membrane localization. L, immunofluorescence of endogenous CD38 obtained with anti-mouse CD38 monoclonal antibody in NGM1. Note that fluorescence is seen in cell membrane and processes. (magnification, $\times 1512$).

Fig. 6. Time course of ADP-ribosyl cyclase activity in membranes prepared from various types of NG108-15 cells. Aliquots (20 μ l) containing 1.5 - 4.5 μ g of membrane protein of NG (A), NGM1 (B), NGM1-V (C), NGM1-mCD38a (D), NGM1-mCD38b (E), and NGM1-mCD38c (F) cells were incubated for the indicated time (0 - 2 min). Radioactivity in spots corresponding to authentic cADPR (filled circle) and ADPR (open square) on TLC sheets was measured. Each data point represents the mean of two independent experiments.

Fig. 7. M-current suppression in two NGM1-mCD38b cells. Top, inward current response to focal application of CCh (3 μ l, 10 μ M), being associated with (A) or without (B) initial outward current. Repetitive hyperpolarization step pulses (-20 mV, 1 s) were applied to estimate the conductance and M current from a holding membrane potential of -20 mV (Middle). Bottom, Expanded records of current transients evoked by ach -20 mV step pulse. They were obtained at the time gap in the top trace. Left (a), before applying CCh; centre (b), during the inward current; right (C and D), after full or over recovery.

Fig. 8. ACh-induced outward and inward currents and inhibition of M current in various types of cells derived from NGM1 cells. Histograms show the average amplitude of outward (A) and inward (B) currents induced by focal application of 100 μ M CCh (3 μ l) in cells indicated on left. C shows the average amount of inhibition of $I_{K(NG, M)}$ recorded in cells as

described in Fig. 4D. Number of cells tested ranged from 5 to 6. Bars, S.E.M. * $p < 0.001$ from two control cells.

Fig. 9. The concentration-dependent inhibition of M-current by acetylcholine in NGM1 (filled circle), NGM1-mCD38b (open square) and NGM1-mCD38c (open diamond) cells. The relative inhibition of the M-current deactivation relaxations produced by the increasing concentration of ACh in three different subclones was plotted. Relative $I_{K(M,ng)}$ was calculated by dividing the amplitude prior to application with that obtained after the perfusion of the indicated concentrations of ACh. Each point represents the mean (\pm S.E.M.) of 4-5 cells.

Figure 1

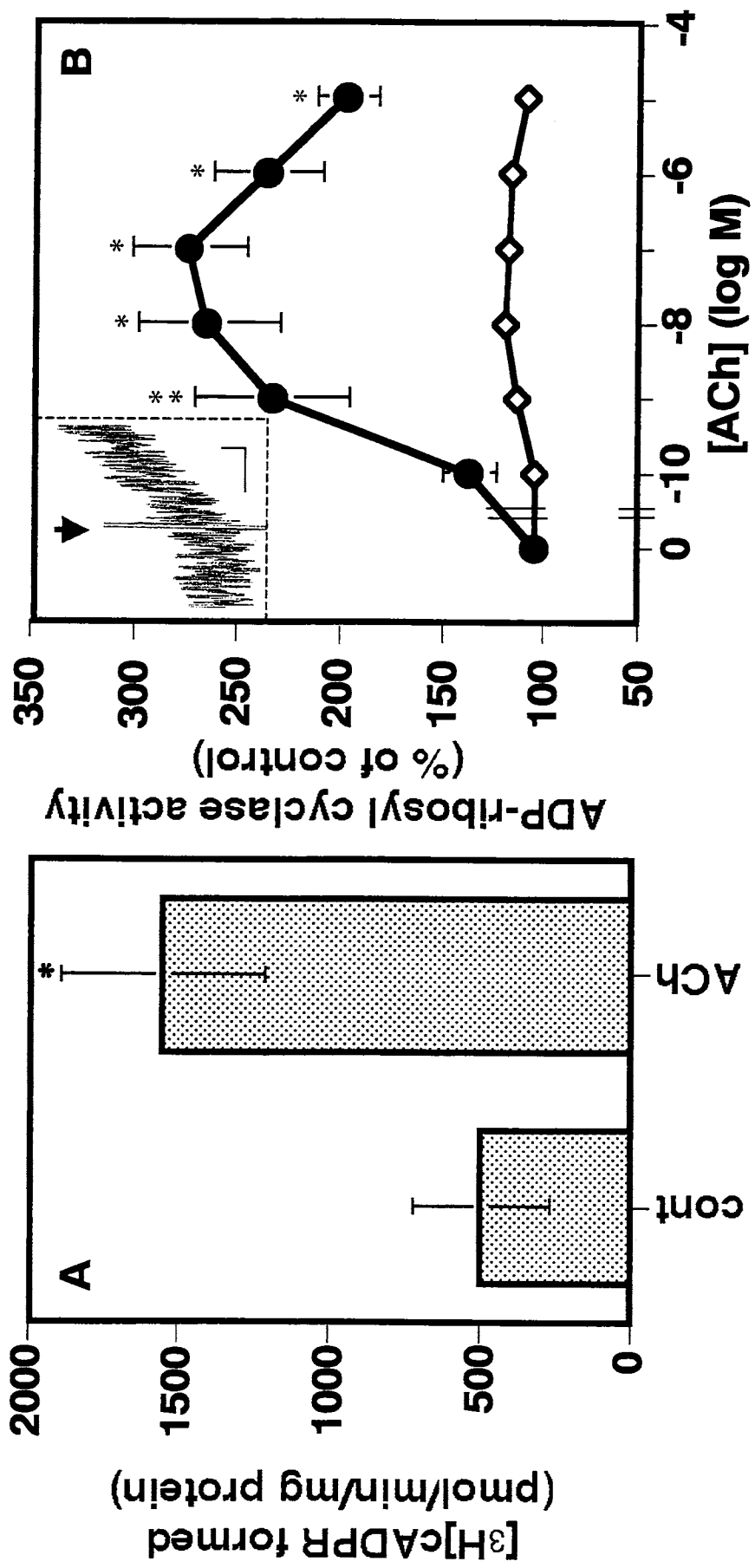


Figure 2

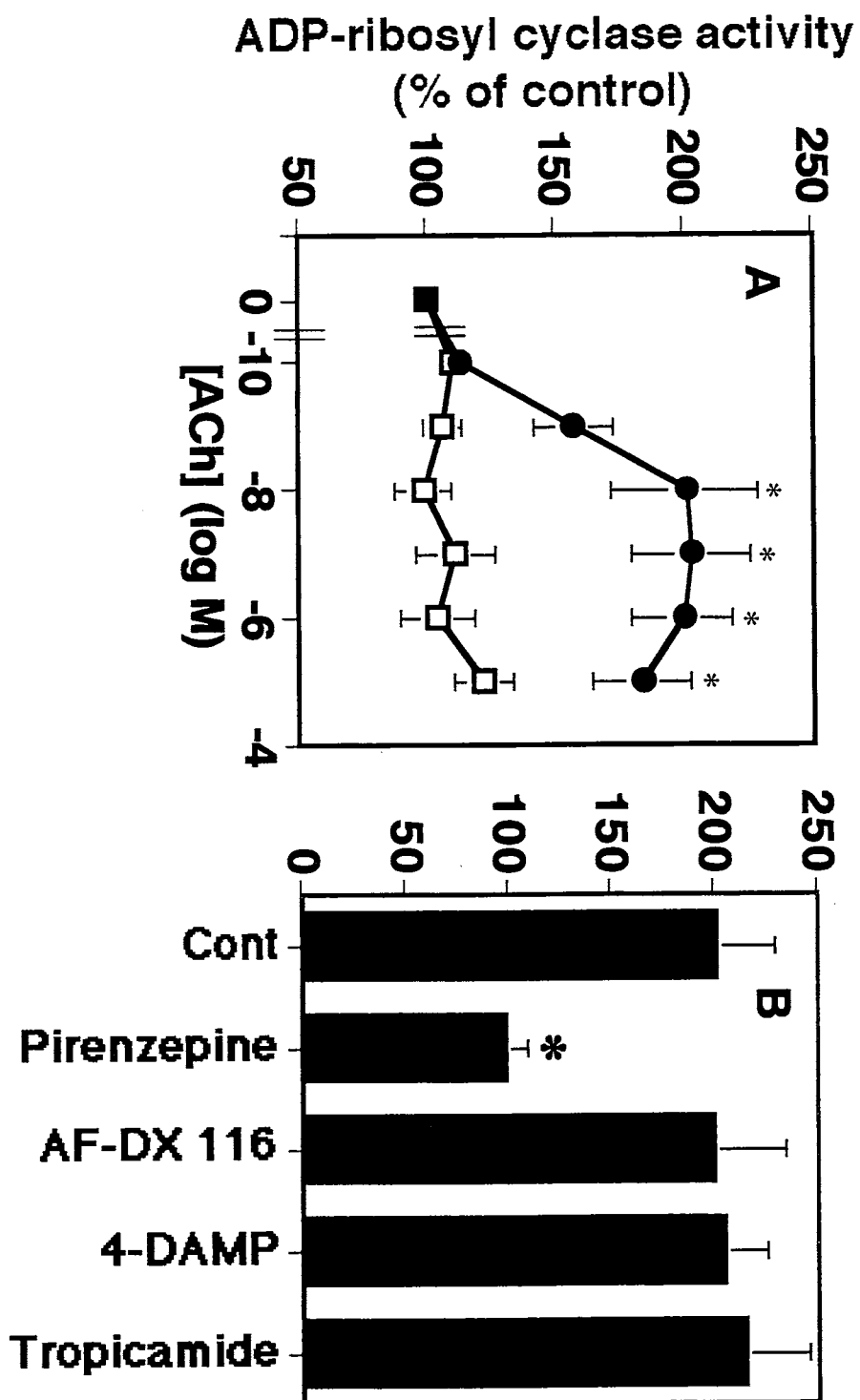
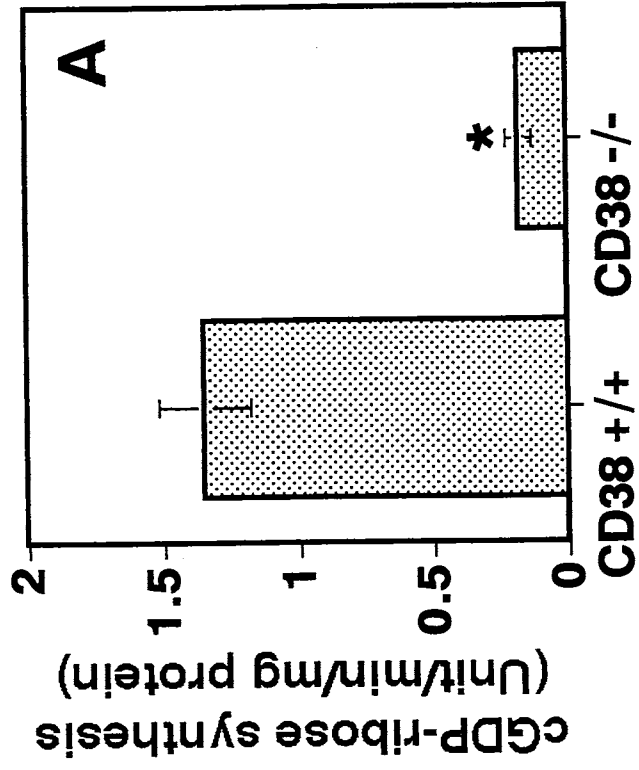
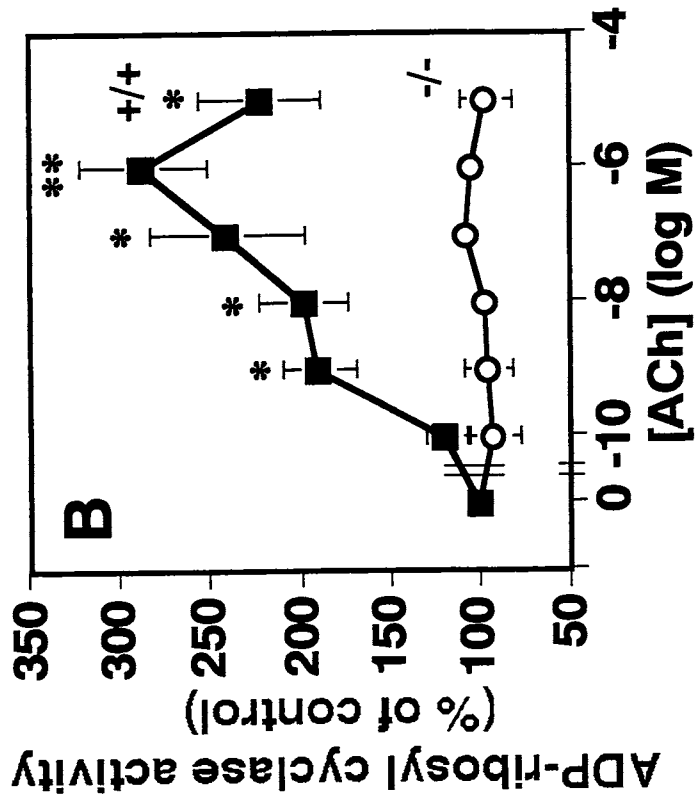
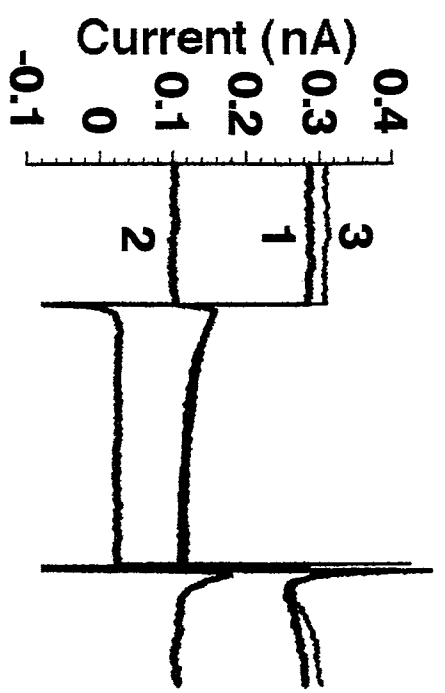


Figure 3



A CD38^{+/+}



B CD38^{-/-}

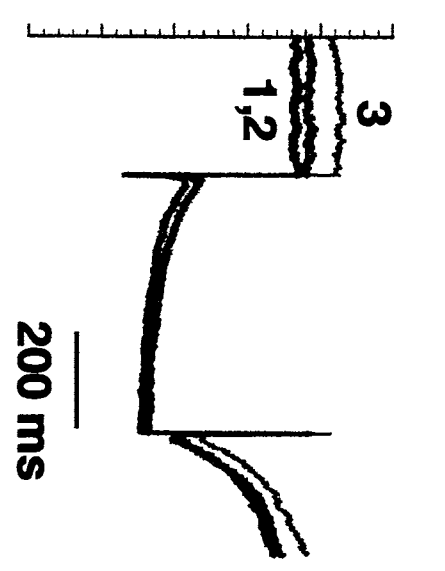


Figure 4

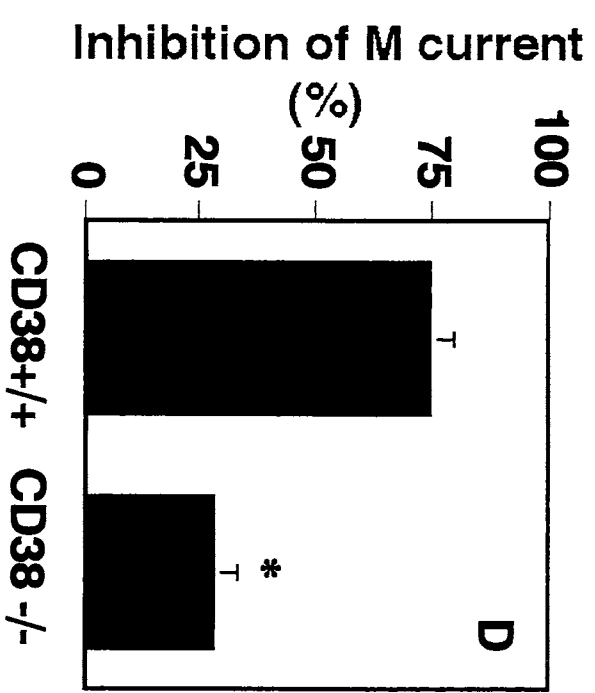
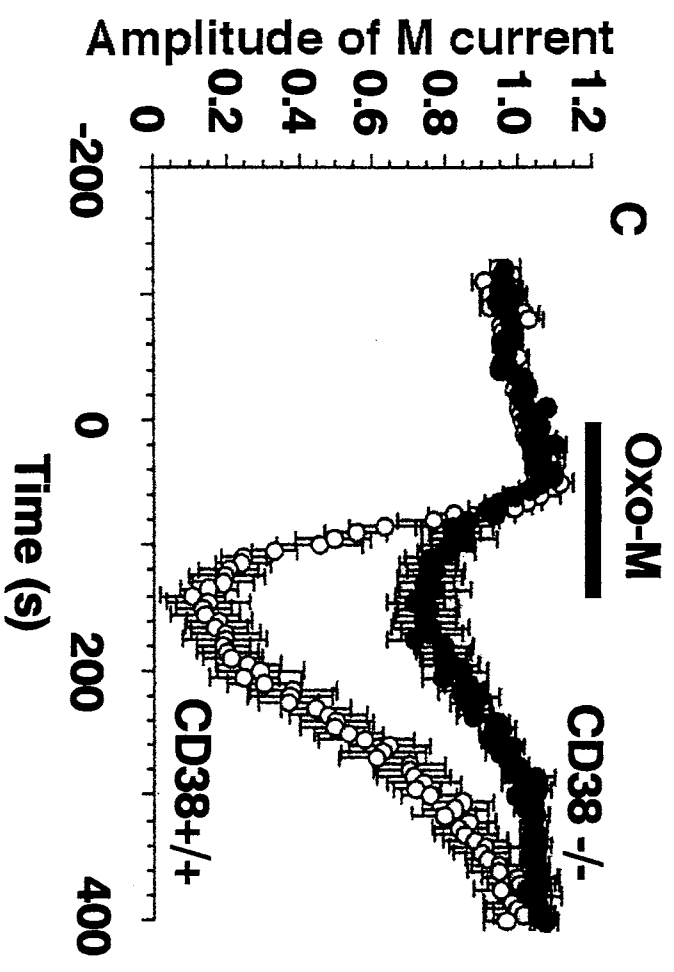


Figure 5

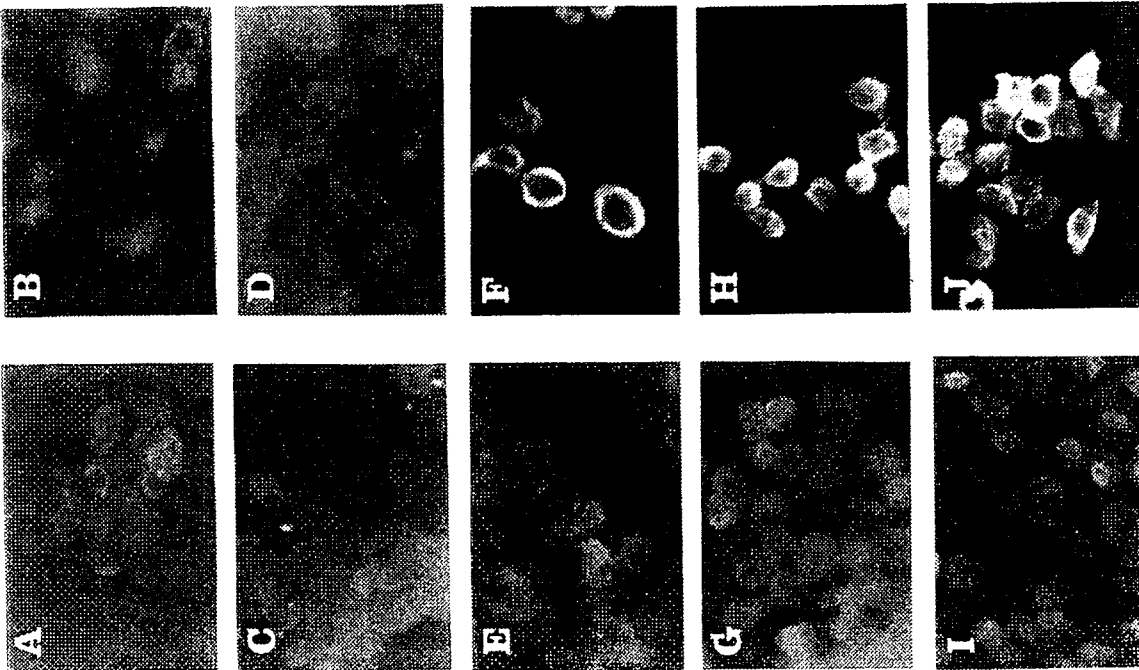
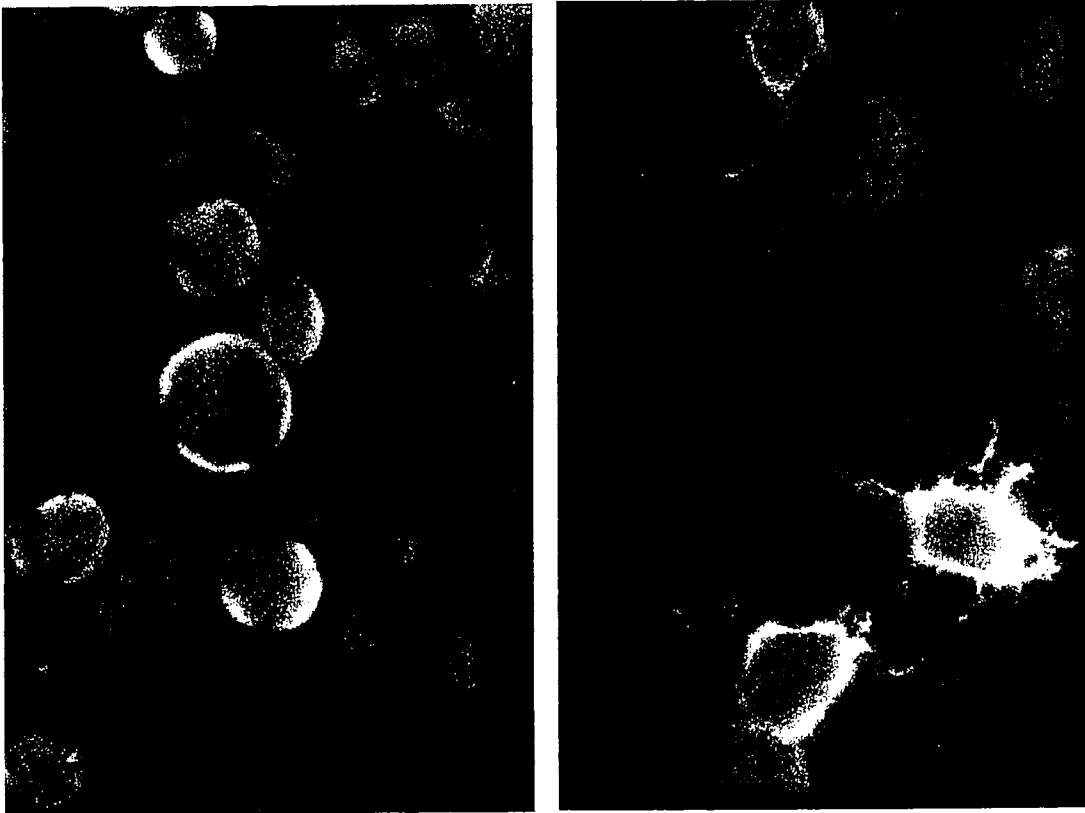


Figure 6

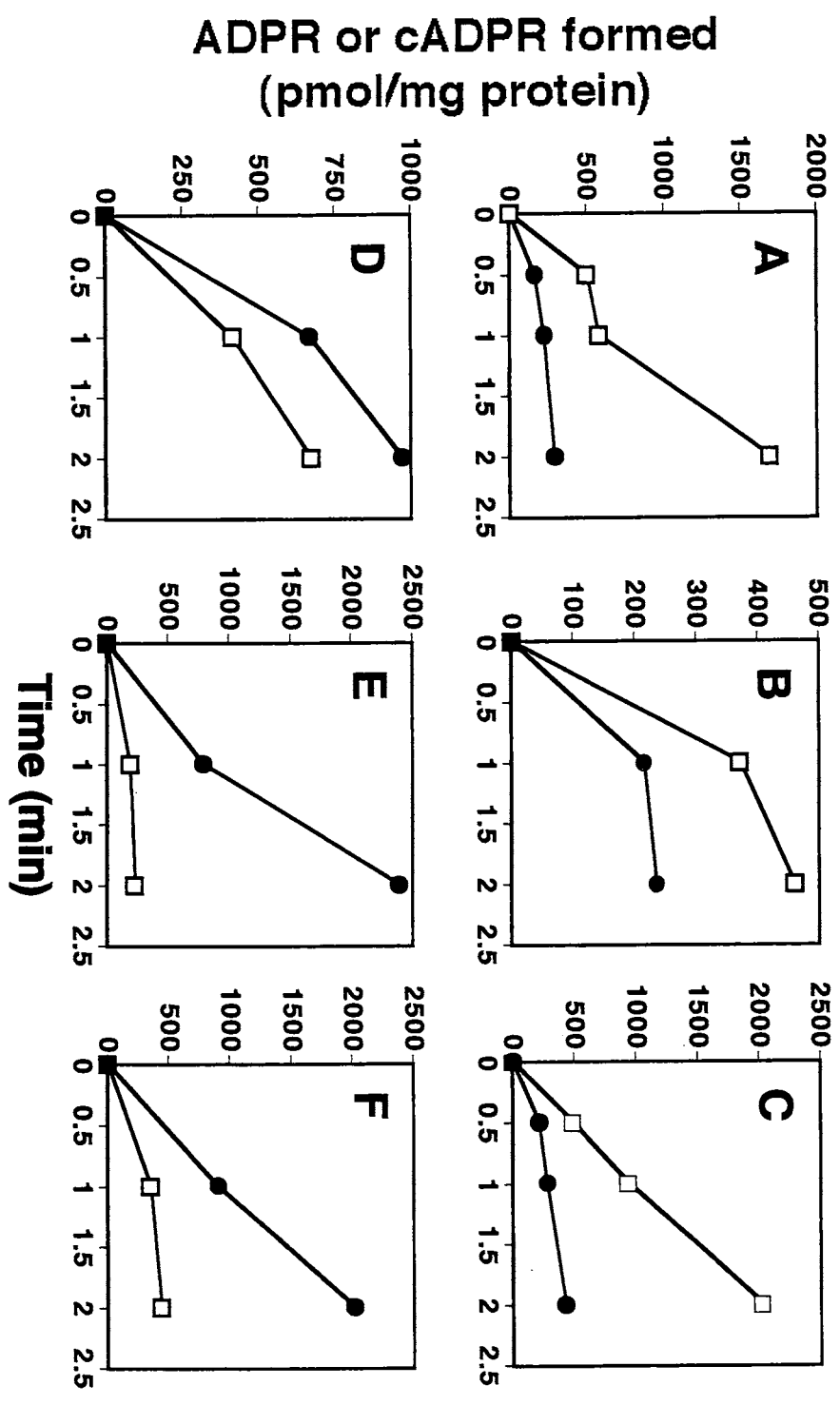


Figure 7

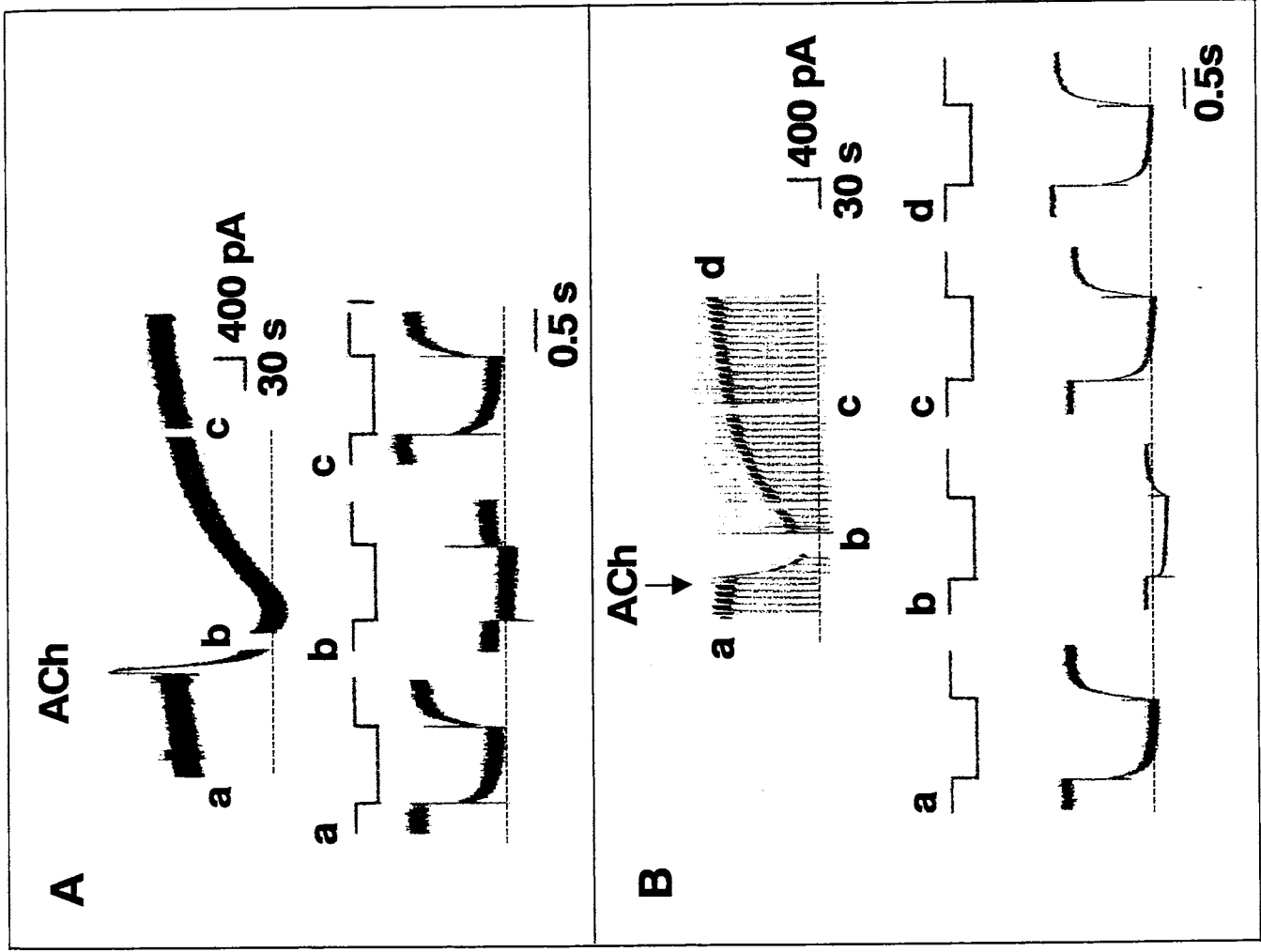


Figure 8

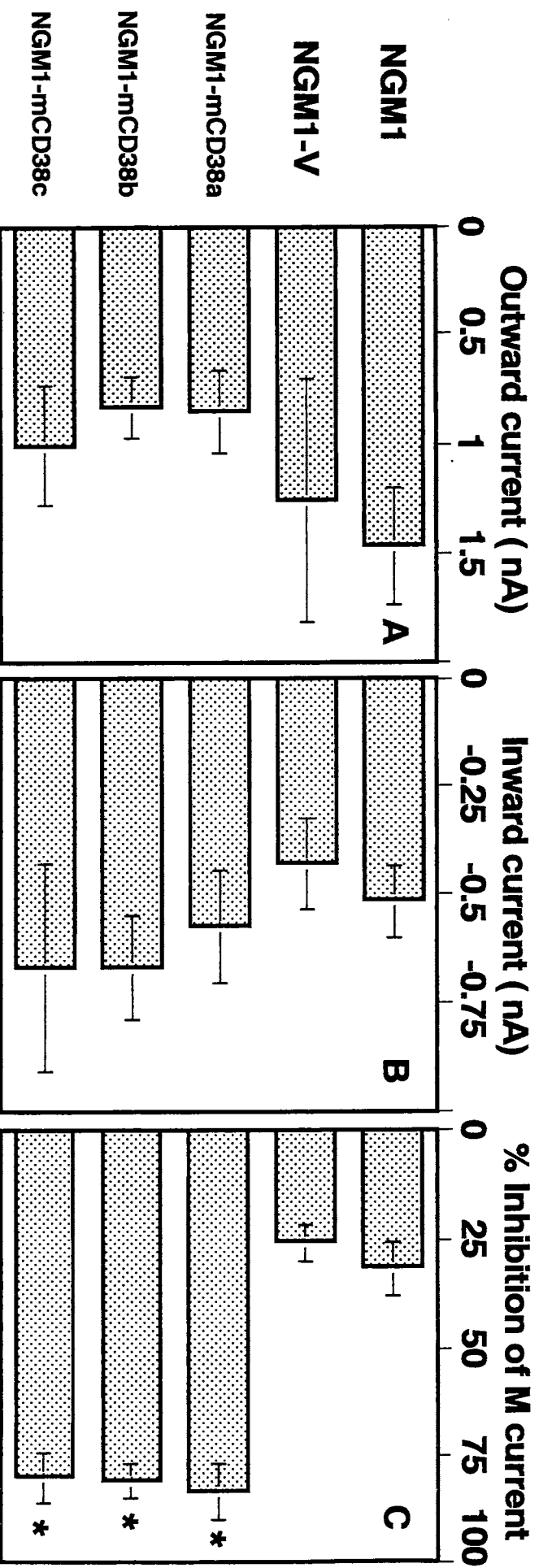
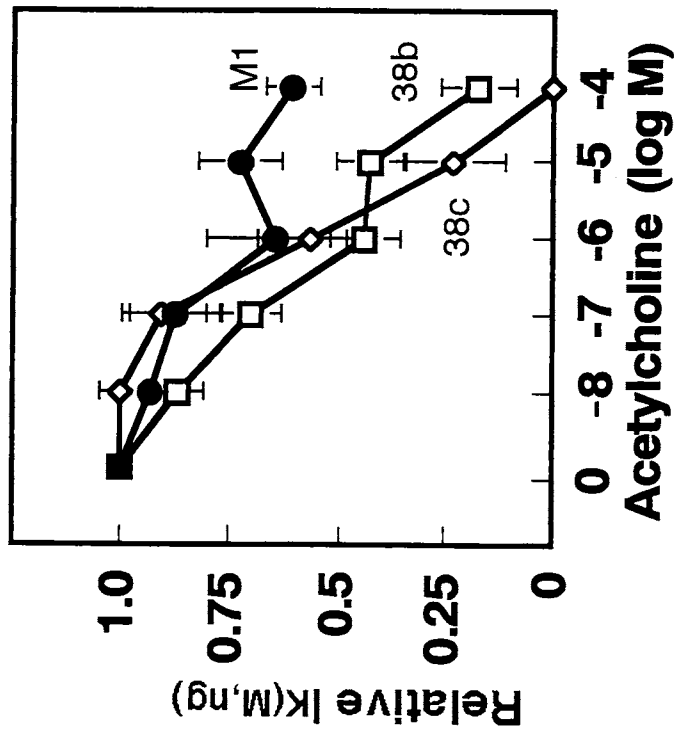


Figure 9



7. 謝辞と展望

本研究は、京都大学中西重忠教授、大阪大学福田 淳教授、生理学研究所重本隆一教授、東北大学岡本 宏教授、九州大学野田百美助教授、ロンドン大学 DA Brown 教授・Robbins 講師等の御支援を賜った。深くお礼申し上げます。

前回の重点 B の報告においても小冊子を作り、日本中広く頒布することを取り止め、その代わり英語で総説を書くと宣言した。その結果、前回の成果を *Journal of Neurochemistry* 76 巻、321–331 頁（2001 年）にまとめて、より広く世界中へ発信した。今回の成果も、一流の Review 誌への掲載を行う覚悟である。