

The G_q and G₁₂ Families of Heterotrimeric G Proteins Report Functional Selectivity

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ABSTRACT

Receptors coupled to the G_q and G₁₂ families of heterotrimeric G proteins have surfaced rarely in the context of functional selectivity and always indirectly. We explore here the differential engagement of G_q and G₁₃ (of the G₁₂ family) by the thromboxane A₂ receptor α (TP_α), via agonist-effected [³⁵S]-guanosine 5'-O-(3-thio)triphosphate binding when the G proteins themselves are used as reporters. We find for TP_α introduced into human embryonic kidney 293 cells and for the receptor expressed normally in human platelets an agonist-selective engagement of G_q versus G₁₃. Pinane thromboxane A₂ (PTA₂) activates G_q in preference to G₁₃, whereas 8-iso-prostaglandin F_{2α} activates G₁₃ in preference to G_q. 9,11-

Dideoxy-9α,11α-methanoepoxy-prosta-5Z,13E-dien-1-oic acid (U46619), in contrast, exhibits no preference. Reserve of receptor in relation to G protein and of G protein in relation to downstream events is apparent in some instances but does not have a bearing on selectivity. Activation of G proteins by PTA₂ is right-shifted from binding of the ligand to receptor, a manifestation of which is a bimodal action: PTA₂ is an antagonist at low concentrations and an agonist at higher concentrations. We posit two populations of TP_α, or two intrinsic sites of ligand binding, with selectivity evident not only in terms of the G proteins activated but properties of antagonism versus agonism.

Functional selectivity refers to changes in the conformation of a receptor that are uniquely induced and perceived, respectively, by ligands and effectors specific to the receptor (for reviews, see Perez and Karnik, 2005; Urban et al., 2007). Selectivity is most often evident as differences in the rank-order of ligands in relation to efficacy and/or potency for distinct downstream events. In the case of 7-transmembrane domain receptors, events of interest are those set into motion by heterotrimeric G proteins, alone or in combination with those engaged by arrestins (Violin and Lefkowitz, 2007).

A great deal of information regarding selectivity has been gained through studies of receptors for serotonin (Berg et al., 1998, 2001; Kurrasch-Orbaugh et al., 2003; De Deurwaerdere et al., 2004), dopamine (Lawler et al., 1999; Kilts et al., 2002; Mottola et al., 2002; Gay et al., 2004; Ryman-Rasmussen et al., 2005), and opiates (Keith et al., 1996, 1998; Whistler et al., 1999; Alvarez et al., 2002), wherein measurements of selectivity as they pertain to G

proteins are based on increases and decreases in cAMP, stimulation of phosphoinositide metabolism and consequent increases in intracellular calcium, and release of arachidonic acid. In some cases (for example, changes in cAMP and sometimes phosphoinositide metabolism), the G proteins employed are obvious; in others, they are not. Inferences of selectivity through second messengers are blurred to some extent by activities shared among G proteins, nonlinearity in signal transmission, and the potential in intact cells for cross-regulation of signaling pathways. Several studies have explored, consequently, functional selectivity more directly through measurements of G protein activation, notably using [³⁵S]guanosine 5'-O-(3-thio)triphosphate ([³⁵S]GTPγS) (Cordeaux et al., 2001) (Cussac et al., 2002) (Lane et al., 2007). Studies using G protein activation are important in that they provide a framework for inferences of receptor conformation that are proximal to the receptor yet operational in nature.

The G₁₂ family of heterotrimeric G proteins in vertebrates consists of G₁₂ and G₁₃. The family has received considerable attention in studies of cell function, especially cell contractility and adhesion, with targets comprising the monomeric G

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ABBREVIATIONS: GTPγS, guanosine 5'-O-(3-thio)triphosphate; TXA₂, thromboxane A₂; U46619, 9,11-dideoxy-9α,11α-methanoepoxy-prosta-5Z,13E-dien-1-oic acid; SQ29548, [1S-[1α,2α(Z),3α,4α]]-7-[3-[[2-[(phenyl amino)carbonyl]hydrazine]methyl]-7-oxabicyclo[2.2.1]hept-2-yl]-5-heptenoic acid; PTA₂, pinane-thromboxane A₂; 8-iso-PGF_{2α}, 8-iso-prostaglandin F_{2α}; HEK, human embryonic kidney; TP, receptor(s) for thromboxane A₂; CGP12177, 4-[3-[[1,1-dimethylethyl]amino]-2-hydroxypropoxy]-1,3-dihydro-2H-benzimidazol-2-one.

protein Rho, cadherins, and Tec tyrosine kinases, among others (for review, see Kelly et al., 2007). The biochemical properties of the G_{12} family are notable (Singer et al., 1994; Kozasa and Gilman, 1995). One of the most striking aspects of receptors coupled to G_{12} and/or G_{13} is that they are invariably coupled to other G proteins as well, almost always G_q (Riobo and Manning, 2005). Events elicited through the G_{12} family are perceived, therefore, to be integrated with events achieved through additional families. In platelets, for example, both thrombin and thromboxane A_2 (TXA_2) act on receptors coupled to the G_{12} and G_q families. The initial response of platelets to either of the ligands is a change in shape (rounding) that requires G_{13} (Offermanns, 2006). The change is a prerequisite to degranulation and aggregation, which are achieved subsequently through G_q . The G_q -dependent aggregation is partly mediated through release of ADP by degranulation and subsequent engagement of G_i . Efficient activation of platelets by thrombin or TXA_2 , therefore, requires direct engagement of G_q and G_{13} and indirect engagement of G_i .

Given that receptors coupled to G_{12} and/or G_{13} invariably couple to other G proteins, and that the integration of signaling is important, a question of considerable interest is whether agonists working through these receptors exhibit functional selectivity. No studies toward this end have been conducted. A large part of the difficulty is the absence of quantifiable enzymes and second messengers uniquely regulated by the G_{12} family. In this study, we used [35 S]GTP γ S-binding to evaluate functional selectivity for agonists operating through the thromboxane A_2 receptor $TP\alpha$. The data reveal, for the first time, selectivity in agonists that engage the G_q and G_{12} families of G proteins and reveal as well an unusual bimodal action of a supposed antagonist.

Materials and Methods

Materials. U46619, SQ29548, pinane thromboxane A_2 (PTA_2), and 8-iso-prostaglandin $F_{2\alpha}$ (8-iso-PGF $_{2\alpha}$) were purchased from Cayman Chemical Company (Ann Arbor, MI). [35 S]GTP γ S (1250 Ci/mmol) and [3 H]SQ29548 (44 Ci/mmol) were purchased from PerkinElmer Life and Analytical Sciences, Inc. (Waltham, MA). Aprotinin, apyrase, normal rabbit serum, GDP, GTP, protein A-Sepharose, and Sepharose 2B were purchased from Sigma-Aldrich (St. Louis, MO). Pansorbin cells and Nonidet P-40 were purchased from Thermo Fisher Scientific (Waltham, MA). Rabbit antisera for G_{α_q} and $G_{\alpha_{13}}$ were produced using peptides corresponding to the C-terminal 10 residues of the two proteins (Butkerait et al., 1995; Windh et al., 1999).

Cells and Membrane Preparation. Human embryonic kidney (HEK) 293 cells stably expressing $TP\alpha$ [4.8 pmol receptor per mg of membrane protein (Wilson et al., 2004)] were maintained in Dulbecco's modified Eagle's medium with 10% fetal bovine serum and 0.5 mg/ml G418 at 37°C with 5% CO_2 . Cells were harvested, washed three times with phosphate-buffered saline, and lysed in 20 mM HEPES, pH 8.0, 1 mM EDTA, 0.1% aprotinin, 0.02% leupeptin, and 0.1% phenylmethylsulfonyl fluoride by repeated passage through a 26-gauge needle. The homogenate was centrifuged at 660g for 5 min, and the resultant supernatant was centrifuged at 20,800g for 30 min at 4°C. The pellet (membrane) was resuspended at ~3 mg/ml protein.

For platelets, human blood was obtained from healthy donors who denied ingestion of aspirin or any other drug for at least 1 week. The donated blood was collected into a one-sixth volume of a solution containing 65 mM trisodium citrate, 60 mM citric acid, and 100 mM

dextrose, pH 4.4. Platelet-rich plasma was isolated by centrifugation of the citrated blood at 180g for 20 min at room temperature, and platelets were obtained thereafter by sedimentation at 880g for 15 min. Platelets were resuspended in a small volume of 10 mM triethanolamine and 5 mM EDTA, pH 6.8, and lysed by immersion in liquid nitrogen then thawing. The lysate was centrifuged at 20,800g for 30 min at 4°C to obtain a membrane pellet, which was resuspended at approximately 3 mg/ml protein in 10 mM triethanolamine, pH 6.8.

[35 S]GTP γ S-Binding. The assay for agonist-promoted binding of [35 S]GTP γ S to G_{α_q} and $G_{\alpha_{13}}$ was performed essentially as described previously (Windh et al., 1999; Zhang et al., 2006). Membranes (20 μ g of protein/assay point) were resuspended in 50 mM Tris-HCl, pH 7.5, 2 mM EDTA, 100 mM NaCl, 20 mM $MgCl_2$, 0.1 μ M GDP, and 5 nM [35 S]GTP γ S in 1.5-ml microcentrifuge tubes on ice. Ligands, if any, were added, and the tubes were transferred immediately to a 30°C water bath for 2 (G_q) or 1 (G_{13}) min. The incubation was terminated by adding 600 μ l of ice-cold 50 mM Tris-HCl, pH 7.5, 20 mM $MgCl_2$, 150 mM NaCl, 0.5% Nonidet P-40, 0.33% aprotinin, 0.1 mM GDP, and 0.1 mM GTP. The extract was transferred to a microcentrifuge tube containing 2 μ l of nonimmune serum preincubated with 100 μ l of a 10% suspension of Pansorbin cells. Nonspecifically bound proteins were removed after 20 min by centrifugation. The extract was incubated for 1 h at 4°C with 10 μ l of a $G\alpha$ -directed antiserum or nonimmune serum, both of which had been preincubated with 100 μ l of a 5% suspension of protein A-Sepharose. Immunoprecipitates were collected and washed three times in the extraction buffer, then once in the buffer without detergent, and then boiled in 0.5 ml of 0.5% SDS followed by addition of 5 ml of Ecolite+ (MP Biomedicals, Irvine, CA). The samples were analyzed directly by scintillation spectrometry. Counts obtained with nonimmune serum, representing nonspecifically bound radiolabel and generally in the range of 50 to 200 cpm, were subtracted before portrayal of the data. In experiments involving inhibition of U46619's actions by PTA_2 , the two ligands were added simultaneously before incubation at 30°C.

Platelet Shape-Change and Aggregation. Platelet-rich plasma was incubated with 1 mM aspirin and, for experiments relating to shape change (rounding), 0.1 unit/ml apyrase. Platelets were then isolated by gel filtration on Sepharose 2B using modified Tyrode's buffer (137 mM NaCl, 2.7 mM KCl, 1 mM $MgCl_2$, 11.9 mM $NaHCO_3$, 3.6 mM NaH_2PO_4 , 10 mM HEPES, 0.2% bovine serum albumin, and 5.5 mM glucose, adjusted pH to 7.4). Agonist-induced platelet shape-change and aggregation were determined in separate experiments by measuring the transmission of light through a 0.3-ml sample of the aspirin-treated, gel-filtered platelets (2×10^8 cells/ml) with stirring in a lumi-aggregometer (CHRONO-LOG CO., Havertown, PA) at 37°C (Prevost et al., 2002). The baseline was set using 0.5 ml of Tyrode's buffer as a blank. Shape-change in response to agonists was evaluated as a decrease in transmission, which was scored as percentage maximum attained; 1 μ M tirofiban was included in these experiments to prevent aggregation. Aggregation was evaluated (without tirofiban) as an increase in transmission in the presence of 1 mM $CaCl_2$. Aggregation was scored as incidence because of its steeply graded nature.

Measurement of Cytosolic Calcium. Platelet-rich plasma was incubated with 1 mM aspirin and 5 μ M Fura-2/acetoxymethyl ester for 1 h at 37°C. Platelets were then isolated by gel filtration as described under *Cells and Membrane Preparation* and, after adjusting to 2×10^8 cell/ml, were placed into a luminescence spectrometer (PerkinElmer Life and Analytical Sciences). Excitation wavelength was set to 340 nm, and emission was evaluated at a wavelength of 510 nm. Calcium was expressed as concentration using a K_d for Fura-2 of 224 nM (Gryniewicz et al., 1985).

[3 H]SQ29548 Displacement. Membranes (20 μ g of protein/assay point) were incubated with 20 nM [3 H]SQ29548 in the presence or absence of other ligands at specified concentrations in 20 mM HEPES, pH 7.4, 2 mM EDTA, and 5 mM NaCl for 30 min at 30°C.

The incubation volume was 0.1 ml. Reactions were terminated by dilution with 10 mM HEPES, pH 7.4, and 0.01% BSA at 0°C and rapid filtration over Whatman GF/C filters presoaked in the same buffer. The filters were washed three times with the same buffer at 0°C and dried. Filter-bound radioactivity was determined by scintillation spectrometry. Nonspecific binding, generally less than 10% at K_d, was defined as the binding of radioligand in the presence of 100 μM SQ29548.

Miscellaneous. Data were analyzed using Prism Software (Graph Pad Software, San Diego, CA). [³⁵S]GTPγS-binding was evaluated by nonlinear regression analysis using a logistic equation and a Hill slope of 1. Displacement of [³H]SQ29548 was fit to a one-site competition mode (a two-site model provided no better fit). Statistical differences in EC₅₀ or K_i values were determined using a two-tailed Student's *t* test, *p* < 0.05 signifying a difference.

Results

Most receptors that are coupled to G proteins of the G₁₂ family couple to those of the G_q family as well. Despite the importance of these receptors and the general interest in functional selectivity, differences in engagement of the two G protein families have not been investigated. Studies were initiated here with TP_α, a receptor for TXA₂, and the G proteins G_q and G₁₃. G₁₃ is a member of the G₁₂ family, the actions of which are essential to those of TP_α in platelets. The ligands employed were U46619, PTA₂, and 8-iso-PGF_{2α}.

The activation of G_q and G₁₃ through TP_α was evaluated first with HEK 293 cells in which TP_α was introduced and stably expressed (Wilson et al., 2004). Activation was evaluated as ligand-promoted binding of [³⁵S]GTPγS to endogenous G_α subunits. U46619, PTA₂, and 8-iso-PGF_{2α} all activated G_q and, as we reported previously (Zhang et al., 2006), G₁₃ (Fig. 1). U46619 was the most potent, exhibiting an EC₅₀ of approximately 80 nM for both G proteins (Table 1). PTA₂ was more potent than 8-iso-PGF_{2α} in activating G_q, whereas the two were similar in activating G₁₃. It is noteworthy that PTA₂ activated G_q in preference to G₁₃ (EC₅₀ = 1 μM for G_q versus 4.5 μM for G₁₃, *p* = 0.03), whereas the converse was true for 8-iso-PGF_{2α} (EC₅₀ = 3.4 μM for G₁₃ versus 12 μM for G_q, *p* = 0.04). Activation in membranes of cells where TP_α was not introduced was negligible. These data, in which opposite preferences for effectors (G proteins) by two agonists were observed, connote functional selectivity.

We turned to human platelets, for which TP_α and the G proteins are both endogenous (the existence of TP_β as well in platelets is a possibility, hence the term 'TP' hereafter, but see Habib et al., 1999). U46619, PTA₂, and 8-iso-PGF_{2α} again activated G_q and G₁₃ (Fig. 2). U46619 remained the most potent of the three and did not distinguish between the two G proteins (EC₅₀ = 0.3–0.4 μM). PTA₂ was more potent than 8-iso-PGF_{2α} in activating G_q, although it was less efficacious. PTA₂ and 8-iso-PGF_{2α} activated G₁₃ with similar potency, but here 8-iso-PGF_{2α} was the less efficacious. Selectivity referenced to potency was again evident: PTA₂ activated G_q in preference to G₁₃ (EC₅₀ = 3 μM for G_q versus 14 μM for G₁₃, *p* = 0.008), whereas 8-iso-PGF_{2α} activated G₁₃ in preference to G_q (EC₅₀ = 35 μM for G₁₃ versus 210 μM for G_q, *p* = 0.003). Activation of the two G proteins by all three ligands was blocked by the TP antagonist SQ29548 (data not shown). The selectivity noted upon overexpression of TP_α in HEK 293 cells, therefore, was corroborated in a setting where receptor and G proteins are expressed normally.

One of the reasons for measuring selectivity using G proteins is that the phenomenon is less likely to be obscured by idiosyncrasies in signal propagation and regulation attending downstream events. With this in mind, we

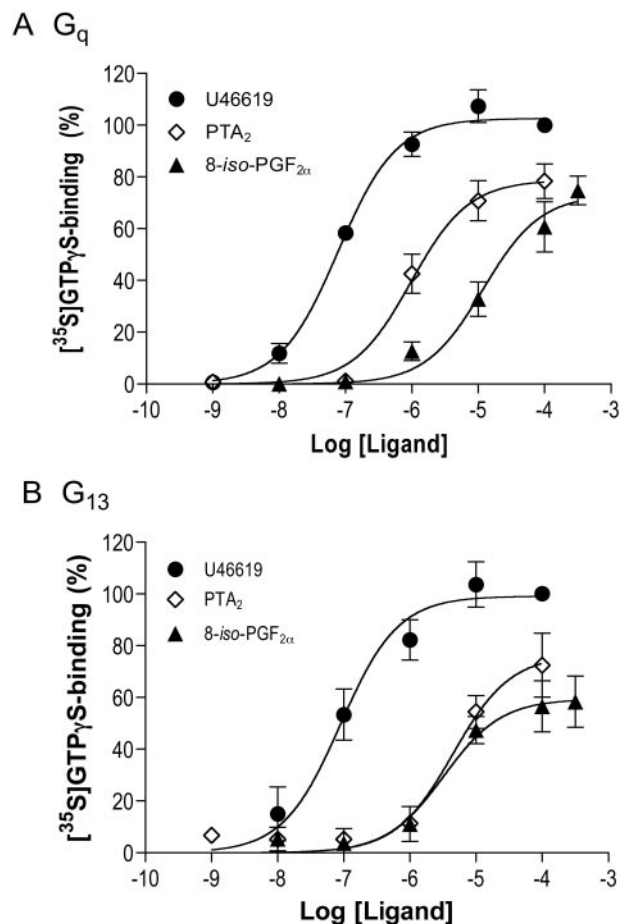


Fig. 1. Activation of G_q and G₁₃ through TP_α in HEK 293 cell membranes. Membranes were prepared from HEK 293 cells stably expressing TP_α and incubated with U46619, PTA₂, and 8-iso-PGF_{2α} at the concentrations indicated, together with [³⁵S]GTPγS. Binding of [³⁵S]GTPγS to G_α subunits of endogenous G_q (A) and G₁₃ (B) was determined by immunoprecipitation with G_α-selective antibodies and scintillation spectrometry. The data are expressed as a percentage binding obtained with 100 μM U46619, equivalent to approximately 3000 cpm for G_q and 500 cpm for G₁₃, minus binding in the absence of agonist, approximately 150 cpm. Each point represents the mean ± S.E. of three to five independent experiments.

TABLE 1

Potency of ligands in promoting activation of G_q and G₁₃ through TP_α. The activation of G_q and G₁₃ was evaluated in membranes of HEK 293 cells stably (over)expressing TP_α and membranes of human platelets. EC₅₀ values for U46619, PTA₂, and 8-iso-PGF_{2α} were determined by nonlinear regression on concentration-response data. Each value is the average of three to five experiments, with 95% confidence intervals (C.I.) given in parentheses. *P* values for comparisons between G_q and G₁₃ for individual ligands are also noted.

	EC ₅₀ (95% C.I.)		<i>P</i>
	G _q	G ₁₃	
	μM		
HEK 293/TP _α			
U46619	0.078 (0.058–0.10)	0.089 (0.049–0.16)	0.8
PTA ₂	1 (0.53–1.8)	4.5 (1.9–10)	0.03
8-iso-PGF _{2α}	12 (5.9–24)	3.4 (1.7–7.2)	0.04
Platelets			
U46619	0.39 (0.35–0.45)	0.31 (0.24–0.39)	0.1
PTA ₂	3.1 (1.7–5.6)	14 (8.1–25)	0.008
8-iso-PGF _{2α}	210 (110–400)	35 (20–64)	0.003

examined platelet rounding and aggregation, respectively, as correlates of G_{13} and G_q activation. All three ligands, as expected, caused rounding (Fig. 3, top). The EC_{50} values were shifted leftward 10- to 20-fold from those for G_{13} activation, suggesting either a surplus of G_{13} in relation to downstream events (analogous to receptor reserve) or a better coupling of receptor and G protein in the intact cell. The maximal effects of the three ligands were similar. Rounding caused by the ligands was inhibited by SQ29548. Activation of G_{13} through one or more forms of TP by all three ligands translates, therefore, into a physiological response. This was not the case for aggregation.

U46619 elicited, as expected, aggregation (Fig. 3, middle), which was steeply graded and inhibited by SQ29548. The EC_{50} for aggregation (135 nM) was close to that for activation of G_q in membranes, suggesting little if any G_q reserve. PTA_2 , however, did not effect aggregation at any but the highest concentration tested (300 μ M), and the aggregation was not inhibited by SQ29548, indicating the effect to be nonspecific. 8-iso-PGF $_{2\alpha}$ was without effect altogether. The activation of G_q noted for PTA_2 and 8-iso-PGF $_{2\alpha}$ through TP, therefore, was inconsequential to aggregation. The ability of

U46619 to effect aggregation was paralleled by its ability to effect a substantial increase in intracellular calcium (Fig. 3, bottom panel). We suspect that the inability of PTA_2 and 8-iso-PGF $_{2\alpha}$ to effect aggregation or large increases in calcium through one or more forms of TP is due to a required threshold in G_q activation. The selectivity noted for G protein activation was, nonetheless, not evident in downstream events, underscoring the problems inherent to these events as measures of the phenomenon.

Selectivity can explain preferential engagement of one G protein or another by a ligand, but it has the ability to explain other actions as well. Contrary to the above-noted properties of PTA_2 as an activator of G proteins through TP α , PTA_2 is most often viewed to be an antagonist of the receptor (for example Nicolaou et al., 1979; Nie et al., 2008). We

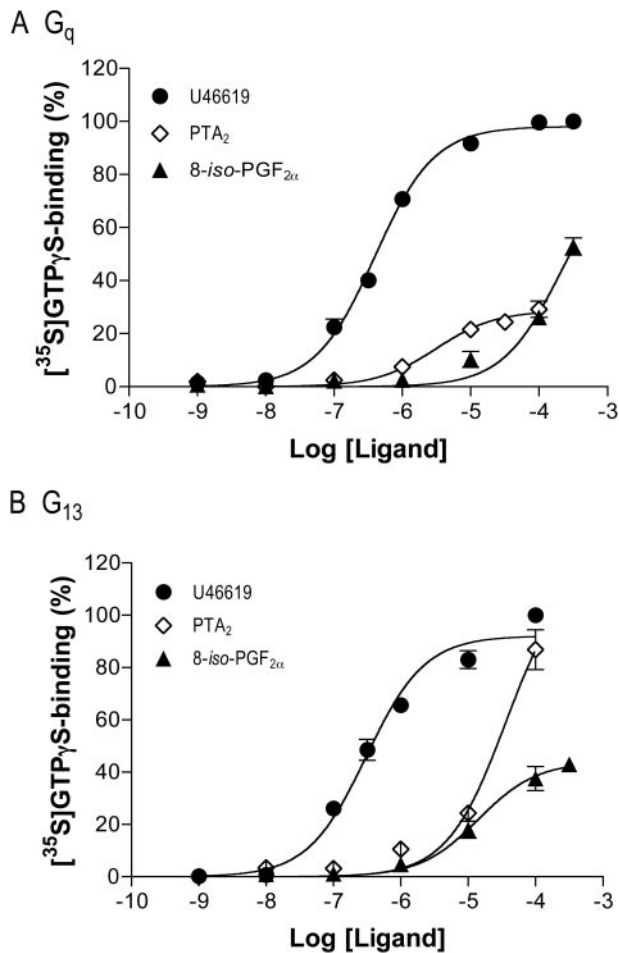


Fig. 2. Activation of G_q and G_{13} through TP in platelet membranes. Membranes were prepared from human platelets, and the activation of G_q (A) and G_{13} (B) in response to U46619, PTA_2 , and 8-iso-PGF $_{2\alpha}$ was evaluated as described in the legend to Fig. 1. The data are expressed as a percentage binding obtained with 100 μ M U46619, equivalent to approximately 8000 cpm for G_q and 1500 cpm for G_{13} , minus binding in the absence of agonist, approximately 500 cpm. Each point represents the mean \pm S.E. of three to five independent experiments.

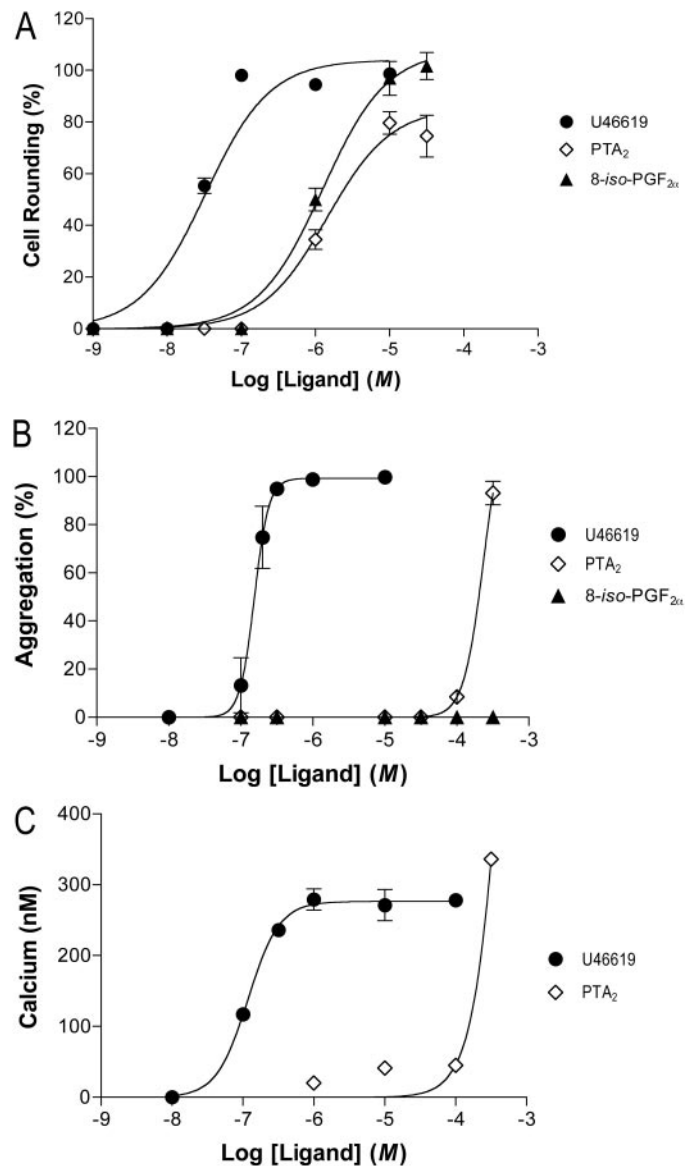


Fig. 3. Response of intact platelets to ligands for TP. The response of human platelets to U46619, PTA_2 , and 8-iso-PGF $_{2\alpha}$ was evaluated in terms of cell rounding (top), aggregation (middle), and intracellular calcium (bottom; data for U46619 and PTA_2 only). Rounding and aggregation are expressed as a percentage of that obtained with 10 μ M U46619. Each point represents the mean \pm S.E. of three to five independent experiments.

wondered whether the difference between our results and cited actions of PTA₂ as an antagonist might rest with a confusion of weak agonism for antagonism, at least at the level of G_q, or with functional selectivity of some form. We found that PTA₂ inhibited the activation of G_q by U46619 (Fig. 4, top), and that the activity at the highest concentration of PTA₂ (10 μ M) was that of PTA₂ alone. This finding conforms to what one would expect for a partial agonist. Of interest, however, was what seemed to be the unusual potency of PTA₂ with respect to inhibition: the IC₅₀ was 0.5 μ M when PTA₂ was tested against 1 μ M U46619, implying an affinity for receptor slightly greater than that of U46619 itself. This finding was in distinction to the data for G protein activation, in which the potency of PTA₂ was 8-fold less than that of U46619. The distinction was even more striking when the activation of G₁₃ was evaluated. PTA₂ is a full agonist with regard to G₁₃, hence suppression of U46619-promoted activation by PTA₂ should not be discernible. This was not the case. PTA₂ inhibited U46619-effected activation of G₁₃ in platelet membranes was shifted rightward from binding with an IC₅₀ of 0.7 μ M (Fig. 4, bottom), well below the EC₅₀ (14 μ M) for its activation of G₁₃. The effects of PTA₂ in the presence of U46619 were biphasic (not shown), with the inhibition at low concentrations followed by an activation analogous to that noted previously for PTA₂ alone at higher concentrations. PTA₂, therefore, displayed concentration-

dependent opposing actions at the level of each of the two G proteins.

To help clarify the behavior of PTA₂, we evaluated the affinity displayed by U46619 and PTA₂ for receptor(s) on platelet membranes by displacement assays using [³H]-SQ29548. The K_i for SQ29548 with platelet membranes was 40 nM (Table 2). Both U46619 and PTA₂ displaced the radiolabel with K_i values of 0.7 and 0.4 μ M, respectively (Fig. 5 and Table 2). In the case of U46619, therefore, binding was roughly coincident with activation of G_q and G₁₃ (EC₅₀ = 0.3–0.4 μ M, as noted above). For PTA₂, binding was consistent with its antagonism of U46619 but *not* with its activation of G proteins—the K_i values for PTA₂ were 5- and 30-fold lower than EC₅₀ values for G_q and G₁₃ activation (3 and 14 μ M, respectively). Variations in Mg²⁺ and/or the inclusion of guanosine 5'-(β , γ -imido)triphosphate or GTP γ S had no impact. Viewed from the perspective of concentration-response relationships, therefore, the activation of G proteins by PTA₂ in platelet membranes was shifted rightward from binding. The Hill slopes for U46619 and PTA₂ were -0.85 ± 0.10 ($n = 3$) and -1.25 ± 0.16 ($n = 6$), respectively. These values were not statistically different from each other or from 1; however, the power of the analysis was not so high as to completely rule out a difference in the two slopes that might connote differences in cooperativity.

A rightward shift in an effect is almost always due to the relevance of two receptors differing in affinity for the ligand, in which the effect is achieved through the one having a lower affinity, but the binding is evident only for the one having the higher affinity for reasons of selectivity in radiolabeling or disproportionate levels of receptor. We turned to HEK 293 cells, therefore, to evaluate TP α in isolation, aware that the receptor reserve implied above and elsewhere (Zhang et al., 2006), which would cause leftward shifts, might complicate the analysis. The binding data were quite similar to those of platelets. U46619 and PTA₂ were comparable with each other in terms of affinities for TP α , having K_i values of 0.8 and 0.7 μ M, respectively (Table 2). As anticipated for receptor reserve, the activation of G_q and G₁₃ by U46619 (EC₅₀ \sim 0.08 μ M in HEK 293 cell membrane) was left-shifted from binding. Despite the reserve, the activation of G proteins by PTA₂ (EC₅₀ values = 1 μ M for G_q and 4.5 μ M for G₁₃) was not left-shifted; in fact, the activation of G₁₃ remained right-shifted from binding. Therefore, two “different” receptors can be entertained as a basis for differences in binding and activation; however, the two would originate with TP α , presumably through different conformations. An alternative is that two sites for PTA₂ exist on TP α , accounting for the duality in the actions of PTA₂.

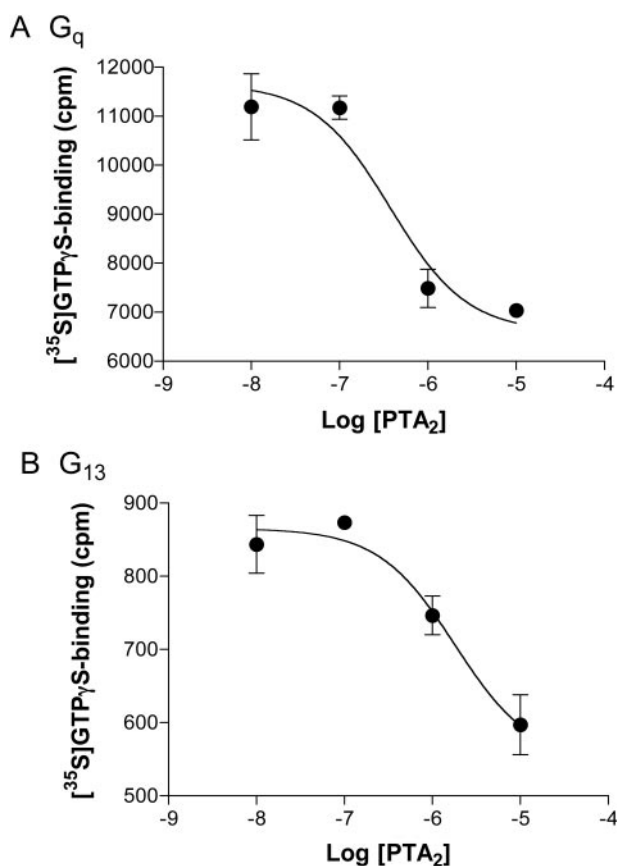


Fig. 4. Inhibition by PTA₂ of U46619's activation of G_q and G₁₃ in platelet membranes. Membranes from human platelets were incubated with 1 μ M U46619, the indicated concentrations of PTA₂, and [³⁵S]GTP γ S. Binding of [³⁵S]GTP γ S to G α_q and G α_{13} was evaluated as described in the legend to Fig. 2. The data are from single experiments performed in duplicate, which are representative of two others each for G_q and G₁₃, and are expressed as counts per minute.

TABLE 2
Binding of ligands to TP

K_i values were determined for SQ29548, U46619, and PTA₂ in displacement assays with 20 nM [³H]SQ29548 using membranes prepared from human platelets and HEK 293 cells, in the latter case stably (over)expressing TP α . Each assay was conducted with five or six concentrations of ligand in duplicate. Each reported value is the mean; 95% confidence intervals, obtained by nonlinear regression on data accumulated from three or four experiments, are in parentheses.

	K _i	
	Human Platelets	HEK 293 α TP α
	μ M	
SQ29548	0.04 (0.03–0.06)	0.07 (0.05–0.10)
U46619	0.7 (0.5–1.0)	0.8 (0.7–1.1)
PTA ₂	0.4 (0.3–0.6)	0.7 (0.5–1.1)

Discussion

The G_{12} family has surfaced only rarely in the context of functional selectivity and always indirectly. The family is possibly involved in the selectivity exhibited through the 5-hydroxytryptamine_{2C} receptor in reference to phosphoinositide accumulation and arachidonic acid release (Berg et al., 1998), because RNA-edited forms of the receptor, which are uncoupled from the G_{12} family (Price et al., 2001), no longer exhibit selectivity (Berg et al., 2001). The family might also underlie the selectivity exhibited through the 5-hydroxytryptamine_{2A} receptor, presuming the family again to be involved in arachidonic acid release (Berg et al., 1998; Kurrasch-Orbaugh et al., 2003). We are aware of no other possible examples. The paucity of information regarding the G_{12} family is not entirely unexpected, in that the actions of the family are not easily separated from those of other signaling entities nor easily quantified through regulation of second messengers, proximate enzymes, or ion channels. However, receptors coupled to the G_{12} family are basic to cell function and, because they are invariably coupled to G proteins of other families, provide ample potential for selectivity.

We approached the need for uniquely linked and quantifiable effectors by measuring G protein activation directly, using agonist-promoted exchange of GDP for [³⁵S]GTP γ S. The assay's utility in measurements of potency and efficacy has already been documented (Barr and Manning, 1997; Windh et al., 1999; Windh and Manning, 2002; Zhang et al., 2006), and extension to selectivity was made most recently for the dopamine D2 receptor in relation to subtypes of the G_i family (Lane et al., 2007). The use of proximal effectors (i.e., G proteins) and cell membranes avoids pitfalls of nonlinearity and convergence in downstream signaling and superimposed forms of regulation that confound results in the intact cell. The choice in this study of G_q and G_{13} for analysis was based on the large number of receptors that engage these two G proteins specifically (Riobo and Manning, 2005) and the relevance of both to platelet function (Offermanns et al., 1997; Moers et al., 2003; Offermanns, 2006).

We found that G_q and G_{13} were, in fact, differentially engaged by agonists for TP α . PTA₂ activated G_q at lower concentrations than it did G_{13} , whereas the converse was true for 8-iso-PGF_{2 α} . Selectivity for the two agonists was discerned in membranes of

both HEK 293 cells overexpressing TP α and platelets. U46619, in contrast, exhibited no selectivity. The opposite (PTA₂ versus 8-iso-PGF_{2 α}) and nonselective (U46619) patterns of G protein engagement are important in precluding receptor reserve alone as a basis for the phenomenon.

Platelet rounding, a measure of G_{13} action, was elicited by all three ligands; however, aggregation, which requires G_q as well, was elicited by U46619 alone through TP. This observation is consistent with our assertion that functional selectivity cannot always be detected through events downstream of G proteins. The more proximal the effector being measured to receptor, we believe, the more likely the measure is to be faithful to receptor conformation. We suspect for platelets that aggregation requires a threshold level of activated G_q , noting that neither PTA₂ nor 8-iso-PGF_{2 α} can achieve full activation of this G protein. It is noteworthy that the EC₅₀ for U46619 in aggregation was congruent with the EC₅₀ in activation of G_q and the K_i for binding of U46619 to platelet membranes, suggesting little or no reserve of receptor or G protein. This finding was in distinction to rounding, for which the concentration-response relationship appeared to be substantially left-shifted from G protein activation and binding, suggesting a reserve of receptor and/or receptor-activated G_{13} . This observation is important—any nonselective ligand such as U46619 would trigger rounding at concentrations much lower than aggregation, an action that might be confused with true (functional) selectivity. Again, the choice of G proteins to evaluate selectivity is essential. The reserve here has nothing to do with, and does not impede, the deduction of selectivity using G proteins as endpoints.

In evaluating the properties of PTA₂, we noted in platelets an antagonistic action for the ligand that did not conform to what would be expected for even a weak agonist; i.e., PTA₂ blocked the actions of U46619 at concentrations well below those anticipated from the affinity suggested by its EC₅₀ in G protein activation. We found that the EC₅₀ values for activation of G_q and G_{13} by PTA₂ were, in fact, *right*-shifted from the apparent binding of the ligand: the EC₅₀ for activation of G_q was shifted from the K_i by approximately 8-fold, whereas that for activation of G_{13} was shifted by approximately 35-fold. Differences between the [³⁵S]GTP γ S-binding and displacement assays are unlikely to account for the shift, because buffer constituents in the activation assay (for example, Mg²⁺ and/or GTP γ S) had no impact on displacement of [³H]SQ29548. More compellingly, no meaningful shift was evident for U46619; the shift was unique to PTA₂.

The almost universal explanation for a rightward shift of function from binding is the existence of two different receptors. The relevance of any receptor apart from TP α itself, however, is doubtful. The receptor labeled with [³H]SQ29548 in platelet membranes is mostly if not solely TP α (Habib et al., 1999). The rightward shift for activation of G_{13} , moreover, is also noted in HEK 293 cells made to express TP α . The possibility of TP β might be entertained in platelets, but it cannot account for the results in HEK 293 cells. We suggest as one possibility, therefore, that TP α itself resolves into two populations. PTA₂ would bind to one as an antagonist with relatively high affinity and to the other as an agonist with lower affinity. The latter population would not be recognized by [³H]SQ29548 binding because of low density (i.e., the population is beneath the threshold of detection by radiolabeling) or low affinity of the population for the radioligand,

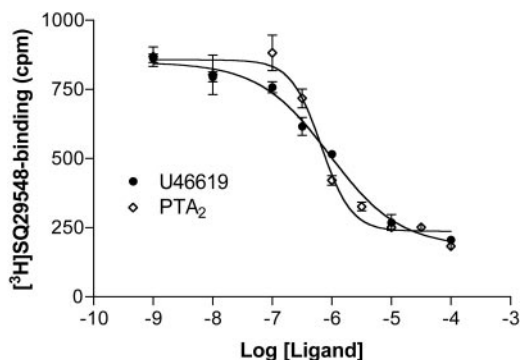


Fig. 5. Displacement of [³H]SQ29548 from platelet membranes. Platelet membranes were incubated with 20 nM [³H]SQ29548 in the absence or presence of increasing concentrations of U46619 and PTA₂ for 30 min at 30°C. Bound radiolabel was determined subsequently by filtration. Shown is a single experiment, performed in duplicate for each displacing ligand, which is representative of several others.

such that binding is not stable in the filtration assay. The existence of different populations of TP α is easily posited. Populations can arise through heterodimerization, other forms of oligomerization, or free receptor versus that complexed to the assayed G protein. Populations might also arise by virtue of the receptor interacting with other proteins, perhaps "silent" G proteins (such as G_i or G₁₂), regulators of G protein signaling, or β -arrestin, or as a function of location within membrane (Zheng et al., 2008). Our data might also be explained by two sites for binding of ligands on TP α , one used by U46619 as an agonist and PTA₂ as an antagonist, and the other by PTA₂ as an agonist. We note the resemblance of our data for PTA₂ with TP α to those obtained by Baker et al. (2003) for CGP12177 with the β_1 -adrenoreceptor, where distinct conformations or activation sites were similarly posited. Functional duality speaks to the possibility of selectivity apparent not so much at the level of one G protein or another but at basic distinctions between agonism and antagonism.

The data obtained here for TP α underscore functional selectivity in relation to the G_q and G₁₂ families, the potential for reserves of receptor and/or G protein that can mimic selectivity, and populations of a single receptor that relate to the agonistic properties of a ligand (PTA₂). All of these introduce a largely unrecognized and important complexity to the coordination of signaling noted for the G_q and G₁₂ families. Remaining questions pertain to the basis of underlying conformations of receptor, specific links to events downstream of the G proteins, and extension to other receptors coupled conjointly to the two families.

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