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Research Article

Inflammation

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Airway smooth muscle prostaglandin-EP₁ receptors directly modulate β_2 -adrenergic receptors within a unique heterodimeric complex

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Multiple and paradoxical effects of airway smooth muscle (ASM) 7-transmembrane-spanning receptors activated during asthma, or by treatment with bronchodilators such as β_2 -adrenergic receptor (β_2 AR) agonists, indicate extensive receptor crosstalk. We examined the signaling of the prostanoid-EP₁ receptor, since its endogenous agonist prostaglandin E₂ is abundant in the airway, but its functional implications are poorly defined. Activation of EP₁ failed to elicit ASM contraction in mouse trachea via this G_{αq}-coupled receptor. However, EP₁ activation markedly reduced the bronchodilatory function of β_2 AR agonist, but not forskolin, indicating an early pathway interaction. Activation of EP₁ reduced β_2 AR-stimulated cAMP in ASM but did not promote or augment β_2 AR phosphorylation or alter β_2 AR trafficking. Bioluminescence resonant energy transfer showed EP₁ and β_2 AR formed heterodimers, which were further modified by EP₁ agonist. In cell membrane [³⁵S]GTPγS binding studies, the presence of the EP₁ component of the dimer uncoupled β_2 AR from G_{αs}, an effect accentuated by EP₁ agonist activation. Thus alone, EP₁ does not appear to have a significant direct effect on airway tone but acts as a modulator of the β_2 AR, altering G_{αs} coupling via steric interactions imposed by the EP₁: β_2 AR heterodimeric signaling complex and ultimately affecting β_2 AR-mediated bronchial relaxation. This mechanism may contribute to β -agonist resistance found in asthma.

Introduction

The 7-transmembrane-spanning (7-TM-spanning) receptors represent the largest signaling family in the genome. We estimate that the lung expresses 25–50 7-TM receptors in airway epithelial cells, airway smooth muscle (ASM), pulmonary vasculature, alveolar walls, and resident immune cells (1). In regard to asthma, several 7-TM receptors play established roles in bronchoconstriction (e.g., M₃-muscarinic receptor) and bronchodilation (e.g., β_2 -adrenergic receptor [β_2 AR]). Despite identification of the endogenous ligands and receptor localization, there are a number of 7-TM receptors expressed in the airway whose functions are unknown, or appear to function paradoxically, based on recombinantly expressed receptors in model cell systems. This lack of understanding of receptor function has impeded our ability to ascertain the role of these ligands (some of which are markedly increased in asthma) in the relaxation/contraction of ASM; thus the mechanistic basis of bronchial hyperreactivity and bronchoconstriction in asthma remains only partially understood. In many cases the basis for incomplete mechanistic information can be attributed to the nature of recombinant expression systems, which may not take

into account multiple receptor subtypes, receptor/receptor interactions, crosstalk via second messengers, or subcellular localization of receptors and specific effectors.

These issues are particularly obvious when one considers the biologic actions of prostaglandin E₂ (PGE₂), which, via at least 4 receptor subtypes (EP₁–EP₄), can evoke multiple signals. By activation of the G_s-coupled EP₂ receptor, PGE₂ can evoke ASM relaxation (2). However, blockade of PGE₂ production leads to partial reversal of β_2 AR desensitization in the context of inflammation (3), and thus PGE₂ may promote bronchoconstriction via desensitization of β_2 AR function or by activation of the G_q-coupled EP₁ receptor. PGE₂ is produced by ASM, airway epithelial cells, and inflammatory cells and is the most abundant prostanoid in epithelial lining fluid (4–6). However, EP₁-mediated signal transduction and its effect on airway responsiveness remains unclear. We show here a unique function of the EP₁ receptor in ASM, which to our knowledge has not been previously described for 7-TM receptors. The EP₁ receptor itself, despite its coupling to inositol 1,4,5-trisphosphate (IP₃) and intracellular calcium ([Ca²⁺]_i), failed to exert the expected bronchoconstrictive response. Instead, EP₁ receptor activation modulated β_2 AR function via formation of a heterodimeric complex that directly decreased β_2 AR coupling to G_{αs} in an EP₁ receptor, agonist-dependent, fashion.

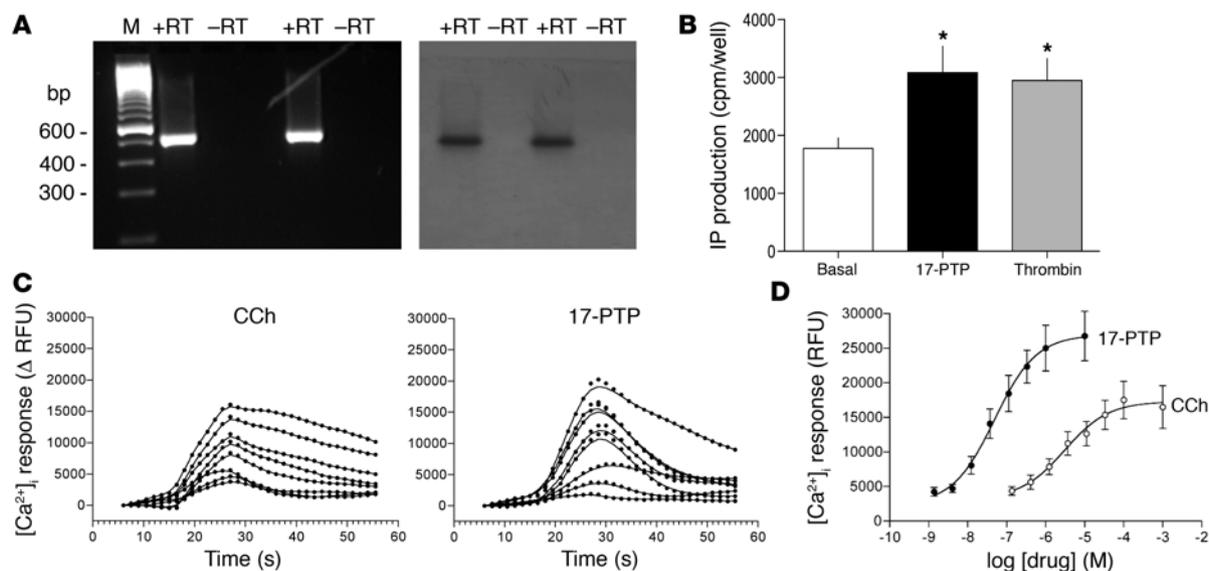
Results

EP₁ receptor activation in isolated and intact ASM. The biologic actions of PGE₂ are potentially mediated by multiple receptor subtypes that differentially couple to G_q, G_i, and G_s. Although a relaxant

Nonstandard abbreviations used: AR, adrenergic receptor; ASM, airway smooth muscle; BRET, bioluminescence resonant energy transfer; [Ca²⁺]_i, intracellular calcium; GABA_B R2, γ -aminobutyric acid receptor subtype bR2; HEK, human embryonic kidney; IP₃, inositol 1,4,5-trisphosphate; PGE₂, prostaglandin E₂; 17-PTP, 17-phenyl trinor-PGE₂; Rluc, Renilla luciferase; 7-TM, 7-transmembrane; YFP, yellow fluorescent protein.

Conflict of interest: The authors have declared that no conflict of interest exists.

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**Figure 1**

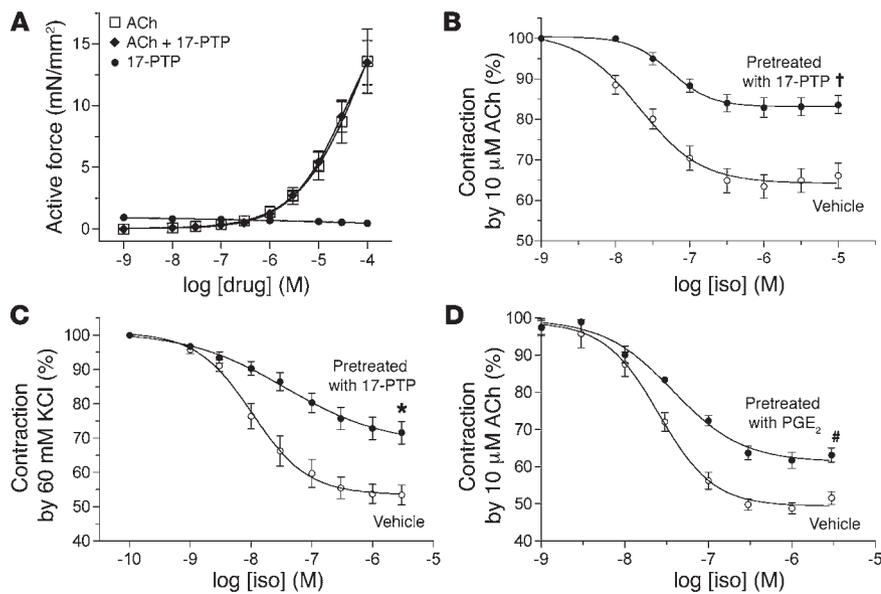
EP₁ receptor expression and signal transduction in murine ASM cells. (A) EP₁ gene expression was detected in cultured murine ASM cells by RT-PCR. The 501-bp product (left) corresponding to sequence from the EP₁ receptor open reading frame was only detected in reactions that included reverse transcriptase (+RT) and hybridized by Southern blots to an EP₁-specific probe (right). (B) The EP₁-selective agonist 17-PTP (1 μM) stimulated inositol phosphate (IP) production in murine ASM ~70% above basal levels (*n* = 4). Thrombin-stimulated inositol phosphate production was used for a positive control. **P* < 0.05 versus basal. (C) Both 17-PTP and acetylcholine-stimulated [Ca²⁺]_i transients in ASM cells. Shown is a representative experiment using the doses shown in D. (D) Maximal agonist-stimulated [Ca²⁺]_i transients in ASM cells exposed to the indicated doses of agonists. Data are mean ± SEM of 5 experiments. Cch, carbachol; RFU, relative fluorescence units.

action of PGE₂, mediated via the EP₂ isoform in ASM, has been documented (7), the extent to which PGE₂ couples to the other subtypes in ASM, particularly the potentially probronchoconstrictive G_q-coupled EP₁ receptor, is not established. EP₁-specific RT-PCR was carried out on murine ASM cells in culture (Figure 1A). A single product of the expected molecular size was obtained, and Southern blots with an EP₁-specific oligonucleotide hybridized to the product. Sequencing further confirmed the product as the murine EP₁ receptor (GenBank accession no. NM_013641). With regard to signal transduction, the EP₁-specific agonist 17-phenyl trinor-PGE₂ (17-PTP) was found to stimulate inositol phosphate production (Figure 1B) and [Ca²⁺]_i transients (Figure 1C). Of note, maximal activation of the EP₁ receptor by 17-PTP stimulated [Ca²⁺]_i to a greater extent, and with greater potency, than the M₃-muscarinic agonist carbachol (Figure 1D). These findings confirmed EP₁ receptor expression in ASM cells and that its activation by agonists resulted in IP₃ and [Ca²⁺]_i turnover as assessed in whole cells. Because these effectors mediate 7-TM receptor contraction of ASM, the data suggested that the EP₁ receptor would likely promote airway contraction.

Excised mouse tracheal ring preparations were utilized to ascertain the effects of EP₁ activation on active force generation in the intact airway. However, despite its ability to stimulate IP₃ and [Ca²⁺]_i turnover in ASM, 17-PTP caused no detectable contraction of tracheal rings even at concentrations up to 10 μM (Figure 2A). Since 17-PTP did not directly effect ASM tone, we next investigated whether EP₁ receptor activation could modify the response to another contractile agonist acting through G_{αq}. For these studies, tracheal rings were pretreated with 1 μM 17-PTP for 15 minutes prior to contraction with various concentrations of acetylcholine (Figure 2A). The maximal acetylcholine response in the absence

(26 ± 5.3 mN) or presence (26 ± 4.8 mN) of EP₁ coactivation was not different, nor was the EC₅₀ for the response (91 ± 11 versus 82 ± 15 μM, respectively). We next considered whether EP₁ receptor activation alters ASM relaxation. The efficacy and potency of the βAR agonist isoproterenol to relax acetylcholine-constricted ASM in the presence and absence of EP₁ receptor activation was ascertained. Tracheal rings were first treated with vehicle or 1 μM 17-PTP for 15 minutes, contracted continuously with 10 μM acetylcholine, and exposed to increasing concentrations of isoproterenol. In the presence of vehicle, isoproterenol caused a dose-dependent relaxation with an EC₅₀ of 23.4 ± 4.8 nM and a maximal reduction in force of 36.6 ± 2.9% (Figure 2B). However, following pretreatment with 17-PTP, a marked reduction in βAR-mediated relaxation was observed, with isoproterenol evoking only 17.1 ± 2.4% relaxation (*n* = 6; *P* < 0.001; Figure 2B). Of note, the EC₅₀ for isoproterenol was also increased under these conditions to 58.2 ± 6.1 nM (*P* < 0.001). As shown in Figure 2C, this same βAR desensitization effect of 17-PTP was observed when tracheal rings were contracted by 60 mM KCl. The endogenous EP₁ receptor agonist is PGE₂, which activates other EP receptors and thus under physiologic conditions could evoke an effect that dominates airway responses over the EP₁-specific phenotype we observed. We therefore studied the effect of pretreatment with 10 nM PGE₂ (which evokes only ~15% relaxation) using the same protocol as for 17-PTP. As shown in Figure 2D, PGE₂ also evoked a loss of isoproterenol-mediated relaxation, amounting to an approximate 25% desensitization. This lower amount of desensitization (compared with 17-PTP pretreatment) may be due to the combined relaxant effect of PGE₂ and isoproterenol.

Given that mouse trachea has a mixed population of β₁AR and β₂AR subtypes (8), the 17-PTP effect was also assessed using the

**Figure 2**

Effect of EP₁ receptor activation on ASM contraction and relaxation. (A) Concentrations of 17-PTP up to 10 μM caused no significant contraction of murine tracheal rings and did not enhance acetylcholine-mediated (ACh) contraction. Data are mean ± SEM of 4 experiments. (B) Pretreatment of tracheal rings with 1 μM 17-PTP for 15 minutes reduced both the potency (EC₅₀ increased) and efficacy of isoproterenol-mediated (Iso) relaxation of tracheal rings contracted with 10 μM acetylcholine. Data are mean ± SEM of 6 experiments. **P* = 0.001 versus vehicle-treated rings. (C) The inhibitory effect of 17-PTP on isoproterenol-mediated relaxation was also present when the tracheal rings were contracted with 60 mM KCl. Data are mean ± SEM of 8 experiments. **P* < 0.05 versus vehicle-treated rings. (D) Pretreatment of tracheal rings with 10 nM PGE₂ evokes desensitization of isoproterenol-mediated relaxation. Data are mean ± SEM of 5 experiments. #*P* < 0.02 versus vehicle-treated rings.

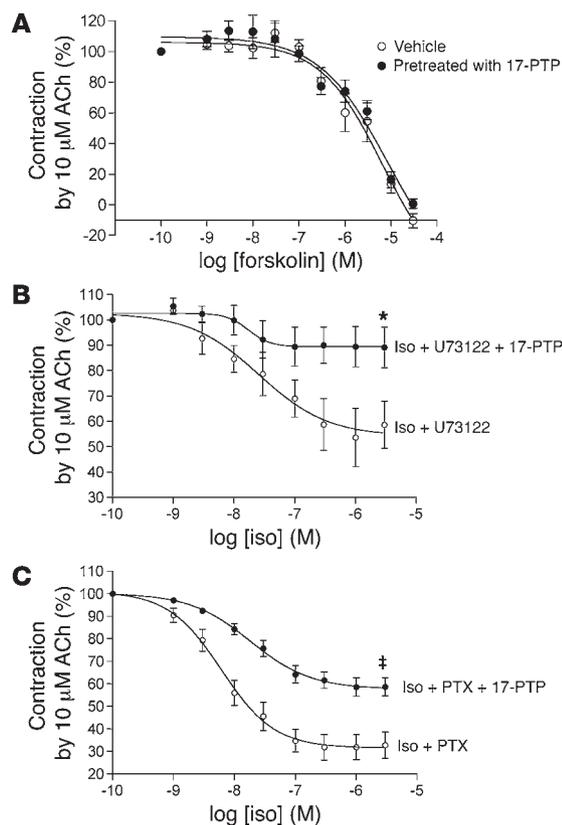
relatively β₂AR subtype-specific agonist formoterol. Under these conditions, 17-PTP evoked a 55% ± 5% desensitization of formoterol-mediated relaxation, indicating a 17-PTP effect on β₂AR (the predominant βAR subtype of human tracheal ASM). To determine whether the inhibitory effect of 17-PTP extended to other ASM relaxants, we performed analogous experiments with forskolin, which directly activates adenylyl cyclase, mimicking a G_s-coupled receptor. However, unlike the response with isoproterenol, pretreatment with 17-PTP had no effect on the maximal response or the EC₅₀ (Figure 3A). Given that the EP₁ receptor couples to G_{αq}, we considered that second messengers generated from phospholipase C (PLC) activation may be involved in the apparent crosstalk between β₂AR and EP₁ receptors. However, pretreatment with the PLC inhibitors U73122 (5 μM; Figure 3B) and Et-18-OCH₃ (10 μM; data not shown) failed to alter the 17-PTP effect. β₂ARs have also been shown in some cells to couple to G_{αi} (9), which may act to oppose G_{αs}-mediated signaling. Treatment of tracheal rings with pertussis toxin, however, had no effect on the EP₁-mediated β₂AR desensitization (Figure 3C), indicating that enhanced G_i coupling was an unlikely mechanism. Finally, the β₂AR desensitization phenotype evoked by EP₁ activation was not altered by pretreatment with 100 μM of the NO inhibitor L-NAME (51.2% ± 6.8% desensitization) or 10 μM of the cyclooxygenase inhibitor indomethacin (57% ± 5% desensitization).

Desensitization of β₂AR by EP₁ receptor activation. The inhibitory effect of 17-PTP on isoproterenol-mediated relaxation observed

in the tracheal ring studies indicated functional interplay between the EP₁ and β₂AR signal transduction pathways. Additionally, the absence of a significant EP₁ effect on forskolin-mediated relaxation and no change in phenotype observed with PLC inhibition suggested that this interaction could potentially occur at the level of the β₂AR itself. The effect of 17-PTP on isoproterenol-stimulated cAMP production was determined in isolated ASM cells. In the absence of 17-PTP, 1 μM isoproterenol stimulated cAMP to 88.0% ± 16.3% of the maximum level obtained with 10 μM forskolin. However, when cells were pretreated with 1 μM 17-PTP for 15 minutes prior to isoproterenol stimulation, cAMP production was reduced to 37.0% ± 3.9% (*n* = 4; *P* = 0.02) of the forskolin response (Figure 4A). Taken together with the tracheal ring data, these results indicated an interplay between the EP₁ receptor and β₂AR, with activation of the former resulting in decreased function of the latter. Two processes that we considered potential mechanisms to account for uncoupling of β₂AR signal transduction within the time frame of minutes included (a) enhanced loss of cell surface receptors due to internalization and (b) enhanced isoproterenol-stimulated β₂AR phosphorylation due to EP₁ activation or phosphorylation of the β₂AR by EP₁ activation via PKC. To ascertain whether EP₁ activation altered the extent or rate of agonist-mediated β₂AR

internalization, we performed radioligand binding on intact ASM cells with the hydrophilic radioligand [³H]-CGP-12177. Since this ligand does not enter intact cells (10), a decrease in binding occurs when surface receptors are internalized. Thus, as expected for agonist-promoted internalization of the β₂AR, exposure of ASM cells to 1 μM isoproterenol caused a time-dependent decrease in [³H]-CGP-12177 binding sites of 39.1% ± 5.7% after 30 minutes (Figure 4B). Pretreatment of cells with 1 μM 17-PTP for 15 minutes had no effect on isoproterenol-stimulated internalization (37.4% ± 3.6%; *n* = 5). We next examined the potential for 17-PTP to promote phosphorylation of the β₂AR. For the phosphorylation experiments, human embryonic kidney (HEK) cells transiently cotransfected with plasmids so as to express the EP₁ receptor and FLAG-tagged β₂AR were loaded with ³²P-orthophosphate. Cells were treated with vehicle, isoproterenol, 17-PTP, phorbol-12-myristate-13-acetate (a positive control for PKC-mediated event), or isoproterenol plus 17-PTP. β₂AR was purified by FLAG immunoprecipitation. As shown in Figure 4C, isoproterenol evoked robust phosphorylation of the β₂AR, while no detectable phosphorylation over background was noted with 17-PTP exposure alone. Importantly, the extent of isoproterenol-promoted β₂AR phosphorylation was not altered by 17-PTP (48 ± 4.5 relative units versus 42 ± 4.2 relative units in the presence and absence of 17-PTP, respectively; *n* = 5; *P* = NS).

β₂ARs and EP₁ receptors form heterodimers. Since these well-defined short-term mechanisms of receptor regulation and crosstalk were not at play during EP₁-induced desensitization of β₂AR, we con-



considered an alternative hypothesis, which was that the 2 receptors formed heterodimers. We also considered whether agonist activation of the EP₁ receptor directly affected the receptor/receptor interaction within the complex, altering β_2 AR coupling to G_{as}. As a screen for such a complex in ASM cells, fluorescence microscopy was carried out using β_2 AR- and EP₁-specific primary antibodies. Secondary antibodies provided for red (β_2 AR) and green (EP₁) signals. Merged images revealed yellow signals consistent with areas of colocalization of these 2 receptors (Figure 5A). To rigorously assess the potential for heterodimer formation, bioluminescence resonant energy transfer (BRET) and coimmunoprecipitation studies were undertaken in transfected HEK 293 cells. BRET was carried out in living cells expressing 2 receptors having carboxyterminally fused Renilla luciferase (Rluc) or yellow fluorescent pro-

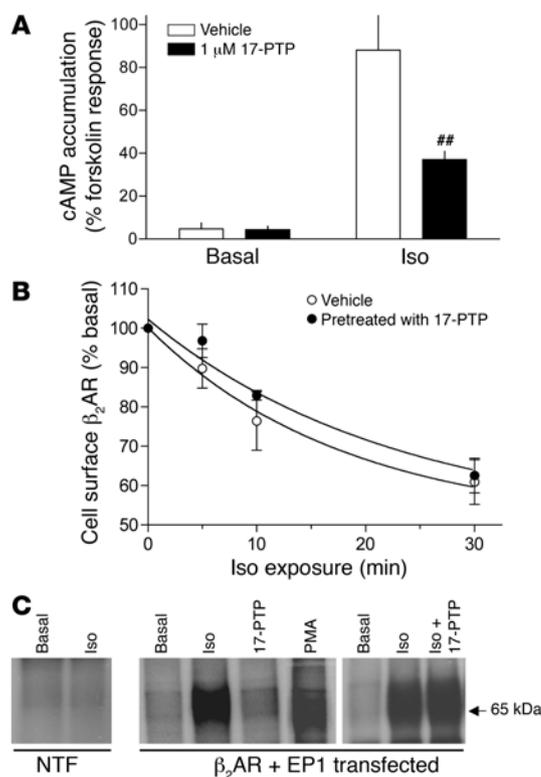
Figure 4

Effect of EP₁ receptor activation on β_2 AR agonist-stimulated cAMP production, receptor sequestration and receptor phosphorylation. **(A)** Pretreatment of ASM cells with 17-PTP (1 μ M) for 15 minutes reduced isoproterenol-stimulated cAMP production. Data are mean \pm SEM of 4 experiments. $^{##}P = 0.02$ versus vehicle-treated cells. **(B)** Isoproterenol caused a time-dependent internalization of β_2 AR in murine ASM cells over 30 minutes as assessed by whole-cell binding using the hydrophilic antagonist [³H]-CGP-12177; pretreatment with 17-PTP (1 μ M for 15 min) had no effect on internalization. Data are mean \pm SEM of 4 experiments. **(C)** EP₁ receptor activation failed to phosphorylate β_2 AR or to alter isoproterenol-mediated β_2 AR phosphorylation in HEK 293 cells transfected to express β_2 AR and EP₁ receptors. Shown is a representative autoradiogram of 4 experiments. PMA, phorbol-12-myristate-13-acetate. NTF, nontransfected.

Figure 3

Characterization of EP₁ receptor-promoted desensitization of β_2 AR-mediated ASM relaxation. **(A)** No effect of 17-PTP pretreatment was observed on forskolin-mediated ASM relaxation, indicating that the site of EP₁ modulation is likely proximal to adenylyl cyclase ($n = 4$). **(B)** Inhibition of phospholipase C (PLC) with U73122 failed to alter the 17-PTP-evoked desensitization of isoproterenol-mediated ASM relaxation. $^*P < 0.05$ versus vehicle-treated rings ($n = 4$). **(C)** Ablation of receptor-G_{ai} coupling with pertussis toxin (PTX) had no effect on the 17-PTP-evoked desensitization of isoproterenol-mediated ASM relaxation. $^{\ddagger}P < 0.01$ versus rings not treated with 17-PTP ($n = 4$).

tein (YFP). As described in detail elsewhere (11–13), this approach takes advantage of a signal produced by the energy donor moiety (Rluc) acting on the energy acceptor (YFP) when the two are in very close proximity. As shown in Figure 5B, studies carried out in HEK 293 cells transfected to express β_2 AR-Rluc and EP₁-YFP (see Methods) indeed revealed a BRET signal, as indicated by the perturbation of the emission spectrum at 530 nm. To verify that this represented oligomer formation, rather than a “pseudo-BRET” signal (11) due to random collisions of overexpressed receptors, studies were carried out with multiple levels of expression (altering the β_2 AR-Rluc/EP₁-YFP ratio). As expected under conditions of dimer formation, the BRET signal saturated, with a calculated maximal ratio (BRET_{max}) of 0.49 ± 0.04 and a BRET₅₀ of 2.97 ± 0.56 (Figure 5C). In contrast, in experiments utilizing cotransfections of β_2 AR-Rluc and a YFP fusion with the γ -aminobutyric acid receptor subtype bR2 (GABA_B R2-YFP) a small signal was observed, as was the case with cotransfections of C-C chemokine receptor 5-Rluc (CCR5-Rluc) and β_2 AR-YFP (Figure 5D). These low-level signals and a lack of saturation (data not shown) were consistent with prior observations that neither the GABA_B R2 nor CCR5 recep-



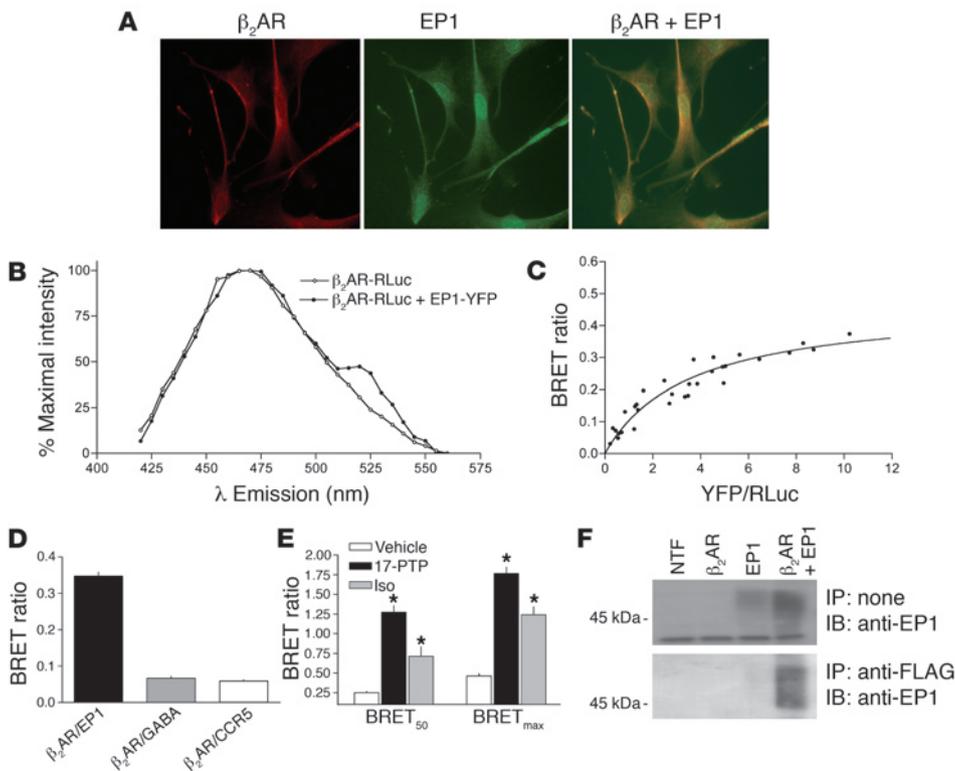


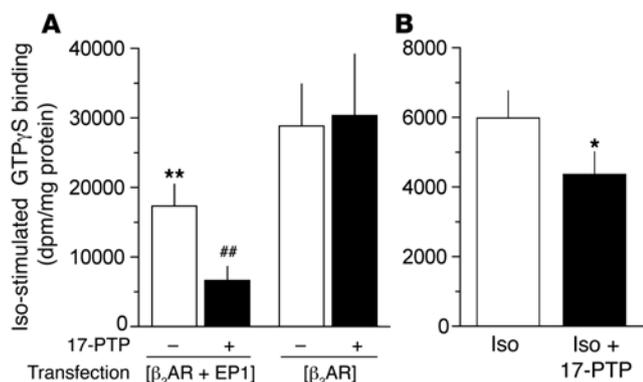
Figure 5
 The EP₁ receptor and β_2 AR form a heterodimer as assessed by BRET and coimmunoprecipitation. **(A)** Immunofluorescence microscopy revealed potential colocalization of EP₁ and β_2 AR in ASM cells. Magnification, $\times 200$. **(B)** The emission spectra of HEK 293 cells transfected with the β_2 AR-Rluc fusion alone showed a single peak with a maximum at ~ 475 nm following addition of coelenterazine h (open symbols). However, a second peak was observed at ~ 530 nm when cells were cotransfected with the EP₁-YFP fusion construct (closed symbols). Shown is a representative experiment of 4. **(C)** The BRET ratios derived from HEK 293 cells transfected with a fixed amount of the donor β_2 AR-Rluc reached saturation with increasing expression of the acceptor EP₁-YFP. Data shown are compiled from 12 experiments. **(D)** GABA_B R2 and C-C chemokine receptor 5 (CCR5) failed to form heterodimers with β_2 AR as indicated by low pseudo-BRET signals. **(E)** The BRET ratios of HEK 293 cells cotransfected with the β_2 AR-Rluc and EP₁-YFP were increased by treatment with either the EP₁ agonist 17-PTP or the β AR agonist isoproterenol. * $P < 0.05$ versus untreated; $n = 4$. **(F)** Coimmunoprecipitation of FLAG-EP₁: β_2 AR receptor complexes. HEK 293 cells were transfected with FLAG- β_2 AR, EP₁ receptor, or both; membranes were solubilized; and proteins were immunoprecipitated with anti-FLAG antibody. Western blots of whole-cell lysates and immunoprecipitates were carried out with polyclonal EP₁ receptor antisera. Shown is a representative blot from 5 experiments.

tors form heterodimers with the β_2 AR (13, 14). We next examined whether activation of the EP₁ receptor or the β_2 AR altered the BRET signal. As shown in Figure 5E, treatment of cotransfected cells with 17-PTP or isoproterenol for 15 minutes prior to the addition of coelenterazine h increased the BRET signal in both cases. Given that most 7-TM dimers appear to be formed intracellularly and then expressed at the cell surface (11, 12, 15, 16), and given the short-term agonist exposure used in these experiments, we do not interpret these results as representing an increased number of dimers per se. Rather, the data are consistent with a conformational change that occurs (potentially in either receptor) when one receptor is bound by its agonist, which causes the Rluc and YFP moieties to move closer together. The magnitude of the agonist-promoted increase in BRET is not trivial, in that the BRET signal is proportional to the distance between the donor and acceptor by a factor of 10^6 (11, 12, 15).

To further corroborate dimer formation, coimmunoprecipitation studies were carried out in HEK 293 cells transfected to express FLAG- β_2 AR and the EP₁ receptor both alone and together. Immunoprecipitation was carried out with anti-FLAG antibody, and the precipitated proteins were fractionated on SDS-PAGE followed by Western blots using a polyclonal EP₁ antisera. As shown in Figure 5F, the EP₁ receptor migrated at approximately 45 kDa in Western blots of cell lysates using the EP₁ antisera. Upon immunoprecipitation with FLAG antibody, protein from cells transfected to express EP₁ or β_2 AR showed no immunoreactivity with the EP₁ antisera. However, when the 2 receptors were cotransfected, a band of the appropriate molecular size was observed. Of note, the high-molecular weight bands, which could represent dimers, are often not resistant to SDS and are not typically visualized (14). These coimmunoprecipitation studies are thus consistent with the BRET results, indicating the presence of the EP₁: β_2 AR heterodimer.

EP₁: β_2 AR heterodimer formation uncouples β_2 AR from G_{α_s} . Taken together, the above results indicated that the EP₁ receptor and the β_2 AR exist as an oligomeric complex, with activation of the EP₁ receptor markedly desensitizing β_2 AR signaling and physiologic function, and that such desensitization is not due to downstream second messenger feedback. This suggested that agonist activation of the EP₁ receptor within the

dimer results in a steric effect on the β_2 AR, altering its conformation in such a way as to directly affect β_2 AR- G_{α_s} interaction. To examine this, we studied agonist-promoted [³⁵S]GTP γ S binding in cell membranes. This provided for direct quantitation of receptor-G protein interaction by measuring GTP turnover and, since it is performed with isolated membranes, was not affected by intracellular factors. In these studies, [³⁵S]GTP γ S binding was carried out by incubating membranes from cells transfected to express G_{α_s} , β_2 AR, and the EP₁ receptor with vehicle, isoproterenol alone, or isoproterenol with 17-PTP. In other studies, G_{α_s} and β_2 AR were expressed without the EP₁ receptor so as to ascertain β_2 AR coupling to G_{α_s} in the absence of the EP₁: β_2 AR dimer. Expression levels of β_2 AR in the β_2 AR and EP₁ (3774 \pm 815 fmol/mg) and β_2 AR transfections (3367 \pm 185 fmol/mg) were comparable. Net isoproterenol-stimulated [³⁵S]GTP γ S binding is shown in Figure 6A. Incubation with 17-PTP resulted in a significant

**Figure 6**

EP₁ receptor activation uncouples the β₂AR from G_s. β₂AR coupling to G_s was assessed by measuring isoproterenol-stimulated [³⁵S]GTPγS binding in membranes from COS-7 cells transfected with G_{αs} and the β₂AR, or with G_{αs} and β₂AR plus EP₁ (A) or mouse primary ASM cells (B). Reactions were performed with partially purified cell membranes. The [³⁵S]GTPγS bound to G_{αs} was recovered by immunoprecipitation with G_{αs} antibody as described in Methods. (A) The presence of 17-PTP in the reaction significantly reduced [³⁵S]GTPγS binding stimulated by isoproterenol in the transfected COS-7 cells, while the β₂AR expressed without the EP₁ had the greatest stimulation by isoproterenol. No effect of 17-PTP was observed on cells transfected to only express β₂AR. ***P* = 0.05 versus β₂AR only; ##*P* = 0.02 versus 17-PTP-untreated cells coexpressing β₂AR and EP₁ receptor. Shown are mean ± SEM from 6 experiments. (B) The presence of 17-PTP in the reaction decreased isoproterenol-stimulated [³⁵S]GTPγS binding in ASM cell membranes. **P* < 0.05 versus isoproterenol in the absence of 17-PTP. Shown are mean ± SEM from 6 experiments. In both types of experiments, basal (non-agonist) binding was ~65% of isoproterenol-stimulated binding.

decrease in isoproterenol-stimulated binding, amounting to an approximately 60% reduction (*n* = 4; *P* = 0.02). Interestingly, the presence of the EP₁ receptor, even in the absence of its activation by agonist, decreased β₂AR coupling to G_s. As shown in Figure 6A, cells expressing β₂AR alone had the highest isoproterenol-stimulated [³⁵S]GTPγS binding, and as expected, 17-PTP had no effect in these cells. Additional studies were also carried out using ASM cell membranes (Figure 6B). Analogous to what was observed in the transfected cell studies, [³⁵S]GTPγS turnover was decreased when membranes were incubated with isoproterenol and 17-PTP compared with isoproterenol alone. These data then confirmed the concept of a direct interaction between the 2 receptors in the cell type of interest when the receptors were expressed at physiologic levels.

Discussion

PGE₂ is a cyclooxygenase-derived product of arachidonic acid metabolism that plays a major role in regulating such diverse physiological processes as smooth muscle tone, inflammation, and pain (reviewed in ref. 17). The biological effects of PGE₂ are often multifaceted and can appear contradictory to traditional signaling paradigms. As introduced earlier, some of these properties of PGE₂ are due to the fact that it is an endogenous agonist for 4 receptor subtypes coupled to several signal transduction cascades. The EP₁ receptor is linked to IP₃ and [Ca²⁺]_i turnover through G_q-mediated signal transduction (18), whereas the EP₂ and EP₄ receptors stimulate adenylyl cyclase and cAMP produc-

tion through G_s (19–21). Splice variants of the EP₃ receptor give rise to multiple isoforms that apparently can couple to G_i, G_s, or G_q (22, 23). Since one or more of these receptors may be expressed by a cell or tissue, the integrated response to PGE₂ often represents the net effect of additive or opposing signal transduction events. Moreover, temporal and spatial changes in patterns of EP receptor expression in response to various stimuli (e.g., injury and inflammation) provide an additional level of dynamic regulation and complexity to PGE₂-regulated processes in the airways. For example, recent studies of the PGE₂-mediated fibroproliferative response due to intratracheal bleomycin have shown an approximately 50% decrease in fibroblast EP₂ receptor mRNA without significant change in the other EP subtypes (24), whereas FITC-mediated fibrosis was associated with decreases in both EP₂ and EP₄ transcript expression (24).

Because PGE₂ is produced endogenously within the lung and activates pathways that may both enhance and inhibit ASM contraction, it may have particular relevance in asthma. PGE₂ is produced by ASM cells, bronchial and alveolar epithelial cells, fibroblasts, and inflammatory cells in the lung, and increased levels have been reported in some studies with human asthma (25, 26). Furthermore, an initial enzyme for prostanoid receptor synthesis (COX-2), as well as PGE₂ precursors, are increased in asthmatic airways (27, 28). Following its discovery, PGE₂ was found to relax ASM and thus has generally been considered to be a bronchoprotective prostanoid. The bronchodilatory effect of PGE₂ appears to be due to activation of the EP₂ receptor subtype, since PGE₂-mediated bronchodilation is present in EP₁-, EP₃-, and EP₄-null mice but absent in EP₂-null mice (29). Interestingly, in nonasthmatic humans, PGE₂ exerts a net bronchodilation; in asthmatics, though, the response is highly variable, sometimes resulting in significant bronchoconstriction (30). The complex nature of PGE₂ action in the asthmatic milieu is further illustrated in that it may serve as an intermediary through which inflammatory cytokines can modulate other receptor systems that govern airway tone. In particular, PGE₂ appears to mediate the inhibitory effects of the inflammatory cytokines TGFβ and interleukin-1 on β₂AR signal transduction in cultured ASM (31–34). These inflammatory cytokines stimulate COX-2 and production of PGE₂ by ASM, and it has been proposed that the resulting increase in cAMP production following EP₂ receptor activation leads to desensitization of the β₂AR (3). However, the mechanism(s) by which this inhibition occurs has not been completely elucidated, nor has the role of other EP receptor subtypes in mediating this effect been explored.

We define here a unique role for the EP₁ receptor expressed on ASM cells using both molecular studies in transfected cells and ASM cells and physiologic studies in trachea. We showed that the EP₁ receptor existed as a heterodimer (or higher-order oligomer) with the β₂AR, and the signaling of this complex was manifested by an altered β₂AR conformation and decreased functional interaction with G_s. EP₁ activation by agonist further enhanced β₂AR desensitization, as shown in the [³⁵S]GTPγS studies. This effect was also readily observed in ASM cells utilizing whole-cell cAMP measurements and membrane [³⁵S]GTPγS binding and manifested at the physiologic level with intact tracheal rings. Interestingly, the extent of β₂AR desensitization evoked by the EP₁ agonist, as assessed using these distinctly different methods, amounted to approximately 50% in all cases. In addition, the EP₁ receptor, while capable of stimulating intracellular IP₃ and [Ca²⁺]_i (as assessed by global whole-cell assays), did not ultimately cause significant ASM



contraction. Thus the dimer may serve not only to form a specialized signaling unit, but could also place the complex in a subcellular compartment that isolates it from activation of the downstream components of contraction. Indeed, β_2 ARs have been shown to be localized to caveolae, which are specialized flask-shaped invaginations of the cell membrane that promote association of the receptor with certain signal transduction elements (35). However, we note that EP₁ receptor-mediated contraction has been observed in guinea pig trachea (36), suggesting that the extent of EP₁-mediated tracheal contraction may depend on the species.

We have thus considered the physiologic role of the EP₁ receptor in ASM as *primarily* one of modulating β_2 AR function, with no observable direct effect on contraction. To our knowledge, this “monitor-like” function has not been previously described for a 7-TM receptor with its heterodimeric partner. However, there are somewhat analogous scenarios with other receptors. For the GABAB R1 and R2 subtypes, it appears that the R1:R2 heterodimer represents a distinct signaling unit, in that the R1 subtype binds the agonist while the R2 subtype couples to G protein activation (37). Furthermore, GABAB R1 cell surface expression requires heterodimer formation with GABAB R2 in the endoplasmic reticulum (38). For the α_{1D} AR, cell surface expression appears to be dependent on oligomeric association with the α_{1B} AR (39). Once so expressed, each subtype can signal independently. Similarly, the β_2 AR can form dimers with the α_{1D} AR and promote its expression (40). Other examples of homo- and heterodimerization function have been recently reviewed (12, 15, 16). We are unaware of a previous report of the unique function of a 7-TM receptor that we describe here.

In the current report, we utilized the physiologic readout of airway contraction and relaxation as the relevant functional signal and found that activation of the EP₁ receptor had no direct physiologic action. However, we found that EP₁ activation resulted in substantial loss of β_2 AR-mediated ASM relaxation. We demonstrated that this interaction of signal transduction networks occurred at the level of the receptors, rather than downstream effectors, and was not due to posttranslational modification of the β_2 AR by phosphorylation or other potential distal-pathway crosstalk mechanisms (41). Rather, formation of the EP₁: β_2 AR heterodimer was a distinct signaling complex, where the EP₁ receptor modulates β_2 AR coupling to G_{as}. Using multiple approaches, the EP₁-agonist dependency of the β_2 AR modulation was demonstrated. However, we cannot exclude the notion that the mere presence of the heterodimer acts to modulate the function of one of its members, the β_2 AR. Indeed, the [³⁵S]GTP γ S binding studies showed that β_2 AR coupling to G_{as} was decreased when coexpressed with the EP₁ receptor. However, the spontaneous switching of the EP₁ receptor to the active conformation may mimic the agonist-occupied form; thus it is not unexpected that coexpression, particularly at relatively high levels, would have an agonist-occupied effect. Regardless, in the native state both receptors were expressed on ASM, and we showed that EP₁ receptor activation decreased β_2 AR function in isolated ASM cells and in intact tissue.

In summary, we define a unique modulatory function of the EP₁ receptor, which acts primarily to regulate β_2 AR function. In asthma, then, local concentrations of elevated PGE₂ could act to quench β_2 AR function, resulting in decreased responsiveness to β -agonist. The mechanism of this receptor interaction appears to be due to the formation of a heterodimer; thus the EP₁: β_2 AR dimer should be considered the functional signaling complex in ASM,

with a unique phenotype based on steric interactions that lead to attenuated β_2 AR-G_{as} coupling.

Methods

Plasmid constructs, cells culture, and transfection. The human β_2 AR cDNA was tagged at the amino terminus with the FLAG epitope and fused at the carboxyl terminus with the cDNA for Rluc, and the cDNA for the human EP₁ receptor was fused at the carboxyl terminus with the cDNA for YFP, using standard techniques. The final constructs representing β_2 AR, FLAG- β_2 AR, and β_2 AR-Rluc were in pcDNA3, and the EP₁-YFP construct was in pEYFP-N1 (BD Biosciences – Clontech). Primary cultures of ASM cells were derived from explants of excised tracheas from the mice as reported previously (42) and maintained in DMEM, 10% FCS, and $\times 1$ antibiotic-antimycotic solution (Invitrogen Corp.) at 37°C in a 95% air, 5% CO₂ incubator. These cells maintain the morphologic and immunohistochemical characteristics of ASM cells for at least 15 passages, although most experiments were performed with cells in passages 4–12. Both COS-7 and HEK 293 cells were similarly maintained as monolayers in DMEM supplemented with 10% FCS, 100 U/ml penicillin, and 100 mg/ml streptomycin. Transfection of these cells for BRET and [³⁵S]GTP γ S binding experiments was performed using Lipofectamine (Invitrogen Corp.) by methods previously described (43).

RT-PCR. Total RNA was extracted from freshly isolated lungs with Tri-Reagent (Molecular Research Center Inc.). An aliquot of RNA was reverse transcribed with random hexamers with murine leukemia virus reverse transcriptase (PerkinElmer) as previously described (43). Transgene cDNA was amplified with 150 nM each of a sense primer (5'-CGTGTCCATTCCTGGGTGGCT-3') and an antisense primer homologous to regions within the mouse EP₁ open reading frame (5'-GTGGCTGTGGCTGAAGTGATG-3'). PCR was at 95°C for 120 seconds followed by amplification for 35 cycles at 95°C for 60 seconds and 56°C for 60 seconds and a final extension at 72°C for 7 minutes. The PCR products were detected in agarose gels stained with ethidium bromide. RNA samples not subjected to reverse transcription were used as negative controls to ensure that the observed PCR product was not the result of contaminating DNA.

Measurement of inositol phosphate production and [Ca²⁺]_i. Formation of [³H]-inositol phosphate was determined in intact ASM cells as previously reported (41). Briefly, near-confluent ASM cells in 12-well plates were incubated with inositol-free media containing [³H]-myo-inositol (5 μ Ci/ml) for 24 hours at 37°C in 5% CO₂ atmosphere. Cells were then washed and incubated with PBS for 30 minutes followed by a 30-minute incubation with 20 mM LiCl in PBS. The cells were treated with either PBS (basal), 1 μ M PGE₂, 1 μ M 17-PTP, or thrombin (a positive control) for 15 minutes, after which total inositol phosphates were extracted by column chromatography with AG1-X8 resin (Bio-Rad).

To measure [Ca²⁺]_i, murine ASM cells were seeded into black-walled, clear-based 96-well plates (Corning) at a density of approximately 80,000 cells per well and cultured overnight in 100 μ l DMEM supplemented with 10% FCS as above. On the following day, the cells were incubated at 37°C for 60 minutes with the Calcium 3 indicator (Molecular Devices) that was dissolved in Hank's balanced salt solution containing 20 mM HEPES, pH 7.4, and 2.5 mM probenecid in a final volume of 200 μ l/well. The cell plates were then placed into the FlexStation II scanning fluorimeter/luminometer (Molecular Devices) to monitor fluorescence before and after the addition of 50 μ l vehicle or a $\times 5$ concentration of the indicated drug (Figure 1, C and D) using an excitation wavelength of 485 nm, emission wavelength of 525 nm, and emission cutoff of 515 nm. Responses were measured as peak fluorescence intensity minus basal fluorescence intensity, with each data point in an individual experiment representing the mean of 4 replicates. Final data reported represent the mean of 4 different experiments using ASM cells derived from 2 different mice.



Radioligand binding and cAMP production. The expression of β_2 ARs in membranes prepared from transiently transfected HEK 293 and COS-7 cells was quantified by radioligand binding assays using the antagonist [125 I]-iodocyanopindolol as previously reported (44). To assess the rate of β_2 AR sequestration in ASM cells, whole-cell binding assays were performed using the hydrophilic, membrane-impermeant radioligand [3 H]-CGP-12177 to determine the amount of cell-surface β_2 AR in the untreated cells as well as in cells treated with either isoproterenol or isoproterenol with 17-PTP (1 μ M each) for periods up to 30 minutes as previously described in detail (45).

To measure cAMP production, ASM cells were seeded onto 24-well plates and maintained overnight in DMEM supplemented with 10% FCS to allow adherence. The cells were then switched to serum-free DMEM for an additional 24 hours. The cells were loaded with [3 H]-adenine (2 μ Ci/well) for 2 hours and then washed twice with PBS to remove excess adenine. Cells were then treated with either vehicle or 17-PTP (1 μ M final concentration) for 15 minutes before stimulation with either 1 μ M isoproterenol or 100 μ M forskolin for 15 minutes. Reactions were terminated by the addition of 50 μ l concentrated HCl. After the samples were frozen and thawed once to ensure cell lysis, cAMP was eluted by alumina column chromatography. Recovered counts were corrected for adenine uptake, and column efficiency ascertained with tracer [14 C]-cAMP. Each data point was determined from the mean of triplicate samples, and expressed as its percentage of the mean forskolin response for that experiment. The results shown represent the mean \pm SEM of 4 separate experiments performed with cell lines derived from 2 different mice.

Tracheal ring contractility studies. Studies of mouse tracheal ring contractility were performed as reported previously in detail (41). Briefly, tracheas were excised and dissected free of surrounding tissues and cut into rings of 5 mm in length. The rings were mounted on stainless steel wires connected to isometric force transducers and immersed in a physiologic saline solution (118 mM NaCl, 11 mM glucose, 4.73 mM KCl, 1.2 mM $MgCl_2$, 0.026 mM EDTA, 1.2 mM KH_2PO_4 , 2.5 mM $CaCl_2$, and 25 mM NaH_2CO_3) at 37°C and bubbled with 95% O_2 and 5% CO_2 , maintaining a pH of 7.40. Each tracheal ring was stretched to a tension of 5 mN, which we have previously determined to be an optimal passive tension for maximizing active force (42). Following a period of equilibration for 30 minutes, the indicated agents (Figures 2 and 3) were added to the bath, and the maximal response over the next 5 minutes was recorded. Cumulative concentration-isometric force curves were generated for the responses using standard curve-fitting techniques. Pretreatments of equilibrated rings included 1 μ M 17-PTP for 15 minutes, 10 nM PGE_2 for 15 minutes, 10 μ g/ml pertussis toxin for 4 hours, 100 μ M L-NAME for 30 minutes, 10 μ M indomethacin for 30 minutes, 5 μ M U73122 for 15 minutes, and 10 μ M Et-18-OCH $_3$ for 30 minutes. All mouse studies were reviewed and approved by the University of Cincinnati College of Medicine Animal Care and Use Committee.

Receptor phosphorylation. Whole-cell phosphorylation experiments were performed as previously described (46), with minor modifications. HEK 293 cells cotransfected with the β_2 AR-FLAG and EP $_1$ -YFP constructs were plated in 100-mm dishes. Forty-eight hours after transfection, the media was changed to serum- and phosphate-free DMEM containing 500 μ Ci/ml ^{32}P -orthophosphate. After incubation for 2 hours at 37°C in 95% air and 5% CO_2 , the cells were treated with vehicle, 1 μ M isoproterenol, 100 nM phorbol-12-myristate-13-acetate, 1 μ M 17-PTP, or 1 μ M isoproterenol plus 1 μ M 17-PTP for 15 minutes. The cells were then washed 4 times with ice-cold PBS and solubilized by rotating for 2 hours at 4°C in solubilization buffer, which consisted of 1% Triton X-100 (Sigma-Aldrich), 0.05% SDS, 0.5 mM EGTA, 1 mM EDTA, the phosphatase inhibitors sodium pyrophosphate and NaF (both at 10 mM), and the protease inhibitors leupeptin, aprotinin, benzamide and soybean trypsin inhibitor (5 μ g/ml each) in PBS. The lysates were centrifuged at 20,000 g for 10 minutes at 4°C

to remove unsolubilized material. β_2 AR-FLAG was immunoprecipitated by incubating the solubilized lysate overnight at 4°C with an anti-FLAG antibody pre-conjugated to agarose beads (Sigma-Aldrich). After washing the beads 4 times with ice-cold solubilization buffer, the tagged receptor was eluted with 3 \times FLAG peptide (Sigma-Aldrich). The eluate was fractionated by 10% SDS-PAGE, and the resultant signal from the dried gel was quantified by a GE Healthcare phosphorimager. Data was expressed as relative units from the pixel densities using ImageQuant software (version 1.2).

BRET measurements. BRET assays were carried out with the Flexstation II scanning fluorimeter/luminometer (Molecular Devices) using methods similar to that previously reported by Bouvier et al. (14). HEK 293 cells either transfected with β_2 AR-Rluc alone or cotransfected with receptor-YFP fusion constructs were detached 48 hours after transfection and seeded onto 96-well plates at a density of approximately 80,000 cells/well in PBS containing 0.1% glucose. To determine the full spectrum of emission, light-emission acquisition (420–560 nm) was started 5 minutes after coelenterazine h (5 μ M final concentration) was added. Emission values recorded at 475 nm and 530 nm after addition of coelenterazine h were used to determine the BRET ratio for each well. In some experiments, the transfected cells were treated with either isoproterenol or 17-PTP for 15 minutes prior to addition of coelenterazine h. The net BRET signal for each condition, performed in triplicate, was calculated by dividing the 530-nm emission of receptor-YFP by the 475-nm emission of the receptor-Rluc after subtracting the background signal determined from cells transfected with the receptor-Rluc construct alone using the formula $[(Em_{530} - Em_{475}) \times CF]/Em_{475}$, where CF represents Em_{530}/Em_{475} , the ratio of the signals from cells expressing receptor-Rluc alone. Maximal BRET levels (BRET $_{max}$) were obtained by cotransfecting increasing levels of receptor-YFP constructs with a constant quantity of the β_2 AR-Rluc construct. BRET $_{max}$ and BRET $_{50}$ were derived from fitting these data to a 1-site hyperbolic function.

Immunoprecipitation and Western blotting. HEK 293 cells expressing the EP $_1$ receptor, FLAG-tagged β_2 AR, or both were solubilized for 2 hours at 4°C in RIPA buffer (1 \times PBS, 1% IGEP, 0.5% sodium deoxycholate, and 0.1% SDS) containing protease inhibitors. The solubilized lysates were cleared by centrifugation at 10,000 g, and a portion of the cleared sample was removed for protein determination and Western blot analysis. The remainder was incubated with M2 anti-FLAG affinity matrix (Sigma-Aldrich) overnight at 4°C. The resin was then collected by centrifugation, washed 3 times with Tris-buffered saline, and eluted with 2 \times Laemmli loading buffer. For Western blot analysis, whole cell lysates and immunoprecipitated samples were fractionated on 8% polyacrylamide gels and transferred to PVDF membranes. Blots were blocked with 5% nonfat milk and incubated overnight at 4°C with a polyclonal anti-EP $_1$ antibody (Cayman Chemical Co.) diluted 1:200. The blots were then washed, incubated with anti-rabbit IgG antibody diluted 1:6,500 at room temperature for 1 hour, and detected by enhanced chemiluminescence.

[^{35}S]GTP γ S binding. [^{35}S]GTP γ S binding assays were performed as described previously (47) with modifications that include a $G_{\alpha s}$ immunoprecipitation step (48). HEK 293 cells were cotransfected with plasmid vectors encoding human $G_{\alpha s}$ and either β_2 AR alone or β_2 AR plus the EP $_1$ receptor. Studies with ASM cells were carried out without transfections utilizing endogenous expression of the receptors and $G_{\alpha s}$. For the binding reaction, partially purified cell membrane protein (~75 μ g) was incubated for 15 minutes at 30°C in a binding buffer (10 mM HEPES, pH 7.4, 3 mM $MgCl_2$, and 50 mM NaCl) that included 50 nCi [^{35}S]GTP γ S and 10 μ M guanosine diphosphate in a total volume of 100 μ l. Basal (vehicle) and agonist-stimulated (1 μ M isoproterenol) conditions were each determined in the absence and presence of 1 μ M 17-PTP. Parallel assays containing unlabeled GTP γ S (100 μ M) were used to define nonspecific binding.



The reactions were stopped by addition of cold solubilization buffer (10 mM HEPES, pH 7.4, 50 mM NaCl, 5 mM EDTA, and 0.5% Triton-X) and rotated for 2 hours at 4°C. After centrifugation at 20,000 g for 15 minutes, the supernatant was incubated overnight with 1 µg G_{αs} antibody (Santa Cruz Biotechnology Inc.) and 20 µl protein A-conjugated agarose beads. The beads were washed 3 times with phosphorylation solubilization buffer, and the bound radioactivity was measured in a liquid scintillation counter. Under these conditions, nonspecific binding was typically <10% of the total. Agonist-stimulated binding was derived by subtracting the counts of basal samples from the counts of the isoproterenol-treated samples. Each data point within an experiment was determined from triplicate samples.

Immunofluorescence microscopy. Murine ASM cells were passed to Lab-Tek chamber slides (Nunc) and grown overnight. After the media was aspirated, the cells were washed with PBS and then fixed with cold methanol for 4 minutes. Cells were then placed in blocking buffer composed of PBS containing 10% donkey serum and 1% Triton-X for 30 minutes at room temperature. The blocking buffer was removed, and cells were incubated overnight at room temperature in PBS containing 1% Triton-X, a 1:50 dilution of a rabbit anti-EP1 antibody (Santa Cruz Biotechnology Inc.), and a 1:200 dilution of a chicken anti-β₂AR antibody (Abcam). Primary antibody was not added to the negative controls. After incubation, the buffer containing the primary antibodies was aspirated, and the cells were washed with PBS containing 0.1% Triton-X. Cells were then incubated with secondary antibodies for 2–4 hours with 0.1% Triton-X in PBS containing a 1:200 dilution

of a Cy2-conjugated donkey anti-rabbit antibody and a 1:800 dilution of a Cy3-conjugated donkey anti-chicken antibody. Cells were then washed with PBS and mounted with Gelmount (Biomedica) to reduce fading. Fluorescent images were captured electronically using the appropriate filter sets on a Zeiss Axioplan II Microscope System at a magnification of ×200.

Statistics. Concentration-response and BRET data were fit by nonlinear functions using the Prism software program (version 4.0; GraphPad Software). Statistical comparisons were by paired or unpaired Student's *t* tests as appropriate, with *P* < 0.05 considered significant. Data are represented as mean ± SEM of the indicated number of experiments.

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