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Ligand regulation of green fluorescent protein-tagged forms of the human β_1 - and β_2 -adrenoceptors; comparisons with the unmodified receptors

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- 1 Stable clones of HEK293 cells expressing either FLAGTM epitope-tagged, wild type human β_1 and β_2 -adrenoceptors or C-terminally green fluorescent protein (GFP)-tagged forms of these receptors were established.
- 2 The binding affinity of [3H]-dihydroalprenolol and other ligands was little affected by addition of GFP to the C-terminal of either receptor.
- 3 Isoprenaline induced the internalisation of both β_1 -adrenoceptor-GFP and β_2 -adrenoceptor-GFP and following removal of the agonist both constructs were able to recycle to the cell surface.
- The extent of internalisation of β_2 -adrenoceptor-GFP produced by isoprenaline was substantially greater than for β_1 -adrenoceptor-GFP.
- 5 C-terminal addition of GFP slowed markedly the rate of internalization of both the β_1 adrenoceptor and the β_2 -adrenoceptor in response to isoprenaline.
- 6 Sustained exposure to isoprenaline (24 h) produced substantially greater levels of downregulation of native β_2 -adrenoceptor compared to β_2 -adrenoceptor-GFP although both were equally effectively removed from the plasma membrane.
- 7 Sustained exposure to isoprenaline resulted in a large fraction of β_2 -adrenoceptor-GFP becoming trapped in internal vesicles/lysosomes but not degraded.
- 8 Even after sustained exposure to isoprenaline a significant fraction of β_1 -adrenoceptor-GFP remained at the cell surface.
- 9 These results indicate that although GFP tagging of β -adrenoceptors can provide qualitative visual patterns of agonist-induced receptor trafficking and regulation in HEK293 cells the quantitative details vary markedly from those obtained with the unmodified receptors. British Journal of Pharmacology (2000) 130, 1825-1832

 β -adrenoceptor; green fluorescent protein; internalization; trafficking; HEK293 cells

Abbreviations: DHA, dihydroalprenolol; EBP50, ezrin radixin-moesin (ERM)-binding phosphoprotein-50; GFP, green fluorescent protein; GPCR, G protein-coupled receptor; KRH, Krebs-Ringer-HEPES; PBS, phosphate-buffered

Introduction

A desire to visualize the cellular location and trafficking of G protein-coupled receptors (GPCRs) following their activation has produced a number of studies utilising forms of GPCRs which have been altered by addition of the Green Fluorescent Protein (GFP) from the jellyfish Aequorea victoria to their Cterminal tail (see Milligan, 1999 for review). Such studies have been highly informative, providing information about the mechanism and kinetics of internalization of a range of GPCRs. However, in most cases there have not been direct comparisons with effects of agonists on the unmodified GPCR. C-terminal addition of GFP to a range of GPCRs does not prevent agonist-induced activation of G proteins and downstream effectors nor does it prevent agonist-induced phosphorylation of GPCRs or their interactions with members of the arrestin family of proteins involved in their desensitization and internalization via clathrin-coated vesicles (Milligan, 1999).

Using co-immunoprecipitation and co-localization approaches, for both the β_1 -adrenoceptor (Tang et al., 1999), and the β_2 -adrenoceptor (Hall et al., 1998a,b; Cao et al., 1999;

Luttrell et al., 1999), a number of other proteins, apart from G proteins, G protein-coupled receptor kinases and arrestins, have been shown to interact directly with these GPCRs (see Hall et al., 1999 for review). Interestingly, a number of these proteins have been shown to interact with the β_2 -adrenoceptor via amino acids at the extreme C-terminus of the GPCR. These include a protein named EBP50 (for ezrin radixin-moesin (ERM)-binding phosphoprotein-50) (Cao et al., 1999) which has been shown to play an important role in determining whether internalized β_2 -adrenoceptor is recycled to the plasma membrane or targeted to a separate set of vesicles which may lead to lysosomal degradation. As little as a single amino acid alteration at the C-terminal tail of the β_2 -adrenoceptor is sufficient to disrupt such interactions and thus potentially to modify the trafficking of the GPCR (Hall et al., 1998a; Cao et al., 1999). Addition of GFP to the C-terminus of the GPCR would be anticipated to prevent such interactions. We have thus generated clones of HEK293-cells stably expressing either wild type or C-terminally GFP-tagged forms of the β_1 - and β_2 adrenoceptors to explore the basic characteristics and kinetics of internalization and recycling of these GPCRs using combinations of confocal microscopy and intact cell [3H]-

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ligand binding studies. The addition of GFP to the C-terminus of both GPCRs substantially slowed their rate of internalization and the capacity of the cell to dispose of the proteins during long-term exposure to agonist. Some of this data has previously been presented in preliminary form (McLean & Milligan, 1999)

Methods

Materials and methods

[³H]-DHA (64 Ci mmol⁻¹), [³H]-CGP-12177 (44 Ci mmol⁻¹), [³H]-adenine and [³H]-cyclic AMP were purchased from Amersham International. All reagents for cell culture were from Life Technologies (Paisley, Strathclyde, U.K.). Receptor ligands were from RBI. All other general laboratory reagents were from Sigma or Fisons and were of the highest purity available.

Construction of GFP tagged forms of the β_1 - and β_2 -adrenoceptors

Human β_1 -adrenoceptor in pcDNA3 was amplified by PCR using a HindIII-FLAG forward primer, 5'-AAAAAAAAGC-TTGCCAC CATGG ACTA CAAGGACGAC GATGATAA-GGGCGCGGGGGTGCTC-3', and a BamHI reverse primer, 5'-AAAAAGGATCCTCCCGCCACCTTGGATTCCGAGG-C-3'. This removed the stop codon and the initiating methionine (start codon) of the β_1 -adrenoceptor, with an initiator ATG being present in the N-terminally added FLAGTM epitope tag (ATG GAC TAC AAG GAC GAC GAT GAT AAG). The PCR product was digested with HindIII and BamHI and the resulting fragment ligated into pcDNA3. A modified form of GFP (Zernicka-Goetz et al., 1997) was also amplified by PCR using a BamHI forward primer, 5'-AAAAA GGATCC AGT AAA GGA GAA GAA CTT TTC-3', and an XbaI reverse primer, 5'-TGCTCTA-GATTATTTGTATAGTTCATCCATGCCATG-3'. This removed the initiating methionine of GFP and the resulting PCR product was digested and linked inframe to generate the FLAGTM β_1 -adrenoceptor-GFP construct. Construction of the human β_2 -adrenoceptor-GFP construct has been described previously (McLean et al., 1999).

Transient and stable transfection of HEK293 cells

HEK293 cells were maintained in Minimum Essential Medium (MEM, Sigma) supplemented with 0.292 g l⁻¹ L-glutamine, and 10% newborn calf serum at 37°C. Cells were grown to 60–80% confluency prior to transient transfection. Transfection was performed using LipofectAMINE reagent (Life Technology, Inc.) according to the manufacturers' instructions. To generate cell lines stably expressing the various constructs, two days after transfection cells were seeded/diluted and maintained in MEM medium supplemented with 1 mg ml⁻¹ Geneticin sulphate (Life Technology, Inc.). Medium was replaced every 3 days with MEM medium containing 1 mg ml⁻¹ Geneticin sulphate.

Confocal laser scanning microscopy

Cells were observed using a laser scanning confocal microscope (Zeiss Axiovert 100) using a Zeiss Plan-Apo 63×1.40 NA oil immersion objective, pinhole of 20, and electronic zoom 2 or 3. The GFP was excited using a 488 nm argon/krypton laser and

detected with 515–540 nm band pass filter. The images were manipulated with Zeiss LSM or MetaMorph software. When examining the time course of internalization and recycling cells were grown on glass coverslips and mounted on the imaging chamber. Cells were maintained in KRH buffer and temperature was maintained at 37°C.

[³H]-ligand binding studies

Stably transfected cells were plated into 24 well microtitre plates the day before the experiment. After appropriate drug treatments the cells were washed twice with ice-cold phosphate-buffered saline (PBS (mm): KCl 2.7, NaCl 137, KH₂PO₄ 1.5, Na₂HPO₄ 8, pH 7.4) followed by addition of 200 μl of binding mix ([3H]-CGP-12177 (10 nm) or [3H]-DHA (2 nm) in Krebs-Ringer-HEPES buffer (KRH (mm): NaCl 130, KCl 5, MgSO₄ 1.2, CaCl₂ 1.2, HEPES 20, Na₂PO₄ 1.2, glucose 10, 0.1% BSA; pH 7.4) buffer). Non-specific binding was assessed by the presence of 10 μ M betaxolol (β_1 -adrenoceptor) or propranolol (β_2 -adrenoceptor). [³H]-DHA binding assays were performed at 30°C for 45 min and [3H]-CGP-12177 binding at 4°C for 90 min in KRH buffer. All experiments were terminated by removal of the binding medium and washing of the cells with ice-cold PBS. 0.5 ml of 0.5 mM EDTA in PBS was added to detach cells from the plate and this volume plus a further 0.5 ml wash of the wells of the microtitre plate were counted. For each assay equivalent unlabelled cells were removed and counted to calculate receptor levels.

Intact cell adenylyl cyclase activity measurements

Were performed essentially as described in (Wong, 1994; Merkouris *et al.*, 1997) Cells were split into wells of a 12-well plate and the cells were allowed to reattach. Cells were then incubated in medium containing [3 H]-adenine (1.5 μ Ci well ${}^{-1}$) for 16–24 h. The generation of [3 H]-cyclic AMP in response to treatment of the cells with various ligands and other reagents was then assessed.

Electrophoresis and immunoblot analysis

Membrane samples were resolved by SDS-PAGE. Proteins were then electrophoretically transferred to nitrocellulose. The nitrocellulose filter was blocked for 1 h in 3% fat-free milk in PBS-T buffer (PBS containing 0.1% Tween 20). After a brief wash in PBS-T buffer, it was incubated overnight at 4°C with an appropriate primary antiserum diluted in PBS-T buffer containing 1% fat-free milk. Polyclonal antisera to either the β_1 - or the β_2 - adrenoceptor (Santa Cruz) were used for detection of the constructs. The primary antiserum was then removed and the blot washed extensively in PBS-T buffer. Subsequent incubation with secondary antibody (donkey antirabbit IgG conjugated with horseradish peroxidase, (Scottish Antibody Production Unit, Carluke, Scotland) proceeded for 2 h at RT and after extensive washing in PBS-T buffer the blot was visualised by ECL (Amersham).

Results

A cDNA encoding a modified form of GFP from *Aequorea* victoria was ligated to the 3' end of cDNAs encoding the human forms of each of the β_1 - and β_2 -adrenoceptors using a polymerase chain reaction-based approach which allowed removal of the stop codons from the GPCR cDNAs. Although

not utilized directly within the current studies, both the β_1 - and β_2 -adrenoceptor cDNAs were also modified at the 5' end to introduce the FLAGTM epitope tag sequence (Asp-Tyr-Lys-Asp-Asp-AspAsp-Lys) at the N-terminus of the proteins. cDNAs encoding the FLAGTM-tagged β_1 - and β_2 -adrenoceptors and the GFP-tagged forms of these (Figure 1), ligated to the mammalian expression vector pCDNA3, were transfected into HEK293 cells. This plasmid also expresses a genetecin resistance marker and clones which displayed resistance to genetecin sulphate (1 mg ml⁻¹) were selected and expanded. Initial screening of clones for the expression of the β_1 adrenoceptor-GFP and β_2 -adrenoceptor-GFP was performed by visual examination in a confocal microscope to detect the autofluorescence of GFP. A significant number of such clones were selected based on clear, plasma membrane-delineated, autofluorescence of the GPCR-GFP fusion protein. Initial screening of the non-GFP tagged forms of the GPCRs was performed by monitoring the specific binding of a single, near saturating, concentration of the β -adrenoceptor antagonist [³H]-dihydroalprenolol ([³H]-DHA). Subsequent saturation [3H]-DHA binding studies were performed on membrane preparations from selected clones to obtain both B_{max} values and measures of ligand affinity for the various constructs (Table 1). Although the measured K_d for [3H]-DHA at the β_1 adrenoceptor constructs was approximately twice that at the β_2 -adrenoceptor constructs, the addition of GFP to the Cterminus of either GPCR had little effect on the binding affinity of [3H]-DHA (Table 1). However, clones expressing the GFP-tagged forms of the GPCRs generally expressed considerably higher levels of [3H]-DHA binding sites than those expressing the equivalent untagged GPCRs. We have previously also noted this feature in clones expressing wild type and GFP-tagged forms of the hamster α_{1b} -adrenoceptor (Steven, P.A. and Milligan, G., unpublished) and believe this to reflect the well appreciated long half-life of GFP. These membranes were also immunoblotted with both β_1 - and β_2 adrenoceptor selective antisera to confirm the identity of the correct GPCRs in each clone (Figure 2). The β -adrenoceptor agonist isoprenaline stimulated adenylyl cyclase activity in a concentration-dependent manner in all clones tested, further confirming (Barak et al., 1997; Kallal et al., 1998) that addition of GFP to the C-terminal tail of either the β_1 - or β_2 adrenoceptor did not prevent productive interactions of the GPCRs with G_s (data not shown).

When grown on a glass cover slip and then imaged on a confocal microscope clones expressing the β_1 -adrenoceptor-GFP construct displayed an essentially homogenous plasma

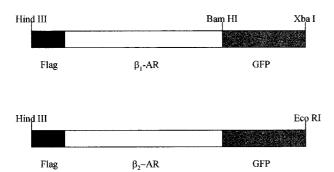


Figure 1 Schematic diagram of the constructs which were stably expressed in HEK293 cells. cDNAs encoding the human β_1 - and β_2 -adrenoceptor (AR) were modified at their N-terminus to encode a FLAGTM epitope. Following removal of the stop codon from the receptor sequences GFP was added in-frame.

Table 1 Expression and ligand binding characteristics of GFP and non-GFP-tagged forms of the human β_1 - and β_2 -adrenoceptors

Construct	$[^3H]$ - DHA K_d (nm)	B_{max} (pmol mg ⁻¹ membrane protein)
$\begin{array}{l} {\rm FLAG^{TM}} \ \beta_{1}\text{-}{\rm AR} \\ {\rm FLAG^{TM}} \ \beta_{1}\text{-}{\rm AR}\text{-}{\rm GFP} \\ {\rm FLAG^{TM}} \ \beta_{2}\text{-}{\rm AR} \\ {\rm FLAG^{TM}} \ \beta_{2}\text{-}{\rm AR}\text{-}{\rm GFP} \end{array}$	$\begin{array}{c} 1.1 \pm 0.2 \\ 1.8 \pm 0.3 \\ 0.6 \pm 0.3 \\ 0.8 \pm 0.2 \end{array}$	2.6 ± 0.4 18.8 ± 2.5 4.1 ± 0.9 8.7 ± 0.7

Data represent means \pm s.e.mean from three independent experiments using different membrane preparations.

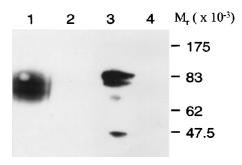


Figure 2 Immunoblots of β₁-adrenoceptor and β₂-adrenoceptor constructs. Membranes expressing FLAGTM β₂-adrenoceptor-GFP (1 and 4) or FLAGTM β₁-adrenoceptor-GFP (2 and 3) were immunoblotted using either anti-β₂-adrenoceptor (1 and 2) or anti-β₁-adrenoceptor (3 and 4) antisera. Similar results were obtained in another set of immunoblots.

membrane distribution of this protein. Addition of isoprenaline (10 μ M) at 37°C for varying times resulted in the plasma membrane signal becoming reduced in intensity and the construct appearing in punctate intracellular vesicles (Figure 3). However, exposure of the cells to isoprenaline for greater than 10 min was required to observe a significant level of internalization. A similar pattern and time-course of internalization was also observed in other clones expressing this construct (data not shown). Betaxolol is a selective β_1 adrenoceptor blocker. Competition binding experiments in membranes expressing either the β_1 -adrenoceptor or β_1 adrenoceptor-GFP with [3H]-DHA confirmed that the high affinity of betaxolol for the β_1 -adrenoceptor was not compromized by addition of GFP (Table 2). Following pretreatment of β_1 -adrenoceptor-GFP expressing cells with isoprenaline (10 μ M, 30 min, 37°C), removal of the agonist and its replacement with betaxolol (10 μ M) allowed recycling of the GPCR construct to the plasma membrane (Figure 4). Equivalent studies were performed with a clone expressing β_2 -adrenoceptor-GFP. Again, isoprenaline (10 μ M) produced internalization of the construct from the plasma membrane into punctate intracellular vesicles and removal of the agonist and replacement with the antagonist alprenolol (10 μ M) allowed efficient recycling of the construct to the plasma membrane (Figure 5).

Despite the ease of visualization of agonist-induced internalization of these constructs it is difficult to use this approach to provide more than a qualitative analysis. [³H]-DHA is a hydrophobic antagonist and thus in intact cell binding studies it can cross the plasma membrane. In this situation, specific binding of this ligand represents a combination of cell surface and internal GPCR. By contrast, [³H]-CGP-12177 is a hydrophilic antagonist which in intact cell

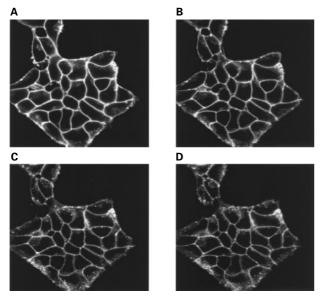


Figure 3 The β_1 -adrenoceptor-GFP construct is internalized following addition of isoprenaline. A group of cells stably expressing the FLAGTM β_1 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for 0 (A), 10 (B), 20 (C) or 40 (D) min during visualization in the confocal microscope. Three separate experiments produced similar results.

Table 2 Ligand binding characteristics of GFP and non-GFP-tagged forms of the human β_1 -adrenoceptor

Construct	Isoprenaline (corrected IC ₅₀) (nM)	Betaxolol (K _i , nm)
FLAG TM β_1 -AR FLAG TM β_1 -AR-GFP	127 ± 10 289 ± 64	16 ± 0.5 25 ± 12

Data represent means \pm s.e.mean from three independent experiments.

A B
C D

Figure 4 Replacement of isoprenaline with betaxolol allows the β_1 -adrenoceptor-GFP construct to recycle to the plasma membrane. Cells stably expressing the FLAGTM β_1 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for 0 (A) or 30 (B–D) min. Following washing in the presence of betaxolol (10 μ M) the distribution of the receptor was visualized 10 (C) and 30 (D) min later. Three separate experiments produced similar results.

binding studies labels only cell surface GPCRs. In preliminary experiments the specific binding at 4°C of [3H]-CGP-12177 to intact cells expressing each of the β_1 - and β_2 -adrenoceptor constructs achieved steady-state within 60 min. All the subsequent binding experiments with this ligand were thus incubated for 90 min at 4°C to ensure maximal binding but to prevent recycling of any internalized GPCRs to the cell surface during the course of the assay as this is an energy and temperature-dependent process. Following incubation of cells expressing either the β_2 -adrenoceptor or the β_2 -adrenoceptor-GFP with isoprenaline (10 μ M) for 30 min at 4°C to allow the agonist to bind but at a temperature at which GPCR internalization would not proceed, the cells were washed with buffer between 0-4 times prior to initiating a specific [3H]-CGP-12177 (10 nm) binding assay. With only a single wash the levels of specific [3H]-CGP-12177 binding achieved was the same as in cells which had not been pre-exposed to isoprenaline, indicating effective removal of isoprenaline. However, two washes of the cells were routinely applied in subsequent studies to ensure this would be the case. Isoprenaline (10 μ M) was then added to cells at 37°C for between 10– 60 min and, following cooling and washing to remove the agonist, the specific binding of either [3H]-CGP-12177 (at 4°C) or [3H]-DHA (at 30°C, to allow ligand to penetrate the cells and bind both cell surface and internal receptors) was determined. For both the β_2 -adrenoceptor and the β_2 adrenoceptor-GFP expressing cells specific [3H]-DHA binding was unaltered over this time period of isoprenaline treatment indicating total cellular levels of the GPCRs were not reduced (Figure 6). However, for both cell lines cell surface levels of the GPCRs, monitored by the binding of [3H]-CGP-12177, were reduced substantially. Although cell surface levels of the β_2 adrenoceptor were reduced by some 40% within 10 min of exposure to isoprenaline, significant loss of the β_2 -adrenoceptor-GFP construct from the cell surface took longer (Figure 7). However, following 60 min exposure to isoprenaline some 60% of both the β_2 -adrenoceptor and β_2 -adrenoceptor-GFP had become internalized (Figure 7). Similar results were

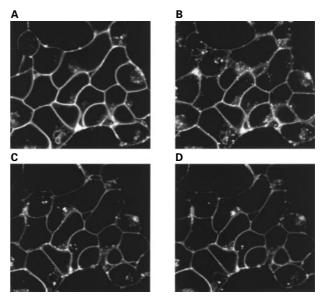


Figure 5 Internalization and recycling of β_2 -adrenoceptor-GFP. Cells stably expressing the FLAGTM β_2 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for 0 (A) or 30 (B–D) min. Following washing with buffer containing alprenolol (10 μ M) the distribution of the receptor was monitored 30 (C) and 40 (D) min later. Three separate experiments produced similar results.

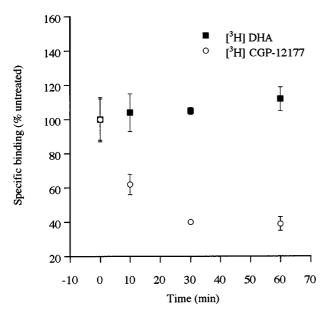


Figure 6 Short-term treatment with isoprenaline internalizes but does not downregulate the FLAGTM β_2 -adrenoceptor-GFP construct. Cells stably expressing the FLAGTM β_2 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for varying times. Specific binding of a single concentration of either [3 H]-DHA (2 nM) or [3 H]-CGP-12177 (10 nM) to intact cells was then measured as described in Methods.

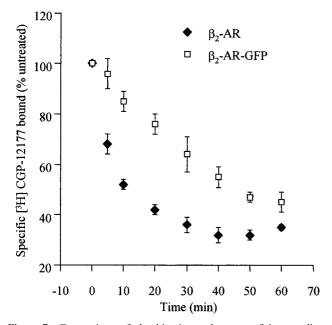


Figure 7 Comparison of the kinetics and extent of isoprenaline-induced internalization of the β_2 -adrenoceptor and β_2 -adrenoceptor-GFP. Cells stably expressing either the FLAGTM β_2 -adrenoceptor or the FLAGTM β_2 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for varying times. Specific binding of a single concentration of [3 H]-CGP-12177 (10 nM) to intact cells was then measured as described in Methods. Results are presented as means \pm s.e.m. for five (FLAGTM β_2 -adrenoceptor) or four (FLAGTM β_2 -adrenoceptor-GFP) experiments.

obtained for the cells expressing the β_1 -adrenoceptor and β_1 -adrenoceptor-GFP (Figure 8). Internalization of the wild type β_1 -adrenoceptor in response to isoprenaline was more rapid than internalization of β_1 -adrenoceptor-GFP, although following a 60 min exposure the two constructs achieved similar (between 30-40%) extents of internalization which was substantially lower than for the β_2 -adrenoceptor constructs.

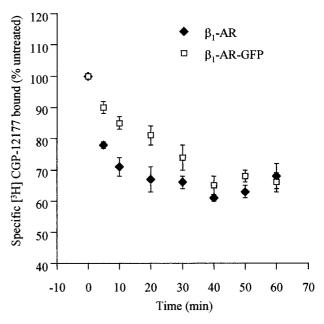
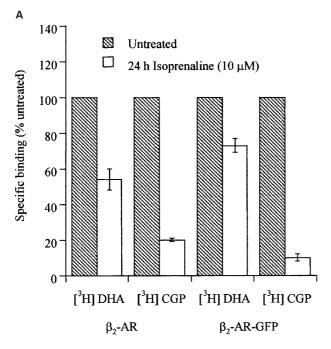


Figure 8 Comparison of the kinetics and extent of isoprenaline-induced internalization of the β_1 -adrenoceptor and β_1 -adrenoceptor-GFP. Cells stably expressing either the FLAG^TM β_1 -adrenoceptor or the FLAG^TM β_1 -adrenoceptor-GFP construct were exposed to isoprenaline (10 μ M) for varying times. Specific binding of a single concentration of [3 H]-CGP-12177 (10 nM) to intact cells was then measured as described in Methods. Results are presented as means \pm s.e.m. for three (FLAG^TM β_1 -adrenoceptor) or five (FLAG^TM β_1 -adrenoceptor-GFP) experiments.

Long-term exposure of cells to agonist often results in a substantial downregulation of total cellular levels of a GPCR. Indeed, following exposure of wild type β_2 adrenoceptor expressing cells to isoprenaline (10 μ M) for 24 h, levels of intact cell [3H]-DHA binding were reduced by 40% and the vast majority of the remaining GPCR was now internalized because specific [3H]-CGP-12177 binding was reduced by 80% (Figure 9A). However, when equivalent experiments were performed on cells expressing β_2 -adrenoceptor-GFP, although very little cell surface GPCR could be measured as [3H]-CGP-12177 binding sites, there was little downregulation of the construct as [3H]-DHA binding was reduced by less than 20% (Figure 9A). To examine this further, membrane fractions were prepared from untreated and isoprenaline (10 µM, 24 h) treated cells expressing either the β_2 -adrenoceptor or β_2 -adrenoceptor-GFP and [³H]-DHA binding monitored. These confirmed that isoprenaline produced a large downregulation of the β_2 -adrenoceptor but not of β_2 -adrenoceptor-GFP (Figure 9B). To examine the basis for this, β_2 -adrenoceptor-GFP expressing cells were grown on glass cover slips and exposed to isoprenaline (10 µM, 24 h). Untreated and agonist-treated cells were then visualized (Figure 10). As anticipated from the intact cell binding studies, although virtually all the construct was intracellular it was still present in high levels. Rather than being present in small, discrete, punctate vesicles as seen with short periods of exposure to isoprenaline, it was predominantly located in large, perinuclear, vesicles. In similar experiments with cells expressing β_1 -adrenoceptor-GFP, 24 h after treatment with isoprenaline (10 μ M) internalization of the construct could be observed but a large amount of cell surface GPCR remained (Figure 11) and parallel ligand binding studies indicated virtually no downregulation of the protein had occurred (data not shown).



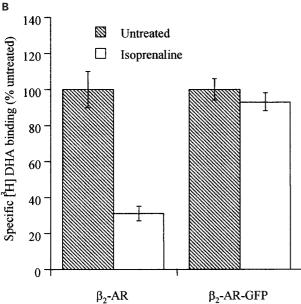


Figure 9 Long-term treatment with isoprenaline. Effects on distribution and downregulation of the $β_2$ -adrenoceptor and $β_2$ -adrenoceptor-GFP monitored by ligand binding studies. (A) Intact Cells. Cells stably expressing either the FLAGTM $β_2$ -adrenoceptor or the FLAGTM $β_2$ -adrenoceptor-GFP construct were either untreated or exposed to isoprenaline (10 μM) for 24 h. Specific binding of a single concentration of either [3 H]-DHA (2 nM) or [3 H]-CGP-12177 (10 nM) to intact cells was then measured as described in Methods. Results are presented as means ±s.e.m. for three separate experiments. (B) Membranes. Cells stably expressing either the FLAGTM $β_2$ -adrenoceptor or the FLAGTM $β_2$ -adrenoceptor-GFP construct were either untreated or exposed to isoprenaline (10 μM) for 24 h. Membranes were prepared and the specific binding of a single concentration of [3 H]-DHA (2 nM) was then measured as described in Methods. Results are presented as means ±s.e.m. for three separate experiments.

Discussion

C-terminal attachment of forms of GFP to the C-terminal tail of GPCRs has recently become a popular strategy to visualize their cellular location to investigate the processes of agonist-induced internalization and recycling to the plasma membrane.

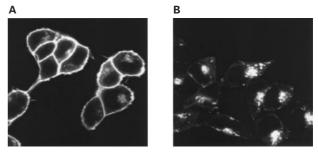


Figure 10 Visualization of the location of β_2 -adrenoceptor-GFP following long-term challenge with isoprenaline. Cells stably expressing the FLAGTM β_2 -adrenoceptor-GFP construct were either untreated (A) exposed to isoprenaline (10 μ M) for 24 h (B). The distribution of the construct was then visualized. Two further experiments produced similar patterns.

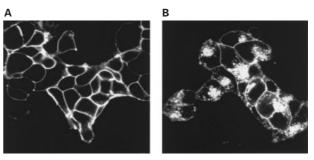


Figure 11 Visualization of the location of β_1 -adrenoceptor-GFP following long-term challenge with isoprenaline. Cells stably expressing the FLAGTM β_1 -adrenoceptor-GFP construct were either untreated (A) exposed to isoprenaline (10 μ M) for 24 h (B) The distribution of the construct was then visualized. Two further experiments produced similar patterns.

GPCRs which have been modified in this manner include the β_1 - (McLean & Milligan, 1999) and β_2 -adrenoceptors (Barak et al., 1997; Kallal et al., 1998; Conway et al., 1999; McLean et al., 1999), the α_{1a} - (Hirasawa et al., 1997) and α_{1b} adrenoceptors (Awaji et al., 1998), the CXCR1 receptor (Barlic et al., 1999), the parathyroid hormone receptor (Conway et al., 1999), the thyrotropin-releasing hormone receptor-1 (Drmota et al., 1998; 1999; Groarke et al., 1999), the Ca²⁺ sensing receptor (Gama & Brietwieser, 1998), the V₂ vasopressin receptor (Schulein et al., 1998) and the cholecystokinin type A receptor (Tarasova et al., 1997). This strategy has many obvious benefits, the most useful being the capacity to monitor the location and movement of the construct in living cells over an extended period of time. However, there are also potential drawbacks. These include that the GPCR is generally increased in size by at least 50% by attachment of the 27 kDa GFP. Furthermore, limitations of resolution of GFP visualization in living cells makes it unclear whether true colocalization rather than overlapping distribution with a second protein is observed (see Drmota et al., 1999; Groarke et al., 1999 for examples). It is generally accepted that addition of GFP does not prevent GPCR interaction with G proteins and this was again confirmed in the present studies for both the β_1 adrenoceptor-GFP and β_2 -adrenoceptor-GFP constructs as both were able to produce a robust, concentration-dependent, stimulation of adenylyl cyclase upon addition of isoprenaline. However, recent studies have indicated the capacity of GPCRs, including the β_1 - and β_2 -adrenoceptors (see Hall et al., 1999 for review) to interact with a range of other intracellular proteins. The C-terminal tail of the β_2 -adrenoceptor appears vital for a number of these interactions including with EBP50 (Cao et al., 1999) which relies on the presence of Ser411. This amino acid is three amino acids from the C-terminal tail of the β_2 -adrenoceptor and is a target for phosphorylation by GRK-5 (Cao et al., 1999). Furthermore, single amino acid alterations at the extreme C-terminus of the β_2 -adrenoceptor have been reported to alter such interactions (Hall et al., 1998a,b). Addition of GFP to the C-terminal tail of a GPCR clearly must modify this region. Indeed, in many cases the sequence of the extreme C-terminal tail is altered during this process, predominantly for expediency and ease of the molecular manipulations, without considering the effects this might have on protein function. Purely in terms of ligand binding, and often when monitoring G protein and effector activation, such sequence alterations are unlikely to be a major issue. Indeed, herein we observed little variation in the binding affinity of either agonist or antagonist ligands following addition of GFP to the C-terminal tail of either the β_1 - or β_2 -adrenoceptor (Table 2). Moreover, isoprenaline was able to stimulate adenylyl cyclase activity via all of the constructs. However, we observed two key alterations in ligand regulation of the C-terminally GFP-tagged form of the β -adrenoceptors compared to the wild type forms. Firstly, isoprenaline-induced internalisation of the GFP-tagged constructs was substantially slower than of the wild type forms of both GPCRs. The molecular basis for this effect has not been established. However, a number of C-terminal truncations of GPCRs are known to slow ligand-induced internalization, e.g. for the rat thyrotropin releasing hormone receptor-1 (Drmota & Milligan, 2000). Indeed, in the case of the gonadotrophin releasinghormone receptors from mammalian species, the absence of a C-terminal tail appears responsible for their very slow rates of agonist-induced internalization (Vrecl et al., 1998). The equivalent GPCR from catfish has a C-terminal tail and both this GPCR and the mammalian version with the tail of the rat thyrotropin releasing hormone receptor-1 appended internalize rapidly in response to agonist when expressed in HEK293 cells (Heding et al., 2000). It is also interesting to note that although C-terminal truncation of the rat thyrotropin releasing hormone receptor-1 slows the rate of internalization it does not affect the recycling rate constant of the GPCR once it has been internalized (Drmota & Milligan, 2000), suggesting that the C-terminal tail of the GPCR is not directly involved in controlling recycling to the cell surface. The current studies indicate that a substantial addition to the C-terminal tail of a GPCR can also markedly slow the rate of internalization. In studies on gonadotrophin releasing-hormone receptors and the thyrotropin releasing hormone receptor-1, the capacity of arrestins to interact with the C-terminal of the GPCR appears intimately involved in the effectiveness of internalization. However, addition of GFP to the C-terminal tail of the rat thyrotropin releasing hormone receptor-1 certainly does not prevent interactions with arrestin-2 (Groarke et al., 1999) but whether it alters the affinity of these interactions has not been explored.

The β_1 - and β_2 -adrenoceptors are highly suitable GPCRs for these type of studies because well characterized radiolabelled ligands are available to measure either total receptor levels or only those at the cell surface in intact cell binding studies. These were then used to provide a quantitative monitor of the rate of internalization of both the GFP-tagged and unmodified GPCRs. Direct visualization of the GPCRs was limited to the GFP-tagged forms and without the use of complex algorithms for analysis (Conway *et al.*, 1999) can provide only qualitative pictures of agonist effects on GPCR localization. However, there are still distinct benefits in being able to directly visualize the presence and distribution of the GFP-tagged forms of the GPCRs.

The second feature of ligand regulation of the GFP-tagged β_1 - and β_2 -adrenoceptors which was altered was the effectiveness of downregulation of the GPCRs during sustained cellular exposure to agonist. Following sustained challenge of the β_2 -adrenoceptor expressing cells with a high concentration of isoprenaline, intact cell ligand binding studies confirmed both physical loss of some 40% of the receptor population and the removal of most of the receptor from the cell surface. However, although a similar removal of β_2 adrenoceptor-GFP from the plasma membrane was evident the protein was apparently not effectively downregulated. Visualisation of such cells confirmed the maintained presence of high levels of β_2 -adrenoceptor-GFP inside the cells and the combination of binding studies and imaging was thus able to confirm that the GFP-linked protein still maintained ligand binding capacity. GFP has a long half-life and it may well be this feature which resulted in the maintained presence of the GFP-tagged β_2 -adrenoceptor in large intracellular vesicles which appear to be lysosomes. It is thus noteworthy that the majority of the clones we were able to isolate following transfection with the β_2 -adrenoceptor-GFP cDNA expressed the protein to relatively high levels. Indeed, clones expressing the protein at less than 3-4 pmol mg⁻¹ membrane protein were not found. This also has been a feature of clones of cells we have isolated expressing GFP-tagged forms of other GPCRs, including the α_{1b} -adrenoceptor. Whilst this is useful in many regards, it may be more difficult to study certain aspects of the regulation of these GPCRs if levels of expression start to outstrip endogenous levels of various, required, assessory proteins. We anticipated that addition of GFP to the C-terminus of the β_2 -adrenoceptor might modulate some aspects of its regulatory cycle as this is a known site of protein - protein interactions which can link this GPCR to the actin cytoskeleton (Cao et al., 1999). The current studies demonstrate again some of the benefits of the use of GFPtagging of proteins but also suggest that a range of comparative studies with the unmodified protein are required in parallel to obtain a useful picture of the kinetics of their regulation and movement.

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