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THEMED ISSUE: GPCR RESEARCH PAPER

Pharmacogenetic characterization of indacaterol, a novel β_2 -adrenoceptor agonist

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Background and purpose: Indacaterol is a novel β_2 -adrenoceptor agonist in development for the treatment of chronic obstructive pulmonary disease. The aim of this study was to investigate the comparative pharmacology of indacaterol in recombinant cells expressing the common polymorphic variants of the human β_2 -adrenoceptor and in human primary airway smooth muscle (ASM) cells.

Experimental approach: Chinese hamster ovarian-K1 cell lines expressing high and low levels of the common human β_2 -adrenoceptor variants were generated [Gly16-Glu27-Val34-Thr164(GEVT), RQVT, GQVT] and also the rare GQVI variant. Human primary ASM cells were isolated from explants of trachealis muscle. Adenosine-3',5'-cyclic-monophosphate production was used as an outcome measure.

Key results: In both the low- and high-expression recombinant GEVT 'wild type' cell lines indacaterol is a high-efficacy agonist. Salmeterol and formoterol were identified as low- and high-efficacy agonists, respectively, and showed similar potencies to indacaterol irrespective of the β_2 -adrenoceptor genotype. The I164 variant cell line was associated with a reduced capacity to generate adenosine-3',5'-cyclic-monophosphate in response to β_2 -adrenoceptor agonist. In the human primary ASM cells indacaterol gave a maximal response intermediate between that of salmeterol and formoterol.

Conclusions and implications: These data demonstrate that indacaterol is a high-efficacy agonist in recombinant cell systems but acts with lower efficacy in human primary ASM cells. No marked genotype-dependent effects were observed for common variants; however, changes in I164 receptor activity were identified, which were dependent on the level of expression of β_2 -adrenoceptors.

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Keywords: asthma; chronic obstructive pulmonary disease; β_2 -adrenoceptor agonist; indacaterol; pharmacogenetics; polymorphism; recombinant; haplotype; cAMP

Abbreviations: ASM, airway smooth muscle; cAMP, adenosine-3',5'-cyclic-monophosphate; CHO, Chinese hamster ovarian; CYP, cyanopindolol; LABA, long acting β_2 -adrenoceptor agonist; SABA, short acting β_2 -adrenoceptor agonist

Introduction

 β_2 -adrenoceptor agonists are the mainstay bronchodilators used in the treatment of asthma and chronic obstructive pulmonary disease. These drugs fall into two main classes,

namely short acting β_2 -adrenoceptor agonists (SABAs) and long acting β_2 -adrenoceptor agonists (LABAs) (Hall and Tattersfield, 1992). The available drugs in clinical usage differ in their basic pharmacology, both in their onset and duration of action, and also in the degree of agonism, with some drugs (e.g. salmeterol) being partial agonists in most cell systems, and other drugs (e.g. formoterol) being full agonists. Recently novel LABAs have been developed for potential clinical usage; one such drug is indacaterol, which in phase 3 trials has been shown to exhibit a once-daily bronchodilator action (Battram *et al.*, 2006; Beeh *et al.*, 2007; Beier *et al.*, 2007; Naline *et al.*, 2007).

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I Sayers et al

The actions of β_2 -adrenoceptor agonists at the human β_2 -adrenoceptor are potentially affected by common polymorphisms within the ADRB2 gene. A number of synonymous and non-synonymous single nucleotide polymorphisms (SNPs) have been described at this locus (Reihsaus et al., 1993). Of these, the two most frequent non-synonymous SNPs result in the presence of either an arginine (R) or a glycine (G) at codon 16 and a glutamine (Q) or a glutamate (E) at codon 27. A third polymorphism potentially results in a methionine (M) substitution for a valine (V) at position 34, but this is extremely rare and hence we have not studied it further. In addition, there is a rare but potentially important non-synonymous SNP that results in either a threonine (T) or an isoleucine (I) at codon 164. The I164 form of the receptor has been shown to be associated with reduced acute responses to most catechol ligands in fibroblast cell lines expressing the β_2 -adrenoceptor (Green et al., 2001). In these Chinese hamster fibroblasts and in human primary cultured airway smooth muscle (ASM) lines, codon 16 or 27 polymorphisms have not been shown to be the acute actions of isoprenaline. However, differences in down-regulation profiles have been identified, at least in transformed cell lines, with reduced receptor downregulation with the E27 form of the receptor and enhanced down-regulation with the G16 form of the receptor (Green et al., 1994; 1995). Codons 16 and 27 reside in the N-terminal extracellular end of the receptor, and the structure-function mechanism underlying altered receptor down-regulation is unclear at this time. Interestingly, the I164 variant resides in the fourth transmembrane domain adjacent to \$165, which is predicted to interact with the β-OH group of adrenoceptor ligands. This may at least in part explain the reduced acute responses of β_2 -adrenoceptor agonists in the I164 variant. The reduced duration of action of the LABA salmeterol in cells expressing the I164 variant has also been rationalized by the concept of a reduced exosite binding (Green et al., 2001). It is important to note that in addition to polymorphic variation, β_2 -adrenoceptor expression and function is also influenced by transcriptional, translational and post-translational mechanisms.

There have been extensive clinical studies that have addressed the possible contribution of polymorphisms to

clinically relevant end points. While it is unlikely these polymorphisms increase the risk of developing asthma (Hall et al., 2006), there are data to suggest they may influence disease progression (Hall et al., 2006) and also response to treatment when SABAs are administered regularly in steroid naïve patients (Israel et al., 2000; 2004). There are conflicting data on the influence of genotype on responses to LABAs, with some studies showing significant reduction in response in individuals homozygous for the R16 form of the receptor (Wechsler et al., 2006), and other studies suggesting maintained bronchodilator responses, at least in subjects taking inhaled steroids (Bleecker et al., 2006). Whether or not the response to indacaterol is altered by genetic variants in the human β₂-adrenoceptor has not to date been studied. Similarly a direct comparison of the ability of indacaterol to stimulate cAMP (adenosine-3',5'-cyclic-monophosphate) production with other clinically relevant compounds on different genetic backgrounds has not been investigated.

The aims of the current study were: (i) to create a series of cell lines expressing the common haplotypes of the human β_2 -adrenoceptor matched for receptor expression; (ii) to investigate the comparative pharmacology of indacaterol in cell lines expressing the 'wild type' form of the β_2 -adrenoceptor; and (iii) to examine the possibility that genetic variation in the β_2 -adrenoceptor alters the response to indacaterol or other LABAs.

Methods

Plasmid construction

PCR was used to amplify the Human *ADRB2* coding region from genomic DNA template using primers designed to engineer a 5'EcoRI site and a 3'XhoI site (See Table 1). PCR conditions included 100–500 ng genomic DNA template, 0.5 U platinum taq high fidelity DNA polymerase, 0.3 mmol·L $^{-1}$ dNTPs, 1 mmol·L $^{-1}$ magnesium sulphate, 0.2 µmol·L $^{-1}$ of each primer in a reaction volume of 50 µL. Cycling conditions were 94°C for 2 min, then 30 cycles of 94°C for 1 min, 60°C for 1 min, 68°C for 2 min. The PCR product was agarose gelpurified, digested with EcoRI and XhoI and cloned into

Table 1 Primers

ADRB2 expression cassette isolation 5'primer GACT-GAATTC-CGCGCCATGGGGCAAC ADRB2 expression cassette sequencing	3'primer GACT- <u>CTCGAG</u> - TTA CAGCAGTGAGTCATTGGTA	Location 214-1461
5'primers	3'primers	Location
TAATACGACTCACTATAGGG		PCDNA3/T7
	CAACAGATGGCTGGCAACTA	PCDNA3/1071
	TCTGCAGACGCTCGAAC	399-415
	GACATGGAAGCGGCCCTCAG	926-945
ATTGCCTCTTCCATCGTCTC		820-839
Site-directed mutagenesis		
5'primer	3'primer	Location
CTTGCTGGCACCCAAT <u>AGA</u> AGCCATGCGCCGG	CCGGCGCATGGCT TCT ATTGGGTGCCAGCAAG	GLY16 > ARG
CCACGACGTCACGCAG CAA AGGGACGAGGTGTG	CACACCTCGTCCC TTG CTGCGTGACGTCGTGG	GLU27 > GLN
GGATTGTGTCAGGCCTT <u>ATC</u> TCCTTCTTGCCCATTC	GAATGGGCAGAGAGGA <mark>GAT</mark> AAGGCCTGACACAATCC	THR164 > ILE

ADRB2 numbering reference sequence NM_000024 (ATG = 220). Underlined are the EcoRI and XhoI restriction enzyme sites. Nucleotides representing codons of interest are shown in bold.

pcDNA3.1 using standard molecular biology techniques. The ADRB2 cassette of the pcDNA3-ADRB2 plasmid was sequenced using BigDye (v3.1) terminator sequencing in conjunction with the ABI-310 DNA sequencer and various sequencing primers (Table 1). Multiple genomic DNA samples were used as template, and plasmids were screened until a Gly16-Glu27-Val34-Thr164 (GEVT) haplotype was obtained. The pcDNA3-ADRB2-GEVT plasmid represents the common haplotype observed in the Caucasian population (Drysdale et al., 2000) and was used as the starting template for sitedirected mutagenesis. The pcDNA3-ADRB2-GQVT, RQVT, GQVI plasmids were generated sequentially by site-directed mutagenesis of the pcDNA3-ADRB2-GEVT backbone using the Quickchange site-directed mutagenesis kit as directed by the manufacturer in conjunction with mutagenic primers (Table 1). All pcDNA3-ADRB2 plasmids were sequenceverified, and large-scale preparations were completed using the Plasmid maxi kit as directed by the manufacturer.

Stable transfection of Chinese hamster ovarian-K1 cells

Chinese hamster ovarian (CHO)-K1 cells were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% foetal calf serum, 2 mmol·L⁻¹ glutamine, $100~U \cdot mL^{\text{--}1}$ penicillin, $0.1~mg \cdot mL^{\text{--}1}$ streptomycin at $37^{\circ}\text{C}/5\%$ CO₂ and passaged by using trypsin/EDTA. For transfection, 10 mL of cells (10⁵ cells·mL⁻¹) were plated in a 10 cm petri dish and grown for 24 h. Six micrograms of pcDNA3-ADRB2 constructs or pcDNA3 empty vector were linearized by using PvuI followed by heat inactivation. CHO-K1 cells were transfected with linearized DNA using Fugene6 as directed by the manufacturer using a 3:1 ratio of Fugene reagent to DNA. Transfected cells were grown for another 24 h followed by passaging into 24-well plates followed by a further 24 h incubation. Complete medium was then replaced by selection medium (complete medium supplemented with 500 µg⋅mL⁻¹ G418), and the cells were maintained for 10–14 days with the addition of fresh selection medium every 2–3 days. Single cell clones of the transfected CHO-K1 cell lines were isolated in 96-well plates by flow cytometry using the cell sorting function (forward/side scatter) of the Coulter ultra flow cytometer. Single cell clones were maintained in complete medium using 20% foetal calf serum to promote survival; medium was replaced every 4-5 days. Clonal cell lines were maintained in selection medium for the rest of the study.

β_2 -adrenoceptor binding studies

Two confluent T162 flasks of cells were used for the preparation of membranes. Medium was aspirated, and cells were washed with ice-cold buffer A (10 mmol·L⁻¹ HEPES, 0.9% NaCl, 0.2% EDTA, pH 7.4) and then incubated with buffer A at 37°C for 10 min. Cells were isolated by using a cell scraper, and a cell pellet was generated by centrifugation at 500× g for 5 min. The cell pellet was resuspended in 2 mL of buffer B (10 mmol·L⁻¹ HEPES, 10 mmol·L⁻¹ EDTA, pH 7.4) and two cell pellets combined and the volume made to 10 mL. The cell suspension was homogenized using an IKA Ultra Turrax T25 polytron using four bursts of 5 s with 20 s intervals between bursts. A further 15 mL of buffer B was added, and the cell suspension was

centrifuged at 39 000× g for 25 min at 4°C. The cell pellet was resuspended in 10 mL buffer C (10 mmol·L⁻¹ HEPES, 0.1 mmol·L⁻¹ EDTA, pH 7.4), and the homogenization procedure was repeated. The cell suspension was centrifuged at 39 000× g for 25 min at 4°C and finally resuspended in 1 mL buffer C. Protein was quantified by using the Bradford protein micro-assay as directed by the manufacturer, and preparations were normalized to 1 mg·mL⁻¹. Samples were snap-frozen in liquid nitrogen and stored at -80°C.

 $[^{125}I]$ -iodo (\pm) -cyanopindolol $([^{125}I]$ -iodo-CYP) saturation binding was used to determine the β_2 -adrenoceptor density in the membrane preparations. Briefly, 100 µL (100 µg) of membrane preparation, 50 µL assay buffer (HBSS with Ca2+ and $Mg^{2+}/0.1\%$ BSA, pH 7.4), 50 µL containing [125I]-iodo-CYP at a range of pmol·L⁻¹ concentrations and a further 50 μL of assay buffer for total counts or 50 µL assay buffer containing 10 μmol·L⁻¹ alprenolol for non-specific binding determination was added to each well of a 96-well round bottom plate. Plates were incubated at room temperature for 2 h. Plates were harvested by using a Packard Filtration harvester onto GF/B filter plates and dried for 2 h at 35°C. Plates were backsealed, and 40 µL Microscintillant 20 was added to each well; the plate was sealed, and c.p.m. per well was determined by using a Packard Topcount. K_d and B_{max} values were generated by using GraphPad Prism v5 (GraphPad, San Diego, CA, USA). All analyses were completed at least in duplicate.

cAMP assav

Cells were seeded in selection medium into 24-well plates at 10⁵ cells per well and grown for 22 h prior to loading of the cells with [3H]-adenine (2 µCi per well) for 2 h. Excess radiolabel was removed by washing with Hanks HEPES buffer, and then 1 mL of Hanks HEPES buffer was added to each well. Plates were incubated at 37°C for subsequent addition of β_2 -adrenoceptor agonists for various time points. Reactions were terminated by the addition of 50 μL of concentrated HCl followed by freezing at -20°C. cAMP production was quantified by column chromatography, and results were corrected for adenine uptake and column efficiency using [14C]-cAMP as described previously (Freyer et al., 2004). Triplicate wells were counted for each condition, and the data are expressed as number of counts (compared with basal counts) or %isoprenaline response (100%). The response to 1 μmol·L⁻¹ isoprenaline was routinely greater than threefold over basal counts in low-expression cell lines except the low-expression GQVI cell line that had a reduced response to all agonists tested (<1.5fold over basal). Typically, the response to 1 μmol·L⁻¹ isoprenaline was 10-fold greater than the basal counts in highexpression cell lines. Mean data from at least three completed in triplicate presented experiments are $(\pm s.e.mean).$

Human primary airway smooth muscle cultures

Human ASM cells were isolated from explants of trachealis muscle obtained from individuals with no history of respiratory disease as described previously (Daykin *et al.*, 1993; Sayers *et al.*, 2006). Briefly, a section of trachealis muscle was isolated by dissection just above the carina and washed in

DMEM supplemented with penicillin (200 U·mL⁻¹), streptomycin (200 µg·L⁻¹) and amphotericin B (0.5 µg·L⁻¹). Explants $(0.2 \times 0.2 \text{ cm})$ were placed in 6-well plates and allowed to adhere. Cells were allowed to grow with fresh complete medium (10% foetal calf serum and 2 mmol·L⁻¹ glutamine) added regularly, after 7-10 days growth cells approached confluency. As cells reached confluency, explants were removed and the cells were harvested by using 5 mg·mL⁻¹ trypsin, 2 mg⋅mL⁻¹ EDTA. Primary ASM cells exhibited >95% cell staining for α -actin. ASM cells were maintained in complete medium and passaged using trypsin/EDTA. Ethical approval for the use of primary cells was obtained from the local ethics committee. Six ASM donors were used in the current analyses with genotypes: (G/R)(E/Q)VT, G(E/Q)VT, G(E/Q)VT, RQVT, GEVT and RQVT. cAMP production in response to β₂-adrenoceptor agonists was measured in human ASM cells in an analogous manner to that described for the recombinant cell lines, except after seeding cells were grown for 2-3 days until ~70-80% confluent prior to [3H]-adenine loading.

Statistical analyses

Differences between the results for the various treatment groups were compared by analysis of variance (ANOVA) in conjunction with Bonferonni's post hoc test or Student's t-test, as appropriate. Figures represent mean values (±s.e.mean). Statistical analyses and curve fitting were completed by using GraphPad Prism v5 (GraphPad, San Diego, CA, USA); a P-value< 0.05 was considered significant.

Materials

All chemicals were analytical grade or higher. Plasticware was from Costar (High Wycombe, UK). All chemicals and reagents were purchased from Sigma-Aldrich (Poole, UK) unless otherwise stated. Platinum taq high fidelity DNA polymerase, pcDNA3.1 and DMEM were obtained from Invitrogen (Paisley, UK); EcoRI, XhoI and PvuI, Promega (Southampton, UK); ABI-310 DNA sequencer, Applied Biosystems (Warrington, UK); Quickchange site-directed mutagenesis kit, Stratagene (Amsterdam, The Netherlands); plasmid maxi kit, Qiagen (Crawley, UK); Fugene6, Roche Diagnostics (Newhaven, UK); Bradford protein micro-assay, Bio-Rad (Hemel Hempstead, UK); [125I]-iodo-CYP, Perkin Elmer (Life and Analytical Sciences, Boston, MA, USA).

Results

Creation of cell lines expressing human β_2 -adrenoceptor variants Initial work was performed to create both low-(<1000 pmol·mg⁻¹) and high-(>2000 pmol·mg⁻¹) expressing CHO-K1 β₂-adrenoceptor cell lines stably expressing the common combinations of ADRB2 polymorphisms. Because of linkage disequilibrium, the combination of arginine 16 glutamine 27 (R16Q27) and glycine 16 glutamate 27 (G16E27) are much commoner than would be expected from the respective allele frequencies of the relevant SNPs. Isoleucine (I) 164 occurs most frequently on the background

Table 2 β₂-adrenoceptor expression levels in recombinant Chinese hamster ovarian-K1 β₂-adrenoceptor cell lines

	Protein (fmol·mg⁻¹)
High level expression	
GEVT	3948
GQVT	2720
RQVT	2745
GQVI	2181
Low level expression	
GEVT	668.1
GQVT	204.9
RQVT	1074
GQVI	302.4
pcDNA(-)	0

Data from [1251]-iodo(±)-cyanopindolol saturation binding assay.

G16Q27, and hence in order to compare responses of different agonists at the commoner haplotypes at this locus the following cell lines were made: G16E27V34T164 (GEVT), R16Q27V34T164 (RQVT), G16Q27V34T164 (GQVT) and G16Q27V34I164 (GQVI). These haplotypes have previously been identified in the Caucasian population with frequencies of GEVT (48.3%), GQVT (13.2%), RQVT (33.0%) and GQVI (1.0%) (Drysdale et al., 2000). In all, 18 clonal cell lines were studied for levels of β_2 -adrenoceptor expression using [125I]-iodo-CYP binding. Cell lines with <100 fmol·mg⁻¹ β_2 -adrenoceptor expression did not generate a measurable and robust cAMP response to 10 μmol·L⁻¹ isoprenaline and were not studied further (data not shown). For each common haplotype, cell lines were chosen for further study where expression was matched as closely as was feasible. No [125I]-iodo-CYP binding was observed for the pcDNA3(-) transfected cell line, confirming that the expression of endogenous β_2 -adrenoceptors could not be detected in this cell line and validating the use of CHO-K1 cells as a host in order to investigate human β_2 -adrenoceptor function. The levels of expression for the different lines chosen for further study are shown in Table 2.

Time course of cAMP formation in 'wild type' (GEVT) cells to a range of β_2 -adrenoceptor agonists

Figure 1 shows the time course of cAMP formation in response to a range of β₂-adrenoceptor agonists in lowexpressing β₂-adrenoceptor GEVT CHO-K1 cells. It can be seen that cAMP formation is essentially linear over the first 60 min stimulation with agonist. Agonists were used at concentrations predicted from previous work to induce near-maximal responses, that is, 1 μmol·L⁻¹. There was a significant difference between the drug treatment groups (two-way ANOVA, P = 0.006). Based on these single dose experiments, it appears that salmeterol (as previously described) is a relatively lowefficacy agonist, whereas indacaterol, like formoterol shows higher efficacy. Similar experiments were performed for all the cell lines expressing different variants of the β_2 -adrenoceptor. In all lines cAMP formation was linear over this time period, and similar findings were observed, that is, that indacaterol and formoterol are higher-efficacy agonists compared with salmeterol. Experiments with CHO-K1 cells transfected with pcDNA3 alone (i.e. without β_2 -adrenoceptor expression inserts) showed no increase in cAMP following exposure to any of the agonists studied complementing the [125I]-iodo-CYP binding data (n = 3, data not shown).

Dose–response relationships of β_2 -adrenoceptor agonists

Figure 2 shows the dose–response relationships for all the β_2 -adrenoceptor agonists studied in the four high-expression cell lines, expressing the three common haplotypes at the *ADRB2* locus and also for the rare GQVI form of the receptor. Figure 3 shows the data from the same set of experiments in the low-expression cell lines. These results are in general similar, although there is somewhat greater variability in responses. EC_{50} and maximal response values derived from these experiments are shown in Tables 3 and 4. However, due to the low response in some cell lines we were unable to

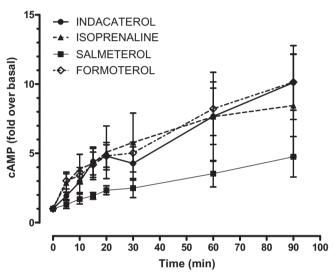


Figure 1 Time course of cAMP (adenosine-3′,5′-cyclic-monophosphate) production in response to 1 μ mol·L⁻¹ β_2 -adrenocepter agonist in the low-expression GEVT 'wild type' cell line. Data are expressed as mean fold increase in cAMP compared with basal level; vertical lines represent s.e.mean (n=3).

calculate accurate values for all the agonists for each cell line combination. This was particularly relevant for the low-expression GQVI cell line that gave a limited cAMP response to agonist (e.g. 1 μ mol·L⁻¹ isoprenaline response was ~1.5-fold over basal) making accurate pEC₅₀ and E_{max} calculations problematic. This low signal to background also gave the impression that salmeterol is a high-efficiency agonist on this haplotype background.

In the low-expression cell lines there were significant differences between the EC₅₀ values generated for the three clinically relevant β_2 -adrenoceptor agonists (pEC₅₀ indacaterol 8.67 ± 0.24 , salmeterol 9.79 ± 0.49 and formoterol 9.81 ± 0.24 , P < 0.001) using the GEVT cell line (Table 3). The significance of the ANOVA was driven by the reduced potency of indacaterol versus salmeterol (P = 0.007) and formoterol (P = 0.0001). Comparison of the EC₅₀ values for indacaterol, salmeterol and formoterol in the high-expression cell lines identified significant differences for the GQVT (pEC50 10.18 ± 0.21 vs. 9.27 ± 0.23 vs. 9.41 ± 0.13 respectively) and RQVT (pEC₅₀ 8.64 ± 0.12 vs. 8.92 ± 0.20 vs. 9.36 ± 0.07 respectively) cell lines (P = 0.004 and P = 0.001 respectively). The significance of the ANOVA was driven by the decreased potency of indacaterol versus formoterol for the RQVT cell line (P = 0.0001) and modest differences between indacaterol and salmeterol (P = 0.020) and formoterol (P = 0.013) for the GQVT cell line (Table 3). Interestingly, there was a trend in both high- and low-expression cell lines for the β_2 -adrenoceptor haplotype to influence agonist potency: for GEVT/RQVT, formoterol > indacaterol and for GQVT, indacaterol > formoterol.

The maximal response at 40 min post β_2 -adrenoceptor agonist confirmed our previous data suggesting that indacaterol and formoterol are higher-efficacy agonists compared with salmeterol (Table 4). Significant differences between the maximal responses were observed, but these were mainly driven by the low-level salmeterol responses. Indacaterol and formoterol overall had similar maximal responses (Table 4).

A comparison of indacaterol responses between cell lines expressing different β_2 -adrenoceptor haplotypes identified a significant difference in EC₅₀ values in the high-expression cell lines (P < 0.0001) potentially reflecting an increased

Table 3 *p*EC₅₀ values

•							
	Indacaterol	Isoprenaline	Salmeterol	Formoterol			
High level expression							
GEVT	8.83 ± 0.23	7.55 ± 0.08	9.00 ± 0.28	9.11 ± 0.08			
GQVT*	10.18 ± 0.21	8.38 ± -0.14	$9.27 \pm 0.23^{\circ}$	$9.41 \pm 0.13^{\circ}$			
RQVT*	8.64 ± 0.12	8.18 ± 0.12	8.92 ± 0.20	$9.36 \pm 0.07^{\$}$			
GQVI	8.87 ± 0.10	7.58 ± 0.05	8.37 ± 0.43	8.67 ± 0.11			
Low level expression							
GEVT*	8.67 ± 0.24	7.73 ± 0.10	$9.79 \pm 0.49^{\circ}$	$9.81 \pm 0.24^{\circ}$			
GQVT	8.85 ± 0.16	7.19 ± 0.11	NC	8.54 ± 0.10			
RQVT	8.33 ± 0.13	7.90 ± 0.18	8.69 ± 0.61	9.15 ± 0.21			
GQVI	NC	NC	NC	NC			
Human airway smooth muscle	8.53 ± 0.16	6.96 ± 0.05	10.00 ± 0.38	8.65 ± 0.17			

Chinese hamster ovarian-K1 β_2 -adrenoceptor data were generated by using 40 min time point (n = 3), human airway smooth muscle by using 10 min time point (n = 12 - 15). Data represent mean \pm s.e.mean. NC = not calculated, experiments were completed; however, due to the low response accurate EC₅₀ values could not be determined. EC₅₀ values for the clinically relevant β_2 -adrenoceptor agonists (indacaterol, salmeterol, formoterol) were compared for each cell line by using the sum of squares F-test (GraphPad Prism v5) *P < 0.004. Statistically significant differences between indacaterol EC₅₀ values and clinically relevant β_2 -adrenoceptor agonists were identified (5P < 0.05).

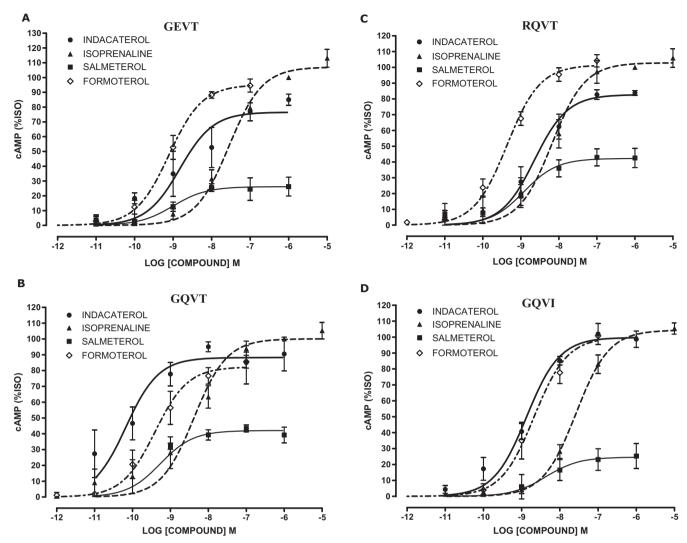


Figure 2 Concentration—response curves for cAMP (adenosine-3',5'-cyclic-monophosphate) production in response to $β_2$ -adrenoceptor agonists (40 min) in the high-expression $β_2$ -adrenocepter cell lines. Variants cell lines (A) GEVT, (B) GQVT, (C) RQVT and (D) GQVI. Data were normalized to response to 1 μmol·L⁻¹ isoprenaline (100%) in each experiment and are presented as mean percentage isoprenaline (%ISO) response; error bars represent s.e.mean (n = 3).

Table 4 E_{max} values

	Indacaterol	Isoprenaline	Salmeterol	Formoterol
High level expression				
GEVT*	77 ± 6	100	26 ± 3 ^{\$}	95 ± 3\$
GQVT*	88 ± 6	100	42 ± 3\$	82 ± 4
RQVT*	83 ± 4	100	42 ± 3\$	$102 \pm 3^{\$}$
GQVI*	100 ± 4	100	25 ± 4 ^{\$}	100 ± 5
Low level expression				
GEVT*	89 ± 3	100	$36 \pm 6^{\$}$	$65 \pm 6^{\$}$
GQVT	83 ± 5	100	NC	94 ± 4
RQVT*	80 ± 4	100	26 ± 6 ^s	$113 \pm 10^{\circ}$
GQVI	NC	NC	NC	NC
Human airway smooth muscle *	48 ± 2	100	10 ± 1\$	79 ± 4 ^{\$}

Chinese hamster ovarian-K1 β_2 -adrenoceptor data were generated using 40 min time point (n = 3), human airway smooth muscle using 10 min time point (n = 12 - 15). Data represent mean \pm s.e.mean. NC = not calculated, experiments were completed; however, due to the low response accurate E_{max} values could not be determined. E_{max} values for clinically relevant β_2 -adrenoceptor agonists (indacaterol, salmeterol, formoterol) were compared for each cell line by using the sum of squares F-test (GraphPad Prism v5, *F < 0.001). Statistically significant differences between indacaterol E_{max} values and other clinically relevant β_2 -adrenoceptor agonists were identified (5P < 0.05).

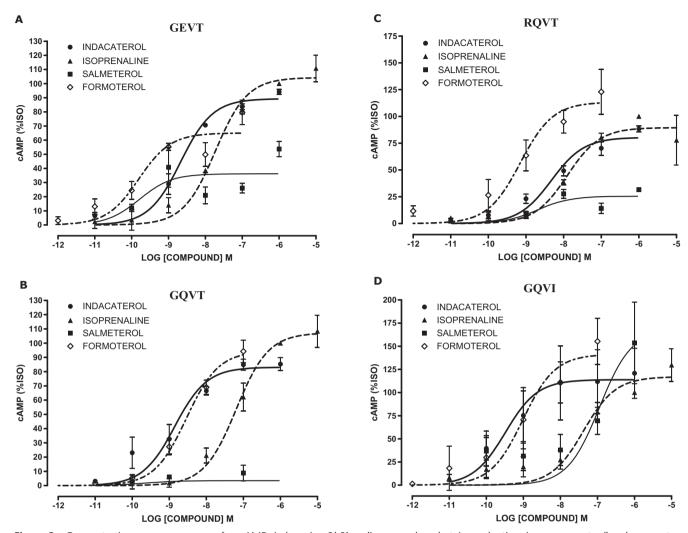


Figure 3 Concentration–response curves for cAMP (adenosine-3',5'-cyclic-monophosphate) production in response to $β_2$ -adrenoceptor agonists (40 min) in the low-expression $β_2$ -adrenocepter cell lines. Variants cell lines (A) GEVT, (B) GQVT, (C) RQVT and (D) GQVI. Data were normalized to response to 1 μmol·L⁻¹ isoprenaline (100%) in each experiment and are presented as mean percentage isoprenaline (%ISO) response; error bars represent s.e.mean (n = 3).

potency of indacaterol on the GQVT background. However, this was not apparent in the more physiologically relevant low-expression cell lines. Neither the low- nor the high-level β_2 -adrenoceptor expression cell lines were matched for receptor expression levels, confounding comparisons of agonist responses on different β_2 -adrenoceptor haplotypes.

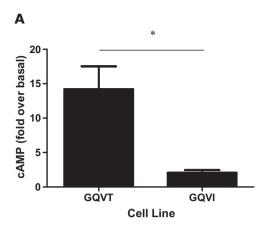
The β_2 -adrenoceptor I164 variant was associated with lower cAMP responses to β_2 -adrenoceptor agonists

The responses to all agonists in the low-expression GQVI β_2 -adrenoceptor cell line were markedly less than in the other cell lines, and too low for accurate EC₅₀ values to be determined for these agonists (Figure 4A). These reduced responses are unlikely to be due to β_2 -adrenoceptor expression differences in the GQVT and GQVI lines as these are well matched for surface receptor expression (204.9 and 302.4 fmol·mg⁻¹ protein, respectively, Table 2). In addition, the ability of forskolin to generate a cAMP response in these two cell lines was similar (Figure 4B). Interestingly, this effect was not apparent

in the high-expression GQVI cell line that had a robust cAMP response to β_2 -adrenoceptor agonist stimulation (16.79 \pm 5.93-fold over basal, 1 μ mol·L⁻¹ isoprenaline, 30 min, n=3, data not shown) suggesting that the genotypic effect is also dependent on the expression level of the β_2 -adrenoceptor. The ability of forskolin to generate cAMP was not significantly different between the four low- and four high-expression cell lines (n=3, data not shown).

Dose–response relationships of β_2 -adrenoceptor agonists in human primary ASM cells

In addition to the recombinant cell lines, we also studied a series of cell cultures of human ASM cells. Because the majority of cell lines studied were heterozygous for the polymorphisms of interest, the data presented were pooled from all the cell cultures established from different individuals (Figure 5). There were no significant differences between the EC_{50} values for the three clinically relevant β_2 -adrenoceptor agonists (indacaterol, salmeterol and formoterol) in the



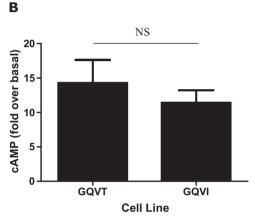


Figure 4 The low-expression lle164 β₂-adrenocepter cell line has a reduced capacity to respond to β₂-adrenocepter agonist stimulation. (A) cAMP (adenosine-3′,5′-cyclic-monophosphate) responses to 1 μmol·L⁻¹ isoprenaline (30 min) in the low-expression GQVT and GQVI cell cells lines. (B) cAMP responses to 10 μmol·L⁻¹ forskolin (30 min) in the low-expression GQVT and GQVI cell lines. Data are expressed as mean fold increase in cAMP compared with basal level; error bars represent s.e.mean (n = 3). *P = 0.01 (t-test). NS, not significant.

human primary ASM analyses (Table 3). There were significant differences between the three agonists with respect to maximal responses (Figure 5, P < 0.0001). As previously demonstrated in the CHO-K1 cell model, salmeterol was identified as a low-efficacy agonist and formoterol as a high-efficacy agonist. Interestingly, in the primary ASM cells indacaterol appears to have a reduced efficacy compared with the CHO-K1 data with a maximal response value intermediate between those to salmeterol and formoterol (Table 4; Figure 5). These data are in contrast to the data generated in the recombinant system, which suggested that indacaterol was a high-efficacy agonist when compared with isoprenaline. This finding was consistent irrespective of the genotype of the human ASM cells examined [e.g. indacaterol (%isoprenaline), GEVT 56 ± 2 (n = 4), RQVT 52 ± 2 (n = 7) data not shown].

Discussion and conclusions

The data presented here make two important points. Firstly, this is the first comprehensive study describing the pharma-

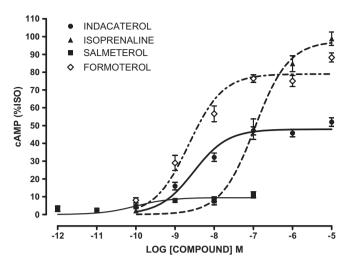


Figure 5 Concentration–response curves for cAMP (adenosine-3′,5′-cyclic-monophosphate) production in response to $β_2$ -adrenoceptor agonists (10 min) in human primary airway smooth muscle cells. Data were normalized to response to $1 μmol·L^{-1}$ isoprenaline (100%) in each experiment and are presented as mean percentage isoprenaline (%ISO); error bars represent s.e.mean (n=12-15).

cology of indacaterol at the β_2 -adrenoceptor in both recombinant cell lines expressing the human β_2 -adrenoceptor and primary cultures derived from human airway myocytes. Secondly, we have been able to study the effects of a range of β_2 -adrenoceptor agonists on cAMP responses in recombinant cell lines expressing the common β_2 -adrenoceptor haplotypes present in the Caucasian population.

Indacaterol acts in both low- and high-expressing CHO-K1 cell lines as a high-efficacy agonist. In this respect it is similar to formoterol, although indacaterol is a slightly less potent agonist in cells expressing the most common Caucasian people β_2 -adrenoceptor haplotype (GEVT). The onset of action in all the cell systems studied is rapid, again being similar to formoterol but producing a significantly greater response than salmeterol in the first 40 min, in keeping with previous studies suggesting that salmeterol is a lower-efficacy agonist. These data are in excellent agreement with previous observations using a recombinant expression system and the guinea-pig isolated trachea (Battram *et al.*, 2006).

The data obtained from the pooled analysis in human primary ASM cells suggest that as predicted, formoterol appears to be a high-efficacy agonist and salmeterol a lowefficacy agonist. Interestingly, in contrast to the recombinant cell line data indacaterol appeared to have a lower efficacy in the ASM with a maximal response intermediate to that observed for salmeterol and formoterol. These findings in part agree with a recent study examining β_2 -adrenoceptor relaxation responses in human isolated bronchi (Naline et al., 2007). The order of maximal response was: formoterol [94 \pm 1 (%)] > indacaterol $(77 \pm 5) \ge$ salmeterol (74 ± 4) at resting tone (Naline et al., 2007). However, it has been noted that the characteristics of these agonists to induce relaxation can be influenced by the tone of the tissue preparation (e.g. salmeterol has low or high intrinsic activity in this system depending on bronchial tone) (Naline et al., 2007). It is important to note that relaxation of ASM (trachealis strips) is not proportional to cAMP levels suggesting cAMP-independent mechanisms are also involved (Kume *et al.*, 1994). cAMP-independent mechanisms may at least in part explain efficacy differences observed in our cell-based/cAMP assay compared with tissue-based approaches using bronchial relaxation (Naline *et al.*, 2007).

In the different cell systems the molecular mechanisms underlying these differences in maximal response remain to be resolved; however, a feature of the CHO-K1 system is the relatively high receptor densities in comparison with that found endogenously, for example, primary bronchial smooth muscle cells in vitro have a β_2 -adrenoceptor density of approximately 70 fmol·mg⁻¹ (Green et al., 1995). In addition, we cannot exclude the possibility that the CHO-K1 signalling machinery influences our key endpoint, that is, cAMP production, and does not adequately represent that observed in human primary cells expressing endogenous β_2 -adrenoceptors. Overall, these data highlight the need to include primary cell and tissue-based analyses in pharmacological characterization of new compounds, but also highlight the utility of recombinant approaches to unmask interesting structure-function mechanisms.

In the next series of experiments we used both high- and low-expressing cell lines with the common haplotypes of the β_2 -adrenoceptor to address the possibility that agonist responses may vary by haplotype. In the Caucasian population the respective minor allele frequencies we have observed in a large population study (Hall et al., 2006) are as follows: R16 36.3% and E27 44.6%. Because of linkage disequilibrium, not all haplotypes occur at the frequency predicted by the allele frequencies per se: in particular the R16E27 combination was not seen in the 1958 birth cohort study (Hall et al., 2006), and the G16Q27 combination is also rare. I164 usually occurs on the G16Q27 background, and hence to study the effects of β₂-adrenoceptor agonists on the most frequent haplotypes seen in the general population we made cell lines, both with high and low expression of the β_2 -adrenoceptor, with all these combinations. In all 18 cell lines were established: these were then screened to try and match expression levels as closely as possible to allow comparisons to be made. In general, cAMP responses to agonists were not detected in cell lines with expression levels <100 fmol⋅mg⁻¹ protein.

In variant cell lines, the overall potency profiles of the agonists studied were similar; however, there was a trend in both the high- and low-expression cells for the receptor haplotype to influence the potency of indacaterol versus formoterol, that is, for GEVT/RQVT formoterol > indacaterol and for GQVT indacaterol > formoterol. The molecular basis of this observation based on the presence or absence of specific amino acid residues is not clear.

However, it is important to note that the genotype effects we observed were present/absent depending on the relative receptor expression, for example, the reduction in β_2 -adrenoceptor agonist responses for the I164 cell line was only observed in the lower-expression cell line. These data suggest that the functional effect of receptor polymorphism may only be unmasked under the appropriate expression level, which has broader implications as β_2 -adrenoceptor

expression is cell-specific, for example, expression on human peripheral blood mononuclear cells is <700–750 receptors per cell compared with 30 000–40 000 on ASM cells (Hayes *et al.*, 1996).

Overall, however, the data suggest that acute responses to these drugs are relatively unlikely to differ between individuals of varying genotype except in those who are homozygous for I164. Such individuals are extremely rare (Hall *et al.*, 2006) because of the low frequency of the I164 allele. We also studied a series of primary cultures derived from human ASM cells from individuals of known genotype: as would be predicted, the number of cultures obtained that were homozygous at all loci was small, limiting our ability to determine whether there are any small differences in response by genotype in this more physiologically relevant system.

Previous functional studies on ADRB2 polymorphisms have addressed the potential to alter down-regulation profiles. In transformed cell lines the G16 genotype appears to show greater down-regulation than the R16 form of the receptor, and the E27 form appears to protect against down-regulation, an effect that is overridden by genotype at codon 16 (Green et al., 1994; 1995). Ideally we would have liked to undertake similar studies in the CHO-K1 cell system to address downregulation profiles with the agonists used in the current study. However, our initial studies demonstrated that while it is feasible to wash out isoprenaline from cultures and observe full recovery of cAMP responses with time [as we have previously described (Hall et al., 1993)], we were unable to wash out some of the other agonists studied. In particular, β_2 -adrenoceptor-mediated cAMP responses could not be recovered following prolonged exposure to indacaterol or salmeterol, suggesting these drugs remain bound to the receptor or its immediate vicinity. The refractory nature of salmeterol's stable activation of β_2 -adrenoceptor to wash out has been described previously (Clark et al., 1996).

We also confirmed that the I164 form of the receptor couples less efficiently with adenylate cyclase in lowexpressing CHO-K1 cells, in keeping with previous work in fibroblasts with other agonists (Green et al., 2001). In the experiments comparing the response to isoprenaline in GQVT and GQVI cells, the reduced cAMP response to isoprenaline is unlikely to be related to differences in receptor expression as these two lines are well matched for expression. In addition, the abilities of both cell lines to generate cAMP responses to forskolin were similar, suggesting that reduced expression of adenylate cyclase in the GQVI cells is unlikely to provide an explanation. Our results show this effect extends to indacaterol as well as the other LABAs studied. These effects were not seen in the high-expressing GQVI cell system, suggesting that high levels of β_2 -adrenoceptor expression can overcome the effect of the I164 polymorphism.

Overall, these data demonstrate that indacaterol is a high-efficacy agonist in CHO-K1 cells transfected with the human β_2 -adrenoceptor. The data from the primary ASM analyses suggest it induces a response intermediate between those observed for salmeterol and formoterol. In keeping with data obtained with a range of other clinically relevant β_2 -adrenoceptor agonists there are no marked genotype-dependent effects on short-term agonist-induced receptor coupling for the common variants of the *ADRB2* gene.

However, for the rare I164 variant we confirmed the reduced activity associated with this variant and identified an additional level of regulation, based on receptor expression, that determines genotype effect.

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Conflict of interest

The authors state no conflict of interest.

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