The utility of *in vitro* data in making accurate predictions of human P-glycoprotein mediated drug-drug interactions:  

A case study for AZD5672

ROBERT ELSBY, MICHAEL GILLEN, CAROLINE BUTTERS, GEMMA IMISSON, PRADEEP SHARMA, VERONICA SMITH & DOMINIC D. SURREY

Clinical Pharmacology and DMPK, AstraZeneca R&D Charnwood, Loughborough, Leicestershire, United Kingdom (RE, CB, GI, PS, VS, DDS), Clinical Pharmacology and DMPK, AstraZeneca R&D Wilmington, Delaware, United States of America (MG).
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Running title: Assessment of P-gp DDI potential in vitro and in the clinic

Corresponding author:

Dr. Robert Elsby
Clinical Pharmacology & DMPK
AstraZeneca R&D Charnwood,
Bakewell Road,
Loughborough,
Leicestershire,
LE11 5RH
United Kingdom
Tel: +44 (0)1509 640000  Fax +44 (0)1509 645576
Email: Robert.Elsby@astrazeneca.com

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Non-standard abbreviations:

DDI, drug-drug interaction; P-gp, P-glycoprotein; C_{max}, observed maximum plasma drug concentration; AUC_{0-72h}, area under the plasma concentration versus time curve from 0 to 72 hours post-dose; FDA, US Food and Drug Administration; ITC,
International transporter consortium; IC$_{50}$, inhibitory concentration that produces 50% inhibition.
ABSTRACT

To support drug development and registration, Caco-2 cell monolayer assays have previously been set up and validated to determine if candidate drugs are substrates or inhibitors of human P-glycoprotein (P-gp). In this study, the drug-drug interaction (DDI) potential of AZD5672 was assessed accordingly and a subsequent clinical digoxin interaction study was performed. AZD5672 (1-500 μM) demonstrated concentration-dependent efflux across cell monolayers, which was abolished in the presence of ketoconazole and quinidine, identifying AZD5672 as a P-gp substrate. Additionally, P-gp mediated digoxin transport was inhibited in a concentration-dependent manner by AZD5672 (IC50=32 μM). Assessment of calculated theoretical gastrointestinal inhibitor concentration [I2], and predicted steady state maximum total plasma inhibitor concentration [I1] indicated the potential for a DDI at the intestinal but not the systemic level following the predicted therapeutic dose of AZD5672 (100mg). A clinical study was performed and the plasma pharmacokinetics (Cmax, AUC0-72h) of orally dosed digoxin (0.5mg) were found to be unaffected by co-administration of AZD5672 (50mg) at steady state. In contrast, a 150mg dose of AZD5672 significantly increased digoxin Cmax and AUC0-72h by 1.82-fold and 1.33-fold, respectively. Concentration-time profile comparisons indicated that digoxin elimination was unchanged by AZD5672 and the interaction was most likely to have resulted from inhibition of intestinal P-gp leading to increased digoxin absorption. The observed dose-dependent clinically significant interaction was accurately predicted utilising calculated I2 and in vitro P-gp inhibition data, confirming AZD5672 to be a P-gp inhibitor in vivo.
INTRODUCTION

P-glycoprotein (P-gp) is ubiquitously expressed in the gastrointestinal tract, liver, kidney and brain and can impact the oral bioavailability, biliary and renal clearance, and/or brain penetration of drugs that are substrates of this efflux transporter (Thiebaut et al., 1987; Schinkel, 1999). If a drug is subject to significant P-gp efflux, its absorption, distribution and elimination could be altered by potent P-gp inhibitors. Equally, drugs that are potent inhibitors of P-gp may modulate the pharmacokinetic and pharmacodynamic parameters of other P-gp substrates resulting in toxicity or altered efficacy. Such drug-drug interactions (DDIs) attributed to P-gp have been observed clinically following co-administration of digoxin and quinidine, digoxin and itraconazole, digoxin and valsopdar, and loperamide and quinidine (Hedman et al., 1990; Jalava et al., 1997; Alderman and Allcroft, 1998; Fromm et al., 1999; Kovarik et al., 1999; Sadaque et al., 2000; Mizuno et al., 2003).

To date the most described P-gp mediated DDIs involve the cardiac glycoside digoxin due to its narrow therapeutic window and ensuing toxicity if plasma levels are elevated above the normal therapeutic range (Tuncok et al., 1997). Since digoxin is a P-gp substrate that is devoid of cytochrome P450 metabolic clearance (Lacarelle et al., 1991), P-gp efflux plays a major role in both the absorption and renal elimination of the compound (Hedman et al., 1990; Cavet et al., 1996; Jalava et al., 1997; Westphal et al., 2000; Igel et al., 2007). Consequently, this makes digoxin susceptible to pharmacokinetic perturbations arising from inhibition of P-gp in the intestinal tract and kidney upon concomitant administration of P-gp inhibitors (Jalava et al., 1997; Westphal et al., 2000; Igel et al., 2007). It is for these reasons that digoxin is considered by both the US Food and Drug Administration (FDA) and the European Medicines Agency (EMA) in their respective draft DDI guidance, and by the
International Transporter Consortium (ITC) in their commentary paper, to be the recommended gold standard probe for studying P-gp inhibition both in the clinic and \textit{in vitro} in order to assess the P-gp mediated DDI potential of new molecular entities.

Such information is required to support drug development and registration and comprises initial \textit{in vitro} studies for the identification of the new molecular entity as a substrate or inhibitor of human P-gp, followed, if warranted, by a clinical interaction study (US FDA draft DDI guidance, 2006; Zhang et al., 2006, 2008; EMA draft DDI guideline, 2010; Giacomini et al., 2010). For P-gp inhibitors, evaluation of the ratios of steady state maximum total (bound plus unbound) plasma inhibitor concentration [$I_1$] and theoretical gastrointestinal inhibitor concentration [$I_2$] to \textit{in vitro} inhibitory potential ($IC_{50}$) are recommended to predict the potential for a systemic (ratio >0.1) or intestinal (ratio >10) P-gp mediated DDI in the clinic, respectively (FDA draft DDI guidance, 2006; Zhang et al., 2008). Calculation of $I_1$ and $I_2$ give “worst-case” concentrations that reflect a conservative approach adopted by the FDA for DDI prediction. Used together the two methods give an accurate prediction rate of approximately 78\% (Zhang et al., 2008). Such predictions facilitate a decision on the need to conduct a clinical interaction study in order to appropriately label a drug for safe and effective use.

AZD5672 is a CCR5 receptor antagonist that was being developed for the oral treatment of rheumatoid arthritis (Figure 1; Gerlag et al., 2010). Patients of this disease population are at risk of DDIs because of the multiple co-medications that might be taken as a consequence of the co-morbid conditions that could be present due to their advanced age (van Roon et al., 2009). Due to a regulatory requirement, and the fact that digoxin might be a likely co-medicant, the first aim of this study was to investigate AZD5672 as a substrate and inhibitor of human P-gp using validated \textit{in vitro} Caco-2
assays (Elsby et al., 2008). Based upon inhibition predictions, the final aim of this study was to conduct a clinical oral pharmacokinetic interaction study in healthy volunteers to establish whether AZD5672 could perpetrate a P-gp mediated DDI with digoxin in the clinic. Two doses (50 and 150mg) of AZD5672 that were close to the predicted therapeutic dose were studied to investigate the dose-dependency of any interaction.
METHODS

Materials

Digoxin, ketoconazole, probenecid, propranolol hydrochloride, quinidine hydrochloride monohydrate and verapamil hydrochloride were purchased from Sigma-Aldrich Inc. \(^{[3}H\)Digoxin (specific activity 870 GBq/mmol, radiochemical purity > 97 %) and UltimaGold scintillation cocktail were purchased from PerkinElmer Life and Analytical Sciences (Buckinghamshire, UK). \(^{[14}C\)D-Mannitol (specific activity 2.15 GBq/mmol, radiochemical purity 97.6 %) and \(^{[3}H\)propranolol hydrochloride (specific activity 740 GBq/mmol, radiochemical purity 99.7 %) were purchased from Amersham Biosciences (Amersham, UK). \(^{[3}H\)AZD5672 (specific activity 415 GBq/mmol, radiochemical purity 99.3 %), non-radiolabelled AZD5672 succinate, \(^{[13}C_6\)AZD5672, and AZD5672 succinate 50 mg tablets were synthesised at AstraZeneca R&D Charnwood (Loughborough, UK). Digoxin 0.25 mg Lanoxin\(^\circledR\) tablets were obtained from GlaxoSmithKline GmbH & Co KG (München, Germany).

Hanks Balanced Salt Solution (HBSS; containing CaCl\(_2\) and MgCl\(_2\)), Dulbecco’s Modified Eagles Medium (DMEM; containing pyruvate, 4500 mg/L glucose and 4 mM L-glutamine) and non-essential amino acids were purchased from Invitrogen (Paisley, UK). Monoflow 2 scintillant was obtained from National Diagnostic (UK) Ltd (Yorkshire, UK). All other chemicals and reagents were purchased from Sigma-Aldrich or Thermo-Fisher Scientific (Loughborough, UK). BD Falcon\(^\text{TM}\) 24-multiwell insert systems with polyethylene terephthalate membranes (1 \(\mu\)m pore size, 0.3 cm\(^2\) surface area) and BD Falcon\(^\text{TM}\) 24-multiwell companion plates were purchased from BD Biosciences (Oxford, UK). Caco-2 cells (HTB37) were purchased from American Type Culture Collection. Phenomenex Luna C18 3 \(\mu\)m column (50 \(\times\) 2.0 mm) was obtained
from Phenomenex Inc. (CA, USA). BDS Hypersil C18 (5 μm; 50 × 3.0 mm) column was obtained from Thermo-Fisher Scientific (MA, USA).

**Preparation of stock solutions and donor and receiver solutions for in vitro assays**

Stock solutions (in dimethyl sulfoxide) and donor and receiver solutions in transport buffer (HBSS containing 25 mM HEPES and 0.1 % (w/v) bovine serum albumin, pH 7.4) were prepared according to the validated methodology described previously (Elsby et al., 2008).

**Caco-2 cell culture**

Caco-2 cells were seeded in cell culture medium (consisting of DMEM supplemented with 20 % (w/v) foetal bovine serum, 1 % (v/v) non-essential amino acids, 98 U/ml penicillin and 98 μg/ml streptomycin) onto polyethylene terephthalate filter membranes (surface area = 0.3 cm²) of 24-well plates and used for transport studies 15-21 days post seeding between passage numbers 25 and 40 as described previously (Elsby et al., 2008).

**Chemical and metabolic stability of [3H]AZD5672 in the Caco-2 test system**

In order to assess metabolic and chemical stability, the radiochemical purity of [3H]AZD5672 (100 μM) in both donor and receiver wells in the bidirectional transport assay, and in the donor working solutions following a 2 h incubation at 37 °C, respectively, was determined and compared to that determined in the donor working solution at t=0 h. Samples (100 μl aliquots) were processed with methanol (200 μl) to
precipitate protein, centrifuged at 13,000 rpm for 5 min and the resulting supernatants analysed by radioflow HPLC as described previously (Elsby et al., 2008).

**Bidirectional transport assay (P-gp substrate assessment)**

The bidirectional apparent permeability of \[^3\text{H}\]AZD5672 (1, 100 and 500 \(\mu\)M) across polarised Caco-2 cell monolayers (n=3) was determined using the previously described validated procedure (Elsby et al., 2008). The P-gp probe substrate \[^3\text{H}\]digoxin (1, 10 and 100 \(\mu\)M) and the high permeability marker \[^3\text{H}\]propranolol (10 \(\mu\)M) were assessed alongside to positively and negatively control the assay, respectively. Briefly, radiolabelled test compounds were incubated in transport buffer (pH 7.4) at 37 °C on either the apical (total volume of 250 \(\mu\)l) or basolateral (total volume of 600 \(\mu\)l) side of the monolayer for 2 hours. Donor compartments also contained the cell monolayer integrity marker \[^{14}\text{C}\]mannitol (50 \(\mu\)M). Corresponding receiver compartments contained transport buffer containing 1 % (v/v) DMSO. The amount of drug appearing in the receiver compartment over time was determined by sampling (50 \(\mu\)l aliquot) at 30, 60, 90 and 120 min time points, quantified by liquid scintillation counting and used to calculate apparent permeability, which was then used to calculate efflux ratio. The sampled volume (50 \(\mu\)l) taken from the receiver compartment was replaced with an equal volume of warmed transport buffer (containing 1 % (v/v) DMSO) after each time point. If efflux of test compound was observed, determination of its bidirectional apparent permeability was repeated, alongside digoxin as positive control, at a single concentration in the presence of the P-gp inhibitors ketoconazole (100 \(\mu\)M) or quinidine (100 \(\mu\)M) to confirm the role of P-gp. Inhibitors were present in both donor and receiver compartments and added at the same time as test compound. Inhibitor was
also present in the transport buffer used to replace sampled volumes from the receiver compartments. Permeability of \( ^{14} \text{C} \)mannitol and transepithelial electrical resistance (TEER) values were measured in all experiments to assess cell monolayer integrity.

**Digoxin interaction (IC\textsubscript{50}) assay (in vitro P-gp inhibition assessment)**

The basolateral-to-apical apparent permeability of \( ^{3} \text{H} \)digoxin (5 μM) across Caco-2 cell monolayers (n=3) was determined over a range of AZD5672 concentrations (0, 0.1, 0.3, 1, 3, 10, 30 and 100 μM), using the previously described validated procedure (Elsby et al., 2008). The P-gp inhibitor verapamil (0.1-100 μM) and probenecid (equivalent to vehicle control, 10 μM) were used to positively and negatively control the assay, respectively. Inhibitors were present on both sides of the monolayer and were added at the same time as digoxin. The degree of inhibition of P-gp mediated digoxin transport was determined by calculation of an IC\textsubscript{50} where appropriate.

\( ^{14} \text{C} \)Mannitol permeability and TEER values were measured in all experiments to assess cell monolayer integrity.

**Data analysis**

For every well the determined concentration (nM; corrected for the volume dilution that occurred during sampling) of substrate appearing in the receiver compartment was plotted against sampling time (min). Curve fitting using standard linear regression parameters (correlation coefficient \( r^2 \geq 0.75 \)) was used to give the rate of drug transport (nM/min). For bidirectional permeability experiments drug transport rates were used to determine apparent permeability (P\textsubscript{app}) values (Elsby et al., 2008). Efflux ratios (ER) were calculated from P\textsubscript{app} basolateral-to-apical / P\textsubscript{app} apical-to-basolateral.
For inhibition studies, calculated digoxin basolateral-to-apical $P_{\text{app}}$ values, in the absence and presence of a range of inhibitor concentrations, were plotted directly against nominal inhibitor concentration and curve fitting was performed using XLfit 3.0 (4 parameter logistic model, equation 201) to determine the concentration that produces half-maximal inhibition of P-gp-mediated digoxin transport (IC$_{50}$). The following equation was used by model 201:

$$Y = A + \left( \frac{B - A}{1 + \left( \frac{x}{C} \right)^0} \right)$$

Where, $A=$minimum y value, $B=$maximum y value, $C=$IC$_{50}$ value, $D=$slope factor.

When fitting, the bottom and top of the curve are locked at values that correspond to the calculated mean digoxin $P_{\text{app}}$ in the presence of 100 μM verapamil (representing complete P-gp inhibition) and the calculated mean digoxin $P_{\text{app}}$ in the presence of probenecid (representing vehicle control in the absence of inhibitor), respectively. Our basolateral-to-apical inhibition and curve-fitting methodology is directly comparable to the recommended “calculation method no.3 (inhibition of corrected basolateral-to-apical permeability of digoxin)” of Balimane et al. (2008) which ensures determination of an accurate IC$_{50}$ value, a prerequisite for accurate prediction of clinical outcome.

Rather than converting $P_{\text{app}}$ to percentage inhibition by correcting it for the (passive) permeability of digoxin observed when P-gp is completely inhibited (Balimane et al., 2008), we are plotting uncorrected $P_{\text{app}}$ directly and are using the (passive) $P_{\text{app}}$ observed when there is complete P-gp inhibition to define where the bottom of the curve should be when fitting.
Mass balance (% recovery) was calculated for all experiments by expressing the sum of the amount of substrate (pmol) found in both the donor and receiver compartments at the end of the incubation (2 h) as a percentage of the initial donor compartment amount (pmol) at t=0 h.

Inhibitor concentration \([I_1]\) represents either predicted (from pharmacokinetic modelling) maximum plasma drug concentration at steady state \(C_{\text{max,ss}}\) or observed \(C_{\text{max,ss}}\). The theoretical gastrointestinal concentration \([I_2]\) was calculated by dividing the oral dose (converted from milligrams to moles) by a volume of 250 ml. Parameters \([I_1]\) or \([I_2]\) were divided by \(IC_{50}\) to give a ratio.

**Clinical interaction study participants**

The clinical study (D1710C00021; Clinical Trial Registration number NCT00723424) was approved by the Independent Ethics Committee (IEC), LaGeSo (Landesamt für Gesundheit und Soziales, Berlin, Germany), and complied with the ethical principles of the International Conference on Harmonization/Good Clinical Practice, in accordance with the Declaration of Helsinki. All subjects provided written informed consent.

Inclusion criteria included the following: male or females, age 18–50 years; body mass index of 19–29 kg m\(^{-2}\); clinically normal physical and laboratory findings, including negative drug test and negative tests of Hepatitis B surface antigen, antibodies to Hepatitis C virus and antibodies to HIV. Exclusion criteria included the following: known allergy to digoxin; QTc interval >450 ms; clinical evidence of Wolff-Parkinson-White syndrome, intermittent complete heart block, second degree heart block or prolonged PR interval of greater than 210 ms on 12-lead ECG; resting heart rate <45 bpm; potassium outside normal range; hypercalcaemia or hypermagnesaemia; abnormal liver function tests; the use of prescription medications within 2 weeks before
study entry; the use of any non-prescription medications (including herbal medications, excluding occasional paracetamol use) within 7 days of first dose of the study medication, unless agreed by the investigator; a history or presence of clinically significant gastrointestinal, hepatic, or renal disease or other condition known to interfere with the absorption, distribution, metabolism, or excretion of the study drugs; and intake of alcohol, grapefruit or caffeine during the study.

Clinical interaction study design

This was a two-period, open-label single-centre study with two dosing groups to determine the pharmacokinetics of a single oral dose of digoxin when administered alone or in combination with AZD5672 at steady-state. The study consisted of a screening period of up to 21 days before the first administration of digoxin. Thereafter, two study periods of 6 and 17 days, respectively, followed. The two study periods were separated by a wash-out phase of at least 7 days (from the digoxin dose of Period I to first dose of AZD5672 in Period II). In Group A, healthy volunteers received a single dose of 0.5mg digoxin on Day 1 of Period I. In Period II, 50mg AZD5672 was given once a day for 13 days. On Day 12 of 13, a further single dose of 0.5mg digoxin was concomitantly administered with AZD5672. Group B was conducted with the same regimen, except that a dose level of 150mg AZD5672 was evaluated. Finally, there was a follow-up visit 4 to 7 days after the last pharmacokinetic (PK) blood sampling in Period II.

Pharmacokinetic assessments

Digoxin plasma samples (2.7 ml blood) were obtained in potassium EDTA tubes at: pre-dose, 10 min, 30 min, 45 min, 1 h, 1.5 h, 2 h, 3 h, 4 h, 6 h, 9 h, 12 h, 24 h, 36 h,
48 h, 72 h, 96 h following each dose of digoxin. Plasma samples for the determination of digoxin were stored at –70 °C and transported to Covance Bioanalytical Services (Indianapolis, IN, USA) where concentrations of digoxin were determined using a commercially available validated and quality-controlled assay using liquid-liquid extraction; after evaporation under nitrogen, the residue was reconstituted, injected onto a BDS Hypersil C18 (5 μm; 50 × 3.0 mm) column, eluted with a gradient mobile phase containing acetonitrile and 5 mM ammonium formate in 0.1 % formic acid, and analyzed using LC-MS/MS. The standard curve range was from 0.0500 to 10.0 ng/ml using a human plasma sample volume of 0.250 ml.

AZD5672 plasma samples (2.7 ml blood) were obtained in potassium EDTA tubes at: pre-dose, 10 min, 20 min, 30 min, 45 min, 1 h, 1.5 h, 2 h, 2.5 h, 3 h, 3.5 h, 4 h, 6 h, 9 h, 12 h, 24 h on Days 10 and 12 of Period II. Plasma samples for determination of AZD5672 were stored at –20°C and transported to Covance Bioanalytical Services. AZD5672 was quantified using a validated, quality controlled bioanalytical method. Briefly, AZD5672 was extracted from human plasma (0.100 ml) by protein precipitation (following addition of acetonitrile containing [13C6]AZD5672 internal standard) and centrifugation. The supernatant was diluted, injected onto a Phenomenex Luna C18 (3 μm; 50 × 2.0 mm) column and eluted using an isocratic mobile phase containing 0.1 % (v/v) aqueous formic acid (77 %) and 0.1 % (v/v) formic acid in acetonitrile (23 %). AZD5672 and internal standard were detected using LC-MS/MS by multiple reaction monitoring of the transitions 640.5 to 399.2 and 646.4 to 399.2, respectively, and quantified with a standard curve of range 0.500 to 500 nM.

**Pharmacokinetic analysis**
The pharmacokinetic variables were determined by non-compartmental analysis using WinNonLin version 5.01 (Pharsight Corporation, Cary, NC, USA). The pharmacokinetic parameters calculated for digoxin were the observed maximum plasma drug concentration (C<sub>max</sub>), area under the plasma concentration versus time curve from zero to 72 h post-dose (AUC<sub>0-72h</sub>) and the time to reach maximum concentration following drug administration (t<sub>max</sub>). The pharmacokinetic parameter determined for AZD5672 for the purpose of this analysis was the observed maximum plasma drug concentration at steady-state (C<sub>max,ss</sub>) on Day 10.

Statistical Analysis

To address the primary objective, a point estimate and 90% confidence intervals for the ratio of digoxin AUC<sub>0-72h</sub> and C<sub>max</sub> (with AZD5672 : without AZD5672) were presented for each dosing group of AZD5672. This was obtained by analysing the log-transformed AUC<sub>0-72</sub> and C<sub>max</sub> for digoxin. No clinically significant drug-drug interaction was concluded if the 90% confidence interval fell within (0.67, 1.5). A similar non-parametric analysis (the Hodges Lehmann 90% confidence interval for the pair-wise treatment difference according to Tukey using the formula for small samples, as outlined in Hollander, Wolfe, Nonparametric statistical methods, 2nd edition, Wiley, 1999) was applied to t<sub>max</sub>.

Safety Assessment

All subjects underwent pre- and post-study medical assessments consisting of a physical examination, routine laboratory tests, vital signs and twelve-lead ECG. Safety parameters monitored throughout the study periods included routine laboratory tests, vital signs, twelve-lead ECG and adverse events.
RESULTS

Assessment of AZD5672 as a substrate and inhibitor of P-gp in vitro

Assessment of cell monolayer integrity and recoveries. All cell monolayers passed acceptance criteria (as described in Elsby et al., 2008) with regard to their TEER (≥ 600 ohms, or 180 ohms × cm²) and determined [14C]mannitol apparent permeability values (≤ 0.953 × 10⁻⁶ cm/s), indicating that the monolayers were acceptable for use.

TEER values ranged from 820-1096 ohms (246-329 ohms × cm²). Apparent permeability values for mannitol ranged from 0.316-0.694 × 10⁻⁶ cm/s. In all experiments conducted, mass balance (% recovery) was acceptable (>70 %). Recoveries ranged from 88-113 %, 89-101 %, 75-91 % and 90-115 % for [³H]digoxin, [³H]AZD5672, [³H]propranolol and [¹⁴C]mannitol, respectively. Occasionally, calculated mass balance values from an individual well exceeded 100 % as the disintegrations per minute value (used to derive pmol) for the volume sampled from the donor compartment at t=2 h was slightly higher than the value achieved for the volume sampled at t=0 h. Whilst these instances can be attributed to the normal minor variability associated with pipetting, which is noticeable because the donor compartment contains such high levels of radioactivity (typically tens to hundreds of thousands of disintegrations per minute), they confirm the absence of non-specific binding for the compound(s) in question.

Chemical and metabolic stability of [³H]AZD5672 in the Caco-2 test system. The determined radiochemical purities of [³H]AZD5672 in donor compartments at the end of the 2 h incubation ranged from 94.2-99.8 %, and for the donor working solution (t=2 h) was 97.2 %, which were qualitatively similar to that determined for the donor
working solution at t=0 h (98.3 %). Additionally, the radiochromatographic traces obtained at the end of a 2-h incubation were qualitatively similar to those obtained at the start of the incubation, i.e. there were no additional peaks (data not shown), demonstrating stability of this compound in the in vitro test system.

Bidirectional transport assay (P-gp substrate assessment). No efflux was observed for the negative control [3H]propranolol (10 μM; Figure 2). The positive control substrate [3H]digoxin (1, 10 and 100 μM) exhibited vectorial efflux (efflux ratios (ER) ranged from 7.1 to 9.2; Figure 2) which was abolished (ER reduced to <2) in the presence of the P-gp inhibitors ketoconazole and quinidine (Figure 3).

The mean bidirectional apparent permeabilities of [3H]AZD5672 determined at 1, 100 and 500 μM are shown in Figure 2; ER ranged from 11 (at 1 μM) to 1.4 (at 500 μM). The efflux of a single concentration of [3H]AZD5672 (1 μM) was abolished (ER reduced from 11 to 0.8 or 1.2, respectively) in the presence of the P-gp inhibitors ketoconazole (100 μM) and quinidine (100 μM, Figure 3).

Digoxin interaction (IC50) assay (in vitro P-gp inhibition assessment). The negative control inhibitor probenecid (10 μM) did not affect digoxin basolateral-to-apical directional apparent permeability (9.58 × 10⁻⁶ cm/s). The positive control inhibitor verapamil inhibited P-gp mediated digoxin transport with a mean (± S.D., n=3) apparent IC₅₀ of 3.1 ± 0.64 μM (Figure 4). AZD5672 inhibited P-gp mediated digoxin transport in vitro with a mean (± S.D., n=3) apparent IC₅₀ of 32 ± 4.2 μM (Figure 4).
Calculation of \([I]/IC_{50}\) ratios for the predicted therapeutic dose of AZD5672, and for the 50mg and 150mg doses of AZD5672 used in the clinical study

The calculated \([I_1]/IC_{50}\) and \([I_2]/IC_{50}\) ratios for the predicted (from pharmacokinetic modelling) therapeutic dose of AZD5672 (100mg) and the actual doses of AZD5672 used in the clinical interaction study are shown in Table 1. The \([I_1]\) value given for the 50 and 150mg dose of AZD5672 is the actual determined total \(C_{max}\) of AZD5672 (at steady state) derived from the plasma concentration data determined in the clinical interaction study.

Effect of AZD5672 on the pharmacokinetic parameters of digoxin in healthy volunteers

The major pharmacokinetic parameters of a single oral dose of digoxin (0.5mg) when administered alone or concomitantly with AZD5672 50mg or AZD5672 150mg at steady state (dosed once daily for 12 days) are summarised and compared in Table 2. The mean plasma concentration-time profiles (0-24 h) are shown in Figure 5. The 50mg dose of AZD5672 at steady state did not appear to alter either the geometric least-squares mean \(AUC_{0-72h}\) nor \(C_{max}\) of digoxin. In contrast, the 150mg dose of AZD5672 at steady state significantly increased the geometric least-squares mean \(AUC_{0-72h}\) and \(C_{max}\) of digoxin by 1.33-fold (90% confidence interval, 1.12-1.57) and 1.82-fold (90% confidence interval, 1.38-2.39), respectively, indicating an interaction. There were no significant differences between digoxin \(t_{max}\) after administration of digoxin alone or in combination with either dose of AZD5672 (data not shown).
DISCUSSION

Inhibition of P-gp can affect the oral bioavailability, biliary and renal clearance, and/or distribution of drugs that are substrates of this transporter, resulting in a drug-drug interaction. To date, the vast majority of DDIs attributed to P-gp have been observed clinically with digoxin since the transporter plays a major role in the drug’s disposition (Hedman et al., 1990; Jalava et al., 1997; Kovarik et al., 1999; Westphal et al., 2000). Resultant safety concerns related to the narrow therapeutic window of digoxin are the reason, in part, that assessment of the potential for P-gp mediated drug-drug interactions in the clinic is required to support drug development and registration (US FDA draft DDI guidance 2006; EMA draft DDI guidelines, 2010). Accordingly, because of the regulatory requirement and the fact that digoxin is likely to be a co-medicant in the target disease population of rheumatoid arthritis, the human P-gp mediated DDI potential of the candidate drug AZD5672 was evaluated.

The validated Caco-2 cell monolayer test system and associated bidirectional transport and digoxin interaction assays were used in this study in strict accordance with the previously defined acceptance criteria for quality control of the assays (Elsby et al., 2008), to evaluate AZD5672 as a substrate or inhibitor of human P-gp in vitro. The demonstration that AZD5672 was both chemically and metabolically stable in the in vitro test system confirmed that the radioactivity quantified by liquid scintillation counting originated from radiolabelled parent compound and not from degradative products/metabolites, and gave confidence in the calculation of apparent permeability. Vectorial efflux of the positive control substrate digoxin, assessed alongside AZD5672 in the bidirectional transport assay, and abolition of this efflux by the potent P-gp inhibitors ketoconazole and quinidine (Elsby et al., 2008) confirmed that the in vitro test system had sufficient P-gp expression for transport experiments and was able to
identify substrates of P-gp. The contribution of an active transporter protein to the apparent permeability of AZD5672 was demonstrated by concentration-dependent vectorial efflux at 1 and 100 μM, which was saturated at 500 μM. This drug transporter was confirmed to be P-gp, as the efflux of AZD5672 was abolished in the presence of ketoconazole and quinidine. A follow-up clinical interaction study between AZD5672 (as a substrate/victim) and a clinical P-gp inhibitor was deemed unnecessary for a number of reasons. Firstly, from an absorption perspective, inhibition of P-gp is unlikely to impact on AZD5672 at the predicted therapeutic dose (100mg) as the primary mediator of absorption at this theoretical intestinal concentration (I2=528 μM, Table 1) would be passive permeability as P-gp is likely to be saturated. Secondly, from a kidney perspective, since a human radiolabelled excretion-balance study indicated that renal elimination of unchanged AZD5672 accounted for only 5% of the administered dose, overall clearance of AZD5672 is unlikely to be significantly affected by inhibition of renal P-gp (unpublished data). Finally, from a CNS-penetration perspective, the clinical impact of P-gp inhibition at the blood-brain barrier remains unclear (Eyal et al., 2009; Giacomini et al., 2010).

In the digoxin inhibition assay, the acceptable basolateral-to-apical apparent permeability of digoxin (~9.5 × 10^{-6} cm/s) in the presence of the negative control inhibitor probenecid, coupled with the determined IC_{50} (3.1 μM) of the positive control inhibitor verapamil, gave confidence in the test system used to generate data for test compound. AZD5672 inhibited P-gp mediated digoxin transport in a concentration-dependent manner with an IC_{50} of 32 μM; identifying AZD5672 as an inhibitor of human P-gp in vitro. Assessment ([I_{1}]/IC_{50} > 0.1 and [I_{2}]/IC_{50} > 10) of the in vitro P-gp inhibitory data generated indicated that for the predicted therapeutic dose (100mg),
AZD5672 had the potential to cause an intestinal ([I₂]/IC₅₀ = 17) rather than a systemic ([I₁]/IC₅₀ = 0.006) P-gp mediated DDI in vivo (Table 1).

Consequently, a clinical interaction study between concomitantly administered AZD5672 (orally dosed to steady state) and digoxin (single oral dose of 0.5mg) was conducted in healthy volunteers, with digoxin plasma pharmacokinetics being determined prior to and following co-administration of AZD5672. Results from the clinical study indicated that the dose selection of AZD5672 mitigated the risk of a DDI with digoxin, as the 50mg AZD5672 dose had no effect on digoxin pharmacokinetics, whereas the 150mg dose elicited a clinically significant DDI resulting in a 1.82-fold and 1.33-fold increase in digoxin C_max and AUC₀-₇₂h, respectively. The observation that C_max was more sensitive to change than AUC for digoxin in this study is in agreement with previous observations evaluating a range of perpetrators (Fenner et al., 2009).

Furthermore, examination of the digoxin concentration-time profiles obtained in the presence and absence of AZD5672 (150mg) indicated that the DDI was attributable to increased digoxin absorption, since the rate of digoxin elimination appeared unchanged as the elimination phases of the profiles were parallel (Figure 5). Whilst P-gp has been demonstrated to be the primary transporter mediating the absorption and disposition of digoxin (Cavet et al., 1996; Jalava et al., 1997; Igel et al., 2007), it has been reported previously that digoxin is also a substrate of the hepatic uptake transporter OATP1B3 (Kullak-Ublick et al., 2001). Nevertheless, since AZD5672 is not a substrate of OATP1B3 (unpublished data), it is unlikely that the observed increase in digoxin absorption (C_max) due to AZD5672 is a consequence of competitive inhibition of OATP1B3. Therefore, the increased absorption of digoxin in this study may be attributed to inhibition of intestinal P-gp by AZD5672. This is further supported by the fact that the observed increase in digoxin AUC in the present study (33%) is close to
the calculated theoretical maximum increase (42%) in digoxin (Lanoxin®) AUC that would be expected if absorption of Lanoxin® increased from the usual 70% to 100% due to intestinal P-gp inhibition (Fenner et al., 2009). Moreover, when the clinically observed AZD5672 steady state $C_{\text{max}}$ and calculated theoretical gut concentration for the 150mg dose of AZD5672 were used to back-extrapolate assessment ([I]/IC$_{50}$) of the $in$ $vitro$ P-gp inhibitory data, the resultant ratios predicted an intestinal-based DDI ([I$_2$]/IC$_{50}$ = 25) rather than an elimination-based (systemic) DDI ([I$_1$]/IC$_{50}$ = 0.009) with 150mg AZD5672.

Interestingly, the back-extrapolated ratios for the 50mg dose of AZD5672 also correctly predicted observed clinical outcome: [I$_1$]/IC$_{50}$ (0.002) and [I$_2$]/IC$_{50}$ (8.3) ratios predicted neither a systemic nor intestinal interaction respectively. These data for both doses of AZD5672 supports the approach proposed by the US FDA for assessing P-gp DDI potential. It was noted that use of the revised [I$_2$]/IC$_{50}$ > 5 ratio suggested by Cook et al. (2010) would have resulted in a false positive DDI prediction for the 50mg dose of AZD5672 if applied to the current study; however, those authors acknowledged that refinement of this ratio to reduce false negatives would inevitably increase the number of false positives.

Since 150mg AZD5672 has been found to cause a clinically significant intestinal P-gp mediated DDI with digoxin, a dose adjustment may be required if this dose of AZD5672 and digoxin (or other drugs whose absorption is primarily restricted by P-gp) are to be co-administered. Clearly no such dose adjustment will be necessary if the therapeutic dose of AZD5672 is determined to be 50mg.

In summary, $in$ $vitro$ studies have identified AZD5672 as both a substrate and an inhibitor of human P-glycoprotein and the data used to support the candidate drug’s
development. In a subsequent clinical study, AZD5672 (150mg) demonstrated a clinically significant DDI with digoxin thereby confirming it to be a P-gp inhibitor in vivo. Finally, the observed clinical interaction was accurately predicted using calculated I₂ and in vitro P-gp inhibition data.
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AUTHORSHIP CONTRIBUTION

Participated in research design: Elsby, Gillen, Imisson, Smith and Surry.

Conducted experiments: Elsby and Smith.

Contributed new reagents or analytic tools: Not applicable.

Performed data analysis: Elsby, Gillen and Imisson.

Wrote or contributed to the writing of the manuscript: Elsby, Gillen, Butters, Imisson, Sharma, Smith and Surry.
REFERENCES


drugs and non-antirheumatic drugs according to rheumatologists and pharmacists.

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LEGENDS FOR FIGURES

Figure 1. Chemical structure of AZD5672.

Figure 2. Determined bidirectional apparent permeabilities (P_{app}) of the positive and negative control substrates [\textsuperscript{3}H]digoxin and [\textsuperscript{3}H]propranolol, and the test compound [\textsuperscript{3}H]AZD5672 across Caco-2 cell monolayers. Data are expressed as the mean ± standard deviation (n=3).

Figure 3. Determined bidirectional apparent permeabilities (P_{app}) of a single concentration of the positive control substrate [\textsuperscript{3}H]digoxin and of [\textsuperscript{3}H]AZD5672 across Caco-2 cell monolayers in the presence and absence of the P-gp inhibitors ketoconazole (100 μM) and quinidine (100 μM). Data are expressed as the mean ± standard deviation (n=3).

Figure 4. IC\textsubscript{50} curves for the inhibition of P-gp mediated [\textsuperscript{3}H]digoxin transport across Caco-2 cell monolayers by the positive control inhibitor verapamil and AZD5672. Data are expressed as the mean ± standard deviation (n=3).

Figure 5. Semi-logarithmic plot of the geometric mean (± standard deviation) digoxin plasma concentration versus time following administration of a single oral dose of digoxin (0.5mg) in the presence and absence of AZD5672 50mg (A) and AZD5672 150mg (B).
**TABLES**

**Table 1. Summary of AZD5672 dose-related concentration and inhibitory kinetic parameters for calculation of [I]/IC50 ratios.**

<table>
<thead>
<tr>
<th>Dose of AZD5672</th>
<th>I1 (μM)</th>
<th>I2 (μM)</th>
<th>IC50 (μM)</th>
<th>[I1]/IC50</th>
<th>[I2]/IC50</th>
<th>Potential for DDI Systemic (&gt;0.1)</th>
<th>Potential for DDI Intestinal (&gt;10)</th>
</tr>
</thead>
<tbody>
<tr>
<td>100mg</td>
<td>0.200a</td>
<td>528</td>
<td>32</td>
<td>0.006</td>
<td>17</td>
<td>No</td>
<td>Yes</td>
</tr>
<tr>
<td>50mg</td>
<td>0.0491c</td>
<td>264</td>
<td>32</td>
<td>0.002</td>
<td>8.3</td>
<td>No</td>
<td>No</td>
</tr>
<tr>
<td>150mg</td>
<td>0.299c</td>
<td>792</td>
<td>32</td>
<td>0.009</td>
<td>25</td>
<td>No</td>
<td>Yes</td>
</tr>
</tbody>
</table>

I1 Mean steady state maximum plasma concentration (Cmax) value for total drug

I2 Concentration in gastrointestinal tract (determined from dose (mol)/250 ml)

IC50 Value determined from P-gp inhibition experiment in Caco-2 cells

a Predicted therapeutic dose from pharmacokinetic modelling

b Predicted value from pharmacokinetic modelling
c Value determined from the clinical interaction study (D1710C00021)
Table 2. Effect of co-administration of AZD5672 at steady state on the geometric least-squares mean (n=12) pharmacokinetic parameters of digoxin.

<table>
<thead>
<tr>
<th>Dose group</th>
<th>Pharmacokinetic parameter</th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>C&lt;sub&gt;max&lt;/sub&gt; (ng/ml)</td>
<td>AUC&lt;sub&gt;0-72h&lt;/sub&gt; (ng·h/ml)</td>
<td></td>
</tr>
<tr>
<td>AZD5672 50mg</td>
<td>Digoxin</td>
<td>2.87 (%CV 29.5)</td>
<td>19.73 (%CV 23.9)</td>
</tr>
<tr>
<td></td>
<td>Digoxin+AZD5672</td>
<td>2.88 (%CV 43.7)</td>
<td>19.55 (%CV 18.9)</td>
</tr>
<tr>
<td>Ratio&lt;sup&gt;a&lt;/sup&gt;</td>
<td></td>
<td>1.00 [0.83, 1.21]&lt;sup&gt;b&lt;/sup&gt;</td>
<td>0.99 [0.92, 1.07]&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
<tr>
<td>AZD5672 150mg</td>
<td>Digoxin</td>
<td>2.58 (%CV 67.0)</td>
<td>19.85 (%CV 40.8)</td>
</tr>
<tr>
<td></td>
<td>Digoxin+AZD5672</td>
<td>4.70 (%CV 37.0)</td>
<td>26.35 (%CV 20.3)</td>
</tr>
<tr>
<td>Ratio&lt;sup&gt;a&lt;/sup&gt;</td>
<td></td>
<td>1.82 [1.38, 2.39]&lt;sup&gt;b&lt;/sup&gt;</td>
<td>1.33 [1.12, 1.57]&lt;sup&gt;b&lt;/sup&gt;</td>
</tr>
</tbody>
</table>

%CV  Coefficient of variation

<sup>a</sup> Calculated from digoxin+AZD5672/digoxin alone

<sup>b</sup> 90% confidence interval
Figure 2

![Graph showing efflux ratios for different concentrations of drugs.](image)

- **Positive/negative controls**
- **A-to-B**
- **B-to-A**

**ER** = efflux ratio

- **Digoxin**
  - 1 μM
  - 10 μM
  - 100 μM

- **Propranolol**
  - 10 μM

- **AZD5672**
  - 1 μM
  - 100 μM
  - 500 μM

**P_app** (x10^-6 cm/s)
**Figure 3**

- Positive control inhibition

**P_app (x10^-6 cm/s)**

- Digoxin 1 μM
- Digoxin 1 μM + ketoconazole
- Digoxin 1 μM + quinidine
- AZD5672 1 μM
- AZD5672 1 μM + ketoconazole
- AZD5672 1 μM + quinidine

**ER (efflux ratio):**
- ER=7.3
- ER=1.1
- ER=1.0
- ER=11
- ER=0.8
- ER=1.2

ER = efflux ratio
Figure 4

B-to-A Digoxin $P_{\text{app}}$ (x10^{-6} cm/s) vs. Verapamil ($\mu$M) and AZD5672 ($\mu$M)
Figure 5

Panel A:
- Digoxin alone
- Digoxin + AZD5672 50mg

Panel B:
- Digoxin alone
- Digoxin + AZD5672 150mg