Clinical investigation of metabolic and renal clearance pathways contributing to the elimination of fevipiprant using probenecid as perpetrator

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Non-standard abbreviations:

ADME: absorption, distribution, metabolism and elimination; AE: adverse events; Ae_{0-24}:

cumulative amount of an analyte excreted in urine from zero to 24 hours; AG: acyl

glucuronide; AUC_{0-24}: area under the plasma (or serum or blood) concentration-time

curve from time zero to 24 hours (mass × time/volume); AUC_{inf}: area under the

concentration-time curve in plasma from time zero extrapolated to infinite time; AUC_{last}:

area under the concentration-time curve in plasma from time zero to time of last

quantifiable concentration; BMI: body mass index; C_{max}: maximum observed

concentration; CI: confidence interval; CL/F: apparent systemic clearance; CLr: renal

clearance from plasma (volume/time); DDI: drug interaction; ECG: electrocardiogram;

FDA: Food and Drug Administration; fu: unbound fraction; GFR; glomerular filtration rate

(approximately 7.5 L/h in a healthy subject); LC-MS/MS: liquid chromatography-mass

spectrometry; MR: metabolite-to-parent drug ratios; OAT: organic anion transporter;

PGD_2: prostaglandin D_2; DP_2: prostaglandin D_2 receptor 2; SAE: serious adverse events;

UGT: UDP-glucuronosyltransferase; Vz/F: apparent volume of distribution

Abstract

Fevipiprant, an oral, non-steroidal, highly selective, reversible, and competitive

prostaglandin D_2 receptor 2 antagonist, is eliminated by glucuronidation, and by direct

renal excretion predominantly via organic anion transporter (OAT) 3. This study aimed

to assess the effect of simultaneous UDP-glucuronosyltransferase (UGT) and OAT 3

inhibition by probenecid on the pharmacokinetics of fevipiprant and its acyl glucuronide

(AG) metabolite to support the dosing recommendation of fevipiprant in the presence of

drugs inhibiting these pathways; however, Phase III clinical trial results did not support

its submission. This was a single-center, open-label, single sequence, two-period,

crossover study in healthy subjects. Liquid chromatography with tandem mass

spectrometry was used to measure concentrations of fevipiprant and its AG metabolite

in plasma and urine. In the presence of probenecid, the mean maximum concentrations

of fevipiprant increased approximately 1.7-fold, and the area under the curve (AUC)$_{last}$

and AUC$_{inf}$ increased approximately 2.5-fold, while the mean apparent volume of

distribution and the AG metabolite-fevipiprant ratio decreased. The apparent systemic

clearance decreased by approximately 60% and the renal clearance by approximately

88% in the presence of probenecid. Using these data and those from previous studies,

the relative contribution of OAT and UGT inhibition to the overall effect of probenecid

was estimated. Furthermore, a general disposition scheme for fevipiprant was

developed, showing how a perpetrator drug such as probenecid, that interferes with two

key elimination pathways of fevipiprant, causes only a moderate increase in exposure,

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and allows estimation of the drug-drug inhibition when only one of the two pathways is inhibited.

Significance statement

In this drug-drug interaction (DDI) study probenecid was used as a tool to inhibit both glucuronidation and active renal secretion of fevipiprant. The combination of plasma and urine pharmacokinetic data from this study with available data allowed the development of a quantitative scheme to describe the fate of fevipiprant in the body, illustrating why the DDI effect on fevipiprant is weak-to-moderate, even if a perpetrator drug inhibits several elimination pathways.

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Introduction

Fevipiprant is an oral, non-steroidal, highly selective, reversible, and competitive antagonist of the prostaglandin D_2 (PGD_2) receptor 2 (DP_2) (Sykes et al., 2016). The DP_2 receptor, a G-protein-coupled receptor, is an important regulator of the inflammatory cascade with a key role in the pathophysiology of asthma (Domingo et al., 2018). Fevipiprant selectively antagonises the DP_2 receptor, thereby targeting and reducing DP_2 receptor-mediated inflammation in the airways of people with asthma (Erpenbeck et al., 2016a; Gonem et al., 2016; Bateman et al., 2017); however, Phase III clinical trial results did not support submission in this indication (Brightling et al., 2020).

Elimination of fevipiprant occurs via glucuronidation, as well as by direct renal and possible biliary excretion (Pearson et al., 2017). Fevipiprant is metabolised to an acyl glucuronide (AG) metabolite (1-O-beta form which can rearrange to isomers), representing the only relevant metabolite in systemic circulation and excreta which is not pharmacologically active. Data from clinical mass balance and drug-drug interaction (DDI) studies revealed that hepatic and renal clearance contribute to the total systemic elimination of fevipiprant and that renal clearance involves an active secretion process (Pearson et al., 2017; Weiss et al., 2020). Because the contributions of both glucuronidation and renal excretion exceed 25% of the clearance of fevipiprant, clinical studies are recommended by health authority guidelines to study the DDI risk in humans (Zhang et al., 2009; European Medicines Agency, 2013; Center for Drug Evaluation and Research (CDER), 2020)

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Fevipiprant was tested *in vitro* as a substrate of the major human UDP-glucuronosyltransferase (UGT) enzymes and drug transporters (Pearson et al., 2017). Fevipiprant is a substrate of the organic anion transporter 3 (OAT3), P-glycoprotein (P-gp), and organic anion-transporting polypeptide 1B3 (OATP1B3) and its metabolism is mediated by the human UDP-glucuronosyltransferase (UGT) enzymes UGT1A3, UGT2B7 and UGT2B17 (Pearson et al., 2017).

A previous DDI study showed that co-administration of cyclosporine, an inhibitor of OATP1B3 and P-gp, increased oral fevipiprant 150 mg area under the curve (AUC) by 2.5-fold and maximum concentration (C_{max}) by 3-fold (Weiss et al., 2020). Our study investigates the effect of inhibition of the other relevant clearance pathways of fevipiprant by probenecid i.e. metabolism by UGTs and OAT3-mediated renal clearance. OAT3 is expressed in proximal kidney tubule cells and plays an important role in the active secretion of low permeable anionic compounds (Wang and Sweet, 2013). UGT1A3 and UGT2B7 are expressed in the liver; UGT2B7 and UGT2B17 are reported to have an important role in the intestine (Strassburg et al., 2001; Kiang et al., 2005; Zhang et al., 2018).

Probenecid was used as a non-selective inhibitor of UGTs (Uchaipichat et al., 2004), and is recommended by the Food and Drug Administration (FDA) to assess sensitivity to DDI for drugs that are metabolised by several UGTs, such as fevipiprant. It is also recommended as an index perpetrator of OAT1 and OAT3 (FDA Center for Drug Evaluation and Research, 2017). The purpose of this study was to assess the effect of concurrent UGT and OAT3 inhibition by probenecid on the pharmacokinetics (PK) of fevipiprant. The assessment of the systemic PK, as well as urinary excretion of both

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fevipiprant and its pharmacologically inactive AG metabolite, allows for some distinction of the metabolic and renal effects of probenecid. The results are discussed further in the context of existing *in vitro* data and clinical results from human mass balance and DDI studies to establish an overall quantitative understanding of the disposition of fevipiprant.

Materials and methods

Regulatory and ethical compliance

The study protocol was reviewed by the Institutional Review Board and the study was conducted according to the ethical principles of the Declaration of Helsinki. Informed consent was obtained from each subject in writing before any study-specific procedures took place.

Study objectives

The primary objective of the study was to determine the effect of probenecid 1000 mg twice daily on the key PK parameters of fevipiprant 150 mg in healthy subjects. The secondary objectives were to assess the safety and tolerability of fevipiprant with and without co-administration of probenecid and to determine the effect of probenecid on the PK of the AG metabolite of fevipiprant.

Study design

This was a single-center, open-label, single sequence, two-period, crossover study in healthy subjects. The study consisted of a 28-day screening period with two baseline visits and two treatment periods, separated by a washout of 7 to 14 days, and an end of

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study assessment (Figure 1). In Treatment Period 1, subjects received a single oral fevipiprant dose of 150 mg on Day 1, followed by collection of reference plasma PK samples over 96 h and urine collection over 24 h. In Treatment Period 2, subjects received oral doses of probenecid 1000 mg twice daily on Days 16 to 21, and a single oral fevipiprant dose of 150 mg (followed by collection of plasma PK samples over 96 h and urine over 24 h) together with the probenecid morning dose on Day 18.

Key inclusion and exclusion criteria

Healthy men and women aged 18–55 years weighing between 50–90 kg were eligible to participate in the study, and were required to have a body mass index (BMI) within the range of 18–30 kg/m^2. Written informed consent was obtained before any assessment was carried out. Subjects using any prescription drugs (with the exception of oral or injectable contraceptives) were excluded from the study.

Pregnant or nursing women and those who smoked were excluded. Those with a history of clinically significant electrocardiogram (ECG) abnormalities, or history or presence of long QT syndrome were also excluded from the study. Full inclusion and exclusion criteria may be found in the Online Data Supplement.

Pharmacokinetic analyses

Blood samples were collected in K3EDTA (anticoagulant) collection tubes. As the AG metabolite is unstable at the physiological pH of plasma and urine (pH> 6.8), both urine and plasma samples were transferred after collection to commercially available Vacuette® tubes (FC) mixture (Greiner, catalogue No. 454513) for stabilization of the AG metabolite. Further details may be found in the Supplement.

The concentrations of fevipiprant and its AG metabolite in plasma and urine were measured using a validated liquid chromatography with tandem mass spectrometry (LC-MS/MS) combo-method (simultaneous quantification of both analytes with same injection), which consisted of a robotized supported liquid extraction (SLE) using Isolute SLE +, 200 µL, 96-well plates for extraction plasma samples, and liquid-liquid extraction for urine samples, respectively, followed by reverse phase LC-MS/MS using ElectroSpray Ionization in the positive ion mode (ESI+) (Erpenbeck et al., 2016b). The plasma method was suitable for quantification of fevipiprant in the range of 1 to 400 ng/mL and in the range of 0.48 to 192 ng/mL for the AG metabolite using 50 µL plasma. The method for urine could quantify fevipiprant in the range of 0.2 to 80.0 µg/mL and its AG metabolite in the range of 0.096 to 38.4 µg/mL using 50 µL urine.

The bioanalytical method for probenecid consisted of protein precipitation followed by reverse phase liquid chromatography with tandem mass spectrometric detection using ElectroSpray Ionization in the negative ion mode (ESI-). The method was suitable for the determination of probenecid in human blood over the range of 1.00 (LLOQ) to 400 µg/mL using 50 µL of human blood (further details may be found in the Supplement).

Fevipiprant PK parameters were determined using the actual recorded sampling times and non-compartmental method(s) with Phoenix WinNonlin (Version 6.4 or higher) from the plasma and urine concentration data.

Key safety and tolerability assessments

All adverse events (AEs) and serious adverse events (SAEs) with their severity and relationship to study drug were collected. Evaluations included: safety assessments

(hematology, clinical chemistry, and urinalysis), pregnancy and assessments of fertility, regular assessments of vital signs, physical condition, body weight and a standard 12-lead ECG.

Statistical analyses

To assess the effect of probenecid on the PK of fevipiprant, log-transformed primary plasma/urine PK parameters of fevipiprant were analysed using a mixed effects model with a fixed effect for treatment (fevipiprant plus probenecid versus fevipiprant alone) and a random effect for subject. Least squares mean differences between treatments and associated 90% confidence intervals (CIs) in the logarithmic scale were back-transformed to produce geometric mean ratio and associated 90% CIs for each PK parameter.

Results

Demographics

In total 16 subjects entered the study, all of whom completed Treatment Period 1, and were included in the safety and PK analysis sets. All subjects except one, who discontinued because of an adverse event, completed Treatment Period 2. Fourteen out of sixteen subjects (87.5%) were female and most were white (68.8%; Table S1 provides additional demographic information). The median age was 27.5 (range, 19–55) years; mean weight was 68.3 ± 10.4 kg and mean BMI was 24.2 ± 2.44 kg/m^2 (Table S1).

Pharmacokinetics of fevipiprant administered with and without probenecid

Co-administration of probenecid increased mean fevipiprant plasma concentrations
(Figure 2a); mean concentration-time data for fevipiprant and its AG metabolite with and
without probenecid are provided in Table S2. PK parameters for fevipiprant with and
without probenecid are summarized in Table 1 and corresponding statistical analysis in
Table 2. Fevipiprant mean C_{max} increased approximately 1.7-fold, and AUC_{last} and
AUC_{inf} increased approximately 2.5-fold in the presence of probenecid (Figure 2a; Table
2). The mean Vz/F decreased from 1470 L to 600 L. Geometric mean ratios showed a
decrease in CL/F of 25.8 L/h (a reduction of approximately 60% with reduction observed
across all subjects, Figure 3) and a decrease in renal clearance (CLr) of 8.5 L/h (a
reduction of 88%) in the presence of probenecid (Table 2; Figure 2). The mean amount
of fevipiprant excreted in urine (Ae_{0-24h}) decreased from 27.7 ± 8.57 mg (approximately
19% of the dose) for fevipiprant treatment alone to 7.08 ± 2.15 mg (approximately 5% of
the dose) for treatment with fevipiprant plus probenecid (Table 1).

*Pharmacokinetics of fevipiprant AG metabolite administered with and without
probenecid*

Although co-administration of probenecid is expected to reduce glucuronidation of
fevipiprant it resulted in an increased plasma exposure to the metabolite (Table 1).
However, the metabolite-to-parent drug ratios (MR; AG metabolite to fevipiprant) for
maximum concentration (MR C_{max}) and overall exposure (MR AUC_{last} and MR AUC_{inf})
decreased in the presence of probenecid (Table 1). The decrease in the MR for
systemic exposure suggests a reduced rate of metabolite formation resulting from
reduced UGT activity in presence of probenecid. The net increase in exposure to the
metabolite results from the 4.3-fold decrease in CLr of the AG metabolite by probenecid

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(Table 1; Figure 2b). The mean amount of the AG metabolite excreted in urine (Ae_{0-24h}) decreased from 36.5 ± 9.62 mg (approximately 17% of the dose considering difference in molecular weight) with fevipiprant alone to 13.0 ± 5.80 mg (approximately 6% of the dose) when probenecid was co-administered.

Adverse events

Headache was the most frequently reported AE (see Table S3 for all adverse events). Two AEs (postural dizziness, headache) reported for one (6.3%) subject were considered to be related to both fevipiprant and probenecid. Eight AEs reported for four (25.0%) subjects were considered to be related to probenecid. All reported AEs were mild in severity (Table S3). One subject discontinued from the study due to an AE. No deaths or serious AEs were reported.

Discussion

In this study we investigated the effect of simultaneous inhibition of UGTs and OAT3 by probenecid on the PK of fevipiprant. The inhibition of two important clearance pathways resulted in a weak (< 2-fold) effect on C_{max} and a moderate (2.5-fold) increase in AUC considering DDI categories also used by the health authorities (European Medicines Agency, 2013; Center for Drug Evaluation and Research (CDER), 2020). The exposure of healthy subjects to 150 mg fevipiprant either with or without co-administration of

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probenecid was found to be safe and generally well tolerated, with no unexpected adverse events reported. This is consistent with previous safety and tolerability findings (Erpenbeck et al., 2016a; Erpenbeck et al., 2016b; Gonem et al., 2016; Bateman et al., 2017).

The metabolic and renal effects of probenecid can be distinguished using the following information: plasma concentration data and the urinary excretion of both fevipiprant and its AG metabolite; complementary literature providing *in vitro* and absorption, distribution, metabolism, and excretion (ADME) data (Pearson et al., 2017); and oral and intravenous (IV) DDI data with and without the OATP1B3 and P-gp inhibitor cyclosporine (Weiss et al., 2020). Based on this, the fractional contribution of OAT and UGT inhibition to the observed effect was estimated and a general disposition scheme for fevipiprant established. This information, in combination with safety and efficacy data from patient trials, would have provided the basis to assess the need for a dose adjustment or contraindication of fevipiprant in presence of co-medications affecting its pharmacokinetics.

Renal elimination of fevipiprant involves glomerular filtration as well as OAT3-mediated active secretion. Fevipiprant has an unbound fraction (fu) in plasma of 0.118 (Pearson et al., 2017); therefore, the clearance by glomerular filtration can be estimated to be approximately 0.9 L/h (fu x GFR = 0.118 x 7.5 L/h ~ 0.9 L/h), assuming a GFR of 7.5 L/h (Davies and Morris, 1993). This suggests a contribution of active secretion to the renal clearance of fevipiprant of approximately 91% (9.0 L/h, Table 3). In the presence of probenecid, the active secretion was reduced to approximately 0.3 L/h (1.2 – 0.9 L/h), indicating near complete (97%) inhibition of active secretion by probenecid. The AG

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metabolite has an unbound fraction in plasma of 0.234 and therefore glomerular filtration can be estimated to be approximately 1.8 L/h (fu x GFR = 0.234 x 7.5 L/h ~ 1.8 L/h), indicating a contribution of active secretion to the renal clearance of approximately 75% (5.3 L/h, Table 3). In the presence of probenecid, urinary secretion decreased to a value close to the estimated glomerular filtration (1.7 vs. 1.8 L/h) indicating near complete inhibition of active metabolite secretion by probenecid. This sensitivity to probenecid suggests that the metabolite is subject to active renal secretion likely involving transporters of the OAT family also.

The systemic clearance (CL) of fevipiprant based on IV data is 19 L/h (Weiss et al., 2020), which allows an estimation of bioavailability (F) of 0.43 using the observed CL/F of 43.7 L/h in this study population. The measured CLr (9.87 L/h) accounts for half of CL (19 L/h) and is comparable to the observed CLr reported previously (9.49 L/h) (Weiss et al., 2020). Inhibition of OAT3-mediated active renal excretion by probenecid reduces CLr by 8.7 L/h (Table 1) and consequently CL to 10.3 L/h (19 L/h - 8.7 L/h) and CL/F to 24 L/h (10.3 L/h / 0.43). Therefore, the expected exposure increase of only inhibiting renal excretion is ~ 1.8 fold (43.7 L/h / 24 L/h), which is lower than the observed ~ 2.5 fold AUC change. Accordingly, inhibiting glucuronidation is expected to have a smaller impact (2.5/1.8 = 1.4 fold). The smaller impact of inhibiting glucuronidation is consistent with findings that the clinical effects of UGT inhibitors on the clearance of UGT substrates are generally weak (Williams et al., 2004). Taken together, inhibitors of a single pathway (OAT3 or UGTs) are expected to cause only a weak increase in AUC of fevipiprant, while simultaneous inhibition results in a weak-to-moderate exposure increase.

While UGT1A3 and UGT2B7 are expressed in the liver, extrahepatic glucuronidation of fevipiprant cannot be excluded. Based on recent protein abundance data, UGT2B7 and UGT2B17 may contribute to the intestinal first-pass and both enzymes are also expressed in the kidney (Margaillan et al., 2015). Probenecid inhibits all three UGT isoforms *in vitro* (Uchaipichat et al., 2004), with highest potency for UGT2B7. Therefore, inhibition of the intestinal UGT2B7 and UGT2B17 could result in a reduced intestinal first-pass metabolism, which is consistent with the reduction in Vz/F seen in this study. In addition, reduced distribution of fevipiprant into the kidney in the presence of probenecid can also contribute to the reduced Vz/F.

Previous *in vitro* and clinical DDI data revealed that OATP1B3-mediated uptake into the liver is a key mechanism of fevipiprant systemic elimination (Weiss et al., 2020). The results from the present study further corroborate that hepatic and renal elimination both contribute approximately 50% to the total systemic clearance of fevipiprant (Figure 4). When fevipiprant was administered together with cyclosporine or probenecid, hepatic clearance and active renal secretion, respectively, were nearly completely inhibited. Hence, the data indicate that transporter-mediated uptake processes (OATP1B3 and OAT3) are the rate-limiting clearance steps in both organs, which is in line with moderate passive permeability of fevipiprant (Pearson et al., 2017).

In summary, results from interaction studies with probenecid and cyclosporine revealed the elimination of fevipiprant is dependent on OATP1B1-mediated hepatic uptake, OAT3-mediated renal excretion and glucuronidation (via UGT1A3, UGT2B7, UGT2B17) (Figure 4). These parallel elimination pathways result in a low risk of major victim DDI, or pharmacogenetic/ethnic variability for this compound (Pearson et al., 2017). This is

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exemplified in the reported study, in which a perpetrator drug interferes with more than

one fevipiprant elimination pathway, but the DDI effect remains weak to moderate.

Acknowledgements

Data sharing statement

Novartis will not provide access to patient-level data, if there is a reasonable likelihood that individual patients could be re-identified. Phase 1 studies, by their nature, present a high risk of patient re-identification; therefore, patient individual results for phase 1 studies cannot be shared. In addition, clinical data, in some cases, have been collected subject to contractual or consent provisions that prohibit transfer to third parties. Such restrictions may preclude granting access under these provisions. Where co-development agreements or other legal restrictions prevent companies from sharing particular data, companies will work with qualified requestors to provide summary information where possible.

Authorship contribution

H Markus Weiss, Thomas Langenickel and Meredith Cain participated in the research design. Bharti Shah, Meredith Cain and Gholamreza Rahmanzadeh conducted the experiments. Bharti Shah, Birk Poller, Swarupa Kulkarni, Janardhana Vemula and Markus Weiss performed data analysis. All authors contributed to the writing of the manuscript.

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References

Bateman ED, Guerreros AG, Brockhaus F, Holzhauer B, Pethe A, Kay RA, and Townley RG (2017) Fevipiprant, an oral prostaglandin DP2 receptor (CRTh2) antagonist, in allergic asthma uncontrolled on low-dose inhaled corticosteroids. *The European respiratory journal* **50:**1-11.

Brightling CE, Gaga M, Inoue H, Li J, Maspero J, Wenzel S, Maitra S, Lawrence D, Brockhaus F, Lehmann T, Brindicci C, Knorr B, and Bleecker ER (2020) Effectiveness of fevipiprant in reducing exacerbations in patients with severe asthma (LUSTER-1 and LUSTER-2): two phase 3 randomised controlled trials. *The Lancet Respiratory Medicine*.

Center for Drug Evaluation and Research (CDER) (2020) In Vitro Drug Interaction Studies — Cytochrome P450 Enzyme- and Transporter-Mediated Drug Interactions Guidance for Industry, US Department of Health and Human Services.

Davies B and Morris T (1993) Physiological parameters in laboratory animals and humans. *Pharmaceutical research* **10:**1093-1095.

Domingo C, Palomares O, Sandham DA, Erpenbeck VJ, and Altman P (2018) The prostaglandin D2 receptor 2 pathway in asthma: a key player in airway inflammation. *Respiratory research* **19:**189.

Erpenbeck VJ, Popov TA, Miller D, Weinstein SF, Spector S, Magnusson B, Osuntokun W, Goldsmith P, Weiss M, and Beier J (2016a) The oral CRTh2 antagonist QAW039 (fevipiprant): A phase II study in uncontrolled allergic asthma. *Pulmonary pharmacology & therapeutics* **39:**54-63.

Erpenbeck VJ, Vets E, Gheyle L, Osuntokun W, Larbig M, Neelakantham S, Sandham D, Dubois G, Elbast W, Goldsmith P, and Weiss M (2016b) Pharmacokinetics, Safety, and Tolerability of Fevipiprant (QAW039), a Novel CRTh2 Receptor Antagonist: Results From 2 Randomized, Phase 1, Placebo-Controlled Studies in Healthy Volunteers. *Clinical pharmacology in drug development* **5:**306-313.

European Medicines Agency (2013) Guideline on the investigation of drug interactions, EMA London, UK.

FDA Center for Drug Evaluation and Research (2017) Clinical Drug Interaction Studies —Study Design, Data Analysis, and Clinical Implications Guidance for Industry.

Gonem S, Berair R, Singapuri A, Hartley R, Laurencin MFM, Bacher G, Holzhauer B, Bourne M, Mistry V, Pavord ID, Mansur AH, Wardlaw AJ, Siddiqui SH, Kay RA, and Brightling CE (2016) Fevipiprant, a prostaglandin D2 receptor 2 antagonist, in patients with persistent eosinophilic asthma: a single-centre, randomised, double-blind, parallel-group, placebo-controlled trial. *The Lancet Respiratory medicine* **4:**699-707.

Kiang TK, Ensom MH, and Chang TK (2005) UDP-glucuronosyltransferases and clinical drug-drug interactions. *Pharmacology & therapeutics* **106:**97-132.

Margaillan G, Rouleau M, Fallon JK, Caron P, Villeneuve L, Turcotte V, Smith PC, Joy MS, and Guillemette C (2015) Quantitative profiling of human renal UDP-glucuronosyltransferases and glucuronidation activity: a comparison of normal and tumoral kidney tissues. *Drug metabolism and disposition: the biological fate of chemicals* **43:**611-619.

Pearson D, Weiss HM, Jin Y, Jaap van Lier J, Erpenbeck VJ, Glaenzel U, End P, Woessner R, Eggimann F, and Camenisch G (2017) Absorption, Distribution, Metabolism, and Excretion of the Oral Prostaglandin D2 Receptor 2 Antagonist Fevipiprant (QAW039) in Healthy Volunteers and In Vitro. *Drug metabolism and disposition: the biological fate of chemicals* **45:**817-825.

Strassburg CP, Barut A, Obermayer-Straub P, Li Q, Nguyen N, Tukey RH, and Manns MP (2001) Identification of cyclosporine A and tacrolimus glucuronidation in human liver and the gastrointestinal tract by a differentially expressed UDP-glucuronosyltransferase: UGT2B7. *Journal of hepatology* **34:**865-872.

Sykes DA, Bradley ME, Riddy DM, Willard E, Reilly J, Miah A, Bauer C, Watson SJ, Sandham DA, Dubois G, and Charlton SJ (2016) Fevipiprant (QAW039), a Slowly Dissociating CRTh2 Antagonist with the Potential for Improved Clinical Efficacy. *Molecular pharmacology* **89:**593-605.

Uchaipichat V, Mackenzie PI, Guo XH, Gardner-Stephen D, Galetin A, Houston JB, and Miners JO (2004) Human udp-glucuronosyltransferases: isoform selectivity and kinetics of 4-methylumbelliferone and 1-naphthol glucuronidation, effects of organic solvents, and inhibition by diclofenac and probenecid. *Drug metabolism and disposition: the biological fate of chemicals* **32:**413-423.

Wang L and Sweet DH (2013) Renal organic anion transporters (SLC22 family): expression, regulation, roles in toxicity, and impact on injury and disease. *The AAPS journal* **15:**53-69.

Weiss M, Umehara KI, Erpenbeck V, Cain M, Vemula J, Elbast W, and Zollinger M (2020) A study of the effect of cyclosporine on fevipiprant pharmacokinetics and its absolute bioavailability using an intravenous microdose approach. *Drug metabolism and disposition: the biological fate of chemicals*.

Williams JA, Hyland R, Jones BC, Smith DA, Hurst S, Goosen TC, Peterkin V, Koup JR, and Ball SE (2004) Drug-drug interactions for UDP-glucuronosyltransferase substrates: a pharmacokinetic explanation for typically observed low exposure (AUCi/AUC) ratios. *Drug metabolism and disposition: the biological fate of chemicals* **32:**1201-1208.

Zhang H, Basit A, Busch D, Yabut K, Bhatt DK, Drozdzik M, Ostrowski M, Li A, Collins C, Oswald S, and Prasad B (2018) Quantitative characterization of UDP-glucuronosyltransferase 2B17 in human liver and intestine and its role in testosterone first-pass metabolism. *Biochemical pharmacology* **156:**32-42.

Zhang L, Zhang YD, Zhao P, and Huang SM (2009) Predicting drug-drug interactions: an FDA perspective. *The AAPS journal* **11:**300-306.

Footnotes

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Financial disclosure statement

All authors apart from TL are employees of Novartis and all hold shares in the company, apart from JV. TL was employed by Novartis at the time the study was conducted and eligible to receive stock.

Figure 1 Study design

Figure 2 (a) Arithmetic mean (± SD) plasma concentration of fevipiprant 150 mg once daily in the presence (black triangles) and absence (black circles) of probenecid over time (0–24 h) on a linear scale. Data presented on a semi-logarithmic scale (0–96 h) are shown as an inset.

(b) Arithmetic mean plasma (± SD) concentration of the AG metabolite of fevipiprant in the presence (black triangles) and absence (black circles) of probenecid over time (0–24 h) on linear scale. Data presented on a semi-logarithmic scale (0–96 h) are shown as an inset.

Figure 3 Fevipiprant individual and geometric mean CL/F by treatment. Individual subjects are shown as an open circle with a line connection for fevipiprant (Day 1) and fevipiprant plus probenecid (Day 18) values. Geometric mean values are represented by closed diamonds

Figure 4. Drug disposition of fevipiprant (blue arrows) and the AG metabolite (AG met, dashed purple arrows) in humans based on *in vitro* phenotyping data (Pearson et al., 2017) and clinical study results. Oral bioavailability (F) was determined in Weiss et al. (Weiss et al., 2020). Hepatic first-pass was calculated with $F_h = 1 - ((CL_h / R_b) / Q_h) = 0.8$, where CL_h represents the hepatic plasma clearance ($CL_h = CL - CL_r = 19 \text{ L/h} - 9.87 \text{ L/h} = 9.13 \text{ L/h}$), R_b is the blood-to-plasma concentration ratio (0.56) and Q_h is the hepatic blood flow (86.9 L/h)(Davies and Morris, 1993). The combined contribution of fraction absorbed and intestinal first-pass ($F_a \times F_g$) was estimated from the equation $F_a \times F_g = F_h / F$. fe (fraction excreted), represents the fractional contribution of each pathway to the extraction of fevipiprant from plasma. Details on active versus filtration clearance in the kidney are provided in the discussion and Table 3. Red lines represent inhibition mechanisms by cyclosporine (OATP1B3) and probenecid (OAT, UGTs). $CL_{r, m}$ represents the renal clearance of the AG metabolite.

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Table 1 Summary of pharmacokinetic parameters for fevipiprant and AG metabolite by treatment

Parameter (unit)	Fevipiprant 150 mg oral	Probenecid + fevipiprant 150 mg oral	AG metabolite (after dosing with fevipiprant 150 mg oral)	AG metabolite (after dosing with probenecid and fevipiprant 150 mg oral)
C_{max} (ng/mL)	812 ± 386 (47.5) [n=16]	1350 ± 682 (50.6) [n=15]	1070± 380 (35.7) [n=16]	1410 ± 703 (49.9) [n=15]
AUC_{last} (ng*h/mL)	3530 ± 945 (26.8) [n=16]	9320 ± 4180 (44.9) [n=15]	6910 ± 2280 (33.0) [n=16]	12600 ± 7560 (60.1) [n=15]
AUC_{inf} (ng*h/mL)	3680 ± 1010 (27.4) [n=15]	9980 ± 4520 (45.3) [n=15]	7220 ± 2340 (32.5) [n=16]	14400 ± 9270 (64.4) [n=12]
T_{max} (h) [†]	1.50 (1.00, 6.00) [n=16]	2.00 (1.00, 6.00) [n=15]	2.50 (2.00, 6.00) [n=16]	2.02 (1.50, 6.00) [n=15)
CL/F (L/h)	43.7 ± 11.7 (26.8) [n=15]	17.8 ± 7.12 (40.0) [n=15]	-	-
Vz/F (L)	1470 ± 926 (63.2) [n=15]	600 ± 299 (49.8) [n=15]	-	-
$T_{1/2}$ (h)	23.5 ± 14.6 (62.0) [n=15]	24.9 ± 11.2 (45.0) [n=15]	24.7 ± 14.5 (58.8) [n=16]	35.2 ± 18.9 (53.8) [n=15]
MR C_{max}	-	-	1.05 ± 0.382 (36.3) [n=16]	0.802 ± 0.329 (41.0) [n=15]
MR AUC_{inf}	-	-	1.41 ± 0.362 (25.7) [n=15]	1.06 ± 0.363 (34.4) [n=12]
MR AUC_{last}	-	-	1.41 ± 0.342 (24.1) [n=16]	0.977 ± 0.354 (36.3) [n=15]
$Ae_{0–last}$ (mg)	27.7 ± 8.57 [n=16]	7.08 ± 2.15 [n=15]	36.5 ± 9.62 [n=16]	13.0 ± 5.80 [n=15]
CLr (L/h)	9.87 ± 1.85 (18.7) [n=16]	1.21 ± 0.378 (31.2) [n=15]	7.14 ± 1.79 (25.0) [n=16]	1.66 ± 0.350 (21.1) [n=15]

Data are arithmetic means ± SD (CV%) [n]. CV% = sqrt(exp(variance for log transformed data)-1)*100.

[†]For T_{max}, data are median (min–max) [n]. MR: Metabolite-to-parent drug ratio.

Table 2 Geometric mean ratio (90% confidence intervals) for fevipiprant 150 mg pharmacokinetic parameters with and without probenecid 1000 mg twice daily (pharmacokinetic analysis set)

Parameter	Treatment	$n^{\dagger}$	Adjusted geometric mean (90% CI)	Comparison	Treatment comparison	
					Geometric mean ratio	90% CI
AUC_{last} (ng*h/mL)	Fevipiprant	16	3412 (2943, 3956)	Fevipiprant + Probenecid vs. Fevipiprant	2.48	2.15, 2.87
	Fevipiprant + Probenecid	15	8466 (7278, 9848)			
AUC_{inf} (ng*h/mL)	Fevipiprant	15	3537 (3022, 4140)	Fevipiprant + Probenecid vs. Fevipiprant	2.55	2.17, 2.99
	Fevipiprant + Probenecid	15	9012 (7701, 10547)			
C_{max} (ng/mL)	Fevipiprant	16	722 (586, 891)	Fevipiprant + Probenecid vs. Fevipiprant	1.67	1.33, 2.10
	Fevipiprant + Probenecid	15	1204 (970, 1494)			
CL/F (L/h)	Fevipiprant	15	42.4 (36.2, 49.6)	Fevipiprant + Probenecid vs. Fevipiprant	0.392	0.335, 0.460
	Fevipiprant + Probenecid	15	16.6 (14.2, 19.5)			
CLr (L/h)	Fevipiprant	16	9.7 (8.6, 10.9)	Fevipiprant + Probenecid vs. Fevipiprant	0.120	0.106, 0.136
	Fevipiprant + Probenecid	15	1.2 (1.0, 1.3)			

Mixed effects model with a fixed effect for treatment and a random effect for subject. Subjects with missing PK parameters for any treatment, but not all treatments, are included in the analysis assuming "missing" as random. Geometric mean ratios and 90% CI are back transformed from log scale.

[†]Number of evaluable subjects.

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Table 3 Renal clearance of fevipiprant and its AG metabolite with and without probenecid

Parameter	Fevipiprant		AG metabolite	
	Without probenecid	With probenecid	Without probenecid	With probenecid
CLr (L/h)	9.9	1.2	7.1	1.7
Glomerular filtration CLr (L/h)	0.9	0.9	1.8	1.8
Active CLr (L/h)	9.0	0.3	5.3	0
Contribution of active CLr to CLr (%)	91%	25%	75%	0

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Figure 1

28 day screening
(Day -28 – Day -2)

7–14 days

(a)

(b)

Figure 2

Fevipiprant (n=15)   **Fevipiprant + Probenecid (n=15)**

Figure 3

Figure 4