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ARTICLE

Understanding the mechanisms of food effect on omaveloxolone pharmacokinetics through physiologically based biopharmaceutics modeling

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Abstract

Omaveloxolone is a nuclear factor (erythroid-derived 2)-like 2 activator approved in the United States and the European Union for the treatment of patients with Friedreich ataxia aged ≥16 years, with a recommended dosage of 150mg orally once daily on an empty stomach. The effect of the US Food and Drug Administration (FDA) high-fat breakfast on the pharmacokinetic profile of omaveloxolone observed in study 408-C-1703 (NCT03664453) deviated from the usual linear correlation between fed/fasted maximum plasma concentration (C_{max}) and area under the concentration–time curve (AUC) ratios reported for various oral drugs across 323 food effect studies. Here, physiologically based biopharmaceutics modeling (PBBM) was implemented to predict and explain the effect of the FDA high-fat breakfast on a 150-mg dose of omaveloxolone. The model was developed and validated based on dissolution and pharmacokinetic data available across dose-ranging, food effect, and drug–drug interaction clinical studies. PBBM predictions support clinical observations of the unique effect of a high-fat meal on omaveloxolone pharmacokinetic profile, in which the C_{max} increased by 350% with only a 15% increase in the AUC. Key parameters influencing omaveloxolone pharmacokinetics in the fasted state based on a parameter sensitivity analysis included bile salt solubilization, CYP3A4 activity, drug substance particle size distribution, and permeability. Mechanistically, in vivo omaveloxolone absorption was solubility and dissolution rate limited. However, in the fed state, higher bile salt solubilization led to more rapid dissolution with predominant absorption in the upper gastrointestinal tract, resulting in increased susceptibility to first-pass gut extraction; this accounts for the lack of correlation between C_{max} and AUC for omaveloxolone.

Scott M. Hynes and Deborah Walker: Employees at the time of development of this publication.

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Study highlights

WHAT IS THE CURRENT KNOWLEDGE ON THE TOPIC?

Omaveloxolone is approved in patients with Friedreich ataxia aged ≥16 years at an oral dosage of 150mg once daily on an empty stomach. Food effect on omaveloxolone pharmacokinetics observed clinically deviates from the usual correlation between fed/fasted maximum plasma concentration (C_{max}) and area under the concentration–time curve (AUC) ratios reported for other oral drugs across food effect studies. Physiologically based biopharmaceutics modeling (PBBM) can predict and explain food effect through integration of drug physiochemical properties, formulation, and metabolism data.

WHAT QUESTION DID THIS STUDY ADDRESS?

A PBBM was developed and validated to predict and explain the effect of the US Food and Drug Administration high-fat meal on the pharmacokinetics of omaveloxolone 150mg.

WHAT DOES THIS STUDY ADD TO OUR KNOWLEDGE?

Consistent with clinical findings, the PBBM predicts the unique pharmacokinetic profile for omaveloxolone administered after a high-fat meal, for which the AUC was only modestly increased $(+15%)$ despite a substantial rise $(+350%)$ in the C_{max} . In vivo, omaveloxolone absorption was solubility and dissolution rate limited in the fasted state. However, in the fed state, bile salt solubilization led to more rapid drug dissolution with predominant absorption in the upper gastrointestinal tract, resulting in increased susceptibility to CYP3A4-mediated first-pass gut extraction; this mechanistically explains the lack of correlation between C_{max} and AUC for omaveloxolone.

HOW MIGHT THIS CHANGE CLINICAL PHARMACOLOGY OR TRANSLATIONAL SCIENCE?

Omaveloxolone pharmacokinetics vary in the presence of a high-fat meal versus the fasted state, reinforcing the importance of physician and patient education on proper administration and dosing. The PBBM shows potential in accurately predicting food-related impacts on drug pharmacokinetics even for outlier trends and potentially reduces the need for clinical pharmacokinetic evaluation.

INTRODUCTION

Friedreich ataxia (FA) is a progressive, autosomal recessive neurodegenerative disorder characterized by difficulty with ambulation, coordination, and speech. $1,2$ In FA, a biallelic trinucleotide (GAA) repeat expansion in the first intron of the frataxin gene leads to impaired transcription and reduced amounts of functional frataxin protein.^{1,2} Frataxin deficiency results in dysregulation of antioxidant defenses, mitochondrial dysfunction, and impaired nuclear factor (erythroid-derived 2)-like 2 (Nrf2) signaling.^{3–5}

Omaveloxolone, an Nrf2 activator, has been shown to improve mitochondrial function, restore redox balance, and reduce inflammation in FA models.^{6,7} In a registrational phase II trial (MOXIe; NCT02255435), omaveloxolone significantly improved neurological function versus placebo, with an acceptable safety profile. $8,9$ Omaveloxolone was approved in the United States and the

European Union for the treatment of FA in patients aged \geq 16 years.¹⁰ The recommended dosage is 150 mg administered orally once daily (QD) in the form of three 50-mg capsules or the entire capsule contents sprinkled on and mixed in 2 tablespoons (30mL) of applesauce, on an empty stomach at least 1h before (United States and European Union) or 2 h after (European Union) eating. $11,12$

Food–drug interactions can potentially affect the release (from a given formulation), solubility, dissolution, absorption, first-pass metabolism, and/or elimination of an oral drug and possibly impact efficacy and safety. 13 Understanding the differences in the pharmacokinetics (PK) of omaveloxolone in various prandial states and the underlying mechanisms explaining these phenomena is essential. Physiologically based biopharmaceutics modeling (PBBM) is an evolving tool that has been widely applied to predict the absorption and PK of oral drug products (DP).[14](#page-11-7) These models integrate the physicochemical and

biopharmaceutics properties of the drug substance (DS), formulation characteristics, and system physiological parameters.¹⁴ By using (biopredictive) dissolution testing as a key input, PBBM enables manufacturing flexibility by delineating a safe space for DP critical quality attributes.^{[15](#page-11-8)} Through virtual population studies, PBBM has served as an alternative to clinical PK trials that evaluate food effects, drug–drug interaction (DDI), or formulation changes.^{[15](#page-11-8)}

Here, a PBBM was developed to predict and explain the effect of a high-fat meal on the PK of a 150-mg dose of omaveloxolone. The model was validated against PK data from dose-ranging, food effect, and DDI clinical studies.

MATERIALS AND METHODS

Overview of modeling strategy

The modeling strategy is illustrated in Figure [1.](#page-3-0) A physiologically based PK (PBPK) modeling absorption baseline model was first established using intravenous and oral data in monkeys to derive relevant PK distribution parameters and then applied to humans. The in vivo capsule opening time and dissolution were based on the DS and DP performances; these together with other biopharmaceutical drug properties such as solubility, passive permeability, and the effect of systemic drug transporter P-glycoprotein (P-gP) on drug efflux were integrated into the PBBM. Metabolic clearance was specified based on in vitro data and verified with DDI studies. The determination of solubility, precipitation rate, dissolution, permeability, drug efflux, distribution, metabolism, and elimination are described in the [Supplementary](#page-12-0) [Materials and Methods](#page-12-0) in Data [S1](#page-12-0) (including Figure [S1](#page-12-0)).

The PBBM was validated using data from nine clinical scenarios from clinical PK studies 408-C-1703 (NCT03664453), thereafter referred to as study 1703, and 408-C-1806 (NCT04008186), thereafter referred to as study 1806, which tested different doses, prandial states, and DDIs. Parameter sensitivity analyses (PSAs) were run on the validated model to identify the main sources of within- and between-participant variability and the factors limiting omaveloxolone absorption. With the validated model, the effect of a high-fat meal on the PK of omaveloxolone was evaluated and explained mechanistically. The DDI and PBPKPlus modules of GastroPlus v9.8.2 (GastroPlus; Simulations Plus) and the ADMET Predictor v10.3 (APv10.3; Simulations Plus) were used.

Clinical studies

The average participant demographic information from two selected studies with comprehensive PK data was

used for validation of the PBPK absorption baseline model. Study 1703 was a phase I, open-label, two-part, food effect (part 1) and dose proportionality (part 2) study of omaveloxolone in healthy adult participants (*N*=34). In part 1 (two-period, fixed-sequence, randomized crossover design), participants were randomly assigned 1:1 to one of the two treatment sequences (sequence 1: period 1 fed and period 2 fasted; sequence 2: period 1 fasted and period 2 fed), each with a 1-week washout period. Participants were administered two single doses of omaveloxolone 150mg at the start of each period. During the fed state, participants were provided the US Food and Drug Administration (FDA) high-fat standardized breakfast (800–1000 calories, with \geq 50% from fat)^{[16](#page-11-9)} prior to dosing. In part 2, participants were randomized 1:1 to receive a single dose of either 50-mg or 100-mg omaveloxolone in a fasted state; data for omaveloxolone 150mg (taken in a fasted state) from part 1 were included in part 2 analyses. The design of study 1703 is further detailed in the [Supplementary Materials and Methods](#page-12-0) (Data [S1](#page-12-0)).

Study 1806 was a phase I, open-label, four-part, DDI study of omaveloxolone in healthy participants $(N=61)$. Participants were treated with omaveloxolone 150mg QD on days 1 and 13, and oral doses of a cytochrome P450 2C8 (CYP2C8) inhibitor gemfibrozil 600mg twice daily (part 2), a strong CYP3A4 inhibitor itraconazole 200mg QD (part 3), or a P-gP inhibitor and a moderate CYP3A4 inhibitor verapamil 120mg QD (part 4), on days 10 to 18. Primary end points were maximum plasma concentration (C_{max}) and area under the concentration–time curve (AUC) from time 0 extrapolated to infinity ($AUC_{0-\infty}$), and time to $C_{\text{max}}(t_{\text{max}})$; AUC from time 0 to the last quantifiable plasma concentration (AUC_{0-t}) was a secondary end point.

Model validation

For model validation, the prediction performance indicators were calculated for the PK parameters and profiles as described below.

Average fold error (AFE) is defined by the following equation:

$$
AFE = 10^{\frac{1}{n} \sum \log \frac{\text{Pred}_i}{\text{Obs}_i}}
$$

The AFE is an indicator of prediction bias. A method that predicted all actual values with no bias would have a value of 1; underpredictions are indicated by an AFE of \leq 1 and overpredictions by AFE of $>$ 1. AFE values generally vary between 0 and infinity; a prediction may be considered satisfactory if the AFE is between 0.8 and 1.2, passable if the AFE is 0.5 to 0.8 or 1.2 to 2, and poor if the

FIGURE 1 Modeling strategy for omaveloxolone PBBM. DS, drug substance; IV, intravenous; *K*p, permeability constant; PBBM, physiologically based biopharmaceutics modeling; PBPK, physiologically based pharmacokinetic; *P_{eff}*, effective permeability; PSD, particle size distribution.

AFE is 0 to <0.5 or >2. A satisfactory AFE is needed for model validation.

Absolute average fold error (AAFE) is defined by the following equation:

$$
AAFE = 10^{\frac{1}{n} \sum \left| \log \frac{Pred_i}{Obs_i} \right|}
$$

The AAFE converts negative log fold errors to positive values before averaging them and measures the spread of the predictions. AAFE values vary between 1 and infinity. A method that predicted all actual values perfectly would have a value of 1; one with predictions that were on average twofold off (above 100% or below

50%) would have a value of 2 and so forth. A prediction may be considered satisfactory if the AAFE is <1.2 , passable if the AAFE is in the range of 1.2 to 2, and poor if the AAFE is >2. A satisfactory AAFE is needed for model validation.

Average absolute prediction error (AAPE%) is defined by the following equation:

$$
AAPE \text{ } (\%) = Average \left(\left| \frac{Pred_i - Obs_i}{Obs_i} \right| \right) \times 100
$$

AAPE is the measurement of prediction error scaled to percentage units. It approximates $(AAFE -1) \times 100$. A model is considered satisfactory if the AAPE is <20%, passable if the AAPE is ≥20 to <50%, and poor if the AAPE $is > 50\%$.

Percent predictions within clinical variability (PPWCV) are defined by following equation:

$$
PPWCV \left(\% \right) = \frac{n_{\text{YES}}}{n_{\text{total}}} \times 100
$$

For each PK sampling time point 1 to n_{total} (apart from pre-dose), a binary criterion yes or no is determined based on whether the predicted concentration falls within the 95% confidence interval of the measured clinical data. The PPWCV calculated for each PK profile was averaged across all the clinical scenarios tested. The same calculations were performed for PK parameters the binary criterion for each clinical scenario was based on whether the predicted PK parameter fell within the 95% confidence interval of the measured average value of that parameter. A satisfactory PPWCV is >80%, a passable PPWCV is in the range of ≥65% to 80%, and a poor PPWCV is <65%.

Parameter sensitivity analyses

The PSA for C_{max} , AUC_{0-t} , and other PK parameters [\(Supplementary Materials and Methods](#page-12-0)) was based on a range of selected DP properties and physiological parameters that could impact omaveloxolone absorption or metabolism by affecting capsule opening time, size of the DS (controlling in vivo dissolution), first-pass gut and liver extraction, and metabolic elimination in vivo (Table [S1\)](#page-12-0). The analysis was performed using omaveloxolone 150mg (target dose) in the fasted state (for increased sensitivity with a lower fraction absorbed vs. the fed state) on a representative population based on the MOXIe registrational study cohort with FA (average age: 26 years; average weight: 69 kg).¹⁰

Simulation design

The PBBM was built using default values based on human fasted or fed physiologies. Since omaveloxolone is not ionized in the physiological pH range, adjustment of surface solubility was not needed. For PBBM validation, populations representative of the clinical trials were created based on the average height and weight of the cohorts. The advanced compartmental and transit model physiologies were adjusted for body weight. All the doses and prandial states tested in the clinical trials were reproduced in the PBBM. The default optimum log D model SA/V 6.1 was applied to calculate absorption scaling factors. Since omaveloxolone is lipophilic with log *p*>5, the amount of absorption scaling factors was increased in the cecum and colon.

To predict food effect in study 1703, the FDA high-fat breakfast option was selected with default zero-order gastric emptying. The gastric emptying time was increased from 3.51 to 5h to match the time needed for the intragastric volume to fall below 100mL after a high-fat meal intake.[17](#page-11-10)

Because CYP3A4, but not CYP2C8, is largely involved in omaveloxolone metabolism, 18 DDI is expected for coadministration of omaveloxolone with CYP3A4 inhibitors. Hence, two DDI simulations based on parts 3 and 4 of study 1806 were performed ([Supplementary Materials and](#page-12-0) [Methods,](#page-12-0) Figures [S2-S5](#page-12-0)). The DDI module of GastroPlus was used, and simpler modeling options based on reported values for inhibition were also tested for model validation. To account for itraconazole 200mg co-administration outside of the DDI module, the maximum rate of reaction (V_{max}) in the gut and liver for CYP3A4 was reduced by a factor of 3.9.^{19,20} The V_{max} for P-gp was set to 0 for verapamil co-administration.

Ethics statement

The clinical studies were designed and monitored in accordance with sponsor procedures, which complied with the ethical principles of good clinical practice and with the Declaration of Helsinki. Each trial was approved by the institutional review board or independent ethics committee at the respective centers. All patients provided written informed consent.

RESULTS

Modeling parameters

The main physicochemical and biopharmaceutical properties of omaveloxolone used for model parameterization are shown in Table [1.](#page-5-0) The derivation of these parameters (solubility and precipitation [Figure [S6\]](#page-12-0), dissolution, permeability [Figures [S7](#page-12-0) and [S8](#page-12-0)], P-gp efflux, and metabolism and elimination [Figure [S9](#page-12-0)]) is shown in the [Supplementary Results](#page-12-0).

Model validation

The PBPK absorption model and PBBM were validated across nine distinct clinical scenarios in studies 1703 and 1806. Profile predictions are shown in Figures [S9](#page-12-0)

TABLE 1 Main physicochemical and biopharmaceutics properties of omaveloxolone used in the model.

Abbreviations: *B*/*P*, blood:plasma concentration ratio; CYP3A4, cytochrome P450 3A4; $D_{(v,0.1/0.5/0.9)}$, 10%/50%/90% of particles are smaller than the stated size; DP, drug product; DS, drug substance; FaSSIF-V2, fasted-state simulated intestinal fluid V2; FDA, US Food and Drug Administration; FeSSIF, fed-state simulated intestinal fluid; fu, fraction unbound; HLM, human liver microsome; IR, immediate-release; IV, intravenous; *K*m, Michaelis–Menten constant; *K*p, permeability constant; log *P*; partition coefficient/lipophilicity; NDA, new drug application; PBPK, physiologically based pharmacokinetic; *P*eff, effective permeability; P-gp, P-glycoprotein; PK, pharmacokinetic; pKa, acid-dissociation constant; PSD, particle size distribution; *V*max, maximum rate of reaction.

(right panel), [S10](#page-12-0), and [S11](#page-12-0); calculations of model prediction performance indicators are shown in Table [2,](#page-6-0) Tables [S2-S4](#page-12-0).

All predicted PK parameters over the nine clinical scenarios were within the observed clinical variability (Table [2,](#page-6-0) Figures [S9](#page-12-0) [right panel], [S10](#page-12-0), and [S11](#page-12-0)). The use of the GastroPlus "controlled release undissolved dispersed" model in conjunction with the DS particle size distribution (PSD) allowed the separation of in vivo drug release (capsule opening) from in vivo dissolution of the capsule contents. This approach provided a satisfactory PK profile prediction, with 98% of the predicted concentrations within the clinical variability at each data point. All model prediction performance indicators on AUC and C_{max} were satisfactory based on predefined criteria, and the PBBM was considered validated.

Parameter sensitivity analyses

The PSA revealed that the main parameters influencing the PK of omaveloxolone 150mg were bile salt solubilization, V_{max} of CYP3A4, DS PSD, drug permeability, and volume in the small intestine (Figure [2](#page-6-1), [Supplementary](#page-12-0) [Results,](#page-12-0) Figure [S12\)](#page-12-0). The model's sensitivity to the DS PSD confirmed the decision to use the measured DS batch particle size for the analysis. Precipitation time, transit times, and P-gp function did not have a substantial effect on omaveloxolone PK. In particular, the stomach transit time from 0.25 to 0.5h or Weibull lag from 0.12 to 0.5h did not affect omaveloxolone PK. The main betweenparticipant sources of variability were drug permeability and CYP3A4 expression and function. The main withinparticipant sources of variability were differences in bile

Note: Bold numbers on the top line represent average values of the simulated clinical scenarios.

Abbreviations: AAFE, absolute average fold error; AAPE, average absolute prediction error; AFE, average fold error.

FIGURE 2 PSA results showing the main parameters that affect AUC_{0-t} (left panel) and *C*_{max} (right panel). In each panel, the first eight most influential parameters are displayed for clarity. $\mathrm{AUC}_{0\text{-}t}$, area under the curve from time 0 to the last quantifiable plasma concentration; *C*max, maximum plasma concentration; CYP3A4, cytochrome P450 3A4; PBPK, physiologically based pharmacokinetic; *P*eff, effective permeability; PSA, parameter sensitivity analysis; V_{max} , maximum rate of reaction.

salt concentration during the day in the fasted and fed states and the type of food.

Prediction of food effect

The effects of a high-fat meal on omaveloxolone 150mg in vivo dissolution, first-pass extraction, and PK are shown in Figure [3](#page-7-0).

The fraction of dose absorbed (Fa) for omaveloxolone and the fraction reaching the portal vein after passing through the gut wall without metabolism ($Fa \times fraction$ of drug escaping first-pass gut metabolism [Fg]) in the fasted and fed states are shown in Figure [4](#page-8-0). In both prandial states, the in vivo dissolution of omaveloxolone was limited by solubility in all compartments. However, the Fa of omaveloxolone increased from 52% in the fasted to 87% in the fed state, attributed to the drug's lipophilicity and strong affinity to bile salt micelles, along with higher concentration of micelles after a high-fat meal. The resulting increased solubility and faster dissolution contributed to the majority of absorption occurring in the upper gastrointestinal (GI) tract in the fed state. In the fasted state, omaveloxolone absorption occurred along the GI tract, with the highest fraction absorbed in the cecum and colon as shown by the secondary peaks in the concentration–time plots (Figure [S12\)](#page-12-0). As there is a lower expression of CYP3A4 in the lower versus upper GI tract, the fraction of omaveloxolone lost by first-pass gut extraction was

FIGURE 3 PBBM prediction of omaveloxolone PK profiles (upper panels) and the amount and percentage of drug dissolved, absorbed, metabolized, and entered into the portal vein or the systemic circulation (lower panels) following a single dose of omaveloxolone 150mg in the fasted (left panels) and fed (right panels) states. In the top panel, the symbols represent the (arithmetic) mean of measured omaveloxolone concentrations and the error bars represent \pm standard deviation. PBBM, physiologically based biopharmaceutics modeling; PK, pharmacokinetic.

FIGURE 4 PBBM prediction of the fraction absorbed (Fa) and the fraction reaching the portal vein after passing through the gut wall without metabolism $(Fa \times Fg)$ at each gut compartment for a single dose of omaveloxolone 150mg administered in the fasted and fed states. Data labels shown are rounded to whole numbers. PBBM, physiologically based biopharmaceutics modeling.

limited in the fasted versus fed state. Fa \times Fg was 36% in the fed state versus 29% in the fasted state (Figure [4\)](#page-8-0), and hepatic extraction was 33% and 28%, respectively. These phenomena explain the negligible increase in AUC (+15%) despite a large increase in C_{max} (+350%) in the fed state for omaveloxolone 150mg observed in the food effect study 1703.

Food effect and dose proportionality results from study 1703

Findings from study 1703 corroborated the results of the PBBM analysis (detailed in the [Supplementary Results,](#page-12-0) Tables [S5–S9,](#page-12-0) Figure [S14\)](#page-12-0).

DISCUSSION

Omaveloxolone is currently indicated to be taken on an empty stomach with capsules either swallowed whole or contents sprinkled on and mixed in 2 tablespoons of applesauce for patients with swallowing difficulties. Omaveloxolone displays a unique PK profile in the presence of standardized FDA high-fat breakfast: AUC was only modestly increased $(+15%)$ despite a substantial rise in *C*max (+350%). The PBBM developed for omaveloxolone provides a mechanistic explanation of this unique food effect.

The PBBM, informed by the DS and DP characteristics and drug metabolism, was validated across the doses, prandial states, and DDI scenarios studied. Consistent with permeability and CYP3A4 activity identified as main between-participant sources of variability from the PSA in this study, Avdeef et al. 21 reported a 60% variability in permeability data obtained from humans across drugs with low and high permeability; furthermore, Lown et al. 22 22 22

reported high interparticipant variability in gut CYP3A4 expression, at 11-fold based on protein, 8-fold based on mRNA, and 6-fold based on catalytic activity. These sources of variability should not exist for cross-over trials. Difference in bile salt concentration was identified as a main within-participant source of variability in this study, in line with a previous report. 23 23 23 The variability in bile salt lumen concentration has also been reported to span several logs, even in the fasted state. 24 24 24 The within-participant variability during the day was reported to be high even in the fasted state, covering 2 log differences. 25

Based on the PBBM prediction, omaveloxolone absorption was mainly limited by drug solubility along the GI tract in the fasted state (Figure [3](#page-7-0)). Omaveloxolone belongs to Biopharmaceutics Classification System class 4 DS, which is characterized by low permeability, restricting absorption in the GI tract, and continuous drug dissolution. This underscores the importance of permeability as a limiting factor for AUC and C_{max} in the fasted state (Figure [2\)](#page-6-1). The Fa of omaveloxolone in the fasted state as predicted by the PBBM was approximately 52%. Consistently, based on the human absorption, metabolism, and excretion study (Study 1805), omaveloxolone was the most abundant component (approximately 40%) in the feces of participants who were administered a 150-mg dose of $[^{14}C]$ omaveloxolone, indicating an Fa of up to 60% ^{[18](#page-11-11)} Under fed conditions, omaveloxolone absorption occurred more rapidly due to the drug's affinity for bile salts, which resulted in faster dissolution in the fed state versus the fasted state.

The PBBM prediction is corroborated by the individual fasted- and fed-state omaveloxolone PK profiles in Study 1703 (Figure [S13\)](#page-12-0). Under the fasted state, in addition to an initial rise in omaveloxolone plasma concentration, multiple peaks were observed in most individual PK profiles, mainly occurring after 4h (when the drug should have reached the lower intestine in humans), suggesting that absorption predominantly occurred in the colon. This supports the model prediction that drug dissolution and solubility were the rate-limiting factors for absorption in the fasted state. Under fed conditions, there was a notably faster drug absorption with t_{max} more frequently observed before omaveloxolone reached the colon (Figure [S13\)](#page-12-0), indicating that drug absorption predominantly occurred in the upper segments of the GI tract. The faster (withinparticipant) absorption in the fed state also suggested that higher concentrations of bile salts in the GI tract lumen could partially overcome the limited solubility (and therefore in vivo dissolution) of omaveloxolone.

The uniqueness of the food effect observed with omaveloxolone, compared with other medications, is shown by the correlation between fed/fasted C_{max} and AUC ratios from 323 food effect studies encompassing a range of compounds and formulations. $26-31$ The linear correlation between $log(C_{\text{max}} \text{ ratio})$ and $log(AUC \text{ ratio})$ for investigated drugs that were proposed by Omachi et al. 29 was applied to the dataset of the present study (Figure [5a\)](#page-9-0).

Based on the linear regression, when $log(AUC \text{ ratio})=0$ (i.e., AUC ratio=1), the corresponding $log(C_{\text{max}}$ ratio) is expected to be slightly negative at −0.0651 (i.e., *C*max ratio = 0.86 on average). This lower C_{max} ratio to AUC ratio (<1) is likely related to prolonged gastric emptying in the fed state, which delays the passage of drug to the small intestine. C_{max} and AUC (ratios) are usually expected to be highly correlated since the AUC is calculated from the integration of concentration–time profiles. In most cases, drugs with solubility-limited absorption and without significant first-pass extraction have increased solubility due to the presence of higher amounts of bile salts in the GI tract lumen under the fed state. This leads to increased drug concentration in close proximity to the absorptive surface of the intestine and a larger amount of drug reaching the systemic circulation (i.e., higher AUC) versus in the fasted state.

There are, however, physiological factors and methodological issues that can explain deviations in the correlation between C_{max} and AUC. PK sampling frequency

FIGURE 5 (a) Correlation between fed/fasted C_{max} ratio versus AUC ratio. (b) σ_i values for each observed C_{max} ratio versus AUC ratio for omaveloxolone and 323 food effect studies. The solid square symbol shows omaveloxolone. A1: CPD3 100mg administered with a high-fat meal³⁸; A2: CPD3 50 mg administered with a high-fat meal³⁸; B1: progesterone-Utrogestran 200mg administered with a 688-calorie meal containing 34% fat content³⁹; B2: progesterone— Utrogestran 100mg administered with an 890-calorie meal containing 55% fat content.^{[27](#page-12-10)} σ_i , mean of sigma; AUC, area under the curve; C_{max} , maximum plasma concentration; CPD3, adenosine monophosphate deaminase inhibitor.

could miss the C_{max} during clinical food effect studies, as the same sampling time points are typically used for the fasted and fed states. In some studies, PK sampling was not frequent enough to capture C_{max} in the fed state due to prolonged gastric emptying, which may lead to underestimation of both C_{max} and AUC and a false-negative finding of food effect. 32 In addition, in both prandial states, gastric emptying of solid phases that comprise the drug might not be happening all at once, and gastric retention could account for a lower C_{max} compared to single-phase emptying. Partial gastric emptying has been reported in the fasted and fed states for various drugs. $32-37$ If first-pass extraction is not limiting drug absorption in a saturable way, a lower C_{max} due to partial gastric emptying could be associated with the same AUC as when gastric emptying happens in a single phase, leading to decorrelation of C_{max} and AUC.

Taken together, a correlation between drug-fed/fasted C_{max} and AUC ratios is expected and justifiable. The presence of outliers in this correlation is particularly intriguing, primarily to understand the mechanism of food effect and to assess the predictive capability of mechanistic models. The correlation depicted in Figure [5a](#page-9-0) could prognosticate measured fed/fasted C_{max} ratios based on corresponding fed/fasted AUC ratios. The statistics associated with using the correlation to predict measured C_{max} ratios would lead to an AFE of 1.00, an AAFE of 1.29, and an AAPE of 27%. The relative standard deviation for the prediction of *C*max based on this correlation was assumed to be 29% from the AAFE value. Assuming that C_{max} ratio observations were randomly spread around the predicted *C*max ratios with a log-normal distribution, the probability of observing any *C*max ratio value is calculated by means of sigma (*σⁱ*) derivation.

$$
\sigma_i = \frac{|P_i - O_i|}{(\text{AAFE} - 1) \times P_i}
$$

In the above equation, the value of σ_i can help estimate the probability of observing a C_{max} ratio (O_i) around the prediction P_i . For $\sigma_i = 2$, the chance to observe (O_i) based on P_i and the spread of the data approximates 1 of 3; for $\sigma_i = 3$, the chance is 1 of 15; and for $\sigma_i = 4$, the chance is 1 of 160. σ_i can be used to identify outliers to the correlation between C_{max} ratio and observed AUC ratio. The σ_i value for omaveloxolone and the 323 food effects studies from the literature are shown in Figure [5b](#page-9-0). The C_{max} ratio versus AUC ratio correlation established is good and roughly what would be expected from a log-normal distribution with 69%, 91%, and 95% of the observed C_{max} ratio within 1, 2, and 3 σ_i of the predictions, respectively. The unique nature of the food effect on omaveloxolone was evident as the σ_i value for omaveloxolone was largest at 11.7 (close

to 12) away from the prediction, while the second largest outlier σ_i value was only at approximately 7. Interestingly, the other four outliers (besides omaveloxolone) in the C_{max} ratio versus AUC ratio correlation in Figure [5b](#page-9-0) corresponded to only two drugs (CPD3 and progesterone).^{27,38,39} These two drugs show similarities with omaveloxolone (e.g., lipophilicity [log P] was reported to be 6.6 and 3.9, respectively).[38,40](#page-12-8) Both CPD3 and progesterone showed secondary absorption phases based on average PK profiles in the fasted state, which was not observed or occurred to a much lower extent in the fed state.^{27,40} Although the detailed evaluation of these effects goes beyond the scope of this article, the effect of food on the PK of CPD3 was correctly predicted using GastroPlus.[40](#page-12-12)

CONCLUSIONS

A PBBM was successfully developed and validated for omaveloxolone. Drug absorption for omaveloxolone is solubility and dissolution rate limited. In the fasted state, omaveloxolone is incompletely absorbed, with absorption predominantly occurring in the lower segment of the GI tract where the CYP3A4-mediated first-pass gut extraction is low. In the fed state, the higher amount of bile salt micelles present led to an increased solubility of omaveloxolone attributed to its weakly acidic and highly lipophilic nature, thereby accelerating in vivo dissolution, resulting in predominant absorption in the upper GI tract. In this region, omaveloxolone is subjected to a more substantial first-pass gut extraction, causing a notable transient surge in *C*max without a correspondingly large increase in AUC. This food effect on the PK of omaveloxolone deviates from that of other drugs for which the fed/fasted ratios for C_{max} and AUC are generally well correlated. These findings point to the impact of fed versus fasted condition on the PK profile of omaveloxolone; hence, reinforcing the importance of physician and patient education on administration and dosing compliance.

In silico PBPK modeling and PBBM tools offer promising platforms that integrate drug dissolution, formulation characteristics, precipitation, degradation, first-pass extraction, and metabolism. These tools accurately forecast the impact of food on oral drugs, such as omaveloxolone, potentially reducing the need for clinical evaluations. Notably, the accurate prediction of a 12-*σ* outlier event of *C*max versus AUC ratio in this case study bolsters the credibility of such models for the prediction of food effects.

AUTHOR CONTRIBUTIONS

X.J.H.P., S.M.H., H.Z., D.W., L.Q.S., and S.S.-S. wrote the manuscript; X.J.H.P., S.M.H., H.Z., and D.W. designed the research; X.J.H.P., H.Z., and D.W. performed the research; **1782** *cc* **e PEPIN** et al.

X.J.H.P., S.M.H., H.Z., D.W., L.Q.S., and S.S.-S. analyzed the data; and H.Z. contributed new reagents/analytical tools.

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CONFLICT OF INTEREST STATEMENT

Xavier J.H. Pepin and Sandra Suarez-Sharp are employees of and hold stock in Simulations Plus, which was commissioned by Reata Pharmaceuticals to conduct the study; Reata was acquired by Biogen in 2023. Scott M. Hynes was an employee and may have held stock in Biogen at the time of development of this publication. Hamim Zahir and Lois Q. Semmens are employees of and may hold stock in Biogen. Deborah Walker was an employee of and held stock and/or stock options in Reata at the time the study was conducted.

DATA AVAILABILITY STATEMENT

Individual participant data collected during the trial may be shared after anonymization and upon approval of the research proposal. Biogen commits to sharing patient-level data, study-level data, CSRs, and protocols with qualified scientific researchers who provide a methodologically sound proposal. Biogen reviews all data requests internally based on the review criteria and in accordance with our Clinical Trial Transparency and Data Sharing Policy. Deidentified data and documents will be shared under agreements that further protect against participant reidentification. To request access to data, please visit<https://vivli.org/>.

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SUPPORTING INFORMATION

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