

# Comparing the Liver Safety Profiles of 4 Next-Generation CGRP Receptor Antagonists to the Hepatotoxic CGRP Inhibitor Telcagepant Using Quantitative Systems Toxicology Modeling

Jeffrey L. Woodhead,<sup>\*,1</sup> Scott Q. Siler,<sup>\*</sup> Brett A. Howell,<sup>\*</sup> Paul B. Watkins ,<sup>†</sup> and Charles Conway<sup>‡</sup>

<sup>\*</sup>DILIsym Services, Inc., A Simulations Plus Company, Research Triangle Park, North Carolina 27706, USA;

<sup>†</sup>Institute for Drug Safety Sciences, UNC-Eshelman School of Pharmacy, University of North Carolina, Chapel Hill, North Carolina 27599, USA; and <sup>‡</sup>Biohaven Pharmaceuticals, Inc., New Haven, Connecticut 06510, USA

<sup>1</sup>To whom correspondence should be addressed at DILIsym Services, Inc., A Simulations Plus Company, 6 Davis Drive, Research Triangle Park, NC 27709, USA. E-mail: jeff.woodhead@simulations-plus.com.

## ABSTRACT

Calcitonin gene-related peptide (CGRP) signaling inhibitors have shown efficacy in both the acute and preventive treatment of migraine. Telcagepant, a first-generation CGRP receptor antagonist, was effective but failed in clinical trials due to hepatotoxicity. Subsequently, although 4 next-generation CGRP receptor antagonists (rimegepant, zavegepant, atogepant, and ubrogepant) were being advanced into late-stage clinical trials, due to telcagepant's failure, more confidence in the liver safety of these compounds was needed. DILIsym v6A, a quantitative systems toxicology (QST) model of drug-induced liver injury (DILI), was used to model all 5 compounds and thus to compare the 4 next-generation CGRP receptor antagonists to telcagepant. *In vitro* experiments were performed to measure the potential for each compound to inhibit bile acid transporters, produce oxidative stress, and cause mitochondrial dysfunction. Physiologically based pharmacokinetic models were produced for each compound in order to appropriately estimate liver exposure. DILIsym predicted clinical elevations of liver enzymes and bilirubin for telcagepant, correctly predicting the observed DILI liability of the first-generation compound. By contrast, DILIsym predicted that each of the 4 next-generation compounds would be significantly less likely to cause DILI than telcagepant. Subsequent clinical trials have validated these predictions for each of the 4 compounds, and all 3 of the compounds submitted to FDA to date (rimegepant, ubrogepant, and atogepant) have since been approved by the FDA with no warning for hepatotoxicity. This work demonstrates the potential for QST modeling to prospectively differentiate between hepatotoxic and nonhepatotoxic molecules within the same class.

**Key words:** liver injury; quantitative systems toxicology; biological modeling; pharmaceuticals.

Inhibition of the calcitonin gene-related peptide (CGRP) pathway has been identified as a potential target for migraine treatment and prevention, which would meet a major unmet medical need (Edvinsson, 2018; Holland and Goadsby, 2018).

While several monoclonal antibody treatments targeting CGRP exist (Berman *et al.*, 2020), small molecules present an advantage in that they may be able to engage the CGRP receptors that exist within the brain, as well as can be dosed orally (Mullin *et al.*, 2020).

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Two of the first-generation small molecules in the CGRP receptor antagonist class, telcagepant and MK-3207, failed during late-stage clinical trials due to liver injury (Hewitt et al., 2011; Ho et al., 2016). Several next-generation drugs have been developed since the failure of telcagepant and MK-3207; 3 compounds, rimegepant (Nurtec ODT), ubrogepant (Ubrovelvy), and atogepant (Qulipta), have been approved by the FDA for the treatment of migraine (Al-Hassany et al., 2022).

The identification of liver injury issues with telcagepant and MK-3207 led to the need to investigate potential differences between those 2 drugs and the other CGRP receptor antagonists. Computer modeling, specifically quantitative systems toxicology (QST) modeling, has been investigated as a way of predicting the toxicity potential of a drug (Watkins, 2020). DILIsym, a QST model of drug-induced liver injury (DILI), has demonstrated the ability to differentiate between hepatotoxic and nonhepatotoxic drugs in the same class (Longo et al., 2016; Woodhead et al., 2014) and to identify mechanistic differences among drugs with DILI liabilities within the same class (Woodhead et al., 2019). DILIsym has also been used to prospectively predict the safety of a drug as compared to another drug within that same class that had DILI liabilities (Woodhead et al., 2020). All of these factors make DILIsym a potentially valuable tool for differentiating hepatotoxic from nonhepatotoxic CGRP receptor antagonists.

In this work, DILIsym was used to predict the hepatotoxicity of telcagepant and compare it to 4 next-generation CGRP receptor antagonists: rimegepant, ubrogepant, atogepant, and zavegepant. *In vitro* hepatotoxicity studies were conducted for all 5 compounds according to standard DILIsym procedure (Longo et al., 2019; Woodhead et al., 2019, 2020). It is worth noting that the simulation work for these compounds was conducted in blinded fashion with assessments of DILI potential made before Phase 3 clinical trial results were completed, making these true prospective predictions.

## MATERIALS AND METHODS

DILIsym v6A was used for all hepatotoxicity simulations. Physiologically based pharmacokinetic (PBPK) modeling for telcagepant, rimegepant, and zavegepant were developed in DILIsym v6A; *in silico* PBPK predictions for ubrogepant and atogepant were generated in GastroPlus 9.5.

PBPK modeling was performed to predict the liver concentration-time profile for each compound. Literature data were used to develop the PBPK model for telcagepant (Han et al., 2010) and to estimate certain PBPK parameters. Clinical trial data were used to develop the PBPK model for rimegepant. The PBPK model for zavegepant was based on animal data and *in silico* predictions and included adaptations for different dosing routes. Further information on the PBPK modeling for the 5 compounds in this study is provided in the [Supplementary Materials](#). Most notably, a custom PK SimPops was created for rimegepant based on the range of plasma concentrations observed in the clinic; the underlying mechanistic susceptibility variation in v4A\_1 was still present in this SimPops. Population variability in PK was not included for the other compounds due to the lack of clinical data for zavegepant, atogepant, and ubrogepant and uncertainty around the variability present in the published data for telcagepant. For atogepant and ubrogepant, no PBPK data were available; as a result, the PBPK for these compounds was developed using the machine learning algorithms in ADMET Predictor (SimulationsPlus, Lancaster, California) as included in GastroPlus 9.5 (SimulationsPlus).

*In vitro* experiments were conducted on each of the 5 compounds to assess the effect of each drug on the 3 main hepatotoxicity mechanisms represented in DILIsym v6A: reactive oxidative stress (ROS), mitochondrial dysfunction, and bile acid transporter inhibition. For oxidative stress assessments, HepG2 cells were cultured for 6 or 24 h with varying drug concentrations and ROS formation was assayed by dihydroethidium (DHE) fluorescence. Three independent studies in triplicate were conducted on HepG2 cells. Assays were performed by Cyprotex (Macclesfield, UK). A detailed description of the protocol for each compound is provided in the [Supplementary Materials](#).

LC/MS/MS analysis was performed on a parallel culture of HepG2 cells to assess the intracellular concentration of each compound; cell lysate concentration was assayed and subsequently corrected for lysate volume and cell volume.

Parameter values for telcagepant-mediated induction of oxidative stress were identified by simulating the experimental data in DILIsym using a DILIsym dosing scheme meant to represent *in vitro* conditions (steady-state liver exposure).

Simulation protocols for each compound were chosen based on clinical experience and proposed clinical doses for each compound. The simulations performed were as follows:

- Telcagepant: 140 and 280 mg BID
- Rimegepant: (1) 75 mg QD, 25 days; (2) 75 mg QD, 25 doses delivered as 5 consecutive days, with one-day rest; (3) 75 mg QD, 14 doses given on alternate days
- Ubrogepant: (1) 100 mg QD, 25 days; (2) 200 mg QD, 25 days; (3) 500 mg QD, 25 days; (4) 1000 mg QD, 25 days
- Zavegepant: (1) 75 mg QD, oral (PO), 25 days; (2) 750 mg QD, PO, 25 days; (3) 2 mg QD, intranasal (IN), 25 days; (4) 20 mg QD, IN, 25 days; (5) 0.75 mg QD, subcutaneous (SC), 25 days; (6) 7.5 mg QD, SC, 25 days
- Atogepant: (1) 60 mg BID, 12 weeks; (2) 120 mg BID, 12 weeks; (3) 300 mg BID, 12 weeks; (4) 600 mg BID, 12 weeks

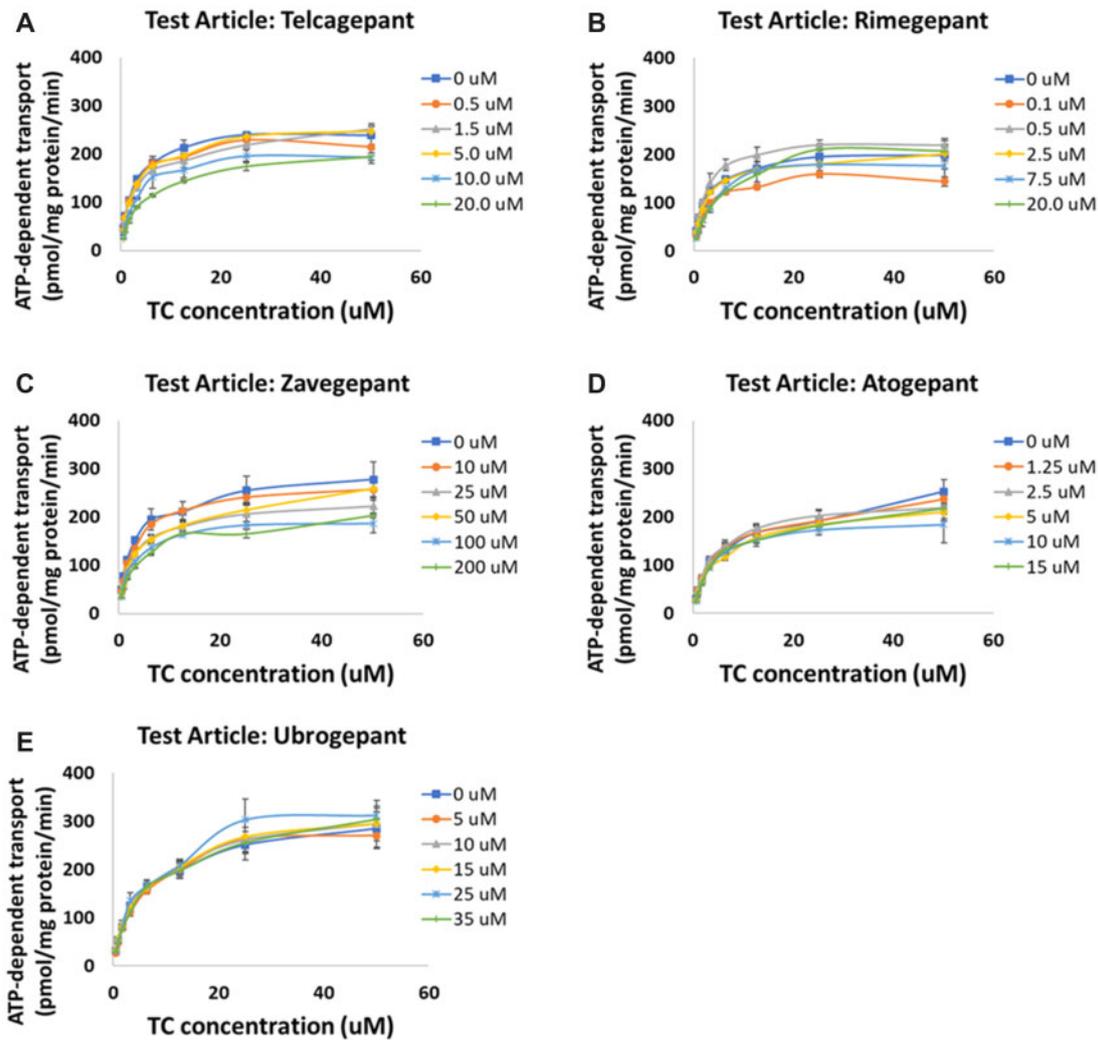
Simulated alanine transaminase (ALT) elevation frequency and severity, as well as several other clinical and mechanistic outputs, were collected for each simulation. Each of the simulation results were compared with clinical experience where applicable (ie, with telcagepant); the other simulation results were treated as a prospective prediction of liver safety.

Mechanistic simulation results were employed as a way of determining the most important mechanism contributing to any predicted ALT elevations for each compound. The general concept of these simulations has been discussed previously (Longo et al., 2019; Woodhead et al., 2017, 2019). Briefly, simulations are performed on a smaller SimCohorts of responder individuals with each potential mechanism of hepatotoxicity turned off in sequence. A decrease in ALT elevation frequency when a mechanism is off demonstrates that the mechanism in question is contributing to the simulated ALT elevations.

## RESULTS

### Results of In Vitro Assays

*In vitro* assays determining the extent of the CGRP signal-blocking compounds' ability to inhibit bile acid transport, induce mitochondrial toxicity, and generate oxidative stress were interpreted and translated into parameters for input into DILIsym v6A. The presence of a parameter for a given mechanism for a compound should not be taken as proof that the mechanism



**Figure 1.** Inhibition of ATP-dependent taurocholate (TC) transport into membrane vesicles mediated by the hepatic bile salt export pump (BSEP) by CGRP receptor antagonist compounds: (A) telcagepant; (B) rimegepant; (C) zavegepant; (D) atogepant; (E) ubrogepant.  $K_i$  studies involving assessment of the transporter inhibition at several different concentrations of taurocholate and the test article were performed, and the data were fit using Michaelis-Menten kinetics to determine the  $K_i$  and inhibition type for each compound. Note: zavegepant top concentration was  $\geq 10\times$  higher than others and it shows negligible BSEP inhibition ( $K_i > 340\mu\text{M}$ , see Table 1).

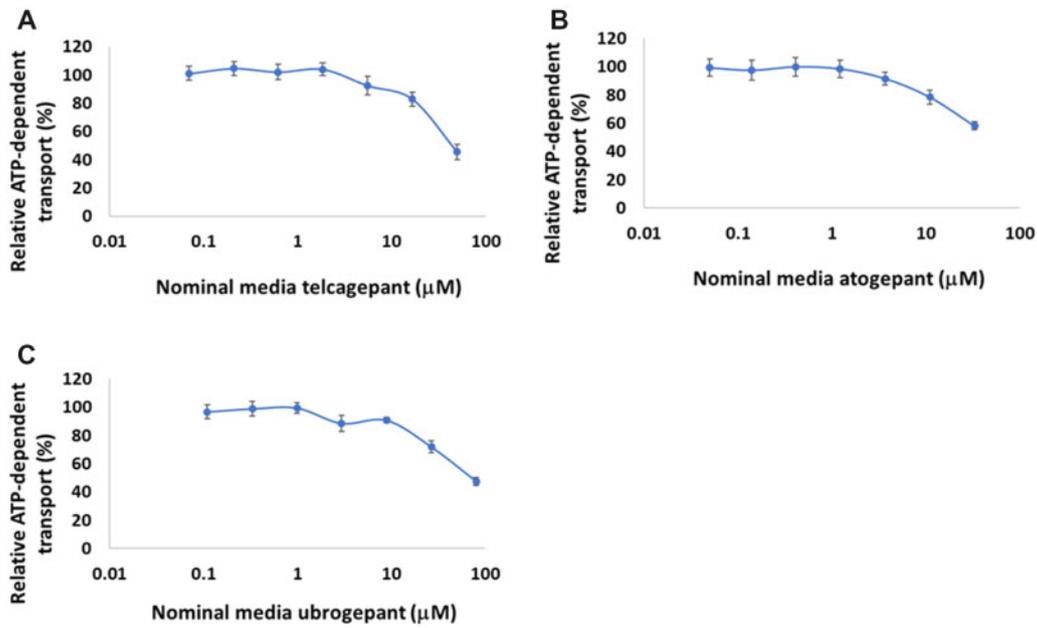
is a potential liability for the compound; the parameters quantify the potential activity of the compound on each mechanism and as such may not be active at the predicted liver concentration range for the compound. DILIsym simulations integrate these parameters with liver exposure predictions to determine the relevance of these mechanisms for potential DILI.

Bile acid transport was inhibited by all compounds to varying degrees. Figure 1 shows the inhibition of the bile salt export pump (BSEP) by each of the CGRP receptor antagonists; Figure 2 shows the inhibition of basolateral bile acid transporters (MRP3 and MRP4). None of the compounds inhibited the sodium taurocholate cotransporting polypeptide (NTCP) (not shown). For each compound, the mode of inhibition for BSEP was assessed; for telcagepant, ubrogepant, and zavegepant, the mode of inhibition was mixed, whereas for rimegepant, the mode of inhibition was competitive. The calculated  $K_i$  values along with the calculated  $\alpha$  values can be found in Table 1. For MRP3/4 and NTCP, the mode of inhibition was assumed to be mixed inhibition with  $\alpha = 5$  (Woodhead et al., 2020).

All the CGRP receptor antagonists except zavegepant inhibited the mitochondrial electron transport chain to

varying extents as well; zavegepant was determined to be a mild uncoupler of the mitochondrial proton gradient. Figure 3 shows the oxygen consumption rate (OCR) versus intracellular compound concentration graphs for each compound and their attendant fits in MITOSym. Each parameter value calculated in MITOSym was translated into a DILIsym parameter (Woodhead et al., 2019; Yang et al., 2014); these parameters are shown in Table 1. For telcagepant, the *in vitro* data lent itself to a range of potential representations rather than to a single representation. As a result, 2 alternate parameterizations were used to bound the potential range of ETC inhibition parameter values that could plausibly represent the *in vitro* data; they are represented as the “high” and “low” parameter sets in Table 1 and Figure 3B. SimPops-based hepatotoxicity simulations were performed with both parameterizations.

Each of the CGRP antagonists also induced oxidative stress *in vitro*. Figure 4 shows the relationship between intracellular compound concentration and ROS production for each compound. These data were fit in DILIsym and the resulting parameters are shown in Table 1.



**Figure 2.** Inhibition of basolateral bile acid transporters by CGRP compounds: (A) inhibition of MRP4 by telcagepant; (B) inhibition of MRP3 by atogepant; (C) inhibition of MRP3 by ubrogepant. Rimegepant and zavegepant did not inhibit basolateral bile acid transporters; telcagepant did not inhibit MRP3; and ubrogepant and atogepant did not inhibit MRP4.

**Table 1.** DILIsym Input Parameters for the CGRP Compounds, Including Parameters Related to Mitochondrial Toxicity, ROS Generation, and Bile Acid Transporter Inhibition

Mechanism	Parameter	Unit	DILIsym Parameter Value <sup>a</sup>					
			Telcagepant— High	Telcagepant— Low	Rimegepant	Zavegepant	Atogepant	Ubrogepant
Mitochondrial dysfunction	Coefficient for ETC inhibition 1	μM	3470	3470	3470	1600	38 170	Not used
	Coefficient for ETC inhibition 3	μM	1.89	Removed	1.89	2	0.1	4,217
	Max inhibitory effect for ETC inhibition 3	Dimensionless	0.45	Removed	0.45	1.5	0.2	0.4
	Uncoupler 1 effect $K_m$	mM	No effect	No effect	No effect		No effect	15 300
	Uncoupler 1 effect $V_{max}$	Dimensionless	No effect	No effect	No effect		No effect	22.5
	Uncoupler 1 effect Hill	Dimensionless	No effect	No effect	No effect		No effect	4.3
	Oxidative stress rate constant 1	ml/nmol/h	$3.5 \times 10^{-4}$	$3.5 \times 10^{-4}$	$3.5 \times 10^{-4}$	No ROS production	$3.41 \times 10^{-4}$	$1.65 \times 10^{-4}$
Bile acid transporter inhibition <sup>b</sup>	BSEP inhibition constant	μM	19.0	19.0	27.2	341	144.2	No inhibition
	BSEP inhibition alpha value	Dimensionless	4.32	4.32	Competitive	1.368	0.64	No inhibition
	NTCP inhibition constant	μM	No inhibition	No inhibition	No inhibition	No inhibition	No inhibition	No inhibition
	MRP4 inhibition constant	μM	42.4	42.4	No inhibition	No inhibition	42	75.3

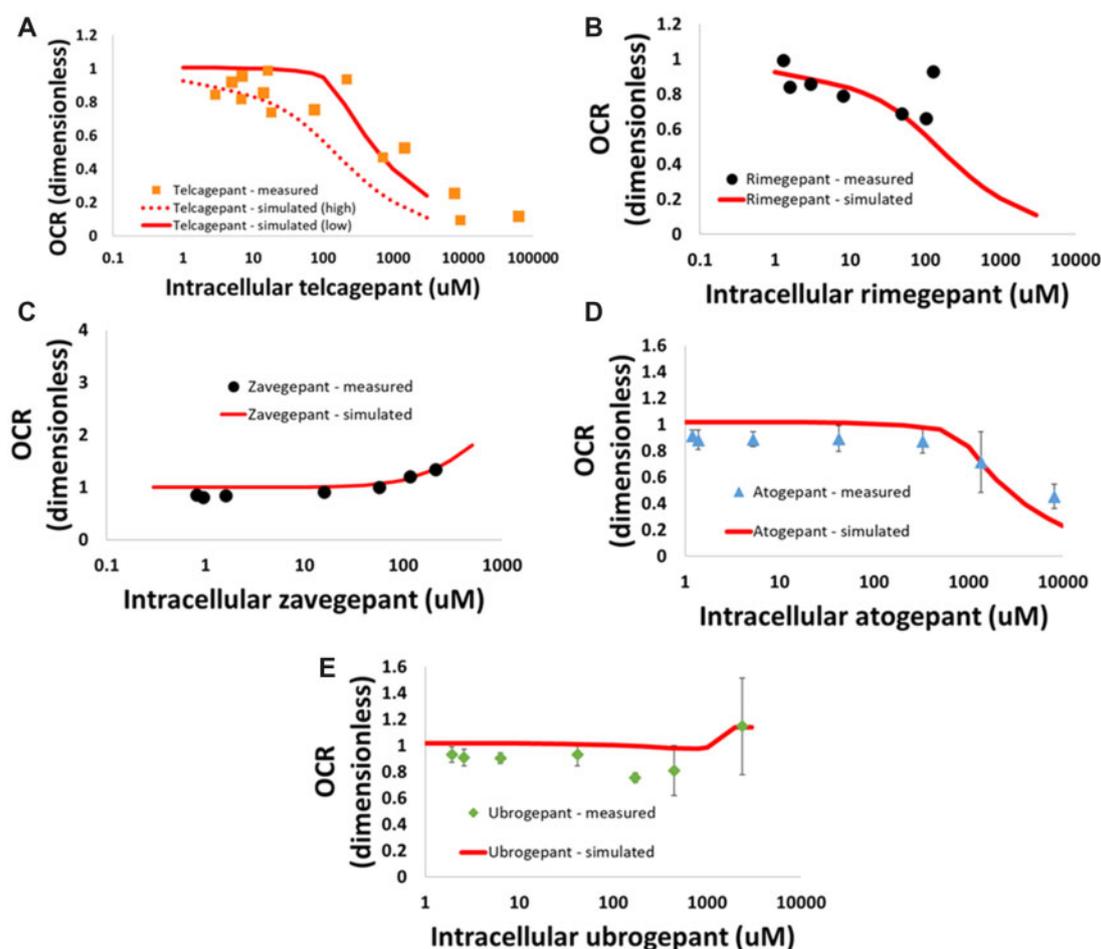
<sup>a</sup>Values shown in the table for DILIsym input parameters should not be interpreted in isolation with respect to clinical implications, but rather, should be combined with exposure in DILIsym to produce simulations that have predictive and insightful value.

<sup>b</sup> $IC_{50}$  values were used for transporters other than BSEP, where  $K_i$  was measured; alpha value of 5 assumed when  $IC_{50}$  measured.

### Simulation Results

Table 2 lists all of the predicted ALT elevation frequencies from the simulations performed in this work. First, DILIsym was used to simulate the telcagepant clinical trials. The 280 mg BID dose

predicted ALT elevations for both parameterizations; ALT elevations were predicted using only the “high” parameterization at the 140 mg BID dose. This is similar to clinical experience, where some ALT elevations occurred at both dosing levels. Given the



**Figure 3.** Oxygen consumption rate (OCR) versus measured intracellular compound concentration after a 24-h incubation in HepG2 cells for each of the CGRP receptor antagonist compounds: (A) telcagepant; (B) rimegepant; (C) zavegepant; (D) atogepant; and (E) ubrogepant. Fits to the data in MITOSym used to parameterize the model are shown along with the *in vitro* data results. For telcagepant, 2 alternate parameterizations were determined to be equally plausible; as a result, both were explored and used in simulations.

uncertainty in the ETC inhibition data and the resulting representation of telcagepant using outer bounds for hepatotoxicity parameter inputs, an underprediction using one parameterization and an overprediction using the other parameterization would naturally be expected. Notably, the simulations do not include adaptation mechanisms, potentially including mitochondrial biogenesis, that would likely reduce the severity of some of the simulated liver injury; this is why the number of Hy's Law cases with telcagepant treatment is overestimated by the simulations (Figure 5).

Rimegepant and zavegepant were both predicted to be relatively safe compared with telcagepant. ALT elevations with rimegepant were simulated in some extreme dosing scenarios, especially when the high-PK representation was used; however, as is seen in the eDISH plots of the simulation results (Figure 5), these were generally mild compared with the severe ALT elevations and Hy's Law cases simulated with telcagepant (Figure 5). For zavegepant, no ALT elevations were predicted at any dose, even those well above the proposed therapeutic doses (Table 2, not shown in Figure 5 because all individuals were in the normal range on the eDISH plot).

Both atogepant and ubrogepant also compared favorably to telcagepant. ALT elevations with both compounds were not simulated at the proposed clinical doses, and only appeared at doses 10-fold higher than the clinical doses (Table 2).

Mechanistic simulation results show that telcagepant hepatotoxicity is driven by ETC inhibition and bile acid transporter inhibition (Table 3). This is distinct from rimegepant, whose predicted minor and mild ALT elevations in the high dosing frequency scenarios were caused by ETC inhibition alone, with only mild contributions from bile acid transporter inhibition and ROS generation (Table 3). Ubrogepant and atogepant simulated ALT elevations at the supratherapeutic doses were largely ROS dependent. For zavegepant, no simulated ALT elevations were observed, so mechanistic simulations were unable to be performed.

## DISCUSSION AND CONCLUSIONS

Telcagepant failed in the clinic due to severe liver injury concerns, and DILISym simulations conducted herein correctly predicted the hepatotoxicity liability of telcagepant. Taken together, the simulation results in this work predicted that the 4 next-generation CGRP receptor antagonists would be significantly safer than telcagepant with respect to liver injury. Neither rimegepant, ubrogepant, atogepant, nor zavegepant were predicted to cause severe liver injury (ie, Hy's Law cases) at the tested clinical dose regimens, even when the worst-case scenarios for clinical exposure were considered. At the actual (rimegepant, atogepant, ubrogepant) or anticipated (zavegepant) FDA recommended

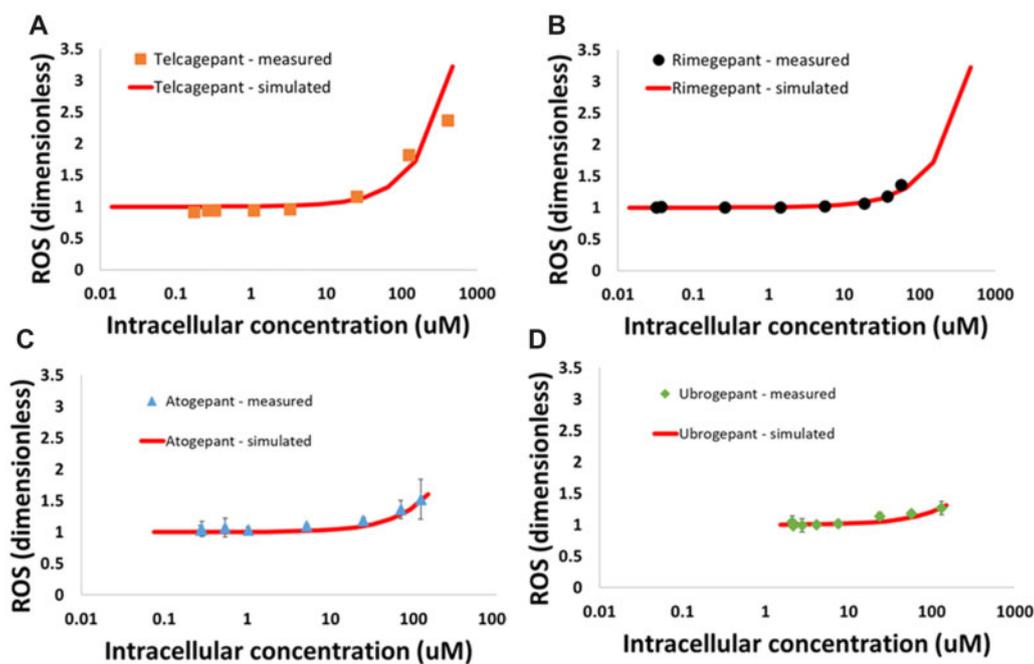


Figure 4. Reactive oxygen species (ROS) as measured by dihydroethidium (DHE) staining in HepG2 cells after a 24-h incubation with CGRP compounds: (A) telcagepant; (B) rimegepant; (C) atogepant; (D) ubrogapant. Zavegepant did not cause an increase in ROS in the experimental system. Fits to the data in DILIsym used to parameterize the model are also shown.

Table 2. Simulated ALT Elevations in the v4A\_1 SimPops for Each of the CGRP Compounds

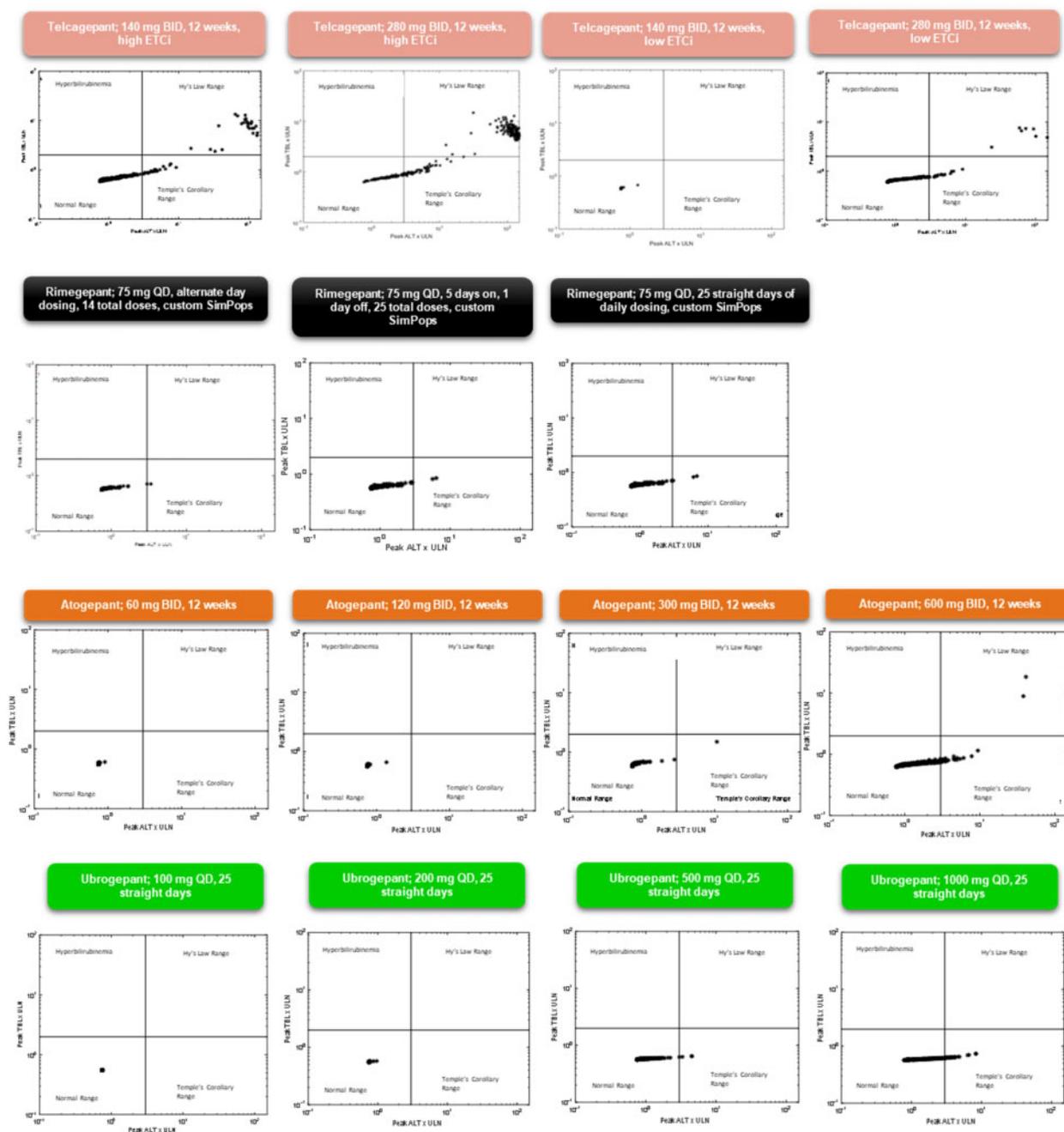
Compound	Oral Dosing Protocol	Simulated ALT > 3X ULN <sup>a</sup>	Observed ALT > 3X ULN in Clinic
Telcagepant—High ETC	140 mg BID, 12 weeks	17.5% (50/285)	1.9% (5/263)
	280 mg BID, 12 weeks	76.1% (217/285)	3.2% (8/265)
Telcagepant—Low ETC	140 mg BID, 12 weeks	0.0% (0/285)	1.9% (5/263)
	280 mg BID, 12 weeks	7.72% (22/285)	3.2% (8/265)
Rimegepant	75 mg QD, alternate day dosing, 14 total doses	0.35% (1/285)	—
	75 mg QD, 5 days on, 1 day off, 25 total doses	0.7% (2/285)	—
	75 mg QD, daily dosing for 25 days, 25 total doses	1% (3/285)	—
Zavegepant	750 mg oral QD, 25 days, 25 total doses	0.0% (0/285)	—
	75 mg oral QD, 25 days, 25 total doses	0.0% (0/285)	—
	20 mg IN QD, 25 days, 25 total doses	0.0% (0/285)	—
	2 mg IN QD, 25 days, 25 total doses	0.0% (0/285)	—
	0.75 mg IV QD, 25 days, 25 total doses	0.0% (0/285)	—
	7.5 mg IV QD, 25 days, 25 total doses	0.0% (0/285)	—
Atogepant	60 mg BID, 12 weeks	0% (0/285)	—
	120 mg BID, 12 weeks	0% (0/285)	—
	300 mg BID, 12 weeks	0.3% (1/285)	—
	600 mg BID, 12 weeks	10.2% (29/285)	—
Ubrogapant	100 mg QD, 15 days	0% (0/285)	—
	200 mg QD, 15 days	0% (0/285)	—
	500 mg QD, 15 days	1.1% (3/285)	—
	1000 mg QD, 15 days	11.6% (33/285)	—
	100 mg QD, 25 days	0% (0/285)	—
	200 mg QD, 25 days	0% (0/285)	—
	500 mg QD, 25 days	1.4% (4/285)	—
	1000 mg QD, 25 days	11.6% (33/285)	—

The alternate parameterizations for telcagepant are included in the table; comparisons to clinical data were only available for telcagepant at the time the simulations were conducted and are included in the table as well.

<sup>a</sup>Upper limit of normal (ULN) for ALT in DILIsym is 40 U/L.

doses and regimens, the vast majority of individuals fall into the normal range for simulated ALT elevations, with for rimegepant a few individuals passing just beyond the normal ALT threshold but with no bilirubin elevations.

Mechanistic simulation results also underscore the difference between telcagepant and the other CGRP receptor antagonist compounds. Telcagepant hepatotoxicity was driven by a combination of ETC inhibition and bile acid transporter



**Figure 5.** eDISH plots demonstrating the severity of simulated ALT elevations for each of the CGRP compounds for which toxicity was predicted at indicated dosing regimens. eDISH plots show ALT fold change on the x-axis and bilirubin fold change on the y-axis. Individuals in the bottom-right quadrant have ALT elevations but no bilirubin elevations; individuals in the top-right quadrant have both ALT and bilirubin elevations and are considered most at risk for developing severe DILI (Hy's Law). The simulations do not include several adaptation mechanisms that would likely reduce the severity of the ALT elevations (though not the frequency).

inhibition; this combination of mechanisms is often responsible for predicted serious hepatotoxicity in DILIsym (Longo et al., 2019; Woodhead et al., 2017). Meanwhile, the other compounds were shown to have only a single mechanism—either ETC inhibition or ROS formation—as their biggest potential liability; this suggests that an adaptive response to ROS that is not included in DILIsym or mitochondrial biogenesis could prevent serious injury from occurring from these other, newer CGRP receptor antagonists. The bile acid transporter mechanism is especially

interesting; whereas rimegepant, atogepant, and ubrogapant all showed signals in a BA transporter  $IC_{50}$  assay, further exploration in the *in vitro* systems and with DILIsym revealed several reasons why those compounds do not have bile acid transport inhibition liabilities; for example, rimegepant is a competitive inhibitor of BSEP whereas telcagepant is a mixed inhibitor of BSEP, and for most individuals the concentration of rimegepant in the liver is well below the measured  $K_i$  while telcagepant liver concentration is well above the measured  $K_i$ . The clinical

**Table 3.** Mechanistic Investigation Simulations for Each of the CGRP Compounds Focused on the Select Specific Scenarios That Did Produce ALT Elevations (ie, to Identify When ALT Elevations Occur, What Underlies It)

Compound and Oral Protocol	Mechanism Off	Mechanisms On	Simulated ALT > 3× ULN
Telcagepant—Original ETC, 140 mg BID, 12 weeks	None	BAi; ETCi; ROS	50/50
	BAi	ETCi; ROS	21/50
	ETCi	BAi; ROS	0/50
	ROS	BAi; ETCi	46/50
Rimegepant, high PK individual, 75 mg QD, daily dosing for 25 days, 25 total doses	None	BAi; ETCi; ROS	21/21
	BAi	ETCi; ROS	14/21
	ETCi	BAi; ROS	0/21
	ROS	BAi; ETCi	18/21
Atogepant, 600 mg BID, 12 weeks	None	All	29/29
	ROS	BAi, ETCi	2/29
	ETCi	BAi, ROS	24/29
	BAi	ETCi, ROS	24/29
Ubrogepant, 1000 mg QD, 25 days	None	All	33/33
	ROS	BAi, ETCi, UC	0/33
	ETCi	BAi, UC, ROS	33/33
	UC	BAi, ETCi, ROS	33/33
	ETCi, UC	BAi, ROS	33/33
	BAi	ETCi, UC, ROS	33/33

Mechanisms were turned off and on to determine which mechanism contributed the most to the observed ALT elevations. Only those individuals who developed ALT elevations in the initial 285-individual SimPops simulation were used in the mechanistic investigation simulations.

Abbreviations: BAi, bile acid transporter inhibition; ETCi, electron transport chain inhibition; UC, mitochondrial proton gradient uncoupling; ROS, reactive oxygen species generation.

importance is 2-fold: (1) a competitive mode of inhibition (rimegepant) can be overcome as bile salt acid levels increase and BSEP continues to provide functional export of bile salts, whereas mixed mode (telcagepant), which includes noncompetitive inhibition, cannot be similarly overcome which leads to the build up of bile salts associated with liver injury; and (2) for most individuals, clinical liver concentrations at  $C_{max}$  are well above the BSEP  $K_i$  (telcagepant) and indicates ongoing inhibition and reduction of the majority of bile salt export function, whereas clinical liver concentrations at  $C_{max}$  for most individuals fall well below the BSEP  $K_i$  (rimegepant) which indicates that the majority of bile salt export function remains intact.

The clinical experience with these 4 next-generation CGRP signal-blocking drugs has been largely in line with DILIsym predictions for these compounds. Rimegepant, ubrogepant, and atogepant have all demonstrated clinical efficacy and safety (Al-Hassany et al., 2022) and have since been approved by the FDA for the treatment of migraine (acute for ubrogepant, preventive for atogepant, and dual-therapy acute and preventive for rimegepant). Meanwhile, clinical trial results for zavegepant which continues to advance in clinical trials for the acute (nasal) and preventive (oral) treatment of migraine without a liver safety signal (Al-Hassany et al., 2022; Goadsby et al., 2020; Moreno-Ajona et al., 2020). These results suggest that the QST approach demonstrated by DILIsym can lead to increased confidence in the ability to differentiate between liver safety profiles of drugs in the same class.

In a previous paper, DILIsym was used to compare telcagepant and ubrogepant to MK-3207, another CGRP receptor antagonist that failed in clinical trials due to liver injury. Telcagepant and MK-3207 were predicted to be hepatotoxic by DILIsym in that work, whereas ubrogepant was predicted to be safe (Yamazaki et al., 2013). These simulations were performed independently from the simulations in this work, with a set of *in vitro* hepatotoxicity assays conducted separately from what was conducted for this report. Although some minor differences between the results from the 2 simulation projects exist, the

striking qualitative similarity between the predictions for telcagepant and ubrogepant (ie, that telcagepant could cause severe hepatotoxicity while ubrogepant would be safe) between the 2 independently conducted simulation projects demonstrates the robustness of the DILIsym QST approach.

In summary, DILIsym prospectively predicted improved liver safety of the 4 next-generation CGRP receptor antagonists rimegepant, zavegepant, atogepant, and ubrogepant relative to the hepatotoxic first-generation molecule telcagepant, and these predictions have been born out in the clinical trials conducted to date. It is worth noting that the same general methods employed in this manuscript have been used to compare hepatotoxic potential among drugs within other compound classes as well (Woodhead et al., 2019, 2020). Our results support the value of QST modeling in drug development.

## SUPPLEMENTARY DATA

Supplementary data are available at Toxicological Sciences online.

## FUNDING

Biohaven, Inc. provided funding for this research.

## DECLARATION OF CONFLICTING INTERESTS

Biohaven Pharmaceuticals is the manufacturer of rimegepant and zavegepant, 2 of the compounds analyzed in this paper. C.C. is employed by Biohaven.

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