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# **Population Pharmacokinetic Analysis of Ensitrelvir, an Inhibitor of 3C-like (3CL) Protease of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) for** patients with SARS-CoV-2 Infection

### **Introduction and Purpose**

- Ensitrelvir is a novel oral inhibitor of 3CL protease of SARS-CoV-2, which is essential for viral replication.
- Ensitrelvir was approved in Japan for the treatment SARS-CoV-2 infection under the Emergency Regulatory Approval System in November 2022.
- Ensitrelvir is administered once daily for 5 days with 375 mg on Day 1 followed by 125 mg.
- The aim of this study is to build a population pharmacokinetic (PK) model to assess covariates on ensitrelvir PK.

# Methods

#### 1) Population PK Analysis

Population PK analysis was performed by using NONMEM ver. 7.4 (ICON Development Solutions, US) with stepwise covariate model (SCM) building tool in Perl-speaks NONMEM ver. 4.9 (developed by Rikard Nordgren).

**Data:** A total of 8034 plasma ensitrelyir concentrations from 2060 healthy participants in Phase 1 studies and patients in Phase 2/3 study were obtained and used for the analysis.

#### **Table.1 Background Characteristics of Participants**

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Study	<b>Background characteristics</b>	Mean (SD)	Medi	
Overall	Body weight (kg)	63.8 (13.6)	62.6 (3	
(N = 2060,	BMI (kg/m²)	23.0 (3.8)	22.5	
8034 points)	Age (years)	36.5 (12.8)	35	
	ALT (U/L)	26.3 (23.4)	19	
[ Phase 1 study	AST (U/L)	25.4 (13.7)	22 (	
(n=175, 4341	BIL (mg/dL)	0.5 (0.3)	0.5	
points)	ALB (g/dL)	4.4 (0.3)	4.4	
Phase 2/3	CrCL (mL/min)	115.3 (30.7)	111.9 (	
study (n=1885,	Scr (mg/dL)	0.76 (0.17)	0.76 (	
3693 points) ]	eGFR (mL/min/1.73 m <sup>2</sup> )	88.1 (19.8)	85.9 (3	
	eGFRabs (mL/min)	84.3 (20.3)	82.5 (3	
	Formulation (Suspension : Tablet) *	62 (3.0%) :	1998 (97	
	Sex (Male : Female) *	1183 (57.4%	b):877 (4	
	Adolescent		2026 (00	
	(12 to < 18 years : >= 18 years) *	34 (1.7%) : 2026		
	Race (Asian : White : Others) *	2030 (98.5%) : 2	25 (1.2%)	
	Country	1/0/ (72 00/) - 120	) (6 20/ ) •	
	(Japan : Korea : Vietnam) *	1484 (72.0%) : 130	) (0.5%) .	
	Health status (Healthy : Infected by SARS-CoV-2) *	175 (8.5%)	: 1885 (9	
ALB = albumin: AI T =	= alanine aminotransferase: AST = aspartate amin	otransferase: BIL = total b	ilirubin: BMI	

index; CrCL = creatinine clearance; eGFR = estimated glomerular filtration rate; eGFRabs = absolute estimated glomerular filtration rate; Scr = serum creatinine

Number of participants (percentage of all participants)

#### **Base model:**

- Structural PK model: Two-compartment model with first-order elimination and first-order absorption.
- Inter-individual variability: exponential error model
- Intra-individual variability: proportional and additive error model

**Covariate model:** Constructed full model by SCM with a forward selection (p<0.01) and inferential assessment. Then assessed the selected covariates with a backward selection (p<0.005). Finally tested omega block and inferential assessment.



**Candidates of covariate** 

- CL/F: body weight, age, adolescent or not, sex, race, country, ALT, ALB, total bilirubin, creatinine clearance, serum clearance, patient or not
- Vc/F: body weight, age, adolescent or not, sex, race, country, ALB, patient
- or not Absorption rate constant (ka): fasted or fed, formulation
- Bioavailability (F1): fasted or fed, formulation, patient or not

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### dian (range)

- (35.0 156.0) 5 (7.0 - 49.8) 5 (12 - 76) 9 (0 - 349) (10 - 272) 5(0.1 - 2.1)(0.5 - 5.8)
- (46.0 354.6) (0.37 - 1.43) (36.4 - 242.9)
- (31.5 278.8) 97.0%)
- (42.6%)
- 98.4%)
- ):5(0.2%)
- : 446 (21.7%)
- 91.5%)
- SMI = body mas

**Model evaluation:** The final model was evaluated by Diagnostic plots, visual predictive check (VPC) and bootstrapping.

#### 2) Post-hoc Exposure Parameters in Patients

Exposure parameters (C<sub>max</sub>, plasma concentration 24 hours post-dose [C<sub>24</sub>] and AUC) on Days 1 and 5 were estimated by Bayesian approach with the final population PK model when ensitrelvir was administered at once daily for 5 days with 375 mg on Day 1 followed by 125 mg (125 mg group) or at once daily for 5 days with 750 mg on Day 1 followed by 250 mg (250 mg group) in Phase 2/3 Study.

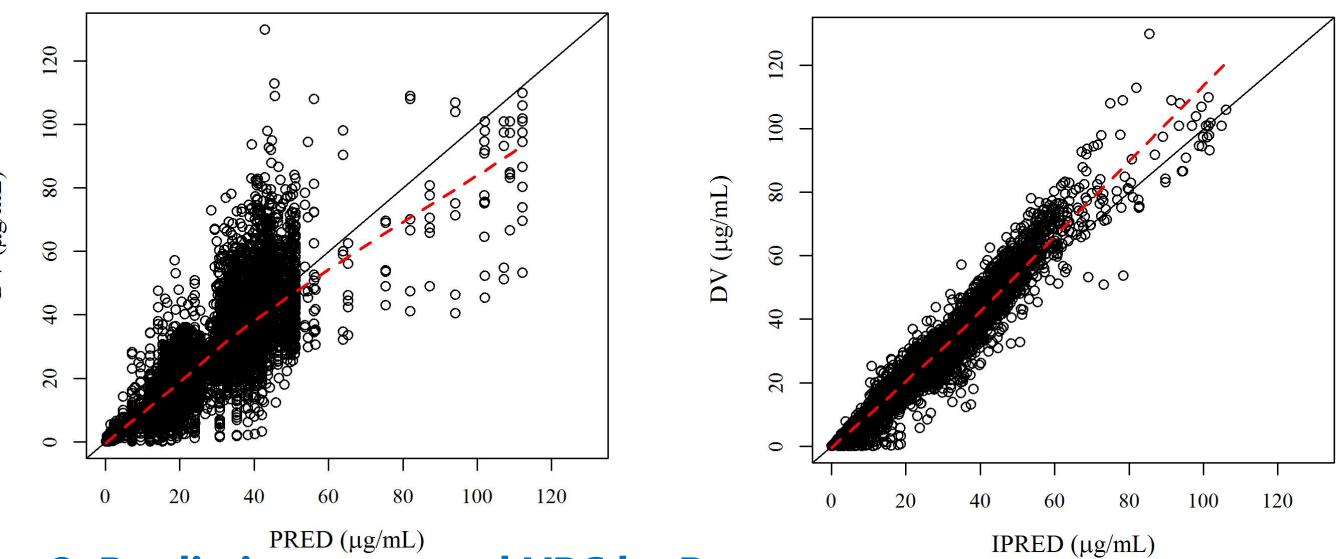
#### 3) PK/Pharmacodynamic (PD) Relationship

PK/PD relationship between change from baseline of SARS-CoV-2 viral RNA load on Day 4 and  $C_{24}$  on Day 1 were assessed in Phase 3 part of Phase 2/3 study. The  $C_{24}$  was categorized by 0 to <10, 10 to < 20, 20 to < 30, 30 to < 40 and  $\geq$  40 µg/mL.

# **Results: 1) Population PK Analysis**

• The final model included the effect of body weight on CL/F, the effect of body weight on Vc/F, the effects of food and formulation on Ka. The final population PK parameters and the results of bootstrapping (rate: 90.2%) are shown in Table 2. Diagnostic plots (goodness-of-fit) are shown in Fig.1. Prediction-corrected VPC by race (Asian/White) for data following multiple doses in Phase 1 and 2/3 studies are shown in Fig.2, indicating good predictive performance of the final model.

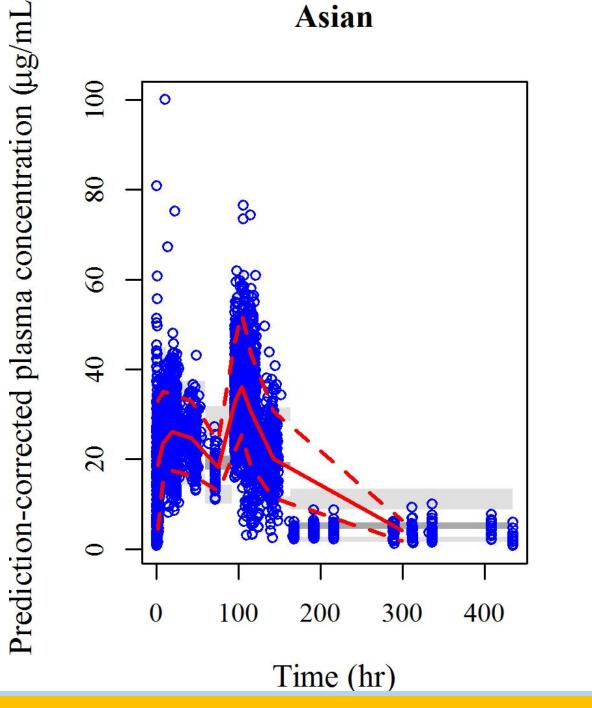
# Fig. 1. Goodness-of-fit Plots for Final Model



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# Fig. 2. Prediction-corrected VPC by Race

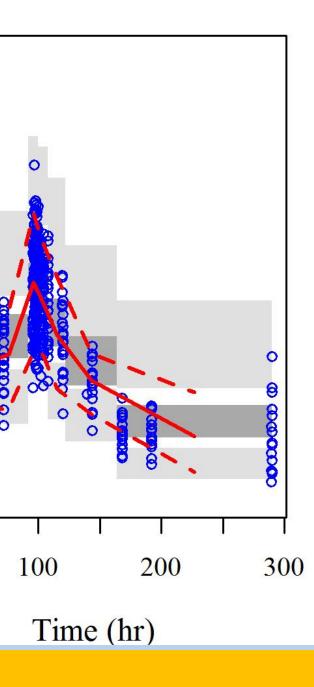


### Conclusion

The population PK model was developed based on the plasma ensitrelvir concentration data from participants including patients infected with SARS-CoV-2, and body weight was the most important covariate on ensitrelvir PK. There was no difference of exposure and dose for anti-viral effect, supporting the selection of the clinical dose (Loading dose 375 mg on Day 1/maintenance dose 125 mg on Days 2-5) regardless body weight.

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# Table.2 Population PK Parameters of Ensitrelvir and Results of Bootstrap

		Final Model					Boot	strap esti	ima	tes
	11		95% Cls			Median	. 95%	95% Cls (lower - upper)		
Pharmacokinetic parameters	Units	Estimates	(lower - upper)			%RSE	S			(lower
Ka	(1/hr)	1.50	1.25	_	1.75	8.6	1.50	1.23	-	1.80
CL/F	(L/hr)	0.211	0.208	-	0.214	0.8	0.211	0.207	-	0.214
Vc/F	(L)	14.7	13.9	-	15.5	2.7	14.8	13.9	-	15.5
Q/F	(L/hr)	0.539	0.321	-	0.757	20.6	0.534	0.331	-	0.823
Vp/F	(L)	2.50	1.80	-	3.20	14.3	2.49	1.74	-	3.26
Effect of food on Ka		0.594	0.398	-	0.790	16.8	0.599	0.447	-	1.00
Effect of formulation on Ka		0.362	0.262	-	0.462	14.1	0.367	0.280	-	0.469
Effect of body weight on CL/F		0.521	0.456	-	0.586	6.4	0.527	0.459	-	0.585
Effect of body weight on Vc/F		1.04	0.960	-	1.12	3.9	1.04	0.959	-	1.11
Inter-individual variability										
Ka	%	72.9	64.3	-	80.6	11.3	72.1	64.4	-	80.4
CL/F	%	21.3	19.7	-	22.8	7.4	21.3	20.0	-	23.0
Covariance between CL/F and Vc/F		0.0216 (R = 0.691)	0.0173	-	0.0259	10.0	0.0217	0.0179	_	0.0263
Vc/F	%	14.7	12.8		16.3	12.3	14.6	12.7	-	16.5
Intra-individual variability										
Proportional residual error	%	19.9	18.5		21.3	3.6	19.8	18.6	-	21.2
Additive residual error	(µg/mL)	0.0317	0.0224	-	0.0410	14.9	0.0331	0.00945	-	0.0570
CI = confidence interval; R = coe	efficient of	correlation; %RSE = ı	relative s	tan	dard erro	or in perc	cent			
Ka = $1.50 \times (0.594 \text{ for food}) \times$ CL/F = $0.211 \times (body weight/62)$ Vc/F = $14.7 \times (body weight/62)$ Q/F = $0.539$	2.6) <sup>0.521</sup>	formulation)						and \ on on		
	stration wi	thin 2 hours after a m	eal, 0 = c	othe	er; formu	lation: 1	= tablet, (	) = suspei	nsic	n]

# **Results: 2) Post-hoc Exposure Parameters in Patients** Table.3 Post-hoc Exposure Parameters in Patients of Phase 2/3 Study

Dose	Day		Day 1			Day 5	
	Parameters	C <sub>max</sub> (µg/mL)	C <sub>24</sub> (µg/mL)	AUC (µg.hr/mL)	C <sub>max</sub> (µg/mL)	C <sub>24</sub> (µg/mL)	AUC (µg.hr/mL)
125 mg Group	n	943	943	943	925	925	925
	Mean	23.4	16.6	437.5	26.6	19.5	578.5
	SD	5.00	3.04	96.53	4.87	3.48	116.7
	CV (%)	21.3	18.3	22.1	18.3	17.8	20.2
250 mg Group	n	942	942	942	920	920	920
	Mean	48.9	34.8	909.5	56.4	41.8	1233
	SD	10.7	6.70	205.1	11.00	8.05	269.5
	CV (%)	21.9	19.2	22.6	19.6	19.3	21.9

# **Results: 3) PK/PD Relationship**

# Fig. 3. Relationship between Change from Baseline of SARS-Cov-2 Virus RNA on Day 4 and C<sub>24</sub> on Day 1

