Integrated Population Pharmacokinetic Analysis of Conjugated and **Unconjugated Payload of Patritumab Deruxtecan in Cancer Patients**

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OBJECTIVES

1. Develop a PopPK model to characterize the PK of anti-HER3-ac-DXd and DXd. 2. Quantify the effects of baseline demographic, laboratory, and disease characteristics on PK in cancer patients.

CONCLUSIONS

- The integrated PopPK model characterized the PK of two analytes in cancer patients treated with Patritumab deruxtecan (HER3-DXd, U3-1402).
- Covariates influencing exposures were identified. Lower baseline eGFR and albumin tended to be associated with lower exposure (Cavgss anti-HER3-ac-DXd) but were still within range of 95% CI of the median projected population exposures.
- The availability of the model will allow sparse sampling in future Phase 3 trials while still characterizing the PK profile.
- This evaluation supports the use of derived individual exposure metrics in exposure response evaluation.

METHODS

 The PopPK analysis included data from 2 phase I/II studies in breast cancer and non-small cell lung cancer (U31402-A-J101 and U31402-A-U102, respectively) treated over the dose range of 1.6 to 8.0 mg/kg administered via intravenous infusion across different dosing regimens.

Table 1. Study Summaries

Study	Dosing Regimen	Objectives
U31402-A-J101	Dose Escalation Part:	• To assess safety and tolerability of U3-1402, to determine the
A Phase 1/2, multicenter, open-label, multiple dose, first-in-human study of patritumab deruxtecan in subjects with HER3-positive metastatic BC.	Q3W IV dosing regimens of:	maximum tolerated dose (MTD) of U3-1402 and to determine the
	1.6 mg/kg, 3.2 mg/kg, 4.8 mg/kg, 6.4 mg/kg, and	The primery objective for deep expansion part was to espece
	8.0 mg/kg	 The primary objective for dose expansion part was to assess safety and evaluate efficacy of U3-1402 at the RDEs in subjects
	Dose Finding Part: 3 cycles of 4.2 mg/kg every 2 weeks, followed by cycles of 6.4 mg/kg Q3W, or	with human epidermal growth factor receptor 2 (HER2)-negative and hormone receptor (HR)-positive breast cancer
	Up-titration regimen of Q3W cycles comprising Cycle 1: 3.2 mg/kg, Cycle 2: 4.8 mg/kg, and Cycle 3 and thereafter: 6.4 mg/kg	
	Dose Expansion Part:	
	4.8 mg/kg or 6.4 mg/kg Q3W IV	
U31402-A-U102 A multicenter, open-label, Phase 1 study of	Dose Escalation Part: 3.2, 4.8, 5.6, or 6.4 mg/kg Q3W IV	 The primary objectives for dose escalation part was to assess the safety and tolerability of U3-1402 in metastatic or unresectable

- The PopPK analysis included 6.869 anti-HER3-ac-DXd and 6.821 DXd observations after IV infusion of HER3-DXd in 401 subjects.
- 182 subjects had BC; 219 with NSCLC.
- Median body weight was 61 kg and median age of 59 years.

Model Development

 The PopPK analysis was conducted via nonlinear mixed-effects modeling using NONMEM software (Version 7.5).

PMX617

- The integrated model characterizes the PK of both anti-HER3-ac-DXd and DXd,
- Integrated model development proceeded as a sequential process:
- 1. Base model for anti-HER3-ac-DXd was developed
- 2. Integrated base model for anti-HER3-ac-DXd and DXd was developed (all parameters estimated simultaneously)
- 3. Covariate effects were estimated for anti-HER3-ac-DXd parameters first, and then for all anti-HER3-ac-DXd and DXd parameters, simultaneously
- 4. Model was evaluated based on objective function value (OFV), plausibility and precision of parameter estimates, and diagnostic plots
- Covariates for anti-HER3-ac-DXd were estimated on clearance and volume terms
- Covariates for DXd were estimated on fractions of formation, since distinguishing effects on systemic PK parameters for DXd and anti-HER3-ac-DXd to DXd formation was not identifiable
- Pre-specified covariates included: hepatic impairment (based on NCIODWG criteria: mild and moderate vs normal), renal function (eGFR), sex, race, weight, age, serum albumin, ECOG performance status (>0 vs 0), tumor type (BC vs NSCLC), and tumor size.
- A full model was implemented providing parameter estimates and confidence interval and the probability distribution without selecting parameters based on a hypothesis testing statistical cutoff

Model Evaluation

INTRODUCTION

• Patritumab deruxtecan (HER3-DXd) is a human epidermal growth factor receptor (HER3)-targeting antibody-drug conjugate with a topoisomerase I inhibitor payload (DXd). The drug-to-antibody ratio is approximately 8. HER3-DXd is currently being investigated as an anticancer agent in several phase 1-3 clinical studies in breast cancer (BC) and nonsmall cell lung cancer (NSCLC) patients.

patritumab deruxtecan in subjects with

metastatic or unresectable NSCLC

Dose Expansion Part: 5.6 mg/kg Q3W IV or up-titration regimen of Q3W cycles of Cycle 1: 3.2 mg/kg, Cycle 2: 4.8 mg/kg, and Cycle 3 and thereafter: 6.4 mg/kg

NSCLC subjects, and to determine the recommended dose for expansion (RDE) of U3-1402 in metastatic or unresectable EGFRm NSCLC subjects who (a) are T790M mutation-negative after disease progression during treatment with erlotinib, gefitinib, or afatinib or (b) develop disease progression while on osimertinib

 The primary objective for dose expansion part was to investigate the antitumor activity of U3-1402

- 500 Monte Carlo simulation replicates of the analysis data were generated using the final PPK model.
- Plots of observed anti-HER3-ac-DXd and DXd were overlaid were corresponding median, 5th and 95th percentiles and stratified by tumor types.

RESULTS

able 2. Summary of continuous baseline covariates by study							
	n	Mean	Median	SD	Min / Max	•	
tudy U31402-J101						ι	
Age (vears)	182	55.6	57.0	12.5	30.0 / 83.0		
Weight (kg)	182	56.7	54.0	13.9	35.1 / 110		
Body surface area (m ²)	182	1.55	1.53	0.181	1.22 / 2.25		
Estimated glomular filtration (eGFR) (mL/min/1.73m ²)	182	98.4	99.8	17.7	37.8 / 134		
Creatinine clearance (mL/min)	182	91.9	89.8	30.1	35.7 / 192		
Albumin (g/L)	182	38.2	39.0	5.10	17.0 / 49.0		
Baseline SLD (cm)	182	7.81	6.55	4.98	1.00 / 31.0		
Baseline H-score	49	157	140	64.8	60.0 / 300		
tudy U31402-U102						•	
Age (years)	219	61.7	63.0	9.93	29.0 / 80.0	(
Weight (kg)	219	64.2	62.0	15.0	32.4 / 113		
Body surface area (m²)	215	1.69	1.66	0.222	1.19 / 2.37		
Estimated glomular filtration (eGFR) (mL/min/1.73m ²)	219	82.1	85.4	18.8	31.9 / 134		
Creatinine clearance (mL/min)	213	76.8	75.4	25.9	30.2 / 174		
Albumin (g/L)	218	38.7	39.0	4.61	22.0 / 46.0		
Baseline SLD (cm)	219	6.31	5.60	3.87	1.00 / 20.5		
Baseline H-score	183	155	170	72.3	0.00 / 300		
II data							
Age (years)	401	58.9	60.0	11.6	29.0 / 83.0	CL_i	
Weight (kg)	401	60.8	57.4	15.0	32.4 / 113	Clim	
Body surface area (m ²)	397	1.63	1.59	0.215	1.19 / 2.37		
Estimated glomular filtration (eGFR) (mL/min/1.73m ²)	401	89.5	91.6	20.0	31.9 / 134	γ: γ:	
Creatinine clearance (mL/min)	395	83.8	80.1	28.9	30.2 / 192	•	
Albumin (g/L)	400	38.5	39.0	4.84	17.0 / 49.0		
Baseline SLD (cm)	401	6.99	6.10	4.46	1.00 / 31.0	Ì	
Baseline H-score	232	155	166	70.6	0.00 / 300	1	

n: number of records summarized; SD: standard deviation; Min: minimum; Max: maximum; Source code: ppk-eda-tables.R; Source file: pk-cont.tex.

Table 3. Summary of categorical covariates by study in PPK analysis

ortions of Total Clearance of anti-HER3-ac-DXd and DXd parate clearance pathways were needed to describe the dynamics of HER3-ac-DXd exposure and can be explained by mechanistic erstanding of ADC disposition and elimination:

- The nonspecific clearance (CLns) pathway was identified as the major clearance pathway and described the mechanism of antibody turnover.
- The CLt pathway was used to describe the initial rapid decrease of clearance within the first cycle, when the antibody quickly binds to the receptors.
- The CLmm clearance pathway mainly affected PK at low anti-HER3-ac-DXd concentrations, when receptors are not saturated, and played relatively a minor role in total elimination of the antibody.
- d exposure also required two clearance pathways to describe PK amics:
- CL_{DXd} described the linear clearance portion and CL_{mm.DXd} described nonlinear portion which accounted for both the nonlinear formation of DXd as well as non-linear elimination.

$$CL_{ns} = CL_{inf} \cdot \left(1 + CL_{inf,EMAX} \cdot \frac{T_{50}^{\gamma}}{T_{50}^{\gamma} + \text{time}^{\gamma}}\right)$$

- non-specific linear clearance at infinity after dosing Q3W. $_{AX}$: max effect of time on CLns. me to half-maximal effect
- coefficient
- CLns pathway was identified as the major clearance pathway initial value at time zero was estimated at 0.0217 L/hr, which dually decreased over time reaching a steady state value of 143 L/hr. CL_{ns} contributed to 23.5% and 77.6% of total clearance at e zero and steady state, respectively.
- CLmm (CLmm = 0.00934 L/h) contributed to 39.5% of the total f total ulation)

Table 4	4. Parameter Estimates	- Structural	Parameters for
anti-H	ER3-ac-DXd		

	Brid			
			Estimate	95% CI
CLt (L/hr)	$\exp(\theta_1)$	Linear transient clearance	0.0858	0.0185, 0.398
V1 (L)	$\exp(\theta_2)$	Central volume of distribution	2.91	2.75, 3.07
Q (L)	$\exp(\theta_3)$	Intercompartmental clearance	0.0221	0.0169, 0.0289
V2 (L)	$\exp(\theta_4)$	Peripheral volume of distribution	3.17	2.59, 3.89
Kdes (1/hr)	$\exp(\theta_5)$	Rate constant of CLt exponential decline	0.217	0.108, 0.439
CLinf (L/hr)	$\exp(\theta_6)$	Nonspecific clearance at infinity	0.0136	0.0106, 0.0173
CLinf _{emax} (-)	exp(θ ₇)	Maximum effect of time on nonspecific time-dependent clearance	0.603	0.207, 1.75
T50 (hr)	$\exp(\theta_8)$	Time to half-maximal time effect	1.38e+03	839, 2.28e+03
Y	$exp(\theta_9)$	Hill coefficient	3.75	0.327, 43.0
Vmax (nmol/L/hr)	$\exp(\theta_{10})$	Maximal MM elimination	2.15	1.72, 2.68
KM (nmol/L)	$exp(\theta_{11})$	MM constant	45.8	27.6, 75.9

Table 5. Parameter Estimates – Structural Parameters for DXd

			Estimat	e 95% Cl
<i>CLt_{DXd}</i> (L/hr)	exp(θ ₁₂)	Linear clearance	4.42	3.83, 5.11
V1 _{DXd} (L)	$\exp(\theta_{13})$	Central volume	5.96	5.40, 6.57

Diagnostics: Goodness of fit plots

Figure 5

 Figure 5: Population predictions (left panels) vs observed, and individual predictions (right panels) vs observed for anti-HER3-ac-DXd (top panels) and DXd (bottom panels). Observed values are indicated by solid black circles. The line of identity (solid grey) is included as a reference(x=y). The dashed blue line represents a LOESS smooth through the data.



Diagnostics: Goodness of fit plots

- Figure 7: Covariates in the full PPK model were varied one at a time from the reference subject, i.e., a 60 kg non-Asian female NSCLC patient with normal hepatic function, ECOG 0, age of 60 year, eGFR of 90 mL/min, albumin of 40 g/L and baseline SLD of 6 cm and used to simulate Cavgss in N=500 parameter sets from the covariance matrix of the fixed effect estimate. The symbols indicate the median and the whiskers show the 95% prediction interval for the simulated Cavgss. The grey shaded area is the reference range with a lower bound of 0.8 and an upper bound of 1.25.
- The magnitude of effect of covariates was evaluated via univariate simulations of DXd exposures. Continuous covariates values were selected that represented the quartiles of the observed range in the analysis. All median point values and 95% CI were in range with 0.8-1.25 of the reference value, indicating none of the values of the covariates assessed within the range included in this analysis warranted a dose adjustment, based on DXd Cavgss.

Figure 7. Effect of Covariates on DXd exposure



	Stu	ıdy	
	U31402-J101 n = 182	U31402-U102 n = 219	Summary n = 401
Sex			
Female	182 (100.0)	139 (63.5)	321 (80.0)
Male	0 (0.0)	80 (36.5)	80 (20.0)
Missing	0 (0.0)	0 (0.0)	0 (0.0)
Kace			
American mulan or Alaska Nativo	1 (0.5)	1 (0.5)	2 (0.5)
Alaska Nalive Asian	142 (78 0)	103 (47 0)	245 (61 1)
Black or African American	4 (2 2)	5 (2 3)	9 (2 2)
Native Hawaiian or Other	· (2.2)	0 (2.0)	0 (2.2)
Pacific Islander	0 (0.0)	2 (0.9)	2 (0.5)
White	34 (18.7)	95 (43.4)	129 (32.2)
Multiple	1 (0.5)	13 (5.9)	14 (3.5)
Other	0 (0.0)	0 (0.0)	0 (0.0)
Hepatic function category (NCI-ODWG)			
Normal	102 (56.0)	186 (84.9)	288 (71.8)
Mild	76 (41.8)	31 (14.2)	107 (26.7)
Moderate	3 (1.6)	0 (0.0)	3 (0.7)
Severe	0 (0.0)	0 (0.0)	0 (0.0)
Missing	1 (0.5)	2 (0.9)	3 (0.7)
Renal function category (CKD-EPI)			
Normal	134 (73.6)	87 (39.7)	221 (55.1)
Mild	41 (22.5)	98 (44.7)	139 (34.7)
Moderate	7 (3.8)	34 (15.5)	41 (10.2)
Severe	U (U.U)	U (U.U)	0(0.0)
IVIISSIIIY FCOG status	0 (0.0)	0 (0.0)	0 (0.0)
0	132 (72 5)	77 (35 2)	209 (52 1)
1	50 (27 5)	142 (64 8)	192 (47.9)
2	0 (0.0)	0 (0.0)	0 (0.0)
Missing	0 (0.0)	0 (0.0)	0 (0.0)
Tumor type			
Breast cancer	182 (100.0)	0 (0.0)	182 (45.4)
NSCLC	0 (0.0)	219 (100.0)	219 (54.6)
Others	0 (0.0)	0 (0.0)	0 (0.0)

Summary is count (percent)

n: number of records summarized; Source code: ppk-eda-tables.R; Source file: pk-cat.tex.

Model Schematic

- The integrated model structure consisted of 3 compartments
- The PK of anti-HER3-ac-DXd was described by a two-compartment model with three elimination pathways:
- *CLt* = *Transient and linear, time-dependent clearance*
- CLns = Nonspecific time-dependent linear clearance
- CLMM = Non-linear Michaelis-Menten clearance

		$CL_t =$	CL_{T} *	exp(-	k _{des} '	* time	e)	
• <i>CL_T</i> wa	as clearanc	e at base	eline	<i>I</i> X	400		,	
• K _{des} ra	te of expor	nential de	cline					
		CLMM	= Vm	ax/(K	m+a	c-DX	ď)	
Vmax : m	aximal MM	l eliminat	ion					
Km : exp	osure eliciti	ing half m	naximur	n effec	t			
Figure	2. Cleara	nce of a	anti-HE	ER3-a	s-DX	d ove	er trea	ıtme
cycles								
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ich were the relative fractions of anti-HER3-ac-DXd CLns, CLt, and CLMM, respectively. • Total DXd clearance at steady-state was 17.3 L/hr and 17.9 for BC and NSCLC populations, respectively.

Figure 3. Formation rate of DXd over treatment cycles

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<i>Vmax_{DXd}</i> (nmol/L/hr)	exp(θ ₁₆)	Maximal MM elimination	6.18	4.58, 8.35
<i>KM_{DXd}</i> (nmol/L)	exp(θ ₁₇)	MM constant	0.483	3 0.342, 0.684
Frac _{ns} (-)	$\exp(\theta_{18})$	Scaling factor for relative conversion fraction from Cl	_ns 1	FIXED
Frac _t (-)	$exp(\theta_{21})$	Scaling factor for relative conversion fraction from Cl	_t 0.272	2 0.0670, 1.11
Frac _{mm} (-)	exp(θ ₂₂)	Scaling factor for relative conversion fraction from CLMM	0.272	2 0.0670, 1.11
Table 6. F	Paramet	er Estimates - Covariate	es	
			Estimate	95% CI
CLt _{type}	$\exp(\theta_{24})$	Tumor type effect on CLt (BC:NSCLC)	0.896	0.455, 1.76
CLt _{s/d}	θ_{25}	Baseline SLD effect on CLt	-0.0758	-0.269, 0.117
V1 _{sex}	$\exp(\theta_{26})$	Sex effect on V1 (male:female)	1.18	1.04, 1.33
V1 _{race}	$\exp(\theta_{27})$	Race effect on V1 (Asian:non-Asian)	0.927	0.851, 1.01
CLinf _{sex}	$\exp(\theta_{28})$	Sex effect on CLinf (male:female)	1.30	1.02, 1.66
CLinf _{race}	$\exp(\theta_{29})$	Race effect on Clinf (Asian:non-Asian)	1.02	0.668, 1.55
CLinf _{ecog}	$\exp(\theta_{30})$	ECOG effect on CLinf (>0:0)	1.04	0.612, 1.76
CLinf _{type}	exp(θ ₃₁)	Tumor type effect on CLinf (BC:NSCLC)	0.937	0.556, 1.58
CLinf _{hep}	exp(θ ₃₃)	Hepatic function effect on CLinf (impaired:normal)	0.906	0.752, 1.09
CLinf _{egfr}	θ_{34}	eGFR effect on CLinf	-0.302	-1.60, 0.999
CLinf _{alb}	θ_{35}	Albumin effect on CLinf	-0.490	-0.939, -0.0401
Kdes _{sld}	$ heta_{36}$	Baseline SLD effect on Kdes	-0.139	-0.391, 0.112
Frac _{nswt}	θ_{37}	Weight effect on Frac _{ns}	0.139	-0.0471, 0.325
Frac _{nssex}	$\exp(heta_{38})$	Sex effect on Frac _{ns} (male:female)	0.848	0.614, 1.17
Frac _{nsrace}	$\exp(\theta_{39})$	Race effect on Frac _{ns} (Asian:non-Asian)	1.06	0.734, 1.53

Diagnostics: VPC

• Figure 8: The lines represent the median (solid) or 5th / 95th (dashed) percentiles of the observed data. The shaded areas represent 95% prediction intervals for median (grey) or 5th / 95th percentiles for data (blue) simulated under the model. The points are observed anti-HER3-ac-DXd concentrations.

• The VPC demonstrated that the model predicted concentrations of both the anti-HER3-ac-DXd and DXd were in reasonable agreement with observed concentrations. In each case, the median of the observed was in close agreement with the distribution summarized from 500 simulated replicates for both studies and across dosing cycles. The simulated 5th percentiles were slightly lower than observed, suggesting a slight overestimation of IIV for both anti-HER3-ac-DXd and DXd, in Study 102

Figure 8.



Diagnostics: VPC

• Figure 9: The lines represent the median (solid) or 5th / 95th (dashed) percentiles of the observed data. The shaded areas represent 95% prediction intervals for median (grey) or 5th / 95th percentiles for data (blue) simulated under the model. The points are observed DXd

- the relative fractions of anti-HER3-ac-DXd CLns, CLt, and CLMM, respectively.
- DXd was described by a one-compartment model with two clearance pathways:
- CLDXd = linear DXd clearance
- CLMM.DXd = non-linear DXd. Michaelis-Menten clearance

Figure 1. Model Schematic

— Rate from CLt ---- Rate from CLns --- Rate from CLMM - - Total rate

Figure 4. Clearance of DXd over treatment cycles

Frac _{nsecog}	$\exp(\theta_{40})$	ECOG effect on Frac _{ns} (>0:0)	1.05	0.818, 1.35				
Frac _{nstype}	$\exp(\theta_{41})$	Tumor type effect on Frac _{ns} (BC:NSCLC)	1.04	0.752, 1.45				
Frac _{nshep}	$\exp(\theta_{43})$	Hepatic function effect on Frac _{ns} (impaired:normal)	1.09	0.904, 1.30				
Frac _{nsegfr}	θ_{44}	eGFR effect on Frac _{ns}	0.0350	-0.999, 1.07				
Frac _{nsalb}	θ_{45}	Albumin effect on Frac _{ns}	-0.271	-0.795, 0.254				
CL _{wt}	$ heta_{46}$	Weight effect on clearance parameters	0.343	0.0521, 0.633				
V _{wt}	$ heta_{47}$	Weight effect on volume parameters	0.475	0.124, 0.826				
Covariate effect parameters are displayed as "(Test:Reference)"								

Table 7. Parameter Estimates – Estimates of Variability

		Estimate	95% CI	Shrinkage	Figu
IIV-CLt	Ω _(1,1)	0.299 [CV%=59.0]	-0.490, 1.09	24.9	
IIV-V1	Ω _(2,2)	0.0205 [CV%=14.4]	-0.0122, 0.0533	7.19	
IIV-Q	Ω _(3,3)	0.431 [CV%=73.4]	-0.236, 1.10	18.7	
IIV-V2	Ω(4,4)	0.112 [CV%=34.5]	-0.218, 0.443	20.2	
IIV-CLinf	Ω _(6,6)	0.127 [CV%=36.9]	-0.143, 0.398	5.17	Baseline
IIV-T50	Ω _(8,8)	0.702 [CV%=101]	-0.166, 1.57	22.4	
IIV-CL _{DXd}	Ω(12,12)	0.108 [CV%=33.7]	-0.208, 0.423	17.8	
IIV-V1 _{DXd}	Ω(13,13)	0.00515 [CV%=7.18]	-0.0626, 0.0729	66.4	Baseline Esti
IIV-Frac _{ns}	Ω _(18,18)	0.0380 [CV%=19.7]	-0.119, 0.195	12.6	
RV on anti-HER3-ac- DXd	Σ _(1,1)	0.0342 [CV%=18.5]	0.0269, 0.0415	6.63	
RV on DXd	Σ (2,2)	0.0814 [CV%=28.5]	0.0612, 0.102	8.78	

Diagnostics: Goodness of fit plots

• Covariates in the full PPK model were varied one at a time from the reference subject, i.e., a 60 kg non-Asian female NSCLC patient with normal hepatic function, ECOG 0, age of 60 year, eGFR of 90 mL/min, albumin of 40 g/L and baseline SLD of 6 cm and used to simulate Cavgss in N=500 parameter sets from the covariance matrix of the fixed effect estimate. The symbols indicate the median and the whiskers show the 95% prediction interval for the simulated Cavgss. The grey shaded area is the reference range with a lower bound of 0.8 and an upper bound of 1.25.

The magnitude of effect of covariates was evaluated via univariate simulations of anti-HER3-ac-DXd exposures. Continuous covariates values were selected that represented the quartiles of the observed range in the analysis. All median point values and 95% CI were in range with 0.8-1.25 of the reference value, indicating none of the values of the covariates assessed within the range included in this analysis warranted a dose adjustment, based on anti-HER3-ac-DXd Cavgss.

Figure 6. Effect of Covariates on anti-HER3-ac-DXd exposure

concentrations.

Figure 9.

Lu D, Lu T, Gibiansky L, Li X, Li C, Agarwal P, et al.. Integrated twoanalyte population pharmacokinetic model of polatuzumab vedotin in patients with non-Hodgkin lymphoma. CPT Pharmacomet Syst Pharmacol (2020) 9(1):48-59.

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