Prediction of Early Onset Cytokine Release Syndrome and Neurologic Events after Axicabtagene Ciloleucel in Large **B-Cell Lymphoma Based on Machine Learning Algorithms**

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BACKGROUND

- Axicabtagene ciloleucel (axi-cel) is an autologous anti-CD19 chimeric antigen receptor (CAR) T-cell therapy approved for the treatment of adult patients with relapsed/refractory large B-cell lymphoma (LBCL) after ≥ 2 prior lines of systemic therapy¹
- In the ZUMA-1 (NCT02348216) multicenter, single-arm, registrational, Phase 1/2 study of axi-cel in patients with refractory LBCL, with a median follow-up of 27.1 months (n=101; pivotal Cohorts 1+2)²: - 83% Objective response rate (58% complete response rate) was seen
- Grade \geq 3 cytokine release syndrome (CRS) and neurologic events (NEs) were reported in 11% and 32% of patients, respectively
- In the long-term follow-up analysis of ZUMA-1 recently presented here at ASH 2021, the 5-year overall survival rate was 43% after a median follow-up of 63 months³
- CRS and NEs associated with CAR T-cell therapy required inpatient management in ZUMA-1, per protocol⁴ • Some axi-cel-treated patients have early, while others have late onset of CRS and NEs,⁵ warranting distinct monitoring and management approaches
- Several exploratory safety management cohorts were added to ZUMA-1 to minimize treatment-related toxicity
- Cohort 4, which evaluated levetiracetam (anticonvulsant) prophylaxis and earlier corticosteroid and tocilizumab (anti-interleukin-6 receptor antibody) intervention, demonstrated a reduced incidence of Grade \geq 3 CRS and NEs⁶
- Cohort 6, which assessed the impact of adding prophylactic corticosteroids to the Cohort 4 toxicity management regimen, demonstrated no cases of Grade \geq 3 CRS, delayed CRS onset, and generally similar NE toxicity compared with pivotal Cohorts 1+2⁷

OBJECTIVE

• To develop a predictive signature for early onset acute toxicities (within 3–4 days after axi-cel) based on machine learning algorithms from ZUMA-1 data, which could facilitate toxicity management in a realworld setting

METHODS

• This post hoc analysis included patients from ZUMA-1 Phase 1 and Phase 2 Cohorts 1, 2, 4, and 6

- Testing and training
- Cohorts 1, 2, and 4 - Validation

pressure.

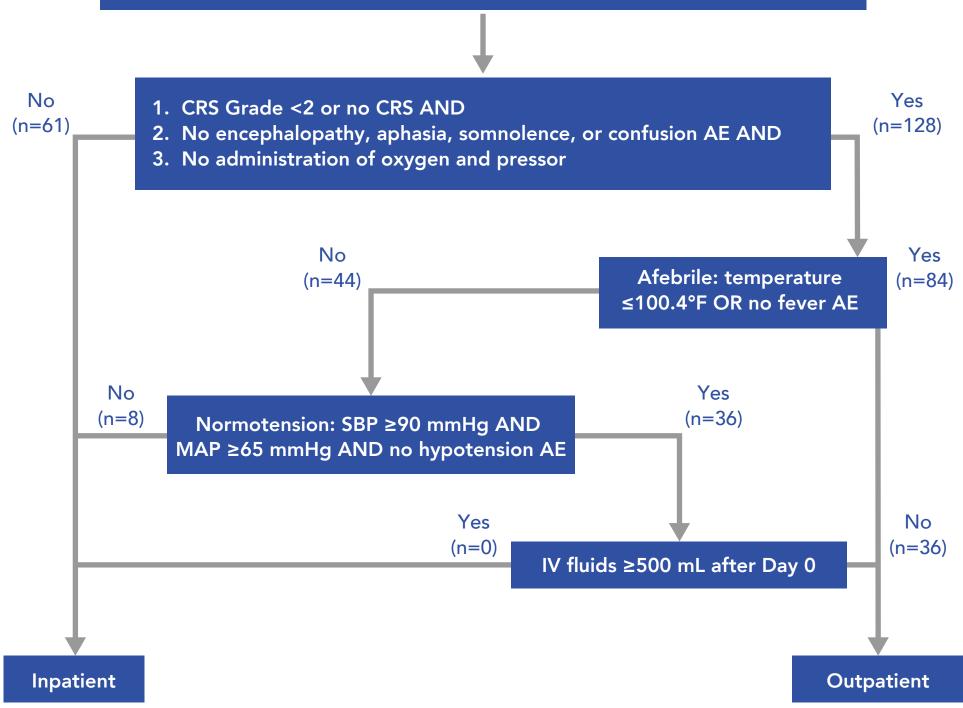
- Data from patients from ZUMA-1 Phase 2 Cohort 6⁷ were included to validate the best-performing model generated using training data
- Patients in Phase 1 and Phase 2 Cohorts 1 and 2 had \geq 2 years of follow-up; patients in Cohort 4 and Cohort 6 had ≥6 months of follow-up

Figure 1. Outpatient Definition

Definition ^a	Early Onset		No. of Patients Meeting Outpatient Definition (n [%])				
	Toxicity Time Point	Acute Toxicity Definition	Ph1 and Ph2 C1, 2, and 4 (N=149)	Ph2 C6 (N=40)			
A2	Day 0 to 2		88 (59)	37 (93)			
A3	Day 0 to 3	Grade ≤ I CRS and no INEs 4	72 (48)	33 (83)			
A4	Day 0 to 4		57 (38)	26 (65)			
B2	Day 0 to 2	5	41 (28)	34 (85)			
B3	Day 0 to 3	Patients with no CRS or NEs	31 (21)	27 (68)			
B4	Day 0 to 4		23 (15)	15 (38)			
C2	Day 0 to 2	Definition C (see schematic)	84 (56)	36 (90)			
C3	Day 0 to 3	Patients identified by AE, vital	64 (43)	32 (80)			
C4	Day 0 to 4	signs, and intervention	50 (34)	24 (60)			

Definition C2^b

Patients during first 72 hours after infusion (N=189^a)



Outpatient definition is based on the acute toxicity definition and the specified early onset toxicity time point. Nine outpatient definitions were assessed. Definition C2 has been provided as an illustration. The criteria within the classification tree are the same for outpatient definitions C3 and C4, apart from the early onset toxicity time point assessed. AE, adverse event; C, Cohort; CRS, cytokine release syndrome; IV, intravenous; MAP, mean arterial pressure; NE, neurologic event; Ph, Phase; SBP, systolic blood

• Three acute toxicity definitions were assessed at 3 different early onset time points for a total of 9 outpatient definitions

- in coculture)

central memory T cells, effector + effector memory T cells.

- to metabolic status)

RESULTS

Patient/Tumor Ch

Baseline

Bulky disease

Outpatient Defi Training AUC

Testing AUC

Covariates that are positively and negatively associated with all 9 outpatient definitions are shown in blue and red, respectively. Covariates that had different association directions across the 9 outpatient definitions are shown in black. Baseline assessments were those last collected prior to conditioning chemotherapy, and Day 0 assessments were those collected prior to axi-cel infusion on the day of infusion. Models that make predictions that are 100% correct have AUC values equal to 1. AUC, area under the curve; LDH, lactate dehydrogenase; RBC, red blood cell; WBC, white blood cell.

Table 2. Summary of Hybrid Covariates With Predictive Potential at Baseline and Day 0 and Hybrid Model Performance

Product Attribut "Fit" Product

Cell viability

lotal cells

Training AUC Testing AUC

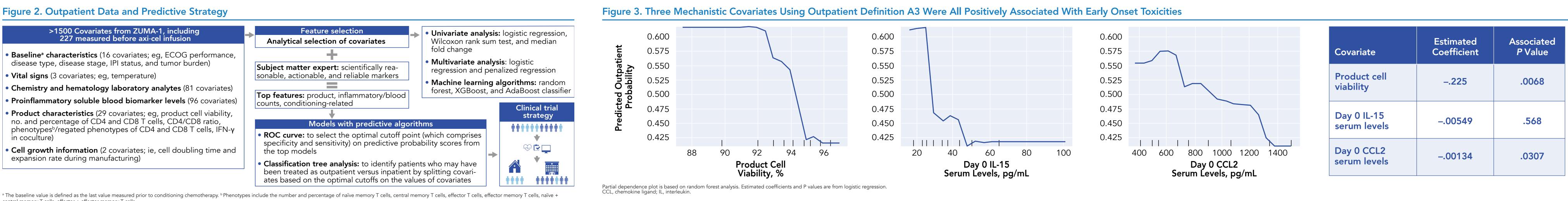
Covariates that are positively and negatively associated with all 9 outpatient definitions are shown in blue and red, respectively. Covariates that had different association directions across the 9 outpatient definitions are shown in black. Baseline assessments were those last collected prior to conditioning chemotherapy, and Day 0 assessments were those collected prior to axi-cel infusion on the day of infusion. Models that make predictions that are 100% correct have AUC values equal to 1. AUC, area under the receiver operating characteristic curve; CCL, chemokine ligand; IL, interleukin; LDH, lactate dehydrogenase; RBC, red blood cell; WBC, white blood cell.

Table 3. Summary of Hybrid Covariates With Predictive Potential at Baseline and Day 0 and Performance of Minimalistic Hybrid Model and Minimalistic Mechanistic Model

Product Attributes	Blood Chemistry		Blood Cells Day 0				Inflammatory Markers Day 0			
Baseline	Baseline									
Coll viobility	Urate		RBCs			IL-15				
Cell viability	Calcium						CCL2			
Model	Outpatient Definition	A2	B2	C2	A3	B3	С3	A4	B4	C4
Minimalistic hybrid model	Training AUC	0.867	0.914	0.839	0.854	0.873	0.785	0.899	1.000	0.903
(6 covariates)	Testing AUC	0.737	0.669	0.633	0.736	0.688	0.770	0.741	0.878	0.638
Minimalistic mechanistic model	Training AUC	0.963	0.864	0.695	0.803	0.998	0.773	0.790	0.859	0.860
(Cell viability + IL-15 + CCL2)	Testing AUC	0.719	0.736	0.609	0.750	0.757	0.766	0.808	0.752	0.620

collected prior to axi-cel infusion on the day of infusion. Models that make predictions that are 100% correct have AUC values equal to 1. AUC, area under the receiver operating characteristic curve; CCL, chemokine ligand; IL, interleukin; RBC, red blood cell.

METHODS (continued)



Axi-cel, axicabtagene ciloleucel; ECOG, Eastern Cooperative Oncology Group; IFN, interferon; IPI, International Prognostic Index; ROC, receiver operating characteristic

• For the model development, patient data were randomly divided into training (70%) and testing (30%) sets

• Machine learning algorithms were applied to 3 categories of covariates:

- Clinical (eg, tumor-related [lactate dehydrogenase level, burden], disease stage, blood cell counts, analytes related to cells [hemoglobin], and analytes related

- Mechanistic (eg, product attributes and inflammatory blood biomarkers) - Hybrid category that integrated both clinical and mechanistic covariates

Table 1. Summary of Clinical Covariates With Predictive Potential at Baseline and Day 0 and Clinical Model Performance

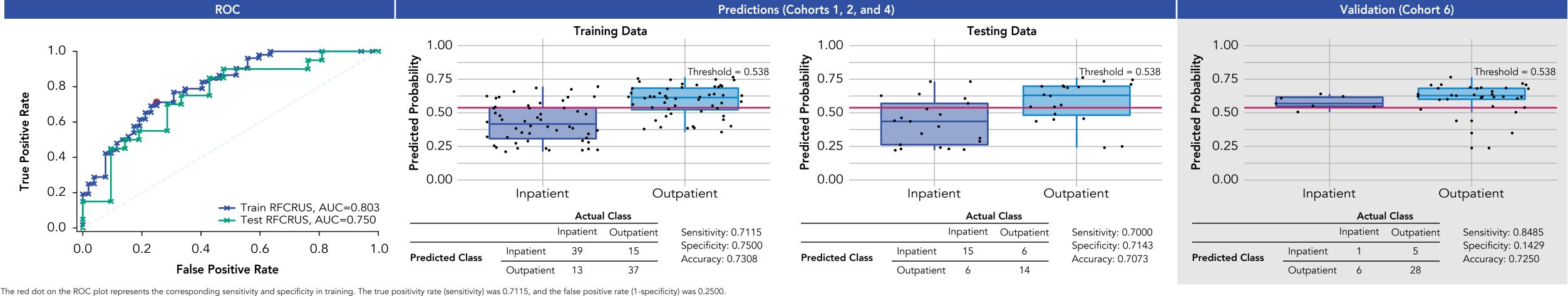
aracteristics	Blood Chemistry				E	Blood Cells	Inflammatory Markers				
	Altered Disease-Related Metabolic and Hematologic State								Inflammatory State		
	Base	line	Day	0	Baselin	le	Day 0	Day 0			
	Ura	te	Albun	nin	RBCs		RBCs				
Calcium Phosphate				WBCs		Hemoglobin					
		hate			Neutrophils						
	Creat	Creatinine				Basophils		Ferritin			
Chloride											
	LDH										
tion	A2	B2	C2	A3	B3	С3	A4	B4	C4		
	0.990	0.887	0.929	0.861	1.000	0.855	1.000	0.978	1.000		
	0.659	0.636	0.657	0.624	0.673	0.583	0.690	0.757	0.699		

	Patient/Tu	mor Characterist	ristics Blood C			nemistry Blood		Cells Inflammatory		ory Markers
		Altered Disease-Related Metabolic and Hematologic State								
		Baseline			eline	Day 0	Baseline	Day 0	Baseline	Day 0
				Ura	ate	Albumin	RBCs	RBCs	IL-17	IL-15
		Bulky disease					WBCs	Hemoglobin		CCL2
	D						Neutrophils			Ferritin
	BI						Basophils			
					ЭН					
on	A2	B2	C2		A3	B3	С3	A4	B4	C4
	0.930	0.948	0.757	7	0.945	0.988	0.879	0.831	1.000	0.897
	0.716	0.779	0.715	5	0.712	0.684	0.647	0.777	0.748	0.668

• Multivariate analysis and machine learning algorithms led to several comparable predictive models for early onset CRS or NEs (best-performing models with receiver operating characteristic (ROC). ROC AUC >0.8 in training and >0.7 in testing; **Tables 1–3**)

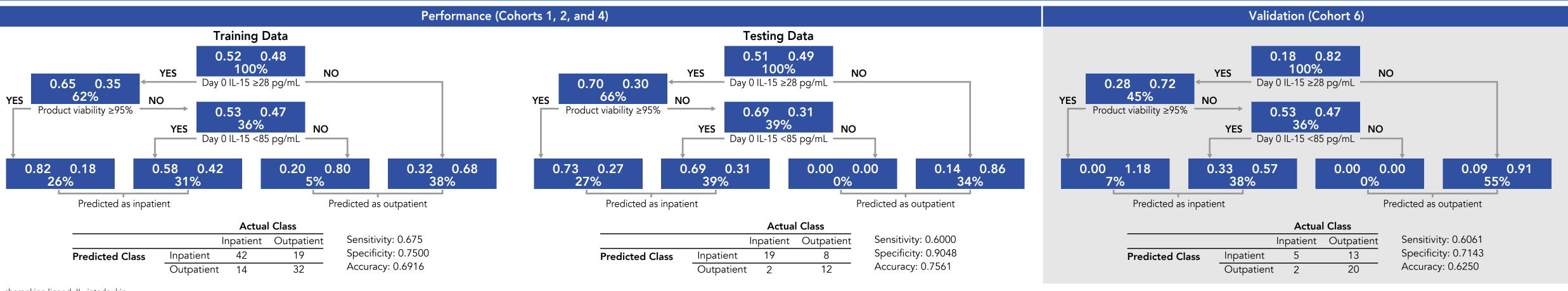
• The covariates in best-performing models included product cell viability, centrally measured Day 0 (pre-axi-cel treatment) IL-15 and CCL2 serum levels and locally measured blood cell counts, blood chemistry analytes, tumor burden, and serum lactate dehydrogenase level (Tables 1–3) • Minimalistic hybrid model (6 covariates) and minimalistic mechanistic model (Cell viability + IL-15 + CCL2) performed comparably with larger models (Table 3)

Figure 4. ROC and Cutoff Selection on Predicted Score From a 3-Covariate Model (Product Cell Viability and Day 0 IL-15 and CCL2 Serum Levels) Using Outpatient Definition A3



CCL, chemokine ligand; IL, interleukin; ROC, receiver operating characteristic.

Figure 5. Classification Tree Analysis From a 3-Covariate Model (Product Cell Viability and Day 0 IL-15 and CCL2 Serum Levels) Using Outpatient Definition A3



CCL, chemokine ligand; IL, interleukin.

• Classification trees of training and testing datasets with splitting based on Day 0 IL-15 and product cell viability showed a potential to categorize patients by early versus late onset of toxicities (specificity ≥0.75 in training and testing) • Models based on data from Cohort 6 did not recapitulate performance of those models optimized using data from ZUMA-1 Phase 1 and Phase 2 Cohorts 1, 2, and 4 (Figure 4 and Figure 5)

CONCLUSIONS

- Machine learning algorithms applied to covariates measured before axi-cel infusion yielded predictive models for early onset CRS or NEs that can be used for toxicity prediction, monitoring, and management
- High performing hybrid (ie, integrated mix of clinical and mechanistic covariates) or mechanistic models corroborated the importance of T-cell viability (product cell fitness) and conditioning-related elevation of factors (IL-15 and CCL2) in influencing toxicities
- Although the validation on Cohort 6 (prophylactic and earlier utilization of corticosteroids and/or tocilizumab) did not entirely recapitulate the performance of the best-performing models generated on previous cohorts where patients were managed differently, the overall analysis confirmed the importance of covariates uncovered in this study, paving the way for predictive algorithms in the real-world setting

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DISCLOSURES

Company, Roche, BMS/Celgene, Novartis, and Miltenyi Biotech; research funding from Kite, a Gilead Company, Roche, Novartis, and Miltenyi Biotech. MST: consultancy or advisory role for Amgen, Kite, a Gilead Company, Celgene, Roche, and Regeneron; and research funding from Amgen, Kite, a Gilead Company; and stock or other ownership in Gilead. KS: employment with Kite, a Gilead Company; and stock

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• A 3-covariate mechanistic model (product cell viability and Day 0 interleukin-15 [IL-15] and CCL2 serum levels) based on outpatient definition A3 performed comparably (ROC AUC > 0.7 in testing) to larger best-performing models

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