Ciltacabtagene **Autoleucel (Cilta-cel) vs Standard of Care (SOC)** in Patients (Pts) With **Relapsed/Refractory Multiple Myeloma (MM): CARTITUDE-4** Survival **Subgroup Analyses**

Yaël C Cohen¹, Joaquín Martínez-López², Abdullah M Khan³, Albert Oriol⁴, Andrew Spencer⁵, Binod Dhakal⁶, Cyrille Touzeau^{7,8,9}, Dominik Dytfeld¹⁰, Hermann Einsele¹¹, Jesús San-Miguel¹², Salomon Manier¹³, Diana Chen¹⁴, Quanlin Li¹⁵, Tzu-min Yeh¹⁶, Katherine Li¹⁷, Vicki Plaks¹⁷, Ana Slaughter¹⁸, Nina Benachour¹⁹, Carolina Lonardi²⁰, Arnab Ghosh¹⁶, Martin Vogel²¹, Nikoletta Lendvai¹⁶, Tamar Lengil¹⁶, Mythili Koneru²², Nitin Patel²², Octavio Costa Filho²², Erika Florendo²², Surbhi Sidana²³, Simon J Harrison^{24,25,26}

al Center, Faculty of Medical & Health Sciences, Tel Aviv University, Tel Aviv, Israel; ⊁hospital 12 de Octubre, Complutense University CNO, MC, Madrid, Spain; ≭The Ohio State Un olumbus, CH, USA, "Institut Català d'Oncologia and Institut Josep Carteras, Hospital Germans Trias i Puijol, Badalona, Barcelona, SpainëAlfred Health-Monash University. Melbourn eaca Longe of Wisconsn, Mwaukee, W, USA; "Service of Hemablogic, Cente Hogbialle Univergiate GHU) Hole Diau, Nartes, France: "Center de Richerche en Candologie et Immurico", Br. NESEM, Center Naliciale de Berderde Solentique, Université d'Anara, Shanes, France; "Site de Richerche Inférgée se la Canochinogie et Immurico", Brick und Polikin, H. 1992barg, Germany: "Canacor Center Ohise Public Method, Narae, CMA, Dissol, Public Method, Narae, CMA, Mick und Polikin, H. 1992barg, Germany: "Canacor Center Ohise Public Method, Narae, CMA, Dissol, "Linkersity of Ludical Sciences, Rozand, Poland," University Mick und Polikin, H. 1992barg, Germany: "Canacor Center Ohise Public Method, Narae, CMA, Dissol, "Linkersity of Ludical Sciences, Rozand, Poland," University Mick und Polikin, H. 1992barg, Germany: "Canacor Center Ohise Public Method, Narae, CMA, Dissol, "Linkersity of Ludical Sciences, Rozand, Poland," University Mick und Polikin, H. 1992barg, Germany: "Canacor Center Ohise Public Method, Narae, CMA, Dissol, "Linkersity of Ludical Sciences, Rozand, Poland," University Janes, Alexand, and Rozand, Poland, "University Alexand, Narae, CMA, Dissol, Telland, Narae, CMA, Dissol, Telland, Narae, CMA, Charles, "Henrico Alexand, Poland," University Alexand, Narae, CMA, Charles, "Henrico Alexand, Poland, "University Alexand, Poland, "University

Key Takeaway

Cilta-cel improved PFS and OS vs SOC across subgroups in CARTITUDE-4, including pts with standard- and high-risk cytogenetics, EMD, and 1 pLOT and beyond

Conclusions



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ITT analysis showed that cilta-cel improved PFS and OS vs SOC across subgroups, including pts with EMD and 1 pLOT and beyond

Compared with SOC, cilta-cel improved PFS and OS in pts with high-risk cytogenetics, suggesting it may overcome the poor prognosis associated with these high-risk features

These data continue to support a positive benefit-risk ratio for cilta-cel in pts with lenalidomide-refractory MM as early as after first relapse

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Introduction

- In CARTITUDE-4 (NCT04181827), a single cilta-cel infusion significantly prolonged progression-free survival (PFS) and overall survival (OS) in pts with lenalidomide-refractory MM after 1-3 prior lines of therapy (pLOT)¹
 - At median follow-up of 33.6 months, PFS (hazard ratio [HR] weighted, 0.29 [95% CI, 0.22-0.39]) and OS (HR, 0.55 [0.39-0.79]; P=0.0009) were significantly improved vs SOC1
- Overall minimal residual disease (MRD)-negative complete response (CR) or better rates (82.1% vs 25.2%) as well as sustained (≥12 months) MRD-negative ≥CR (51.7% vs 9.7%) rates in evaluable pts were higher in the cilta-cel arm vs SOC²
- Here, we report PFS and OS from subgroups of pts with standard-/high-risk cytogenetics, with/without extramedullary disease (EMD), and with 1, 2, or 3 pLOT

Results

Study population

- As of May 1, 2024, the median follow-up was 33.6 months (range, 0.1-45.0
- Cilta-cel vs SOC by cytogenetic risk
- Cilta-cel consistently improved PFS and OS compared with SOC in pts with standard risk and high risk (Figures 2 and 3)
- High-risk cytogenetics was defined as del(17p), t(4;14), t(14;16), or gain/amp(1q) by fluorescence in situ hybridization

Figure 2: Kaplan-Meier analysis of pts with standard- and highrisk cytogenetics



Methods

Treatment and data analysis

- CARTITUDE-4 study design has been described previously3
- Pts (n=208) randomized to the cilta-cel arm underwent apheresis, bridging treatment. lymphodepletion, and then a single cilta-cel infusion (n=176, Figure 1)
- Bridging treatment consisted of either:
- Pomalidomide, bortezomib, and dexamethasone (PVd) or
- Daratumumab, pomalidomide, and dexamethasone (DPd)

cvtogenetics



Α	Subgroup	HR (95% CI)ª	c	Cilta-cel		SOC	
		:	N	Median (95% CI)	N	Media (95%)	
	del (17p)		49	29.9 (13.4–NE)	43	8.7 (5.1–14	
	t(4;14)	н — н	30	37.1 (18.0–NE)	30	6.7 (3.8–13	
	Gain/amp(1q)	щ	89	37.1 (25.9–NE)	107	10.3 (7.5–14	
	≥2 cytogenetic abnormalities		43	29.8 (10.8–NE)	49	6.7 (4.7–10	
	Favors cilta	0 0.5 1	2 → Favors SOC	;			

3	Subgroup	o HR (95% Cl)ª Cilta-cel		ilta-cel	SOC	
			N	Median (95% CI)	N	Media (95% C
	del (17p)		49	NR (NE–NE)	43	NR (17.3–N
	t(4;14)		30	NR (NE–NE)	30	26.8 (20.3–N
	Gain/amp(1q)		89	NR (NE–NE)	107	38.0 (34.0–N
	≥2 cytogenetic abnormalities		43	NR (29.8–NE)	49	23.0 (17.3–N
		0 0.5 1	2			
	Favors cilta	-cel ← 🗕 🗕 🔶	Favors SOC	;		

1. Mateos M-V, et al. Clin Lymphoma Myeloma Leuk 2024;24(suppl 2):S290. 2. Popat R, et al. Blood 2024;144(suppl 1):1032. 3. San-Miguel J, et al. N Engl J Med 2023;389:335-47

Figure 1: Randomization and treatment

As-treated population

ITT population



*32 pts did not receive cilta-cel as study treatment (n=30 due to disease progression; n=2 due to death during bridging therapyllymphodepletion), of which 20 received cilta-cel as subsequent LOT. ITT, intent to treat; LOT, line of the rapy; tx, treatment,

Pts (n=211) randomized to the SOC arm received physician's choice of PVd or DPd until disease progression (n=208)

PFS was assessed using a validated computerized algorithm; HR was analyzed using an unweighted Cox proportional hazards model for the ITT analysis set

Multiple Myeloma

