

Ciltacabtagene Autoleucel in Lenalidomide-Refractory Multiple Myeloma Responding to Bridging Therapy: CARTITUDE-4 Cytogetic Subgroup Analysis

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Key Takeaway



Cilta-cel demonstrated a profound benefit in patients with high- and standard-risk cytogenetics when MM is well controlled at time of infusion

Conclusions



In patients with high- and standard-risk cytogenetics, those with ≥PR to BT had better survival outcomes with cilta-cel compared with those who had <PR to BT



There were no cases of IEC-parkinsonism in patients who had ≥PR to BT



Infections were a key cause of NRM in patients who had <PR to BT

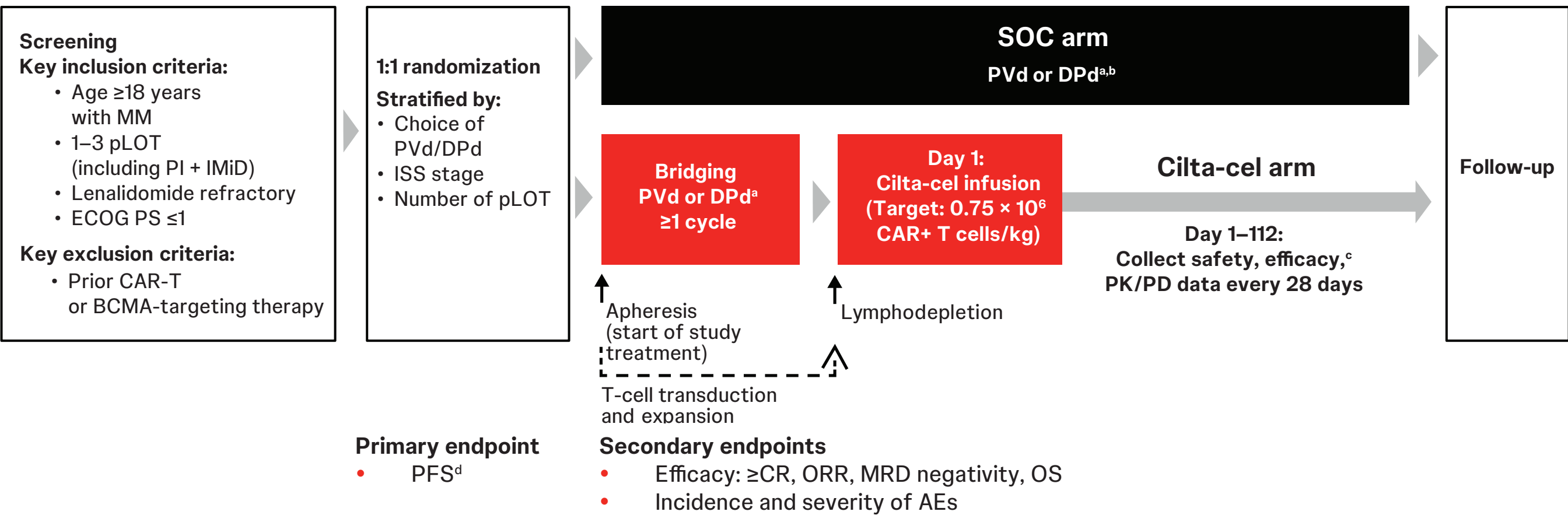
Introduction

- Previous analyses from the phase 3 CARTITUDE-4 study of patients with lenalidomide-refractory multiple myeloma (MM) after 1–3 prior lines of therapy (pLOT) who were treated with ciltacabtagene autoleucel (cilta-cel) have shown that survival and safety outcomes vary by cytogenetic risk status and response to bridging therapy (BT)^{1–3}
 - Patients with high- or standard-risk cytogenetics treated with cilta-cel had improved progression-free survival (PFS) and overall survival (OS) compared with patients treated with standard of care (SOC)¹
 - Partial response or better (≥PR) during BT was associated with better survival and safety outcomes with cilta-cel³
- Emerging real-world data underscore the importance of effective BT before infusion with cilta-cel⁴
- Here, we present efficacy and safety for patients who received cilta-cel as study treatment according to response to BT and cytogenetic risk (median study follow-up, 33.6 months)

Methods

- CARTITUDE-4 study design is shown in Figure 1

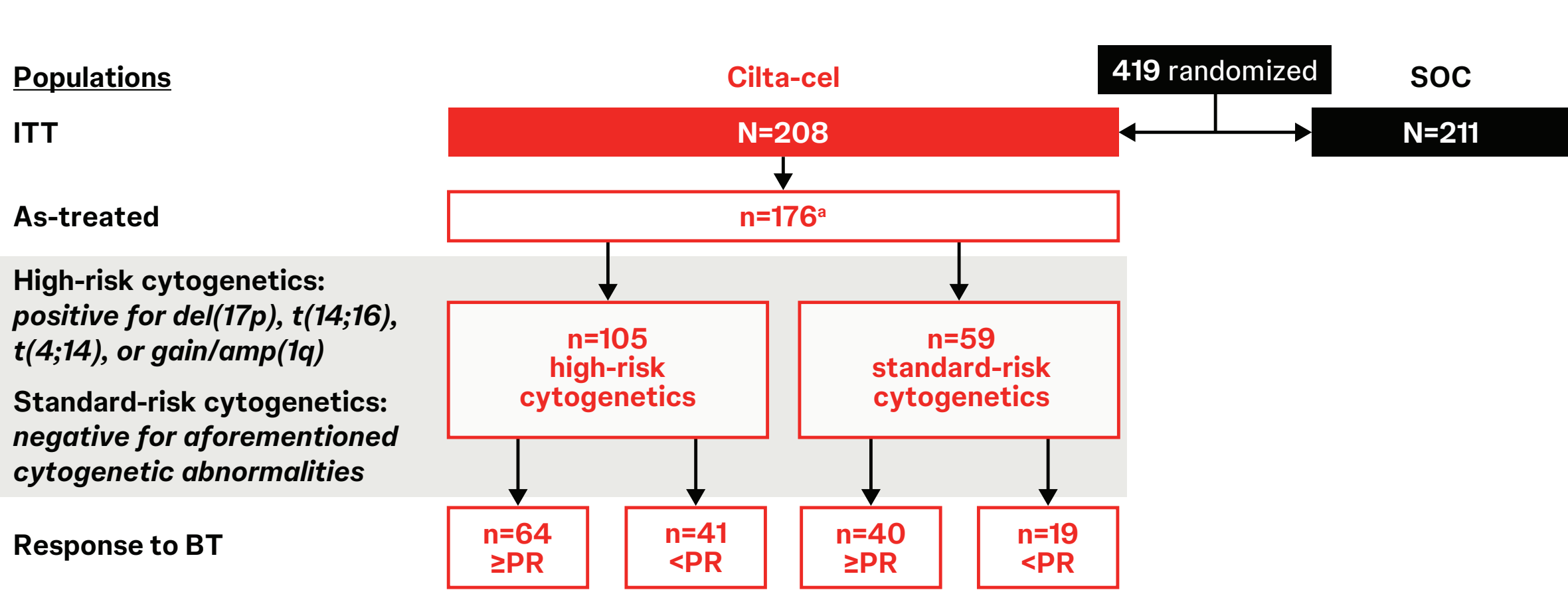
Figure 1: CARTITUDE-4 study design^{2,5}



^aPhysician's choice; ^bAdministered until disease progression; ^cEfficacy data were collected after day 112 every 28 days; ^dTime from randomization to disease progression/death; AE, adverse event; BCMA, B-cell maturation antigen; CAR, chimeric antigen receptor; CR, complete response; DPd, daratumumab, pomalidomide, and dexamethasone; ECOG PS, Eastern Cooperative Oncology Group performance status; IMiD, immunomodulatory drug; ISS, International Staging System; MRD, minimal residual disease; ORR, overall response rate; PD, pharmacodynamics; PI, proteasome inhibitor; PK, pharmacokinetics; PVD, pomalidomide, bortezomib, and dexamethasone.

- Study population is shown in Figure 2

Figure 2: Study population^{3,5}



^a12 patients did not have cytogenetic risk information available. ITT, intent-to-treat.

Kaplan-Meier estimated PFS and OS by cytogenetic risk status and response to BT

- Deeper responses to BT were associated with improved PFS and OS outcomes in the high- and standard-risk cytogenetics subgroups (Figures 3 and 4)

Figure 3A: PFS from cilta-cel infusion in the high-risk cytogenetics subgroup by response to BT

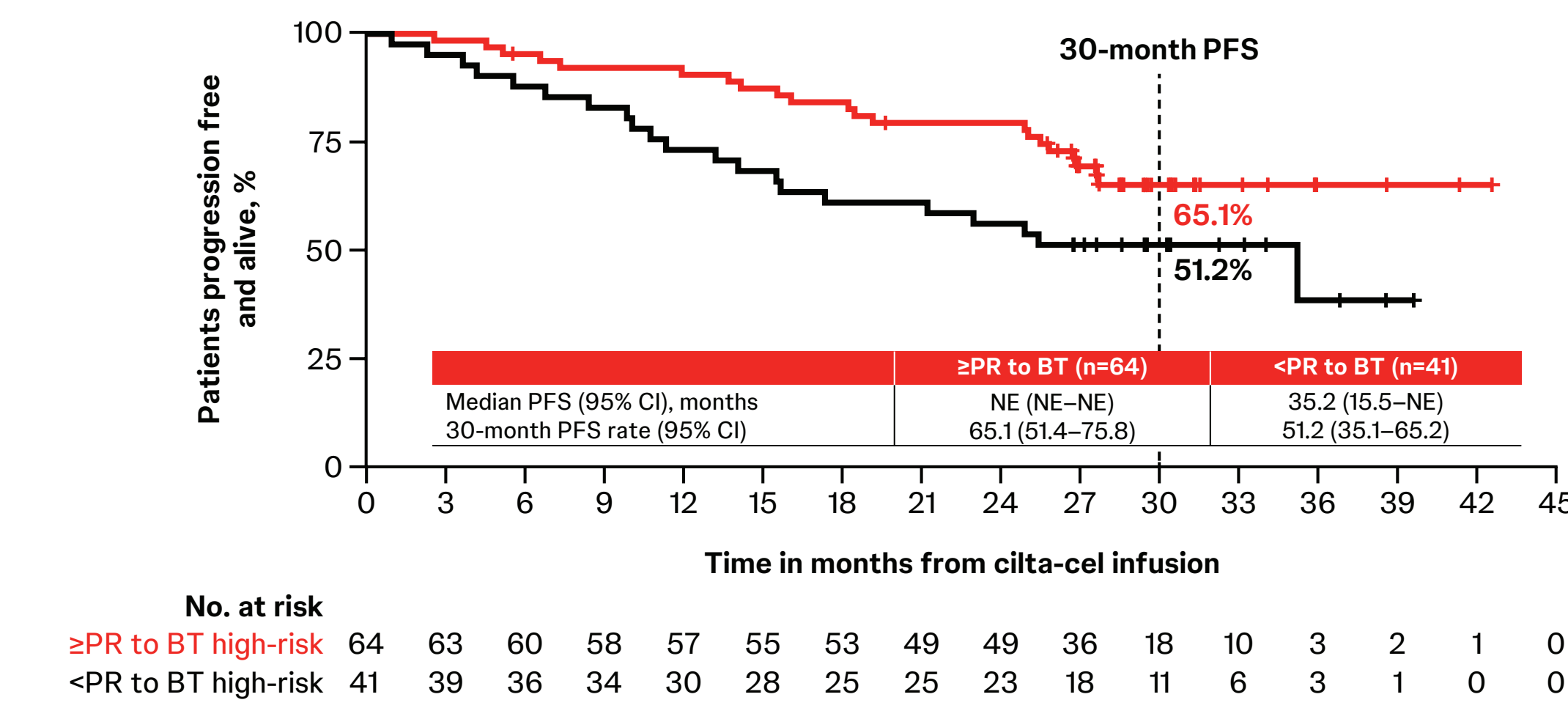


Figure 3B: OS from cilta-cel infusion in the high-risk cytogenetics subgroup by response to BT

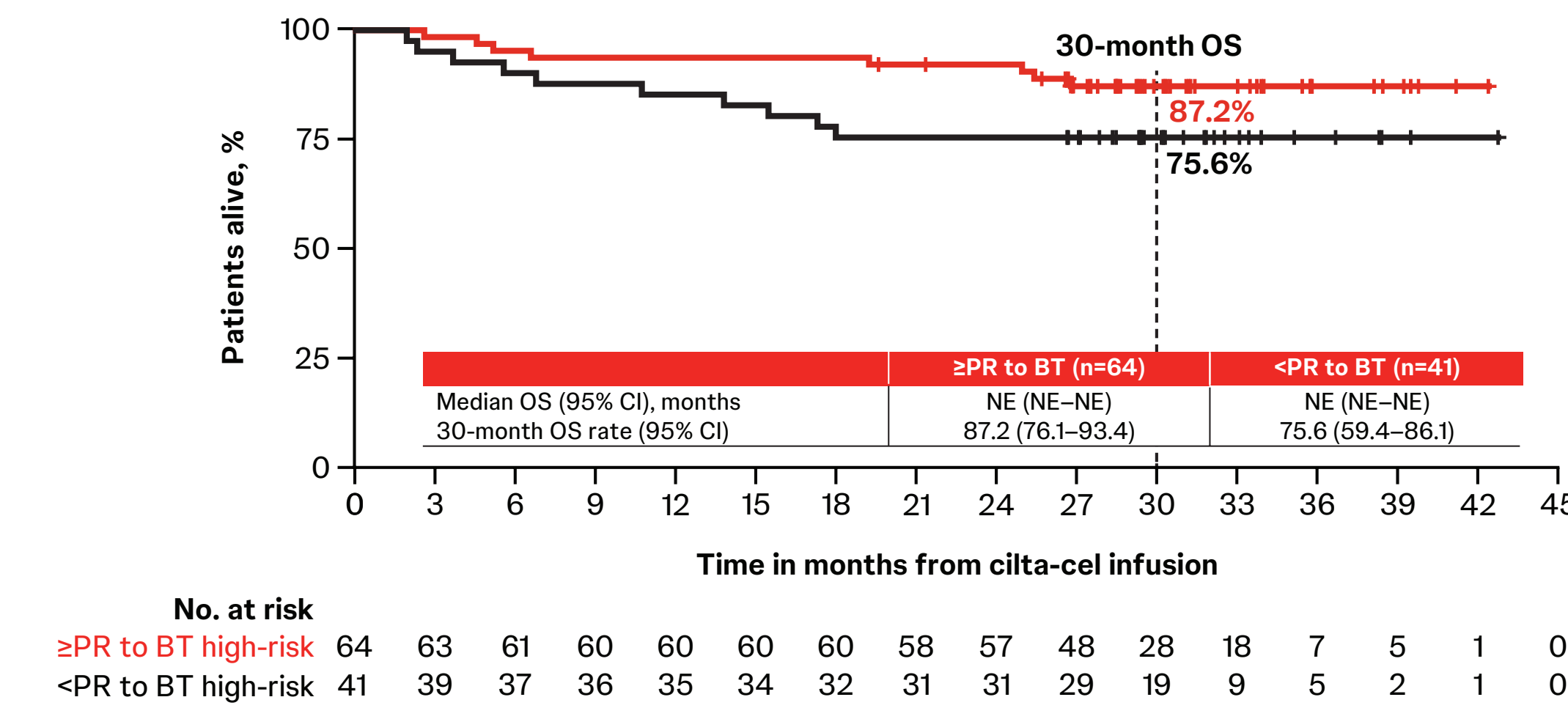


Figure 4A: PFS from cilta-cel infusion in the standard-risk cytogenetics subgroup by response to BT

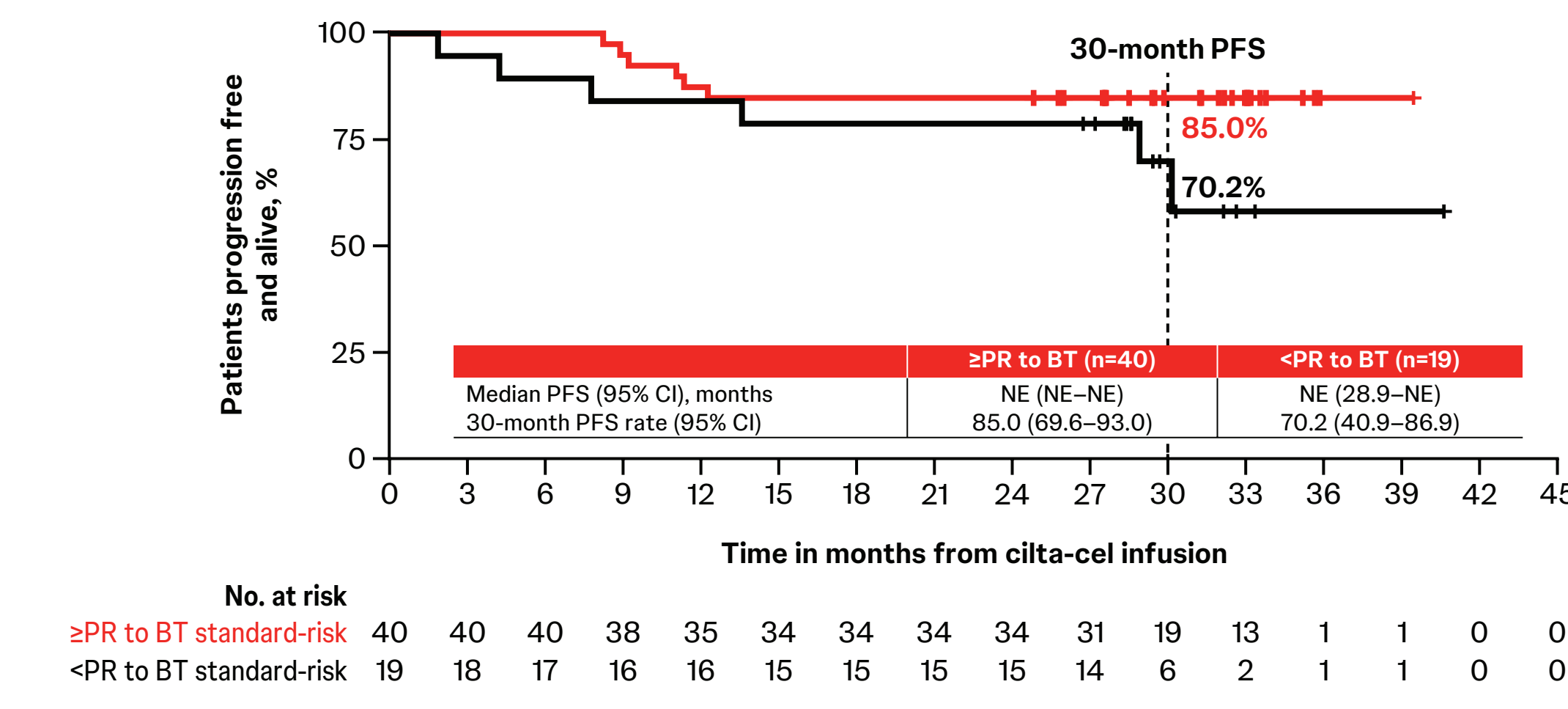
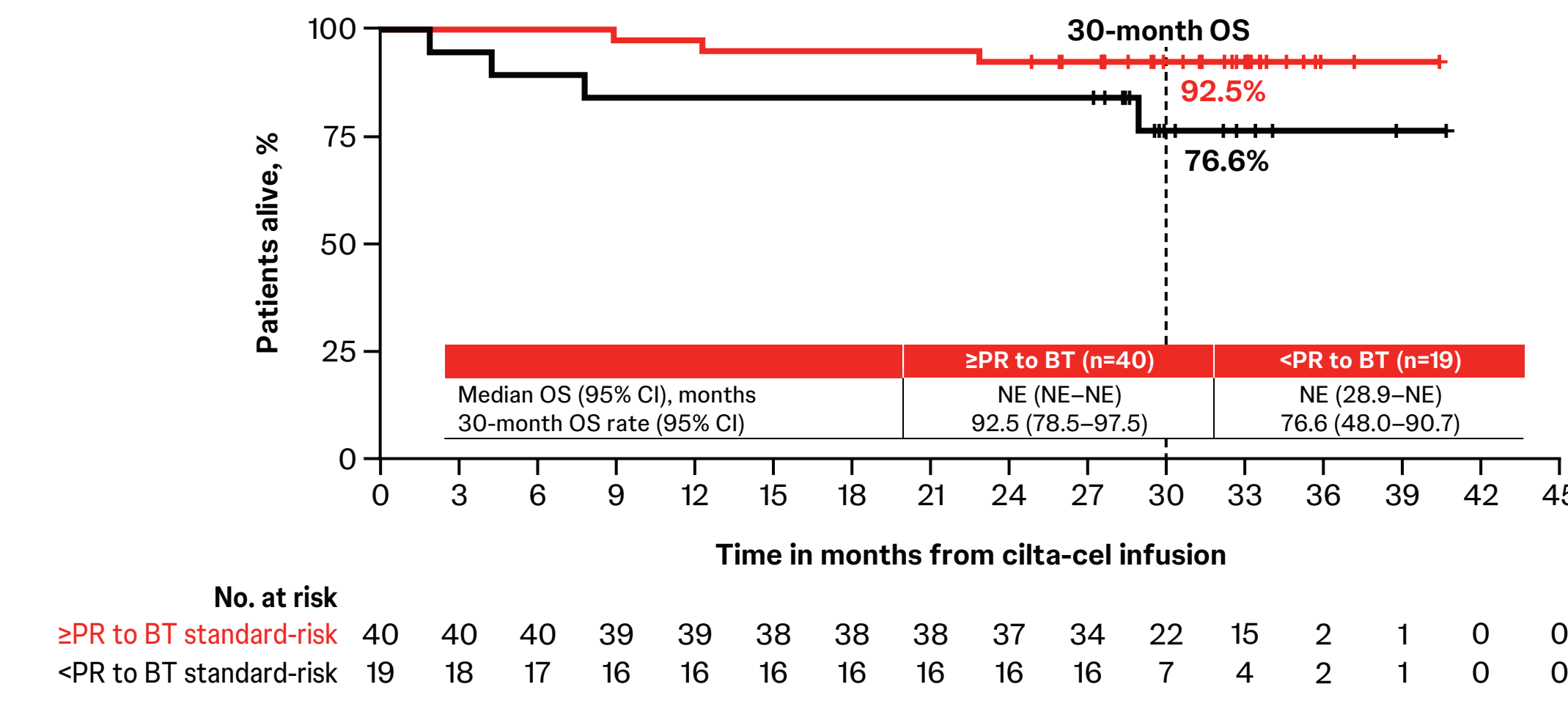


Figure 4B: OS from cilta-cel infusion in the standard-risk cytogenetics subgroup by response to BT



Safety (Table 2)

- Grade 3/4 infections were numerically higher in patients with high-risk cytogenetics (≥PR to BT, 43.8%; <PR to BT, 48.8%) vs standard-risk cytogenetics (≥PR to BT, 30.0%; <PR to BT, 26.3%) subgroups
- Patients with ≥PR to BT experienced fewer fatal infections, irrespective of cytogenetic risk
- There were no cases of immune effector cell (IEC)-parkinsonism in patients who achieved ≥PR to BT
- Nonrelapse mortality (NRM) rates were higher in patients who had <PR to BT, regardless of cytogenetic risk

Table 2: AEs in patients with high- and standard-risk cytogenetics

AE, n (%)	High-risk cytogenetics		Standard-risk cytogenetics	
	≥PR to BT (n=64)	<PR to BT (n=41)	≥PR to BT (n=40)	<PR to BT (n=19)
CRS	47 (73.4)	34 (82.9)	28 (70.0)	16 (84.2)
ICANS	5 (7.8)	2 (4.9)	0	1 (5.3)
CNP	7 (10.9)	4 (9.8)	2 (5.0)	2 (10.5)
IEC-parkinsonism	0	1 (2.4) ^b	0	0
Grade 3/4 infections	28 (43.8)	20 (48.8)	12 (30.0)	5 (26.3)
NRM	7 (10.9)	6 (14.6)	2 (5.0)	4 (21.1)
Infections ^a	4 (6.2)	5 (12.2)	0	2 (10.5)

^a≥PR to BT: COVID-19 infection (n=2), cytomegalovirus colitis (n=1), fungal pneumonia (n=1); <PR to BT: COVID-19 infection (n=5), pneumonia (n=1), *Pneumocystis jirovecii* pneumonia (n=1).

^bOccurred in a patient with unconfirmed PD at cilta-cel infusion.

CNP, cranial nerve palsy; ICANS, immune effector cell-associated neurotoxicity syndrome.

Results

- 176 patients received cilta-cel as study treatment and were divided into 4 subgroups based on cytogenetics and response to BT:
 - High-risk cytogenetics subgroups (n=105): ≥PR to BT (n=64) and <PR to BT (n=41)
 - Standard-risk cytogenetics subgroups (n=59): ≥PR to BT (n=40) and <PR to BT (n=19)
- Baseline characteristics and response to BT by subgroup are summarized in Table 1

Table 1: Baseline characteristics and response to BT by subgroup

Baseline characteristic	High-risk cytogenetics		Standard-risk cytogenetics	
	≥PR to BT (n=64)	<PR to BT (n=41)	≥PR to BT (n=40)	<PR to BT (n=19)
Median age (range), years	63.5 (43.0–78.0)	57.0 (43.0–75.0)	60.0 (27.0–78.0)	65.0 (38.0–78.0)
ISS stage, n (%)				
I	47 (73.4)	23 (56.1)	31 (77.5)	13 (68.4)
II	16 (25.0)	13 (31.7)	7 (17.5)	5 (26.3)
III	1 (1.6)	5 (12.2)	2 (5.0)	1 (5.3)
Higher baseline tumor burden, n (%)	4 (6.3)	10 (24.4)	5 (12.5)	3 (15.8)
pLOT, n (%)				
1	26 (40.6)	9 (22.0)	14 (35.0)	6 (31.6)
2	24 (37.5)	17 (41.5)	14 (35.0)	8 (42.1)
3	14 (21.9)	15 (36.6)	12 (30.0)	5 (26.3)
Triple-class ^a refractory, n (%)	6 (9.4)	6 (14.6)	2 (5.0)	3 (15.8)
Response to BT, n (%)				
sCR	1 (1.6)	–	0	–
CR	3 (4.7)	–	1 (2.5)	–
VGPR	17 (26.6)	–	15 (37.5)	–
PR	43 (67.2)	–	24 (60.0)	–
MR	–	6 (14.6)	–	2 (10.5)
SD	–	32 (78.0)	–	17 (89.5)
NE	–	3 (7.3)	–	0

^aPI + immunomodulatory agent + CD38 antibodies.

MR, minimal response; NE, not evaluable; PD, progressive disease; sCR, stringent complete response; SD, stable disease; VGPR, very good partial response.

High-risk cytogenetics subgroups by response to BT

- In patients with ≥PR to BT, median PFS and OS were not reached; 30-month PFS and OS rates were 65.1% (95% CI, 51.4–75.8) and 87.2% (95% CI, 76.1–93.4), respectively
- Patients with <PR to BT had median PFS of 35.2 months and median OS was not reached; 30-month PFS and OS rates were 51.2% (95% CI, 35.1–65.2) and 75.6% (95% CI, 59.4–86.1), respectively

Standard-risk cytogenetics subgroups by response to BT

- In patients with ≥PR to BT, median PFS and OS were not reached; 30-month PFS and OS rates were 85.0% (95% CI, 69.6–93.0) and 92.5% (95% CI, 78.5–97.5), respectively
- For patients with <PR to BT, median PFS and OS were not reached; 30-month PFS and OS rates were 70.2% (95% CI, 40.9–86.9) and 76.6% (48.0–90.7)

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Disclosures

RM reports a consulting/advisory role or honoraria for Amgen, BMS/Celgene, GSK, Janssen, Menarini-Stemline, Pfizer, Sanofi, and Takeda.

