

# Clinical Pharmacological Characterization of Cilta-cel, a CAR-T Therapy Directed Against BCMA in Adult Patients with Multiple Myeloma in a Multicohort CARTITUDE-2 Study

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### Key Takeaway

Following a single infusion of cilta-cel at a target dose of 0.75x10<sup>6</sup> cells/kg, CAR-T cellular expansion and persistence were observed coinciding with continued reduction of sBCMA, which reflects CAR-T cell mediated pharmacodynamic activity. Although there was evidence of treatment emergent immunogenicity (ADA positive), the impact of ADA on cellular kinetics was not apparent.

### Conclusions

PK exposures assessed using transgene and cellular levels were concordant and showed similar expansion and persistence profiles. High interindividual variability of exposures was observed. The median t<sub>max</sub> of cilta-cel transgene expansion in peripheral blood was approximately 2 weeks post-infusion and the median t<sub>bqcl</sub> of cilta-cel transgene levels was approximately 4-5 months post-infusion.

Coincided with CAR-T cell expansion, continued reduction of sBCMA was observed and reached nadir level around 2 or 3 months, sBCMA levels thereafter showed a gradual increase over time, but these values remained substantially lower than baseline in most participants.

The ADA incidence was 25 to 42%, with the median onset time of 2 to 6 months post-infusion. There was no apparent impact of ADA on cilta-cel cellular kinetics.

### Acknowledgments

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### Disclosures

JMW, YR, HV, YJ, QS, CC, NL and IS are currently employed by Janssen R&D LLC, SC is not currently employed by Janssen. JMW, YR, HV, YJ, QS, CC, NL and IS are employed by Johnson & Johnson and own Johnson & Johnson stock. MA and TY are currently employed by Legend Biotech.

### Introduction

- Ciltacabtagene autoleucel (cilta-cel) is an autologous chimeric antigen receptor T cell (CAR-T) therapy that targets B cell maturation antigen (BCMA) expressed on the surface of mature B lymphocytes and malignant plasma cells.
- Cilta-cel was approved in US and EU based on results of Phase 1b/2 CARTITUDE-1 study (NCT03548207) and Phase 3 randomized controlled trial, CARTITUDE-4 study (NCT04181827). Cilta-cel administration resulted in deep and durable responses in patients with relapsed and refractory (R/R) multiple myeloma (MM), who have received at least 1 prior line of therapy, including a proteasome inhibitor (PI) and an immunomodulatory agent (IMiD), and are refractory to lenalidomide.
- CARTITUDE-2 (NCT04133636) is a phase 2, multicohort, open-label study evaluating safety and efficacy of cilta-cel in various MM patient populations. Cohort A patients had received 1 to 3 prior lines of therapy and were lenalidomide-refractory, Cohort B patients received one line of prior therapy including a PI and an IMiD and had disease progression ≤12 months after ASCT, and Cohort C patients were previously treated with a PI, an IMiD, an anti-CD38 monoclonal antibody and BCMA-directed therapy (excluding cellular immunotherapy) (Figure 1).

### Results

#### 1. Cellular Kinetics:

- Cilta-cel PK measurements by using transgene and cellular levels were concordant. Similar cellular kinetic profiles were exhibited across the 3 cohorts with an initial expansion phase followed by a rapid then a slower decline over months as shown in Figure 2. High interindividual variability was observed.
- Although higher mean C<sub>max</sub> was observed in Cohort A(i) and Cohort B, and lower C<sub>max</sub> in Cohort C, the AUC<sub>0-28d</sub> was comparable across cohorts. The median t<sub>max</sub> ranged 11 to 15 days, and median t<sub>bqcl</sub> (time to reach BQL level) were comparable ranging from 125 to 157 days among Cohorts A(i), B, and C as shown in Table 1.
- There was no clear dose-exposure relationship observed across 3 cohorts as shown in Figure 3.

Figure 2: Mean (+SD) Blood Cilta-cel CAR Transgene Levels versus Time Profile

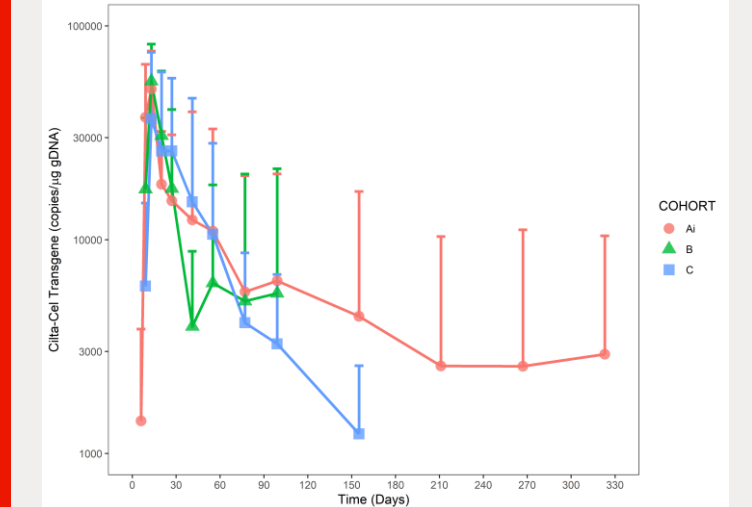


Table 1: Pharmacokinetic Parameters of Cilta-Cel Transgene Levels in Blood After a Single Infusion of Cilta-cel

Pharmacokinetics of Cilta-cel Transgene (mean [SD], t <sub>max</sub> , t <sub>0.5</sub> and t <sub>bqcl</sub> ; median [range])	Cohort Ai	Cohort B	Cohort C
n	20 <sup>a</sup>	19 <sup>b</sup>	18 <sup>c</sup>
C <sub>max</sub> , copies/μg genomic DNA	64039 (28184)	62097 (31062)	47809 (38948)
t <sub>max</sub> , day	10.50 (8.73 - 42.88)	13.08 (8.96 - 209.88)	14.94 (8.92 - 41.02)
C <sub>last</sub> , copies/μg genomic DNA	3421 (6912)	6714 (15138)	4450 (15703)
t <sub>0.5</sub> , day	183.05 (20.97 - 331.92)	96.97 (26.90 - 330.84)	126.71 (8.92 - 323.80)
t <sub>bqcl</sub> , day	153.47 (57.12 - 336.78)	124.81 (40.99 - 221.75)	156.97 (14.91 - 328.88)
AUC <sub>0-28d</sub> , day*copies/μg genomic DNA	601430 (295664)	639474 (389912)	558681 (480470)
AUC <sub>0-6m</sub> , day*copies/μg genomic DNA	1505597 (2190896)	1368431 (1799134)	1286263 (830054)
AUC <sub>0-12m</sub> , day*copies/μg genomic DNA	1712545 (3109405)	1387920 (2055365)	1202972 (1214932)
t <sub>1/2</sub> , day	38.3 (34.8)	11.0 (5.8)	40.8 (29.9)

AUC<sub>0-28d</sub>=area under the analyte concentration-time curve from time 0 to 28 days; AUC<sub>0-6m</sub>=area under the analyte concentration-time curve from time 0 to 6 months; AUC<sub>0-12m</sub>=area under the analyte concentration-time curve from time 0 to the time of last measurable (non-BQL) concentration; BQL=below quantification level; C<sub>last</sub>=last observed measurable (non-BQL) analyte concentration; C<sub>max</sub>=maximum observed analyte concentration; SD=standard deviation; t<sub>0.5</sub>=apparent terminal elimination half-life; t<sub>max</sub>=time of first BQL concentration after reaching C<sub>max</sub>; t<sub>bqcl</sub>=time of last measurable (non-BQL) analyte concentration; t<sub>1/2</sub>=time to reach maximum observed analyte concentration.  
<sup>a</sup> n=19 for AUC<sub>0-28d</sub> and AUC<sub>0-6m</sub>; n=11 for t<sub>1/2</sub> and n=10 for t<sub>bqcl</sub>.  
<sup>b</sup> n=17 for t<sub>0.5</sub> and n=8 for t<sub>1/2</sub>.  
<sup>c</sup> n=16 for AUC<sub>0-28d</sub>; n=13 for AUC<sub>0-6m</sub>; n=9 for t<sub>0.5</sub> and n=8 for t<sub>1/2</sub>.

### References

- Clinical Protocol, A Phase 2, Multicohort Open-Label Study of JNJ-68284528, a Chimeric Antigen Receptor T cell (CAR-T) Therapy Directed Against BCMA in Subjects with Multiple Myeloma. Dated Oct. 23, 2023.
- 2023 ACCP: Pharmacokinetics, Pharmacodynamics, and Exposure- Response for Neurotoxicity Adverse Events in a Phase Ib/II CARTITUDE-1 Open-Label Study. Shih-Yu Chang<sup>1</sup>, Jianmei Wu<sup>1</sup>, Yupeng Ren<sup>1</sup>, Dong Geng<sup>2</sup>, Tairan Yang<sup>2</sup>, Deepu Madduri<sup>1</sup>, Carolyn Jackson<sup>1</sup>, Jordan Schecter<sup>1</sup>, and Xuewen Ma<sup>1</sup>

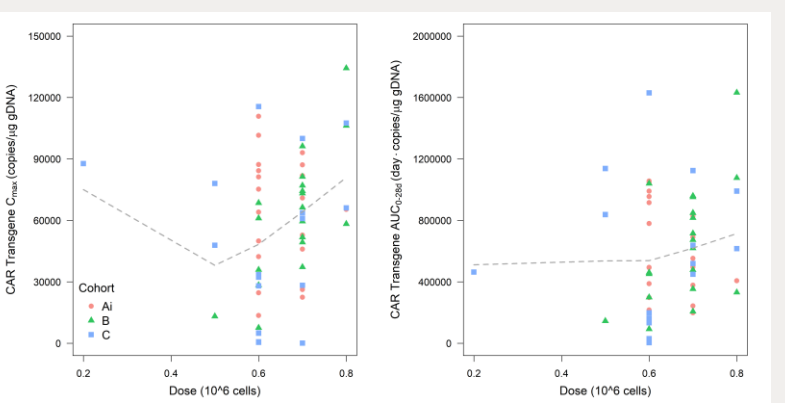
### Objectives

Cellular kinetics, pharmacodynamics, and immunogenicity results of cilta-cel are presented for Cohorts A initial group [A(i)], B and C to improve our understanding of cilta-cel *in vivo* expansion and its relationships with clinical factors in various MM populations.

### Methods

- Cellular kinetics was assessed for Cohorts Ai (first 20 patients who received cilta-cel using the clinical trial process), B and C by using CAR transgene copy level and CAR<sup>+</sup>CD3<sup>+</sup> cell level in blood after a single infusion of 0.75 (range: 0.5-1.0) x 10<sup>6</sup> CAR-positive viable T cells/kg. PK parameters were estimated by NCA using Phoenix WinNonlin (version 8.0).
- Pharmacodynamics: soluble BCMA (sBCMA) level in serum was assessed using a ligand binding assay.
- Serum samples were evaluated for antibodies binding to cilta-cel (anti-drug antibodies or ADA)

Figure 3: Correlation Between Dose and Exposures (C<sub>max</sub> and AUC<sub>0-28d</sub>)



#### 2. Pharmacodynamics:

Following a single cilta-cel infusion, expansion and persistence of CAR-positive T cells coincided with decreases of sBCMA levels. Mean sBCMA levels reached nadir (i.e. LLOQ value <0.250 μg/L) on Day 56 for Cohort A(i) and on Day 100 for Cohorts B and C. Thereafter, sBCMA levels showed gradual increase in some patients; however, the reversed sBCMA levels (up to 1 year post cilta-cel infusion) in most patients were substantially lower than baseline sBCMA as shown in Figure 4.

Figure 4: Mean (+ SD) Serum sBCMA Concentration versus Time Profile

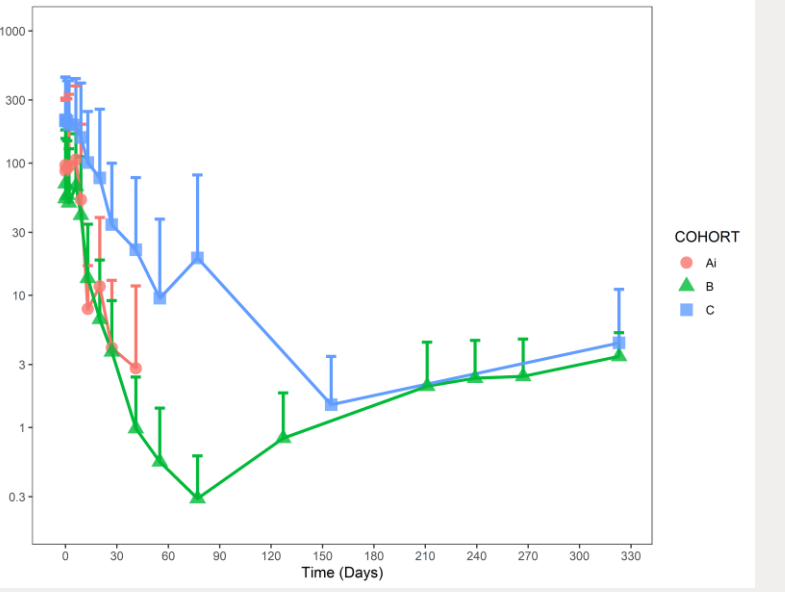
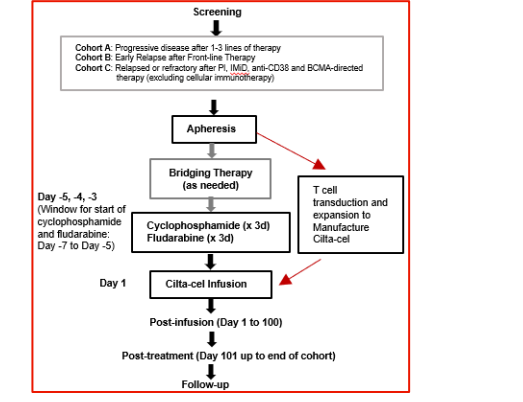


Figure 1. Schematic Overview of CARTITUDE-2 Study (Cohorts A to C)



#### 3. Immunogenicity:

The overall incidence of treatment emergent ADA to cilta-cel were 25.0% for Cohorts A(i) and C, 42% for Cohort B, with median ADA onset time ranging from 57 to 186 days across 3 cohorts.

As shown in Figure 5, given the high inter-individual variability and overlapping range of PK parameters, the overall cellular kinetics looks similar between ADA positive and ADA negative patients across 3 cohorts. In addition, the observed median ADA onset time for each cohort (57 to 186 days) was much later than the reported median t<sub>max</sub> (around 14 days), which also suggests there was no apparent impact of ADA on the cellular kinetic parameters (C<sub>max</sub> and AUC<sub>0-28d</sub>) of cilta-cel.

Figure 5: Correlation Between ADA Status and Exposures (C<sub>max</sub> and AUC<sub>0-28d</sub>)

