

## Expansion, Persistence and Pharmacodynamic Profile of ADI-001, a First-in-Class Allogeneic CD20-targeted CAR Gamma Delta T Cell Therapy, in Patients with Relapsed/Refractory Aggressive B-cell Non-Hodgkin's Lymphoma



Figure 1. ADI-001 is manufactured from qualified donor leukopheresis using a proprietary activating antibody, designed to expan the V $\delta$ 1 subset of  $\gamma\delta$  T cells. CAR expression is achieved following transduction with proprietary vectors in a proprietary scalable

## GLEAN: ADI-001 First-in-Human Study (CD20 CAR+ γδ T cells)

Dose limiting toxicity; DOR= Duration of response; ECOG= Eastern Cooperative Oncology Group; Flu= Fludarabine; GLEAN= Gamma Delta adoptive

therapy for Nhl-1; OS= Overall survival; PFS= Progression-free survival; R/R= Relapsed or refractory; TTP= Time to progression



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DL3 and DL4 showing mean Cmax, Area under the curve (AUC<sub>0-28</sub>), Tmax and Day 28 persistence. Mean DL3 and DL4 Cmax by ddPCR aligns with what was previously reported by internal flow cytometry analysis. AUC<sub>0-28</sub> was calculated using a model-based cellular kinetics analysis for CAR T cells for the first 28 days after infusion (units of days\*CAR copies/ug DNA). CAR cells/µL were derived from ddPCR and consider product VCN. LLOQ (64.5 CAR copies/µg) and LOD (49.5 CAR copies/µg) are shown (y-axis, dashed lines). N= 20 patients were assessed by ddPCR; N = 16 patients were assessed for AUC<sub>0-28</sub>. (C) Quantitative SNP profiling of cell product (AlloCell) was assessed in 24 patients and plotted as Mean ±SEM. (D) Whole blood (WB) from 24 patients across multiple timepoints was evaluated for the presence of CAR+ V $\delta$ 1+ T cells by flow cytometry and expressed as absolute cell counts (cells/µL blood). The lower limit of quantification (LLOQ) for this assay is 3.2 cells/µL blood with a lower limit of detection (LOD) of 0.18 cells/ $\mu$ L blood. CAR+ V $\delta$ 1+ T cells detected below LLOQ were graphed at ½ the LOD of 0.18 cells/ $\mu$ L blood; internal paired flow analysis subject to verification (E) The gross dynamic range of average ADI-001 genomes (measured by AlloCell and expressed as % ADI-001 cells per total nucleated cells) across all dose levels is shown versus the average recovery of patient CD8+ 1 cells by flow cytometry (N = 24). For all measures detection of ADI-001 returns to <LOD by the month 3 assessment.

# **Consolidation Dosing Regimen Results in 2<sup>nd</sup> Distinct Expansion**

Figure 4. Representative data from a single DL3 (3E8) DLBCL patient showing distinction of product expansion and detection for two products generated from two independent donor sources (Donor 1 and Donor 2). Following completion of a first cycle of enhanced lymphodepletion (eLD) and dosing with Donor 1-derived product, the patient received a second cycle of standard lymphodepletion (sLD) followed by dosing with Donor 2derived product. (A) Representative cellular kinetics by ddPCR of consolidation dosing exhibit a notable second expansion and exposure of ADI-001. (B) Results demonstrate unique detection of the first and second donor-sourced cell products using a single nucleotide polymorphism (SNP) 3genome analysis. Both product lots dosed in a single patient were dosed at DL3 (3E8)

DLT evaluable patients across multiple timepoints (Day 1 pre-infusion to Day 28) using flow cytometry and shown as mean ± SEM. (A) Subjects whose BOR was CR or PR (N=17) appear to have higher expansion and stimulation/proliferation of CAR+ cells (%Ki-67+ CAR+ cells) than those subjects with NR/PD as BOR (N=7); Mann-Whitney, p=0.005 for Day 10 and p=0.045 for Day 14. Time-points with less than a total cell count of 10 CAR+ Vd1+ ce were plotted as 0 for % of CAR+ Vd1+ cells and % of Ki67+ cells; internal paired flow analysis subject to verification. (B) Relationship of ADI-001 Cmax at DL2, DL3, DL3x2, and DL4 with serum cytokine/chemokine levels represented as mean ± SEM of the peak fold change over Day 1 pre-infusion Subjects whose Cmax was above 20,000 CAR copies/µg (High Cmax, blue bars, N=14) appear to have a higher mean peak fold change of serum cytokines/chemokines than subjects with Cmax below 20,000 CAR copies/ug (Low Cmax, red bars, N=6); Mann-Whitney, p=0.05 for IL-8, p=0.04 for IL 18, and p=0.02 for MCP-1.

### Production of Polyfunctional Serum Cytokines is Associated with **Clinical Response**



**Figure 7**. Relationship of ADI-001 serum cytokines/chemokines levels with BOR, represented as mean ± SEM of the peak fold change over Day 1 preinfusion. Subjects whose BOR was CR or PR (blue bars, N=17) appear to have higher polyfunctional cytokines/chemokine mean peak fold change than subjects who had a SD or PD as BOR (red bars; N=7).

Shared HLA Alleles	CR/PR Rate n/N (%)
0/8	7/8 (87.5)
1/8	7/11 (63.6)
≥2/8	3/5 (60.0)

- expansion and persistence of an allogeneic CAR  $\gamma\delta$  T cell therapy, ADI-001.
- At DL3 and DL4, ADI-001 Cmax and Tmax were comparable to, or exceeded those demonstrated by approved autologous CD19 CAR T therapies <sup>1-4</sup>. Additionally, DL4 demonstrated a notable mean Day 28 persistent exposure, measured as 16,533 copies/ $\mu$ g or approximately 27 cells/ $\mu$ L
- Higher ADI-001 Cmax and  $AUC_{0-28}$  were associated with clinical response.
- Higher ADI-001 Cmax associated with higher stimulatory and proliferative flux of CAR+ cells and coincided with production of functional serum cytokines/chemokines and increased clinical response.
- Endogenous cytokines, including SCF and IL-15, may contribute to both ADI-001 expansion dynamics and clinical response.
- Degree of shared HLA alleles between patient and ADI-001 product did not associate with differences in ADI-001 exposure or clinical response.
- Taken together ADI-001 demonstrates a robust exposure profile and is positively associated with both pharmacodynamic correlates and clinical response.

### References

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