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### NTLA-5001, a T Cell Product Candidate with CRISPR-Based Targeted Insertion of a High-Avidity, Natural, WT1-Specific TCR, Shows Efficacy in *In Vivo* Models of AML and ALL

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### Conflict of Interest Disclosure and Legal Disclaimer

- D. Liu, A. Prodeus, A. Becker, M. Foisey, I. Balwani, I. Dutta, Q. Zhang, M.S. Arredouani, M. McKee, L. Sepp-Lorenzino, and B.C. Schultes are employees of Intellia Therapeutics, Inc.
- F. Ciceri and C. Bonini are consultants for Intellia Therapeutics, Inc.
- C. Bonini received research funding from Intellia Therapeutics, Inc.

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### NTLA-5001: Engineered T Cell Product Candidate with Site-Specific Insertion of a High-Avidity, Natural TCR to WT1<sub>37-45</sub>

- Initial clinical indication: AML, the most common type of acute leukemia in adults with limited treatment options<sup>1</sup>
- HD1-TCR selected for NTLA-5001 based on high avidity and ability to specifically kill AML blasts



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### CRISPR Editing Overcomes Key Challenges in TCR T Cell Engineering

Traditional tgTCR addition



### CRISPR/Cas9 tgTCR replacement



### **Key Challenges**

- Mutagenesis risk from random lentiviral insertion
- Mixed expression of endogenous, tg and mispaired TCRs
- Unpredictable specificities of mispaired TCRs and GvHD risk
- Lower tgTCR expression per T cell leads to reduced efficacy

### **Our Solution**

- Precise replacement of endogenous TCR with tgTCR
- No insertional mutagenesis risk
- Reduced risk of normal tissue reactivity
- High tgTCR expression per T cell leads to higher efficacy

Proprietary Sequential Editing Process Enables Multiple Edits with High Efficiency and Cell Viability



WT1-TCR T Cells Engineered with Proprietary Process Have Enhanced Potency vs. Standard Methods



- ~90% of T cells carry desirable central and stem cell memory phenotype
- Higher cytokine secretion in response to WT1-presenting tumor cell lines in vitro
- Long-term proliferative capacity in a repeat-stimulation assay with tumor cells in vitro



### Proprietary Process Minimizes Translocations and Gene Expression Changes vs. Standard Electroporation Process

KromaTiD dGH<sup>™</sup> Assay Shows No Detectable Translocations With Sequential Editing



**Proprietary Process** 

# (780 genes) Unedited Proprietary Standard

Gene Expression Profiling<sup>1</sup>

-0.3

<sup>1</sup>NanoString nCounter® CAR-T characterization panel; EP: Electroporation

#### # Genes Differentially Regulated

6h	Std	Proprietary
P<0.05	195	75
>2 FC & P<0.05	34	4

- Standard process leads to gene expression changes in pathways of metabolism, memory and exhaustion
- Proprietary process
  results in minimal gene
  expression changes

### WT1-Specific TCR T Cells Are Efficacious *In Vivo* in Mice; Proprietary Process Enhances Anti-Tumor Activity



\*\*\*\* p<0.0001, WT1 TCR vs. MART1 Control TCR (2-way ANOVA) \* or \*\* p<0.05 or p<0.01, Standard vs. Proprietary Process (2-way ANOVA)

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Better tumor control with WT1 T cells manufactured using the proprietary process

### WT1-Specific TCR T Cells Are Efficacious *In Vivo* in Mice; Human IL2/IL15 Enhances Anti-Tumor Activity

ALL



**Days post Tumor Cell Injection** 



## In NOG mice producing human IL2 or IL15\* WT1 T cells

- Persist longer
- Control tumor burden more effectively

\*IL2 or IL15 levels comparable to levels in patients post lymphodepletion

9

### Conclusions and Next Steps for NTLA-5001

- Proprietary process enables efficient, scalable genome editing
  - ~99% KO efficiency of target genes; 50-70% in locus insertion of tgTCRs
  - Sequential editing with high viability and potential for safer products
  - Faster T cell expansion with favorable T cell memory phenotype leading to potentially reduced vein-to-vein time
  - Enhanced in vitro function and in vivo anti-tumor efficacy in mouse models
- Intellia is continuing to advance NTLA-5001 toward the clinic in 2021
  - Scale-up for clinical process completed

### NTLA-5001: Proposed first-in-human trial evaluating safety and activity in AML



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