

Bill, living with transthyretin amyloidosis, and his wife, Maura



Lipid nanoparticles (LNPs) as a superior CRISPR/Cas9 delivery modality for highly efficient multiplex gene editing of T cells for adoptive cell therapy

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CRISPR/Cas9 Genome Editing Can Power Next-Generation T Cell Therapies



To unlock the full potential of T cell therapies, we need to enable multiple gene edits with high efficiency and cell viability but low translocations



Intellia's proprietary LNP CRISPR/Cas9 delivery technology empowers safe and potent T cell therapies with multiple genome edits

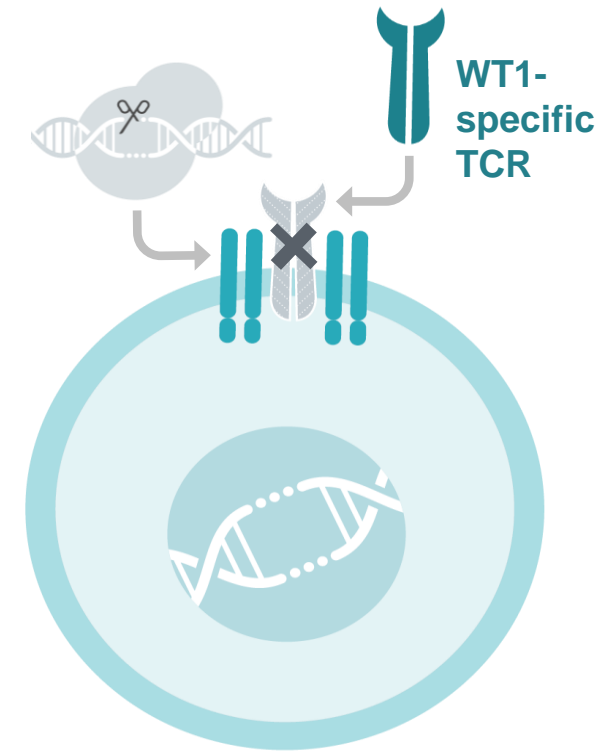
Our Goal: Engineer T Cell Therapies While Preserving Normal Cell Physiology

CRISPR-Based Gene Engineering

- **Precise** gene knockout (KO) and insertion
- **Site-specific insertion: TCR or CAR transgenes**
 - Homogeneous targeted insertion and expression
 - Improved cell drug quality and potency

LNP-based T cell transfection of CRISPR/Cas9

- Improved method for engineering autologous and allogeneic cell therapies
- **Modular** platform for sequential multiplex genome editing with minimal translocation rates
- **Scalable:** manufacturing process suitable for clinical development of autologous and allogeneic therapies

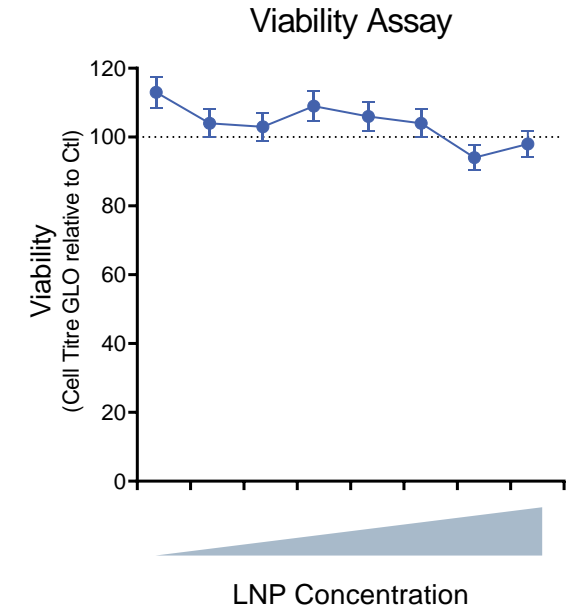
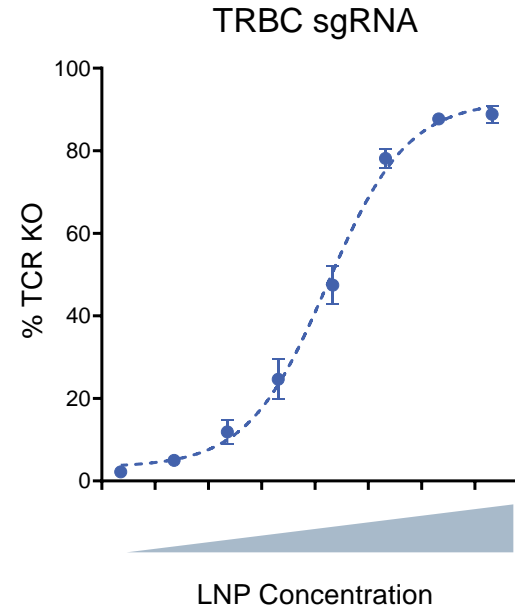
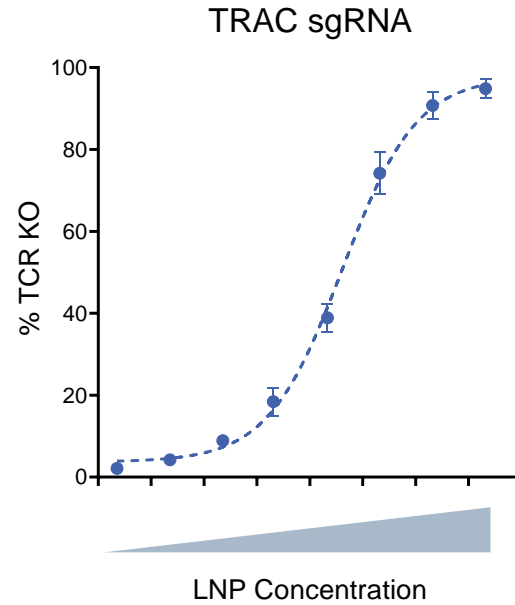
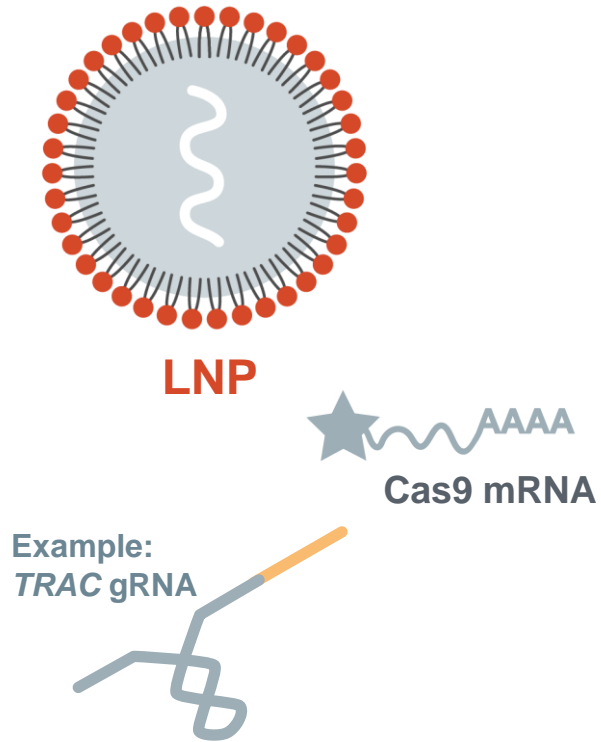


PRECISE • POTENT • PERSISTENT

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LNPs are an improved delivery system for
CRISPR/Cas9 T cell engineering

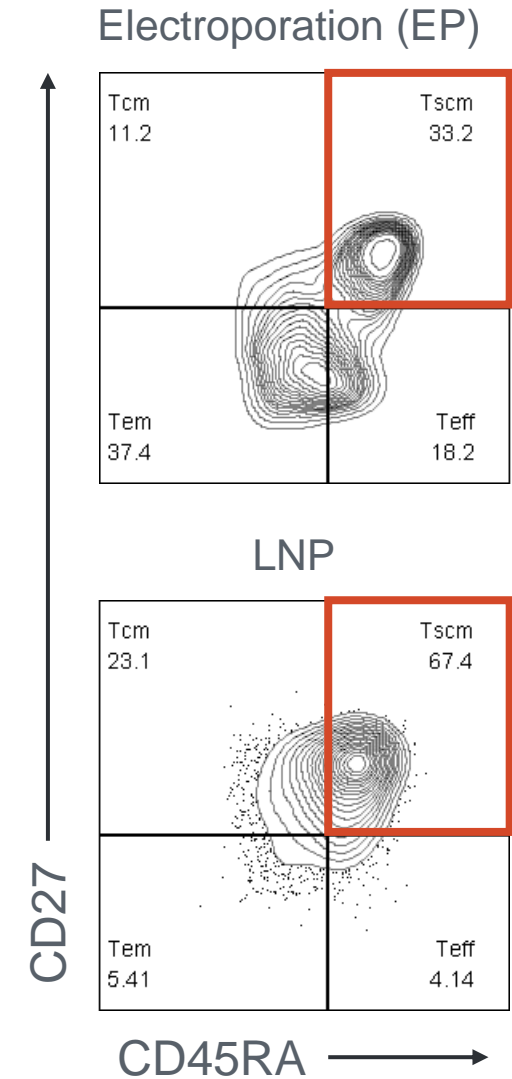
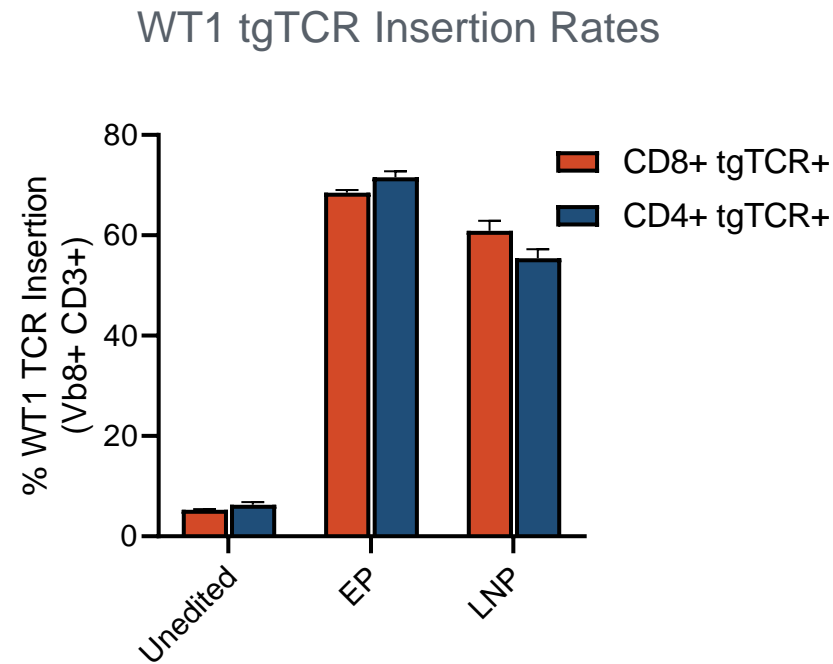
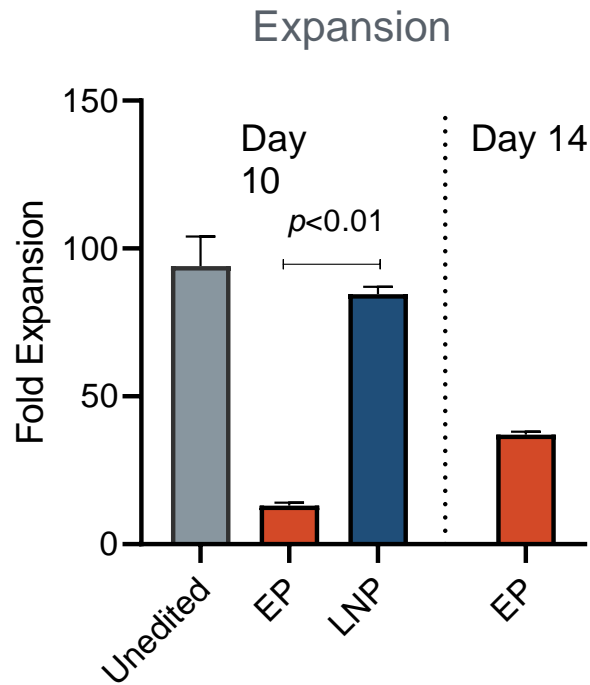
Dose-dependent Knockout of TRAC and TRBC by LNP Transfection of CRISPR/Cas9 in Primary Human T Cells



- LNP formulation designed for *ex vivo* delivery to T cells
- High KO rates without negative impact on T cell viability

Improved Expansion and Memory Phenotype of WT1 TCR-T cells using LNPs

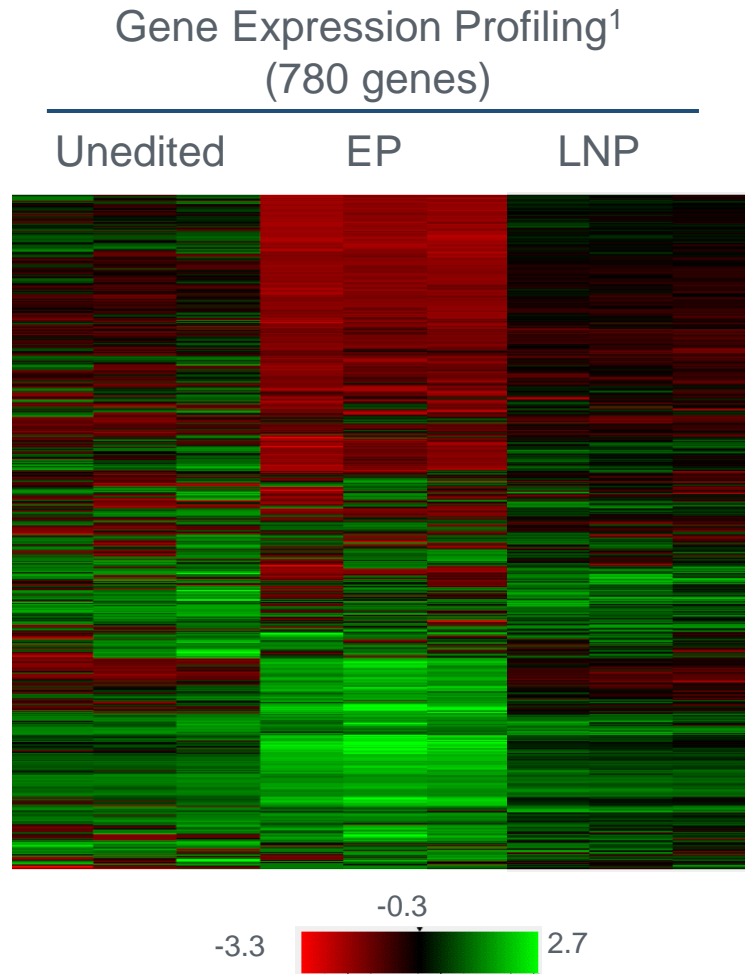
- T cells engineered with WT1 antigen specific TCR insertion in *TRAC locus*, combined with *TRBC KO*



LNP_[CRISPR/Cas9] delivery results in:

- More rapid expansion post-editing
- Favorable CD45RA⁺CD27⁺ memory phenotype
- Comparable editing rates

LNP_[CRISPR/Cas9] Transfection Minimizes Gene Expression Aberrations vs. Electroporation



¹NanoString nCounter® CAR-T characterization panel;
EP: Electroporation

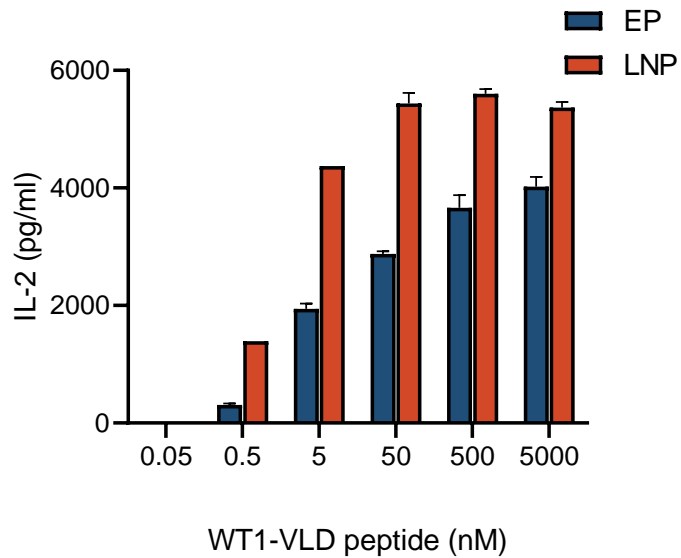
Genes Differentially Regulated

6h	EP	LNP
P<0.05	195	75
>2 FC & P<0.05	34	4

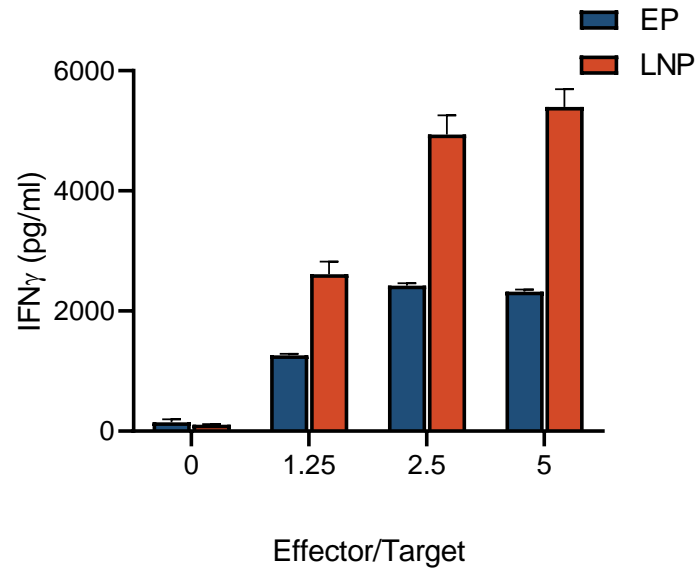
- **LNP_[CRISPR/Cas9] transfection showed minimal gene expression changes**
- **Electroporation results in global transcriptome aberrations in key pathways such as metabolism, memory and exhaustion**

WT1 TCR-T Cells Engineered with LNP_[CRISPR/Cas9] Method Have Enhanced Potency

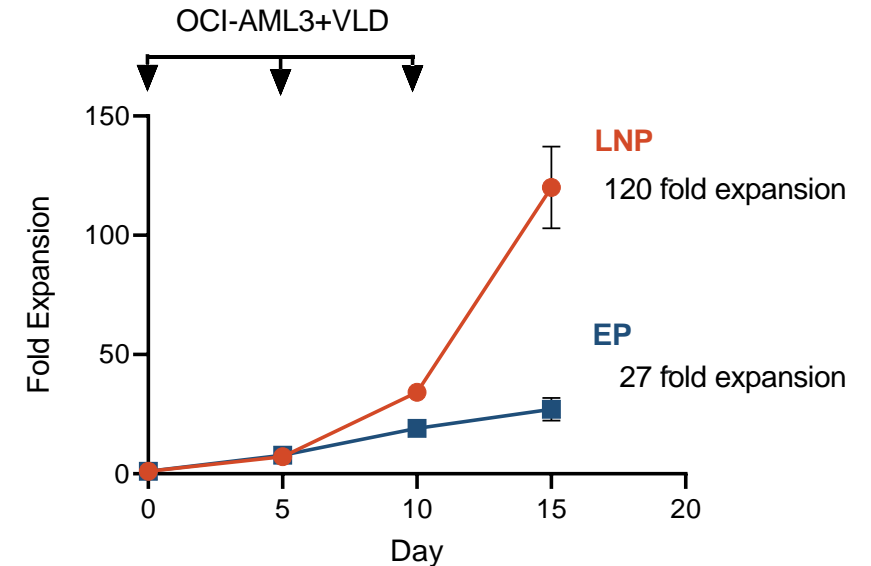
IL-2 Secretion: WT1-VLD Peptide Pulsed OCI-AML3



IFN- γ Secretion: K562-HLA-A*02:01⁺ Cells



Re-stimulation Stress Test: OCI-AML3 pulsed cells

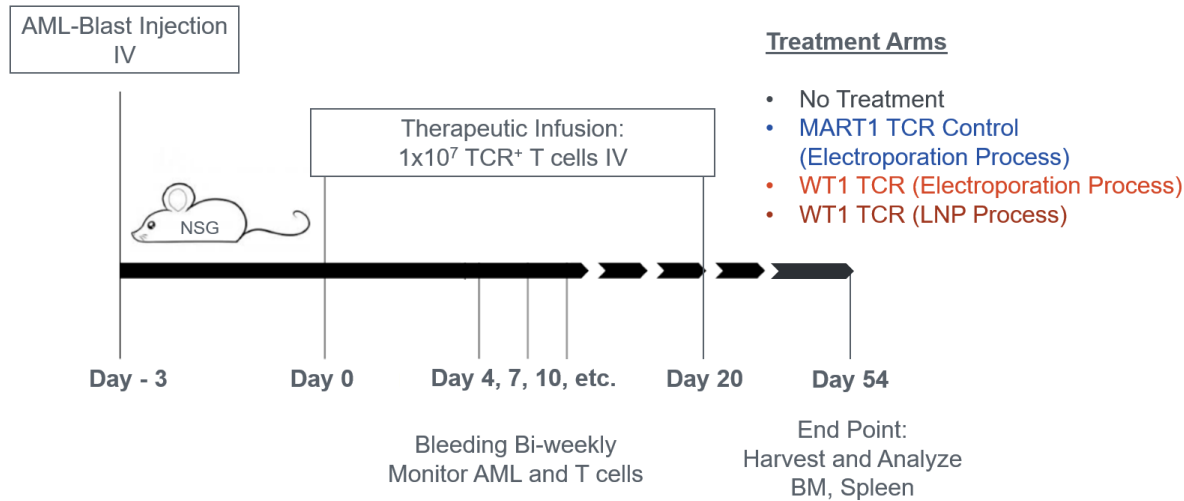


WT1 TCR-T cells engineered with LNP_[CRISPR/Cas9] delivery:

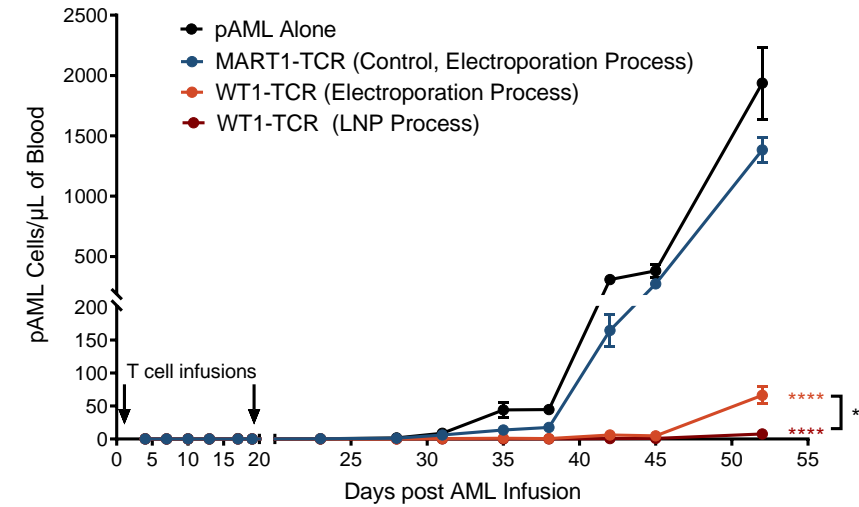
- Secrete more cytokines in response to WT1-presenting tumor cell lines
- Have long-term proliferative capacity in a repeat-stimulation assay with tumor cells

WT1-Specific TCR T Cells Are Efficacious *In Vivo* in Mice; LNP_[CRISPR/Cas9] engineered T Cells Possess Enhanced Anti-Tumor Activity

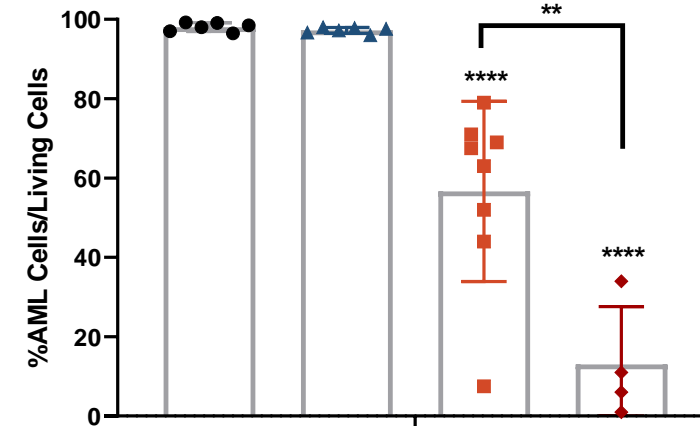
AML-PDX Model



Better tumor control with WT1 T cells manufactured with LNP_[CRISPR/Cas9] delivery vs. electroporation



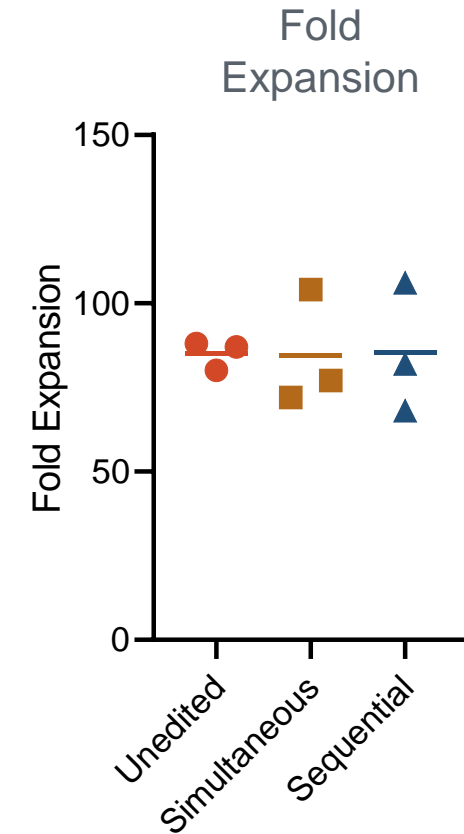
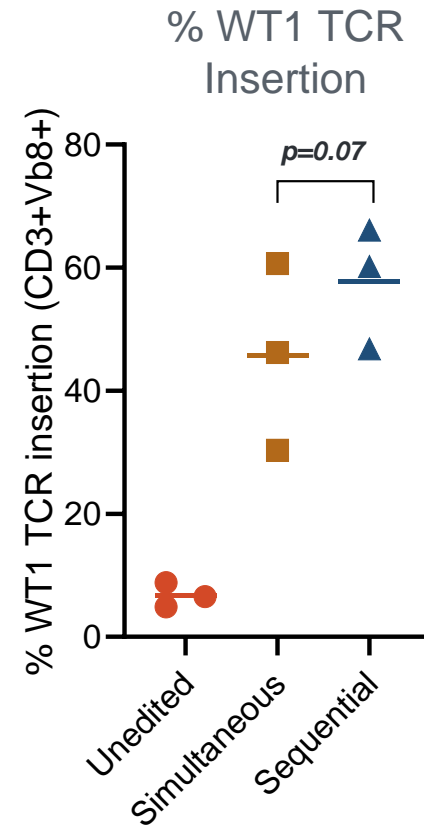
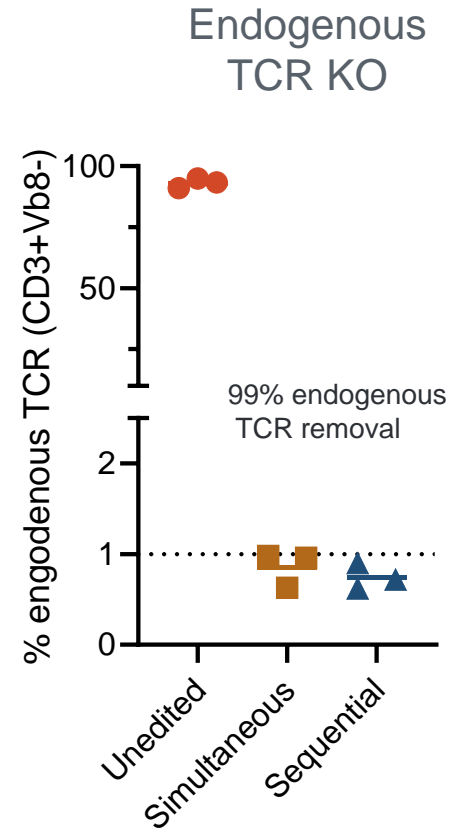
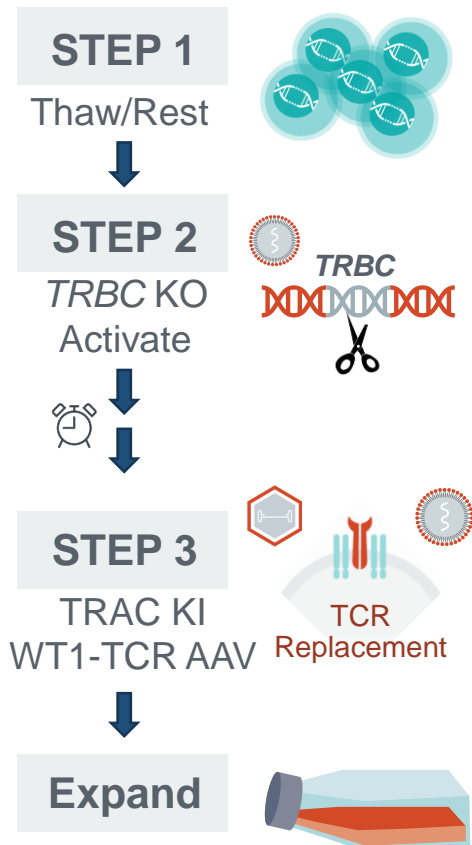
AML in Bone Marrow (Day 54, End of Study)



**** $p < 0.0001$, WT1 TCR vs. MART1 Control TCR (2-way ANOVA)
* or ** $p < 0.05$ or $p < 0.01$, EP vs. LNP Process (2-way ANOVA)



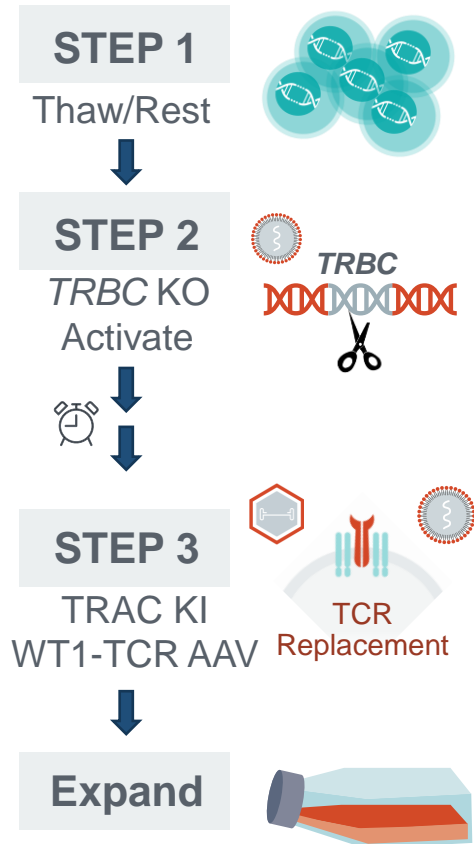
Sequential Editing of Target Genes Reduced Translocations Without Impacting Expansion Rates



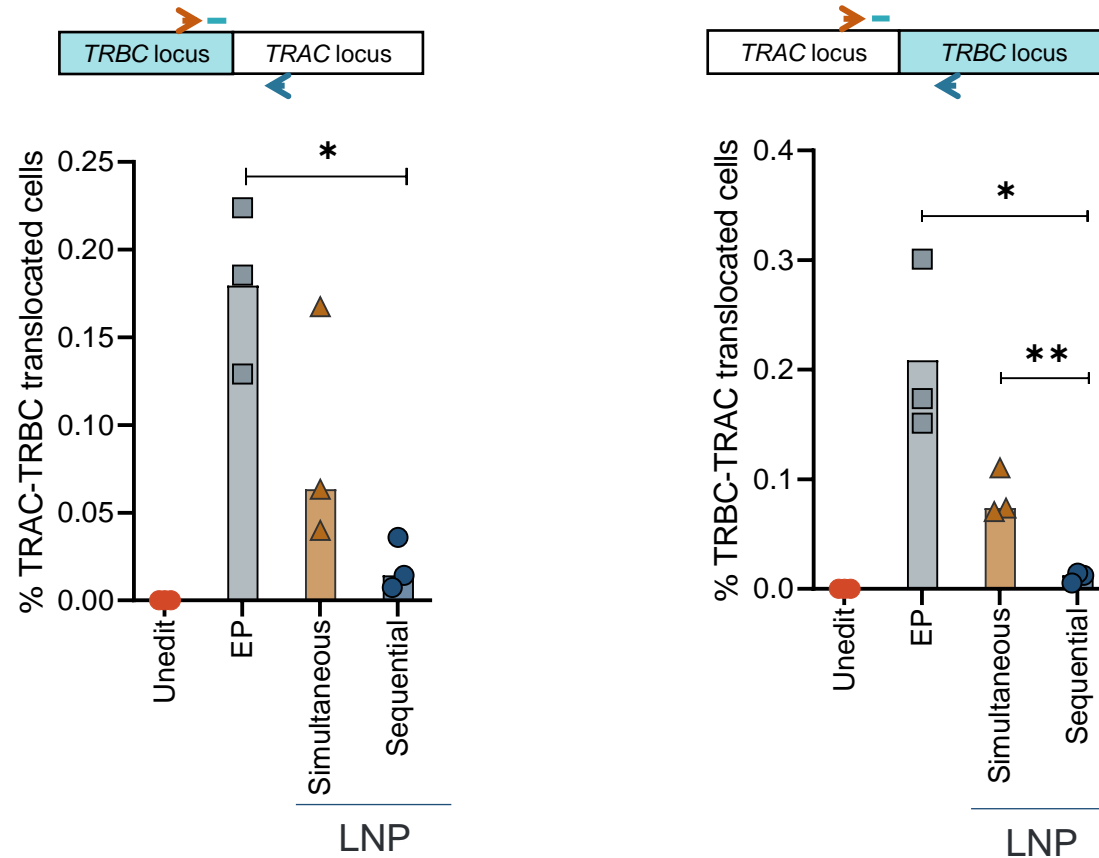
Sequential editing of *TRAC* and *TRBC* results in:

- High endogenous TCR KO (~99%) & insertion rates (~50%)
- High cell expansion

Sequential Editing of Target Genes Reduced Translocations Without Impacting Expansion Rates



ddPCR assay to detect *TRAC-TRBC* translocations



Sequential LNP [CRISPR/Cas9] editing results in a significant reduction in *TRAC-TRBC* translocations

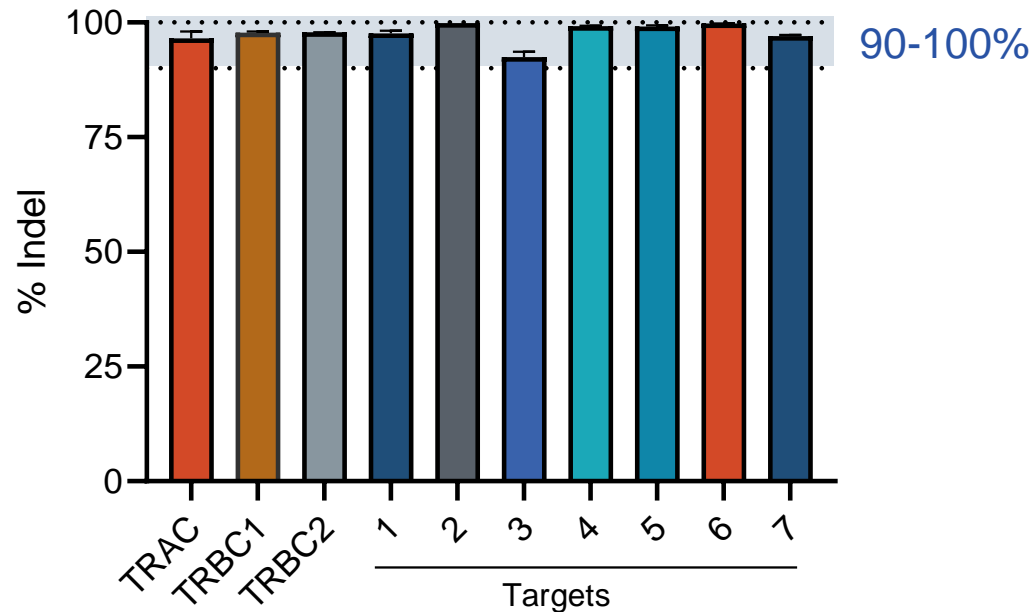
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Modular Processes for Allogeneic TCR and CAR-T cell therapies

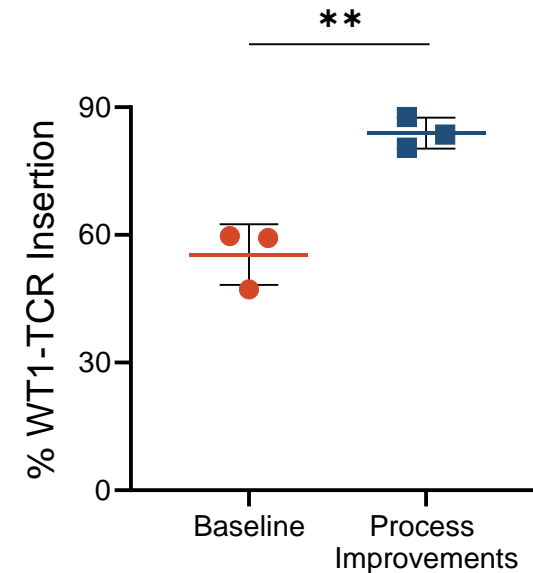
Building a Modular Platform for Next-Generation T Cell Therapies

- Allogeneic and solid tumor T cell therapies will require edits of numerous targets

90+% editing achieved via LNP delivery across numerous targets

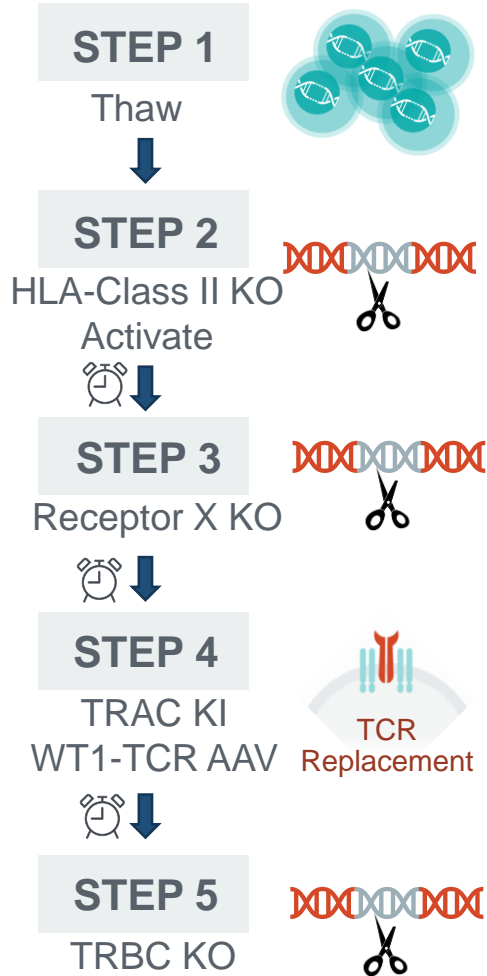


Process Improvements to Enhance Site-Specific Insertion Efficiency

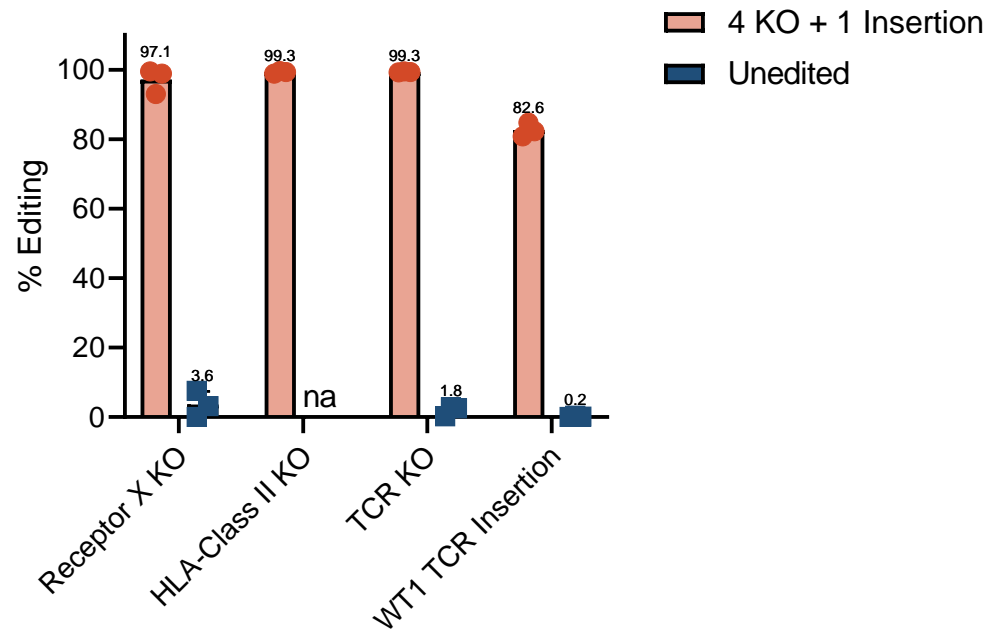


High insertion has been reproduced across several TCRs, CARs, and other transgenes

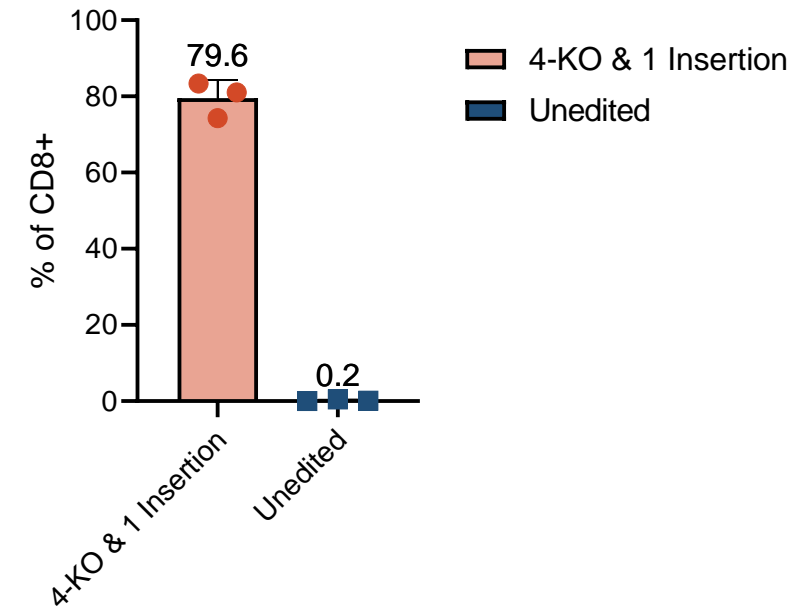
Multiplex CRISPR/Cas9 T Cell Editing: 4 KO Edits with a tgTCR Insertion



% Editing Across Loci



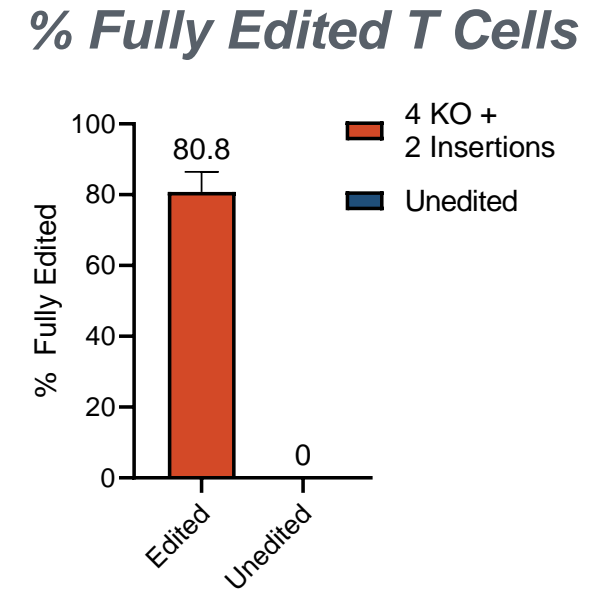
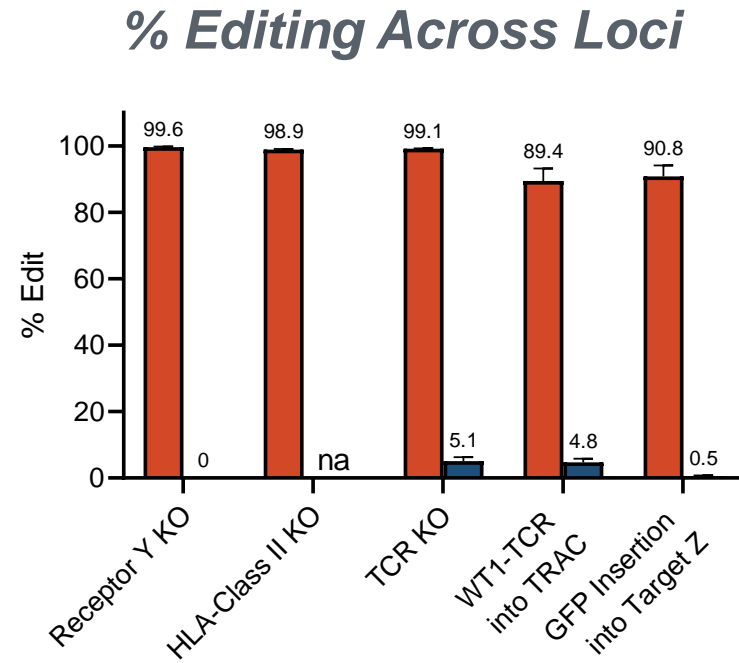
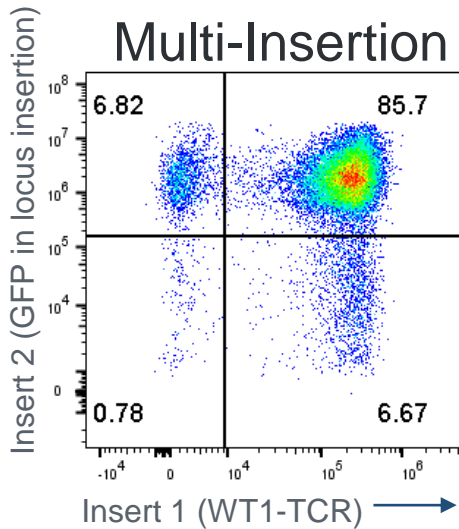
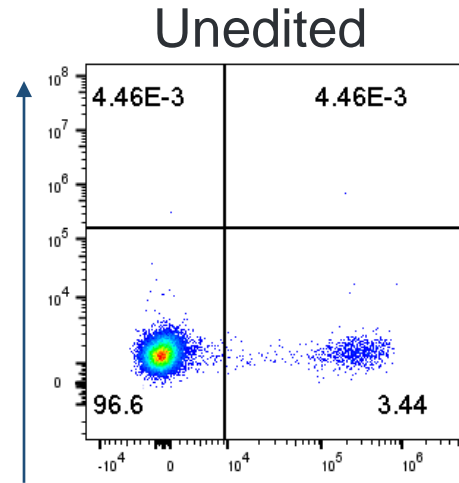
% Fully Edited T Cells



- Highly efficient editing across all targets while retaining high viability & expansion
- Yield of ~80% fully-edited T cells will support allo-T cell therapy candidates
- For more information on Intellia's allogeneic platform see Dr. Yong Zhang's presentation: Session 2c (OR18) Oct. 20th 10:45 CEST

Multiplex CRISPR/Cas9 T Cell Editing: 4 KO Edits with 2 Insertions

Dual site-specific insertion strategy enables co-expression of CAR/TCR construct and immune enhancing transgene



- >80% of cells have insertion of both the TCR and GFP transgene as well as complete editing of 4 other KO targets
- Cells retained high viability
- Modular platform for insertion of T cell supporting transgenes

Key Takeaways

1. LNPs represent a superior CRISPR/Cas9 delivery system for *ex vivo* T cell editing

- Improvements on cell viability and expansion vs. standard electroporation methods
- Improved phenotype and potency of WT1 TCR-T cells
- Sequential editing with LNP transfections minimizes translocations
- NTLA-5001: First-in-human clinical study with LNP-edited CRISPR/Cas9 T cells expected to begin screening in 2021 (NCT05066165)

2. Modular cell engineering platform for highly multiplex editing of T cells

- High editing across numerous targets
- Process improvements led to 80-90+% CAR and TCR insertion rates
- Efficient sequential multiplex editing of up to 5 genes including multiple insertion and KOs

Robust and modular platform technology enables allogeneic and solid tumor candidates with highly multiplex gene edits