



Intellia is Leading the Genome Editing Revolution

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

Intellia
THERAPEUTICS

Corporate Overview
MARCH 2022

Intellia Therapeutics' Legal Disclaimer

This presentation contains “forward-looking statements” of Intellia Therapeutics, Inc. (“Intellia”, “we” or “our”) within the meaning of the Private Securities Litigation Reform Act of 1995. These forward-looking statements include, but are not limited to, express or implied statements regarding Intellia’s beliefs and expectations regarding: the safety, efficacy and advancement of our clinical programs for NTLA-2001 for the treatment of transthyretin amyloidosis, NTLA-2002 for the treatment of hereditary angioedema, and NTLA-5001 for the treatment of acute myeloid leukemia pursuant to its clinical trial applications (“CTA”) and IND submissions, including the expected timing of data releases, regulatory filings, and the initiation and completion of clinical trials; the advancement of development candidates including NTLA-3001 for the treatment of alpha-1 antitrypsin deficiency (AATD)-associated lung disease, NTLA-2003 for AATD-associated liver disease, and NTLA-6001 for CD30+ lymphomas; the ability to generate data to initiate clinical trials and the timing of CTA and IND submissions; the expansion of its CRISPR/Cas9 technology and related technologies, including manufacturing and delivery technologies, to advance additional development candidates; the ability to maintain and expand our related intellectual property portfolio, and avoid or acquire rights to valid intellectual property of third parties; the ability to demonstrate our platform’s modularity and replicate or apply results achieved in preclinical studies, including those in our NTLA-2001, NTLA-5001, and NTLA-2002 programs, in any future studies, including human clinical trials; its expectations of Rewrite Therapeutics, Inc.’s (“Rewrite”) DNA writing to advance additional novel platform capabilities; the ability to optimize the impact of our collaborations on our development programs, including, but not limited to, our collaboration with Regeneron Pharmaceuticals, Inc., including our co-development programs for hemophilia A and hemophilia B, with AvenCell Therapeutics, Inc. (“AvenCell”) for the development of universal CAR-T cell therapies, with SparingVision SAS (“SparingVision”) for the development of ophthalmic therapies, with Kyverna Therapeutics, Inc. (“Kyverna”) for the development of KYV-201, and with ONK Therapeutics Ltd. (“ONK”) for the development of engineered NK cell therapies; and the potential timing and receipt of future milestones and royalties, or profits, as applicable, based on our license, collaboration and, if applicable, co-development agreements with Regeneron, Novartis Institutes for Biomedical Research, AvenCell, SparingVision, Kyverna, and ONK; the timing of regulatory filings and clinical trial execution, including dosing of patients, regarding our development programs; the potential commercial opportunities, including value and market, for our product candidates; our use of capital and other financial results during 2022; and our ability to fund operations beyond the next 24 months.

Any forward-looking statements in this presentation are based on management’s current expectations and beliefs of future events, and are subject to a number of risks and uncertainties that could cause actual results to differ materially and adversely from those set forth in or implied by such forward-looking statements. These risks and uncertainties include, but are not limited to: risks related to our ability to protect and maintain our intellectual property position; risks related to valid third party intellectual property; risks related to our relationship with third parties, including our licensors and licensees; risks related to the ability of our licensors to protect and maintain their intellectual property position; uncertainties related to regulatory agencies’ evaluation of regulatory filings and other information related to our product candidates; uncertainties related to the authorization, initiation and conduct of studies and other development requirements for our product candidates; risks related to the development and/or commercialization of any of Intellia’s or its collaborators’ product candidates, including that they may not be successfully developed and commercialized; risks related to the results of preclinical or clinical studies, including that they may not be positive or predictive of future results; risks related to the Rewrite acquisition, including that it may not result in novel platform capabilities; risks related to Intellia’s reliance on collaborations, including that its collaborations with Regeneron, AvenCell, SparingVision, Kyverna, ONK or its other collaborations will not continue or will not be successful. For a discussion of these and other risks and uncertainties, and other important factors, any of which could cause Intellia’s actual results to differ from those contained in the forward-looking statements, see the section entitled “Risk Factors” in Intellia’s most recent Annual Report on Form 10-K as well as discussions of potential risks, uncertainties, and other important factors in Intellia’s other filings with the Securities and Exchange Commission (“SEC”). All information in this presentation is as of the date of the release, and Intellia undertakes no duty to update this information unless required by law.



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Intellia is Leading the Genome Editing Revolution

Transforming lives of people with severe diseases by developing curative genome editing treatments



▶ **Full-Spectrum Strategy**
Robust R&D engine to develop *in vivo* and *ex vivo* therapies for diseases with high unmet need

▶ **Modular Solutions**
Focused on building differentiated technology with broad applicability that can be applied to future candidates

▶ **Setting the Standard**
Extensive characterization for potent and highly specific editing

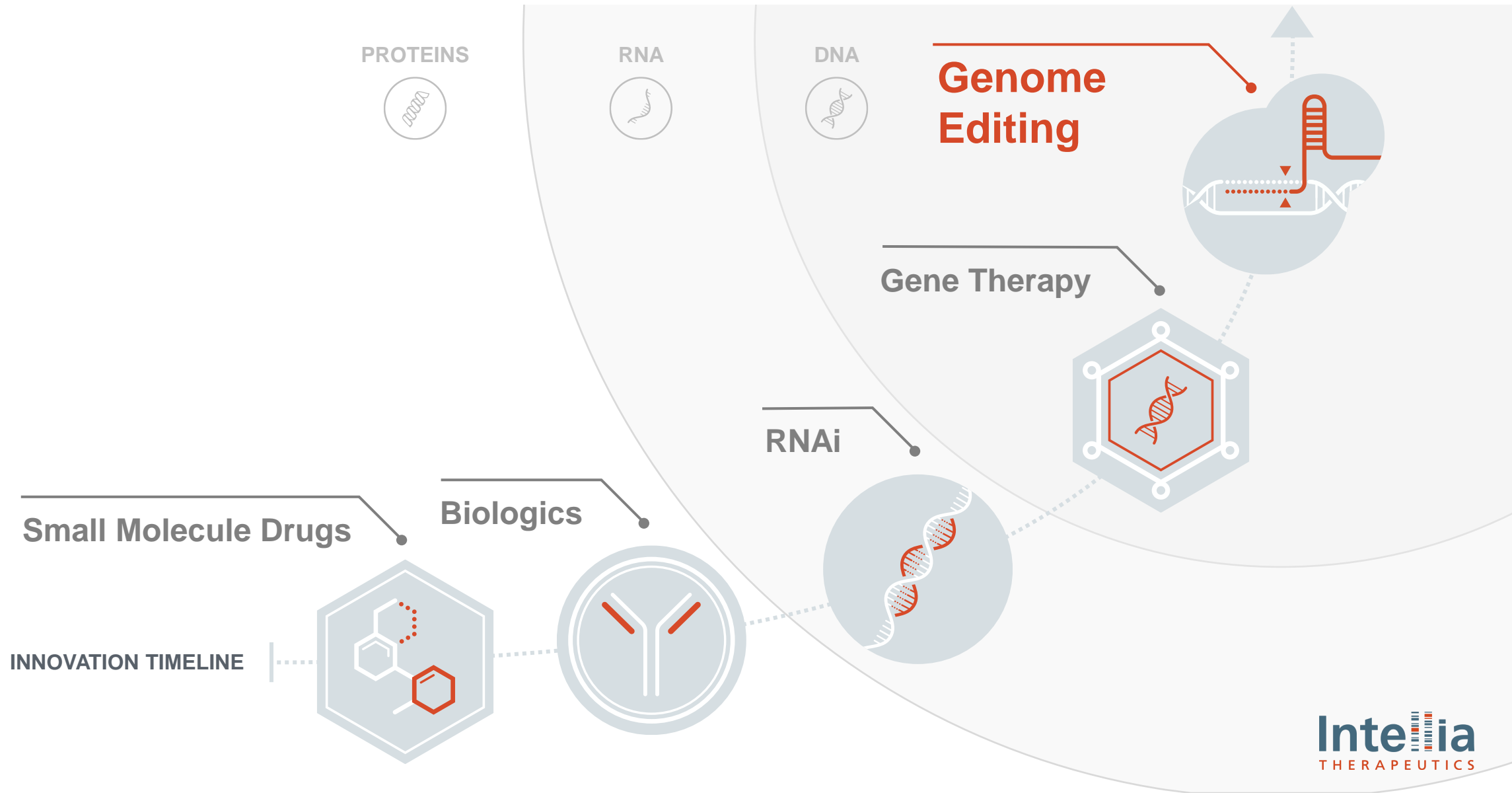
▶ **Applying Novel Tools**
Building an array of editing tools and delivery modalities for therapeutic application

▶ **Leaders of the Field**
First company to demonstrate initial safety and efficacy of *in vivo* genome editing in a clinical study

**Unsurpassed
Genome Editing
Pipeline**

**World-class
Genome Editing
Toolbox**

Therapeutic Strategies to Treat Life-Threatening Diseases Have Advanced Over Time



Power of CRISPR:

Nobel-Prize Winning Genome Editing Technology



Precise and modular approach for editing the genome



Potential for life-long effect following one-time treatment



Locates a genetic sequence to make a permanent change



Overcomes key limitations of gene and RNAi therapies



High level of specificity to make one or multiple edits



Provides foundational capabilities for derivative tools

In Vivo Leader: First to Demonstrate Systemic CRISPR Gene Editing in Humans



The NEW ENGLAND
JOURNAL of MEDICINE

August 5, 2021

CRISPR-Cas9 In Vivo Gene Editing for Transthyretin Amyloidosis

Julian D. Gillmore, M.D., Ph.D., Ed Gane, M.B., Ch.B., Jorg Taubel, M.D.,
Justin Kao, M.B., Ch.B., Marianna Fontana, M.D., Ph.D.,
Michael L. Maitland, M.D., Ph.D., Jessica Seitzer, B.S., Daniel O'Connell, Ph.D.,
Kathryn R. Walsh, Ph.D., Kristy Wood, Ph.D., Jonathan Phillips, Ph.D.,
Yuanxin Xu, M.D., Ph.D., Adam Amaral, B.A., Adam P. Boyd, Ph.D.,
Jeffrey E. Cehelsky, M.B.A., Mark D. McKee, M.D., Andrew Schiermeier, Ph.D.,
Olivier Harari, M.B., B.Chir., Ph.D., Andrew Murphy, Ph.D.,
Christos A. Kyratsous, Ph.D., Brian Zambrowicz, Ph.D., Randy Soltys, Ph.D.,
David E. Gutstein, M.D., John Leonard, M.D., Laura Sepp-Lorenzino, Ph.D., and
David Lebwohl, M.D.

Science
JOURNALS

***“CRISPR injected into the blood treats
a genetic disease for the first time”***

FT
FINANCIAL
TIMES

***“CRISPR gene-editing ‘revolution’
treats internal organ for first time”***

USA
TODAY

***“‘It’s a wow’: New CRISPR gene-editing
success holds promise for treating many
genetic diseases with a single dose”***

nature

***“Landmark CRISPR trial shows
promise against deadly disease”***

Building a Full-Spectrum Genome Editing Company

CRISPR-based Modular Platform

EMPLOY NOVEL EDITING AND DELIVERY TOOLS

In Vivo
**CRISPR is
the therapy**

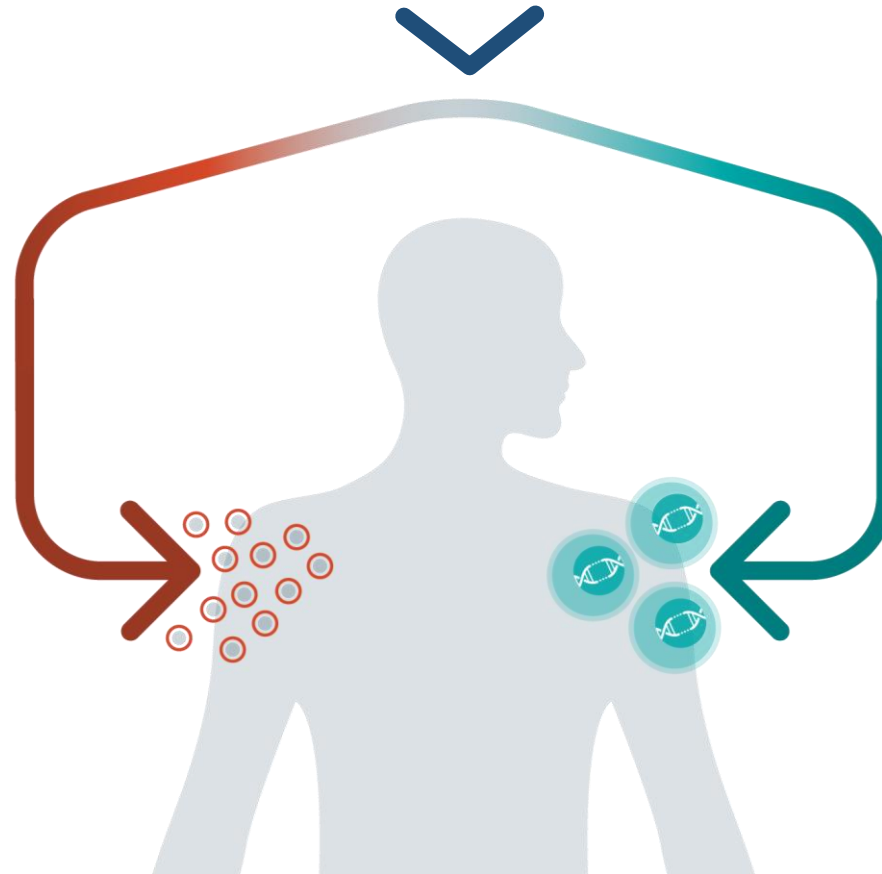
FIX THE TARGET GENE

Genetic diseases

Ex Vivo
**CRISPR creates
the therapy**

REWIRE & REDIRECT CELLS

Immuno-oncology
Autoimmune diseases



2022 and Beyond: Key Expected Milestones

In Vivo

NTLA-2001
ATTR

- ☐ Present additional interim clinical data from Phase 1 study later in 2022
- ☐ Complete enrollment of Phase 1 study for both ATTRv-PN and ATTR-CM in 2022

NTLA-2002
HAE

- ☐ Present interim data from Phase 1/2 study in 2H 2022

NTLA-3001
AATD

- ☐ Plan to file an IND or IND-equivalent in 2023

Ex Vivo

NTLA-5001
AML








- ☐ Continue to enroll patients in Phase 1/2a study in 2022

Platform Innovation














**Research
and Platform
Advancements**

- ☒ Nominated NTLA-6001, an allo-CAR-T cell therapy for CD30+ lymphomas
- ☒ Nominated NTLA-2003, an *in vivo* knockout candidate for AATD-associated liver disease
- ☐ Advance at least one additional *in vivo* development candidate by end of 2022
- ☐ Advance additional novel platform capabilities in 2022¹

In Vivo Development Pipeline Fueled by Robust Research Engine

PROGRAM	APPROACH	Research	IND-Enabling	Early-Stage Clinical	Late-Stage Clinical	PARTNER
<i>In Vivo: CRISPR <u>is</u> the therapy</i>						
NTLA-2001: Transthyretin Amyloidosis	Knockout					LEAD Inteilia* REGENERON
NTLA-2002: Hereditary Angioedema	Knockout					Inteilia
NTLA-2003: AATD-Liver Disease	Knockout					Inteilia
NTLA-3001: AATD-Lung Disease	Insertion					Inteilia
Hemophilia B	Insertion					LEAD Inteilia REGENERON*
Hemophilia A	Insertion					LEAD Inteilia REGENERON*
Research Programs	Knockout, Insertion, Consecutive Edits					Inteilia
Research Programs	Various					Inteilia REGENERON** SPRINGVISION

Ex Vivo Development Pipeline Fueled by Robust Research Engine

PROGRAM	APPROACH	Research	IND-Enabling	Early-Stage Clinical	Late-Stage Clinical	PARTNER
Ex Vivo: CRISPR <u>creates</u> the therapy						
OTQ923 / HIX763: Sickle Cell Disease	HSC					
NTLA-5001: Acute Myeloid Leukemia	WT1-TCR					
NTLA-6001: CD30+ Lymphomas	Allo CAR-T					
Solid Tumors	WT1-TCR					
Allo Undisclosed	Undisclosed					
Research Programs	Allo Universal CAR-T					
Other Novartis Programs	CAR-T, HSC, OSC	Undisclosed				

***Milestones & royalties only



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In Vivo

CRISPR is the therapy

GENETIC DISEASES

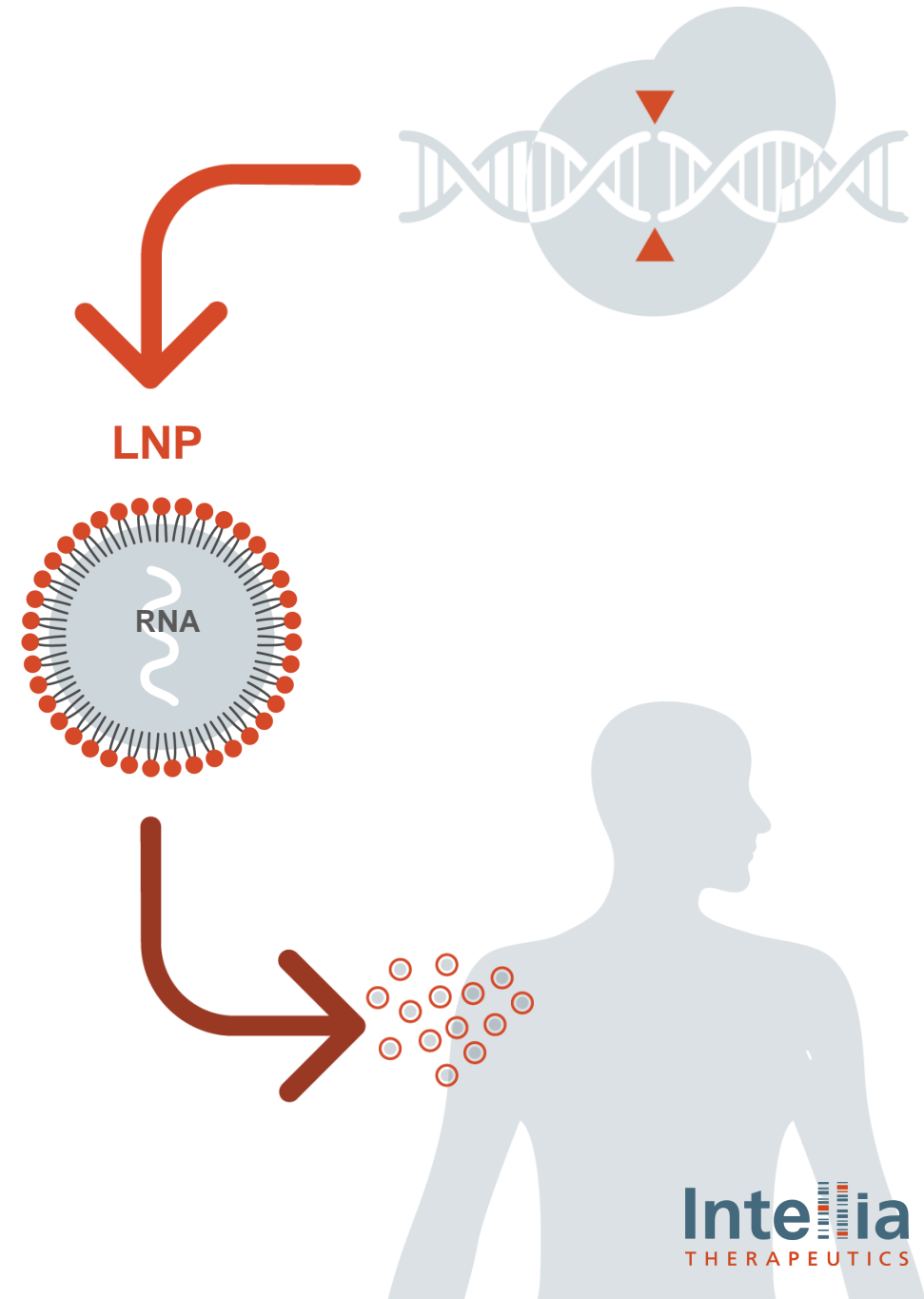
Strategic Advantages:

Potential curative therapy from single dose

Systemic non-viral delivery of CRISPR/Cas9 provides transient expression and potential safety advantages

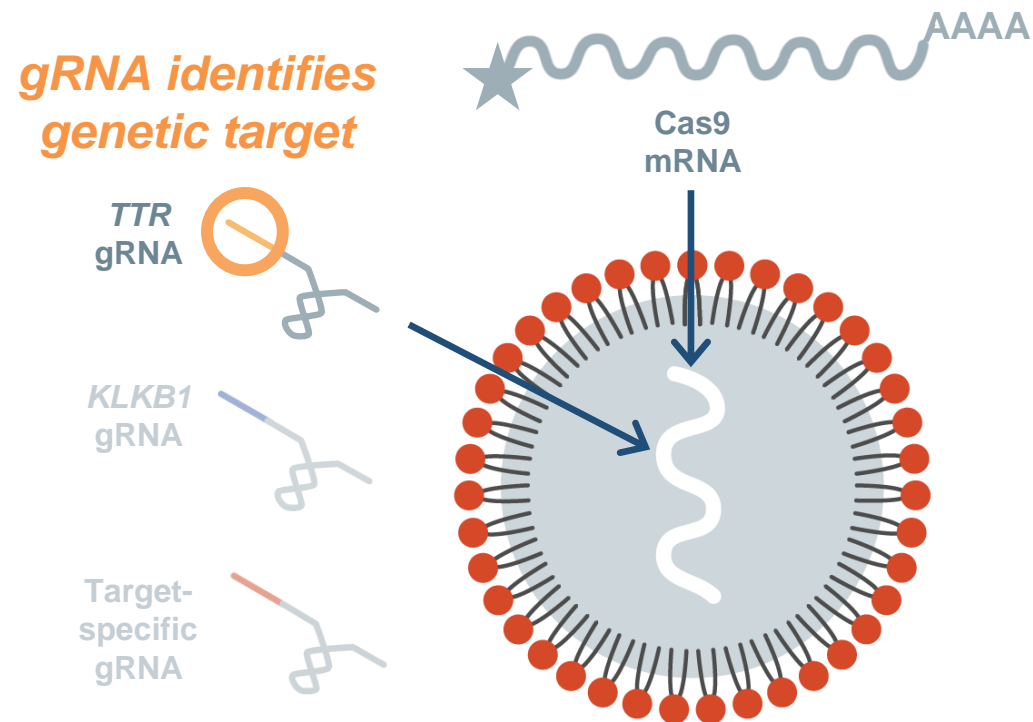
Permanent gain of function with targeted gene insertion

Capable of delivering to multiple tissue types for various therapeutic applications



Modular Delivery Platform Enables Rapid and Reproducible Path to Clinical Development

LNP Delivery System:



Key Advantages of LNP Delivery

- Clinically-proven delivery to liver
- Large cargo capacity
- Transient expression
- Biodegradable
- Low immunogenicity
- Well-tolerated
- Redosing capability
- Scalable synthetic manufacturing
- Tunable to other tissues

NTLA-2001 for Transthyretin (ATTR) Amyloidosis



- Caused by accumulation of misfolded transthyretin (TTR) protein, which affects nerves, heart, kidneys and eyes
- Chronic dosing is required with current treatments

OUR APPROACH

Knock out *TTR* gene with a single dose

- Reduce wild-type and mutant TTR protein
- Aims to address polyneuropathy and cardiomyopathy

KEY ADVANTAGES

- Potential to halt and reverse disease
- Potential “one-and-done” treatment
- Expect lifelong, stable TTR reduction

50K

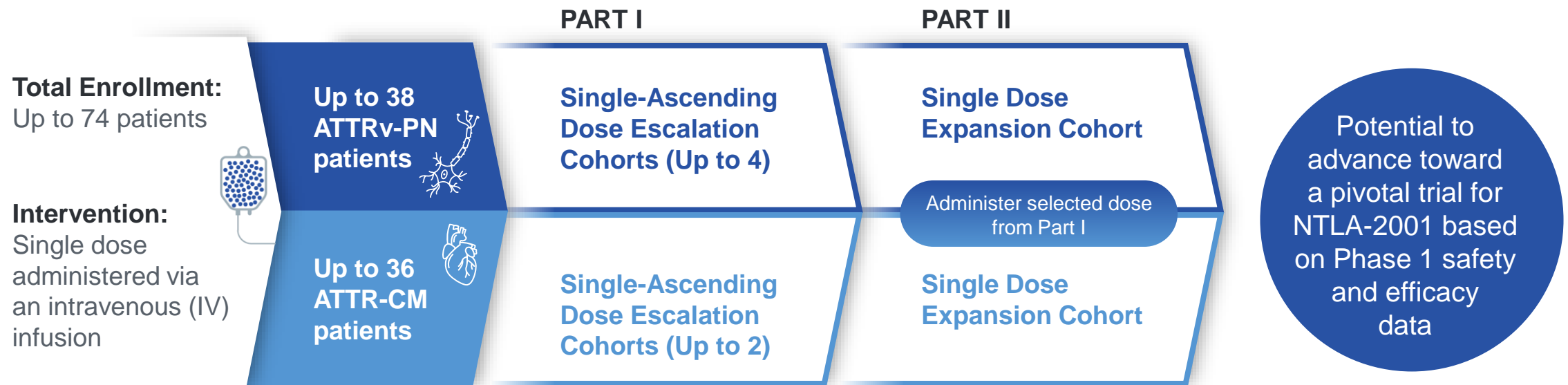
ATTRv patients worldwide¹

~200-500K

ATTRwt patients worldwide²

NTLA-2001 Expanded Phase 1 Study

Two-part, open-label, multi-center study in adults with hereditary ATTR with polyneuropathy (ATTRv-PN) or ATTR amyloidosis with cardiomyopathy (ATTR-CM)



PRIMARY OBJECTIVES

Evaluate safety, tolerability, PK and PD

- Measure serum TTR levels

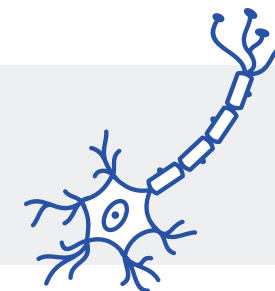
SECONDARY OBJECTIVES

Evaluate efficacy on clinical measures of:

- Neurologic function in subjects with ATTRv-PN
- Cardiac disease in subjects with ATTR-CM

NTLA-2001 Phase 1 Study: Polyneuropathy Arm

Hereditary transthyretin amyloidosis with polyneuropathy (ATTRv-PN)



Intervention:

Single dose
administered via an
intravenous (IV) infusion



PART I – DOSING COMPLETE Single-Ascending Dose

1.0 mg/kg (n=6)

0.7 mg/kg (n=3)

0.3 mg/kg (n=3)

0.1 mg/kg (n=3)

PART II – INITIATE IN Q1 2022 Single Dose Expansion Cohort

N = 8 subjects
Administer 80 mg fixed dose

PRIMARY OBJECTIVES

Evaluate safety, tolerability, PK and PD

- Measure serum TTR levels

SECONDARY OBJECTIVES

Evaluate efficacy on clinical measures of neurologic function

- Neuropathic impairment endpoints include NIS (Part 1 and 2) and mNIS+7 (Part 2 only)

NTLA-2001 Phase 1 Study: Cardiomyopathy Arm

Hereditary transthyretin amyloidosis with cardiomyopathy (ATTRv-CM)
or wild-type cardiomyopathy (ATTRwt-CM), NYHA Class I - III



Intervention:

Single dose
administered via an
intravenous (IV) infusion



PART I Single-Ascending Dose

1.0 mg/kg

0.7 mg/kg

PART II Single Dose Expansion Cohort

N = 12 subjects
Administer dose derived
from Part I data

PRIMARY OBJECTIVES

Evaluate safety, tolerability, PK and PD

- Measure serum TTR levels

SECONDARY OBJECTIVES

Evaluate efficacy on clinical measures of cardiac disease

- Cardiac imaging, biomarkers, cardiopulmonary exercise test, 6MWT

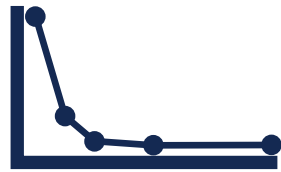
A decorative vertical bar on the left side of the slide, composed of a series of horizontal stripes of varying lengths and colors (white and orange) against a dark blue background.

Interim Clinical Trial Results from Ongoing Phase 1 Study of NTLA-2001 in ATTRv-PN Patients

Data disclosed on February 28, 2022

First-in-human evidence of deep, consistent, and durable TTR reductions following *in vivo* CRISPR-based gene editing

Single systemic administration of NTLA-2001 resulted in deep reductions in serum TTR



**93% mean reduction
at 1.0 mg/kg by Day 28**



**6 out of 6 patients at 1.0 mg/kg
achieved >80% reductions in TTR**

- Durable reductions in serum TTR observed over follow-up period
 - Consistent with animal data supporting potential lifelong TTR suppression
- Generally well tolerated: predominately mild adverse events
- A fixed dose of 80 mg has been selected for evaluation in Part II pending regulatory feedback

These data further support and extend early findings from this pioneering trial demonstrating the promise of CRISPR-based *in vivo* gene editing in humans

NTLA-2001 was generally well tolerated across all dose levels

- **Across all dose levels, the most frequent adverse events* were headache, infusion-related reactions, back pain, rash†, and nausea**
 - Majority of adverse events were mild in severity with 73% (n=11) of patients reporting a maximal adverse event severity of Grade 1
 - All patients received a complete study dose of NTLA-2001
 - All infusion-related reactions were considered mild, resolving without clinical sequelae
- **A single related Grade 3 event (SAE) of vomiting was reported at the 1.0 mg/kg dose in a patient with underlying gastroparesis**
 - 1.0 mg/kg dose level expanded per protocol to 6 patients to further characterize safety and PD
- **No clinically significant laboratory findings observed**
 - Transient Grade 1 liver enzyme elevations observed
- **Maximally tolerated dose was not reached**

Median follow-up for all subjects is 6 months

* Related and unrelated events in more than 2 patients

† Date of onset D6–D145; all mild in severity

PD, pharmacodynamics; SAE, serious adverse event

Data disclosed on February 28, 2022

This slide includes data for investigational products not yet approved by regulatory authorities

Majority of adverse events were mild in severity

Parameter	0.1 mg/kg n=3			0.3 mg/kg n=3			0.7 mg/kg n=3			1 mg/kg n=6			All n=15		
	Gr. 1	Gr. 2	Gr. 3	Gr. 1	Gr. 2	Gr. 3	Gr. 1	Gr. 2	Gr. 3	Gr. 1	Gr. 2	Gr. 3	Gr. 1	Gr. 2	Gr. 3
Patients with at least one TEAE	3	–	–	3	–	–	2	–	1*	3	2	1†	11	2	2
Headache	2	–	–	–	–	–	2	–	–	3	–	–	7	–	–
Infusion-related reaction	1	–	–	–	–	–	2	–	–	4	–	–	7	–	–
Back pain	1	–	–	–	–	–	2	1	–	1	–	–	4	1	–
Rash	1	–	–	–	–	–	–	–	–	3	–	–	4	–	–
Nausea	1	–	–	–	–	–	1	–	–	1	–	–	3	–	–

Adverse events reported in more than 2 patients

Patients counted once per row, per dose level, as highest grade reported

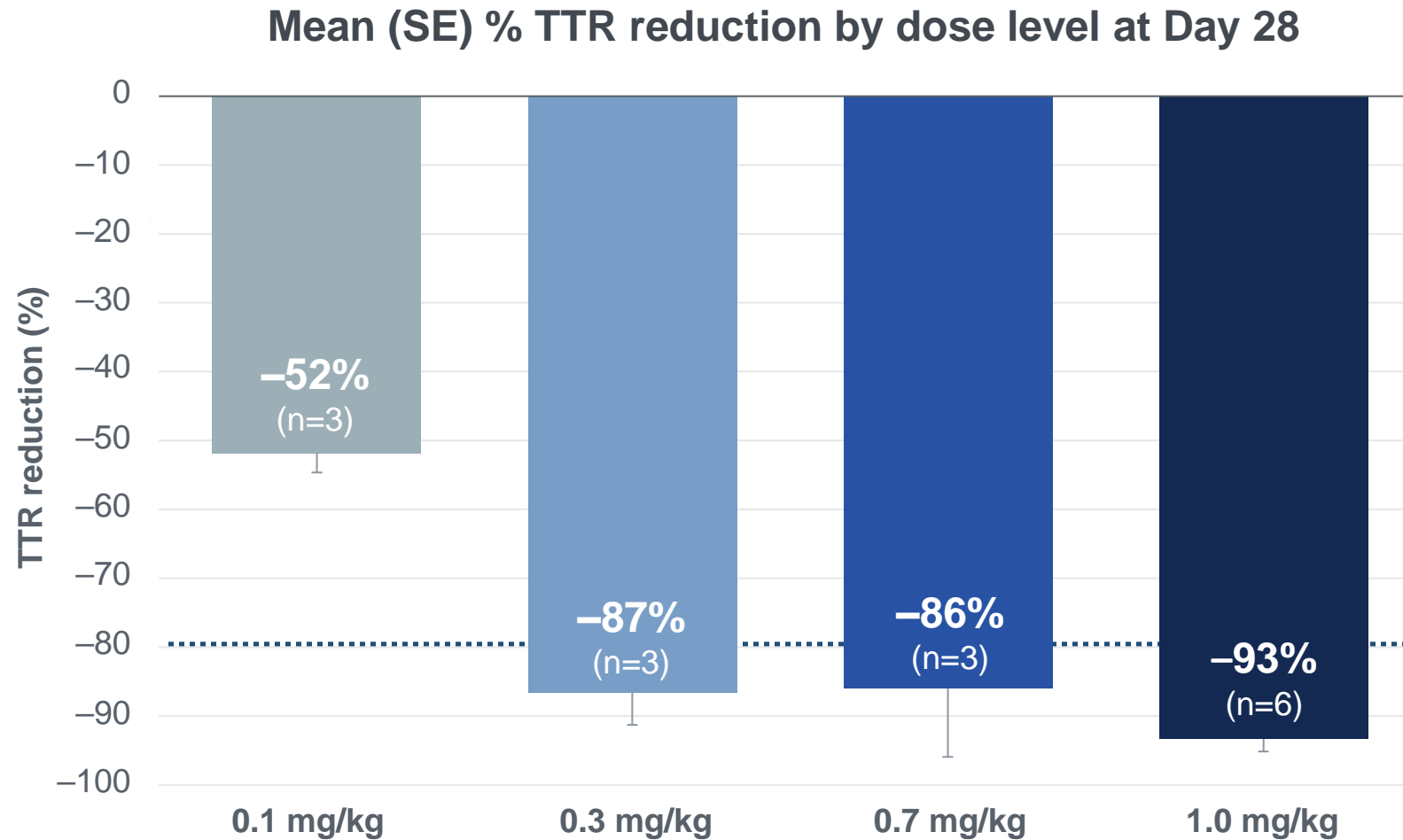
* Unrelated Grade 3 (SAE) of COVID-19 pneumonia

† Related Grade 3 (SAE) of vomiting in a patient with concomitant medical history of gastroparesis

Gr., Grade; SAE, serious adverse events; TEAE, treatment-emergent adverse event

Data disclosed on February 28, 2022

Dose-dependent reductions in serum TTR, reaching a mean reduction of 93% at 1.0 mg/kg



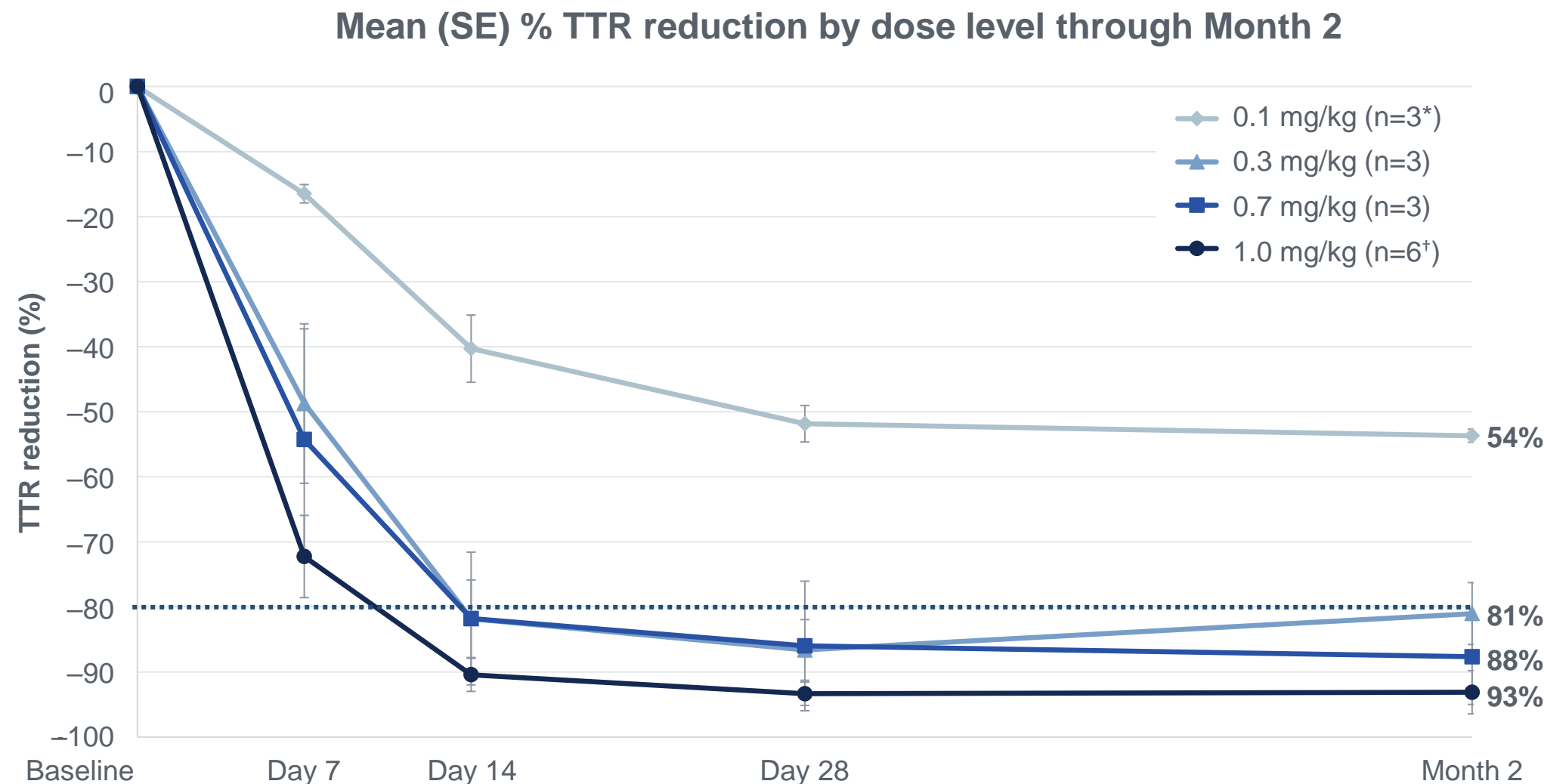
Dashed line represents the targeted minimum reduction

SE, standard error; TTR, transthyretin

Data disclosed on February 28, 2022

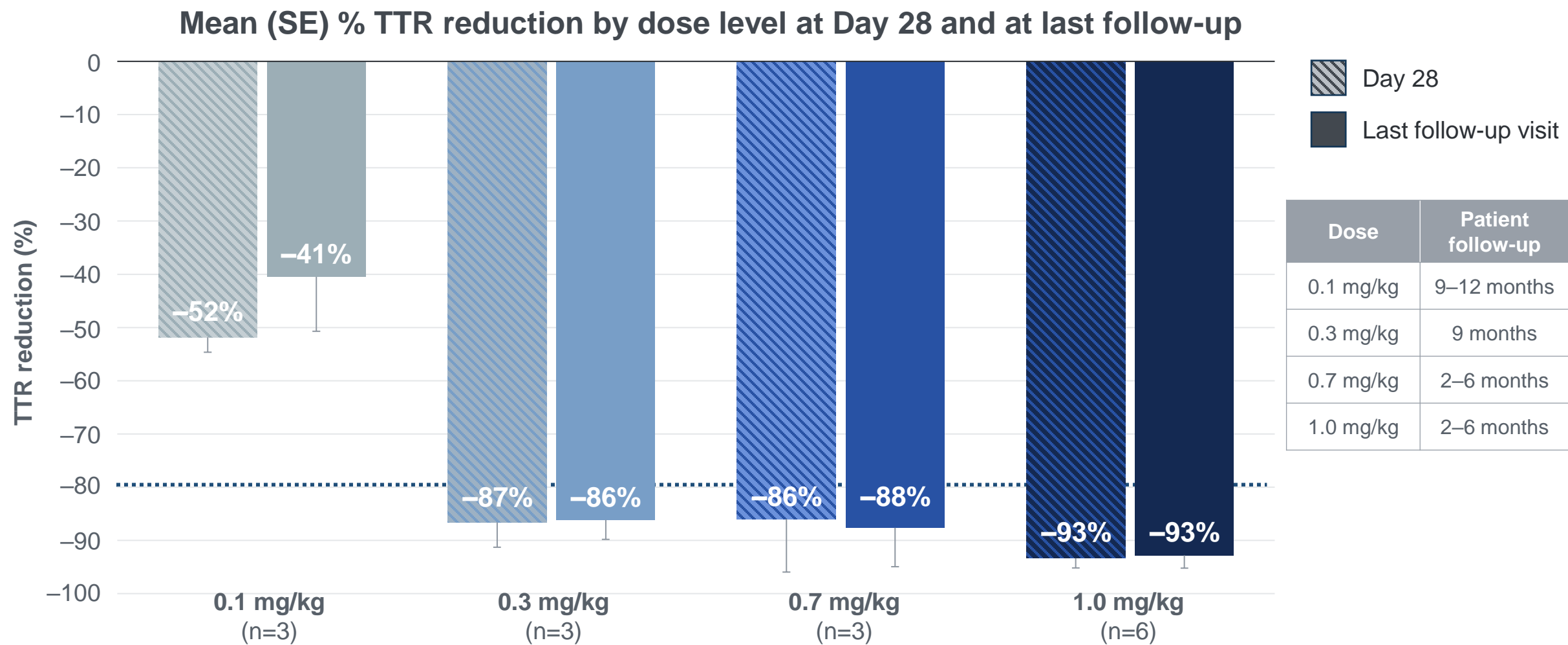
This slide includes data for investigational products not yet approved by regulatory authorities

Rapid reductions in serum TTR, achieving nadir by Day 28



* n=2 at Month 2; † n=5 at Month 2
Dashed line represents the targeted minimum reduction
SE, standard error; TTR, transthyretin
Data disclosed on February 28, 2022
This slide includes data for investigational products not yet approved by regulatory authorities

Durable reductions in serum TTR were observed over the follow-up period



Mean % reduction at last follow-up calculated using TTR value from last available follow-up visit for each patient per dose level





Dashed line represents the targeted minimum reduction

SE, standard error; TTR, transthyretin

Data disclosed on February 28, 2022

This slide includes data for investigational products not yet approved by regulatory authorities

Growing Confidence in NTLA-2001 as Potential Treatment for ATTR Amyloidosis

Key Insights from Ongoing Phase 1 Study	Supported by Interim Data
Generally well-tolerated at all dose levels	
Dose-response relationship with deep reductions at higher doses	
Consistent reductions in serum TTR across ATTRv-PN patients	
Durable response following a single dose	

93% mean serum TTR reduction demonstrated at 1.0 mg/kg by Day 28 (n=6)

NTLA-2002 for Hereditary Angioedema (HAE)



- Genetic disease characterized by recurring, severe and unpredictable swelling in various parts of the body
- Chronic dosing is required with current treatments

~7-14 days

Average frequency of attacks for untreated patients¹

~1 in 50,000

HAE patients worldwide¹

OUR APPROACH

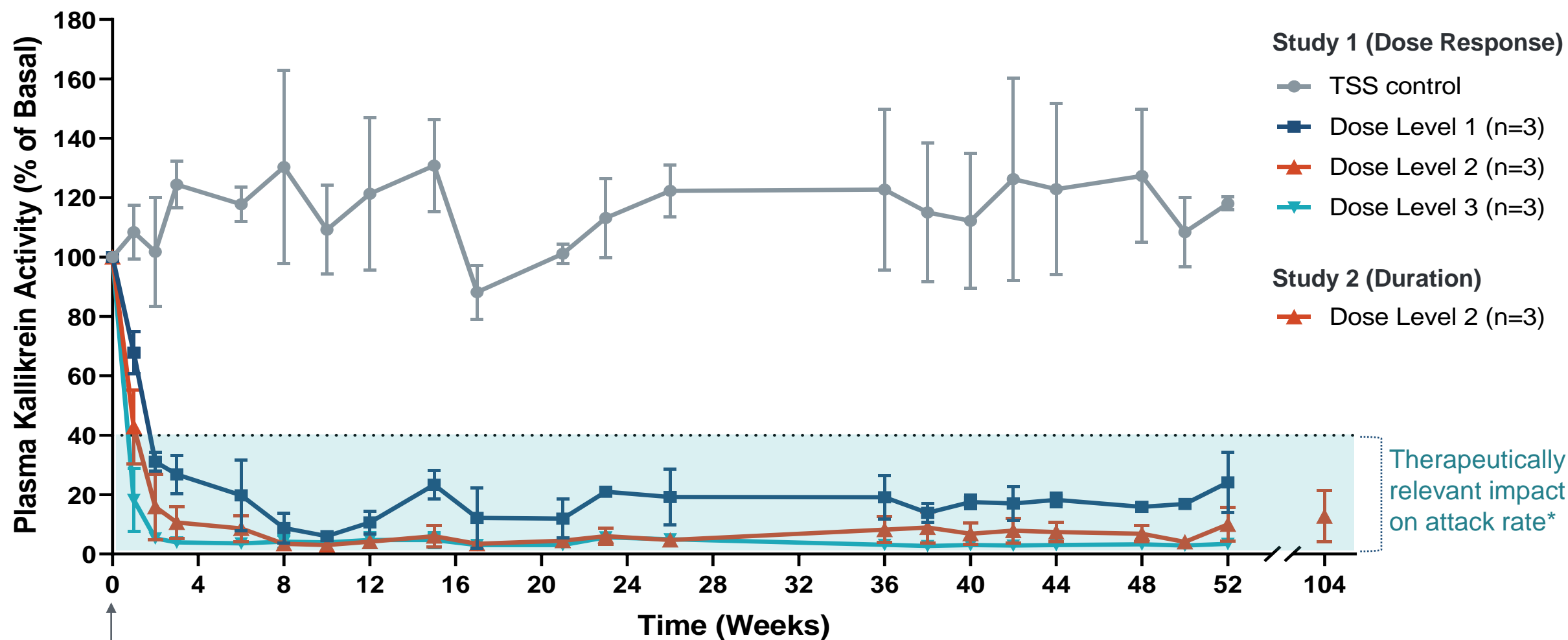
Knock out *KLKB1* gene with a single dose

- Reduce kallikrein activity to prevent attacks

KEY ADVANTAGES

- Potential “one-and-done” treatment
- Expect extensive and continuous reduction in kallikrein activity
 - Intended to minimize the risk of breakthrough attacks
- Potential to eliminate significant treatment burden

Achieved Sustained Therapeutically Relevant Kallikrein Activity Reduction After a Single Dose in NHPs



Single Dose

NTLA-2002 Phase 1/2 Trial Design

International, multi-center study to assess safety, tolerability, PK, PD and effect of NTLA-2002 on attacks in adults with Type I or Type II HAE

Total Enrollment:

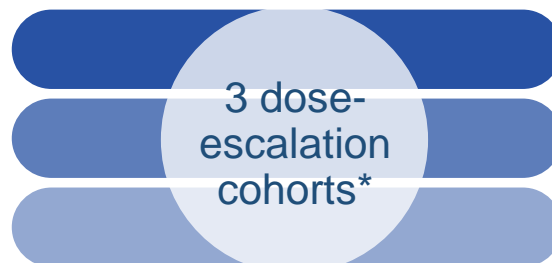
Up to 55 patients,
age 18 and older



Intervention:

Single dose
administered via
an intravenous
(IV) infusion

PHASE 1 Open-Label, Single-Ascending Dose



PHASE 2 Expansion study to confirm recommended dose

Randomized

Dose 1 (N=10)

Dose 2** (N=10)

Placebo Arm (N=5)

KEY ENDPOINTS

- Evaluate safety and tolerability
- Change in plasma kallikrein protein and activity levels
- Change in attack rates (Phase 2)

Clinicaltrials.gov ID: NCT05120830

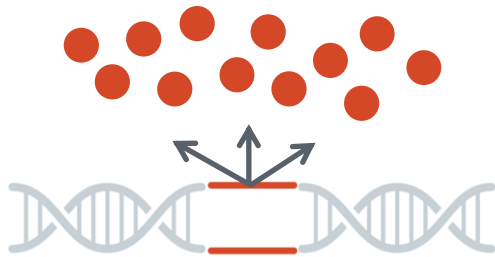
PK: Pharmacokinetics **PD:** Pharmacodynamics

*3 to 6 subjects per cohort; up to 2 additional cohorts, if necessary **Optional cohort

Beyond Gene Inactivation, Intellia is Also Advancing Targeted Insertion Programs

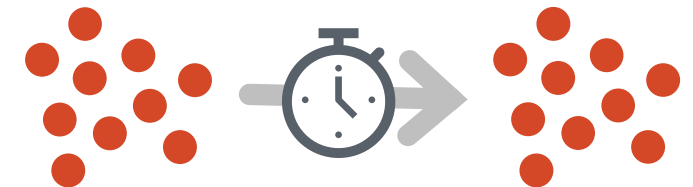
CRISPR-Enabled Targeted Insertion Approach Offers Significant Advantages Over Alternate Gene Therapy Approaches

**High Levels of
Protein Expression**

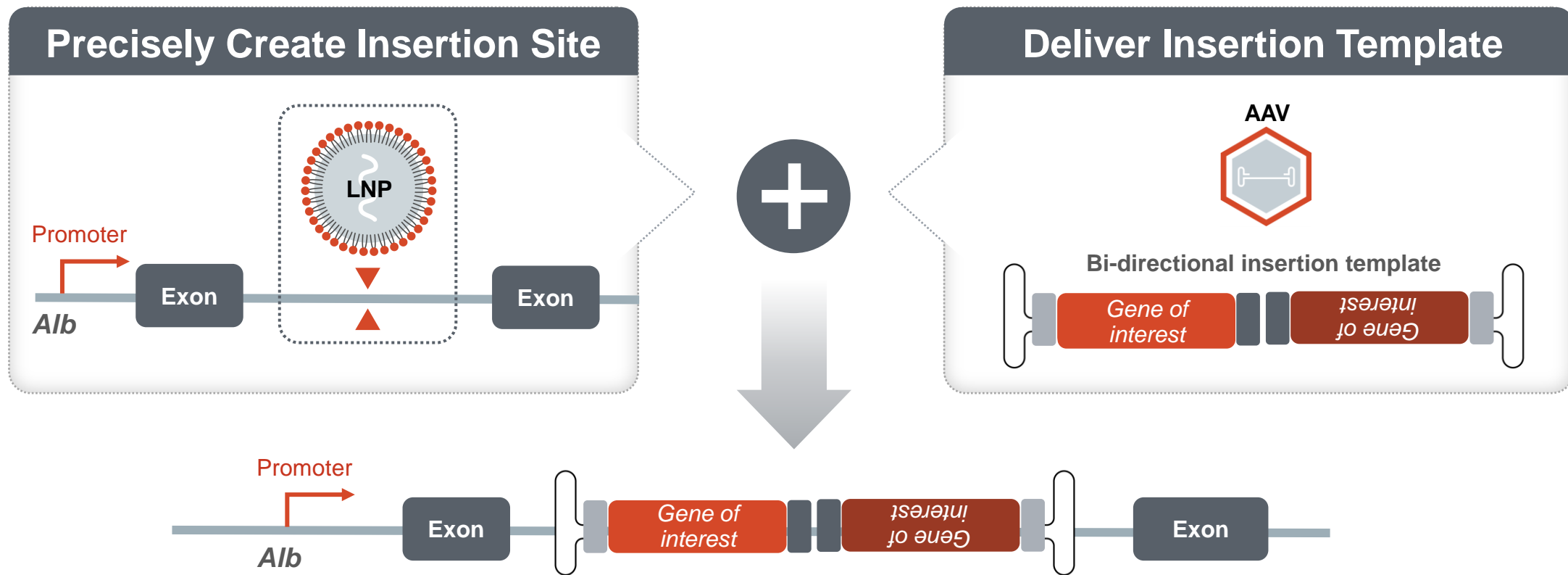


**Potential to
Revolutionize
Gene
Replacement**

**Durable
Protein Expression**



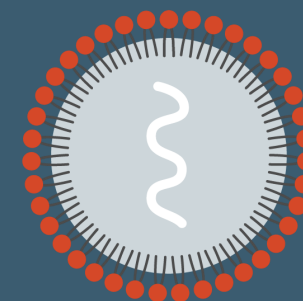
Insertion Technology Enables Production of High Levels of Therapeutic Protein



Targeted, stable gene insertion in the albumin locus

NTLA-3001 and NTLA-2003 for Alpha-1 Antitrypsin Deficiency (AATD)

Genetic disorder leading to progressive lung and/or liver disease¹



AATD patients*

> 60K in the U.S.²

~250K globally³

OUR APPROACH

NTLA-3001: Targeted insertion of a functional *SERPINA1* gene

- Continuous expression of functional A1AT protein at normal levels
- Address AATD-associated lung disease

NTLA-2003: Knockout mutant *SERPINA1* gene

- Reduce and prevent accumulation of mutant A1AT protein
- Address AATD-associated liver disease

KEY ADVANTAGES

- Each candidate is designed as a single-dose treatment
- **NTLA-3001:** Aims to achieve normal human levels of A1AT protein and halt progression of lung disease
- **NTLA-2003:** Aims to halt the progression of liver disease and eliminate the need for liver transplant in severe cases

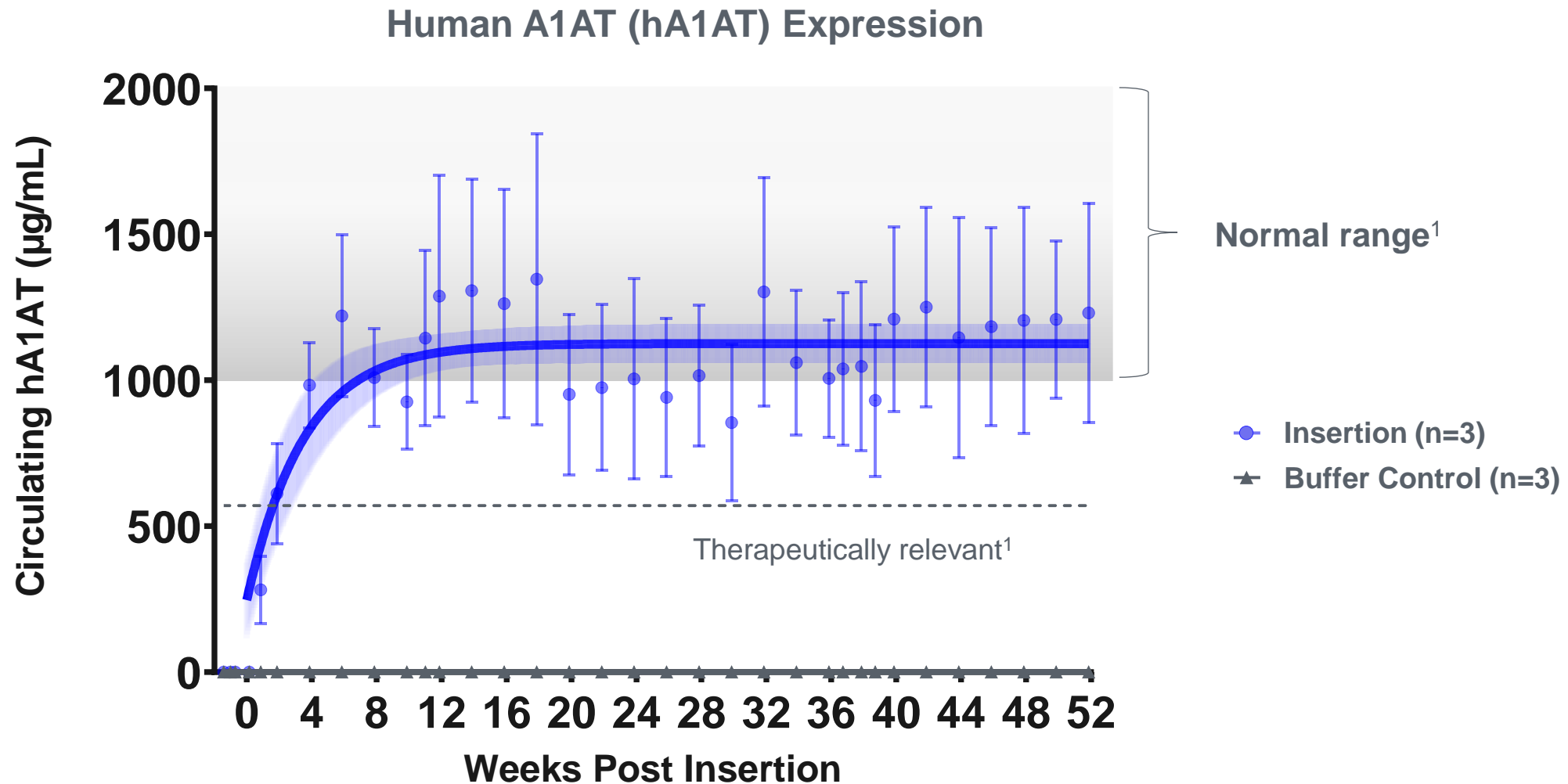
¹<https://www.genome.gov/Genetic-Disorders/Alpha-1-Antitrypsin-Deficiency>

²Brantly M. *Clin Chem*. 2006; 52:2180-2181

³Blanco et al. *Int J Chron Obstruct Pulmon Dis*. 2017; 12:561-569

*In severe AATD patients defined as individuals with Pi*ZZ genotype

Durable Physiologic Levels of hA1AT Maintained Through One Year in NHP



Clinical Validation of LNP Delivery Platform Supports *In Vivo* Pipeline Acceleration

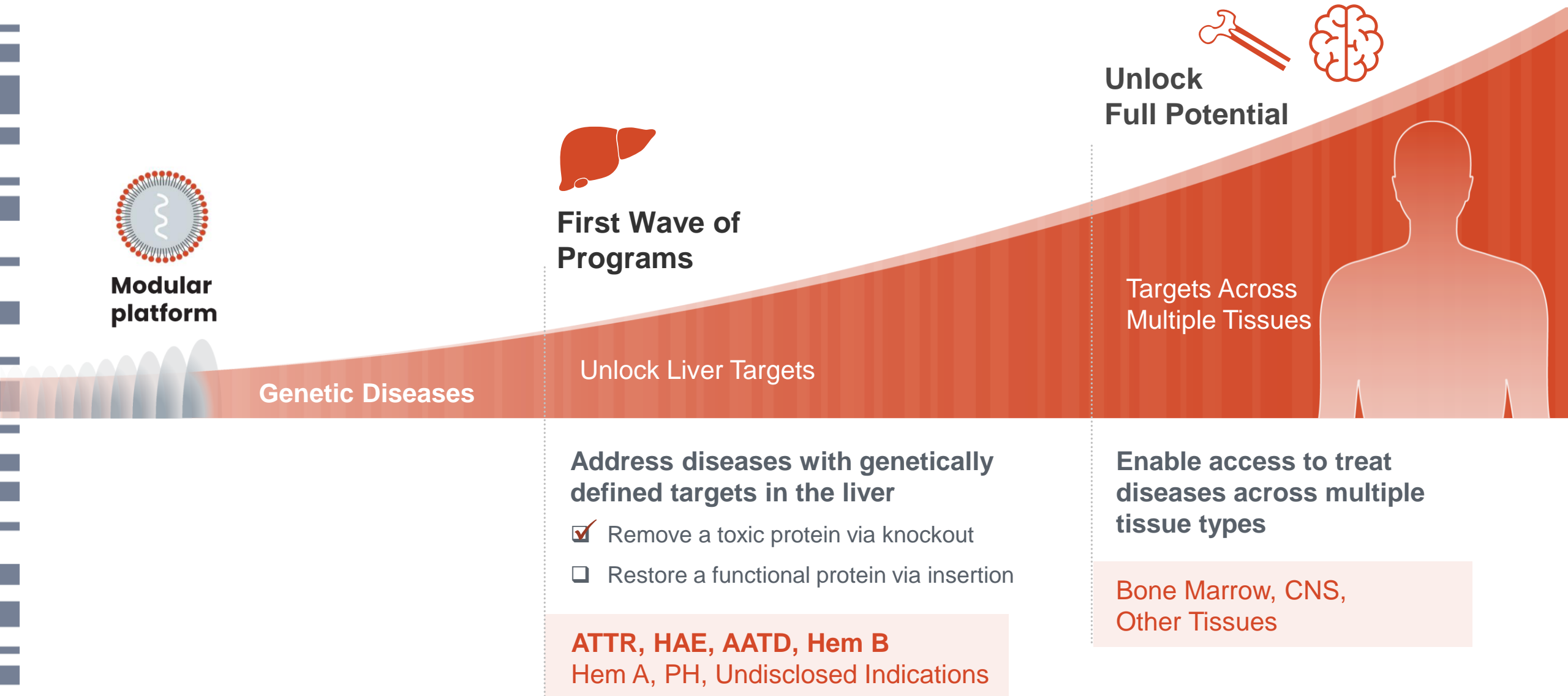




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Ex Vivo

CRISPR creates the therapy

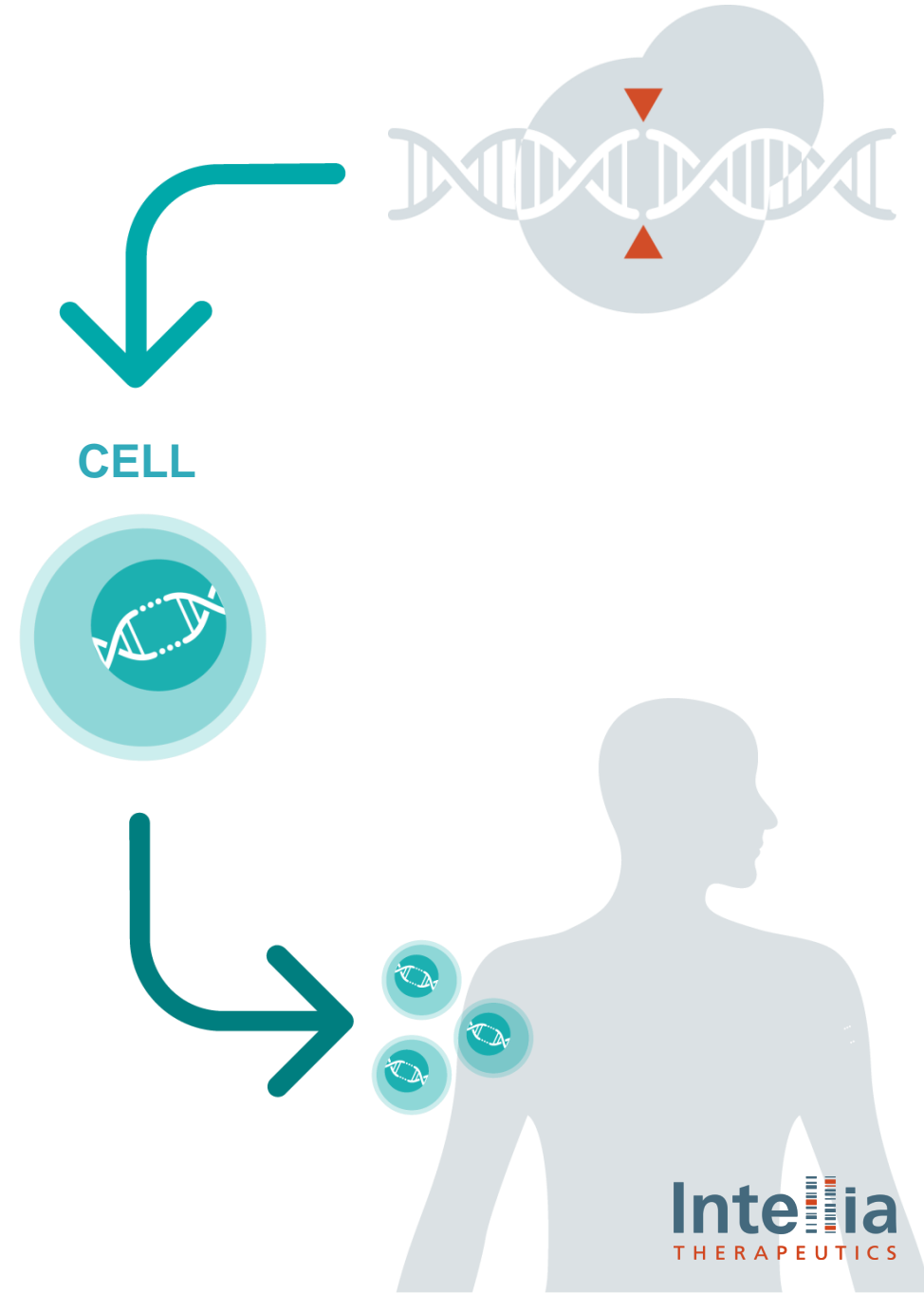
IMMUNO-ONCOLOGY / AUTOIMMUNE DISEASES

Strategic Advantages:

Utilizing proprietary CRISPR engineering platform to create differentiated cell therapies for IO and AI diseases

Targeting modalities, such as TCR, with broad potential in multiple indications

Focused on reproducing natural cell physiology for potential improvements to safety and efficacy in immuno-oncology



Proprietary Engineering Platform to Power Next-Generation Engineered Cell Therapies

LNP-BASED CELL ENGINEERING PLATFORM

Highly efficient sequential editing

Optimal cell performance

Scalable manufacturing process

ENABLES VERSATILE SOLUTIONS BY “MIXING AND MATCHING” INCLUDING:

Cell Type

HSCs, T cells
NK cells, Macrophages



Targeting Modality

TCRs
CAR-Ts, Universal CARs



Rewiring Instructions

Immune-enhancing edits
Novel targets



Differentiated Approach to Cell Therapy Genome Engineering

		Intellia THERAPEUTICS	Other Approaches	
Gene Editing Approach	Delivery	Lipid Nanoparticle	Electroporation	Electroporation
	Editing Mode	Sequential	Simultaneous	Simultaneous
	Knockout (KO)	Cleavase or Base Editor	Cleavase	Base Editor
	Insertion	CRISPR insertion	Lenti/Retroviruses	Lenti/Retroviruses
Key Questions From Preclinical Data	Minimize random DSB?	✓	✗	✗
	Minimize random insertion?	✓	✗	✗
	Minimize genotoxicity risk?	✓	✗	✗

Intellia
THERAPEUTICS

LNP-based,
sequential process



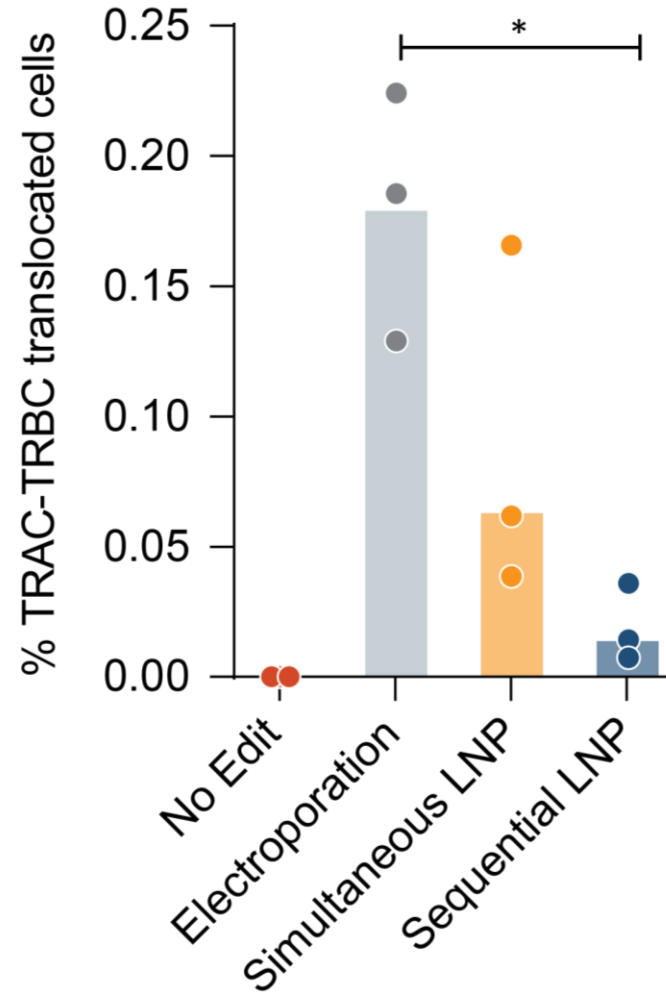
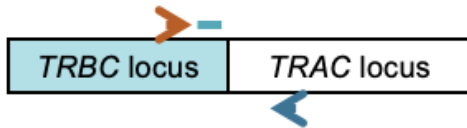
Precise CRISPR
KOs & insertion(s)



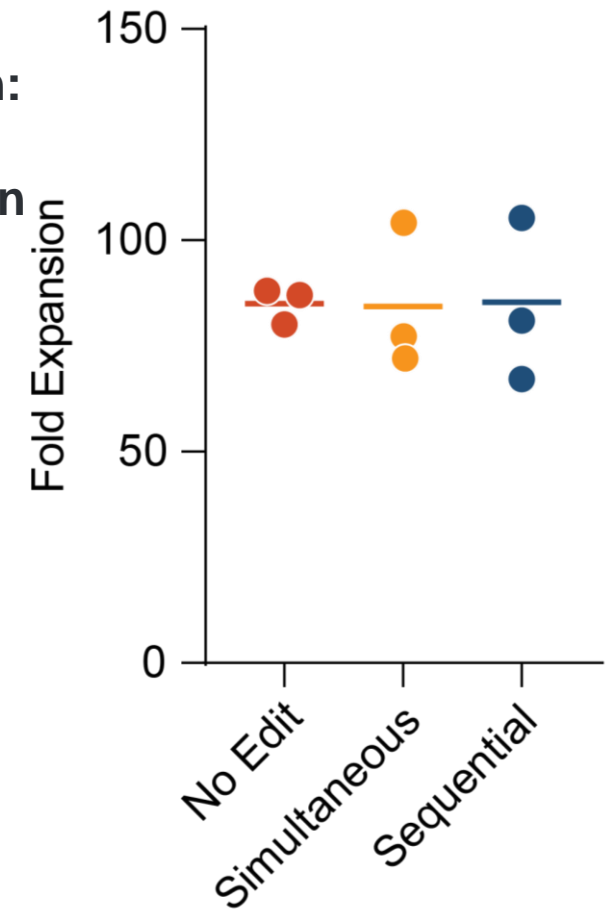
Quality cell
product

Sequential Editing with LNP Approach Minimizes Translocations While Retaining Robust Cell Viability and Expansion

ddPCR assay to detect *TRAC-TRBC* translocations



LNP Approach:
Cell Expansion
at D10



NTLA-5001 for Acute Myeloid Leukemia (AML)



Most common acute leukemia in adults¹

~20K

New cases in the U.S. in 2021¹

> 40K

New cases in the 7 Major Markets in 2020²

< 30%

5-year overall survival¹

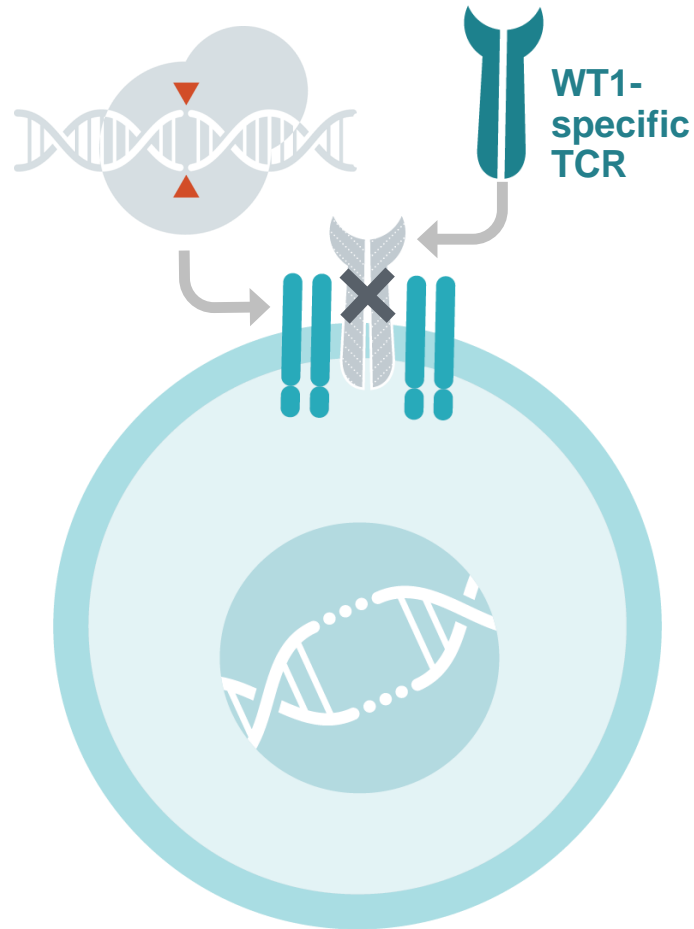
OUR APPROACH

Engineer TCR-T cells directed against Wilms' Tumor Type 1 (WT1) to specifically kill AML blasts

KEY ADVANTAGES

- Potential to address all mutational subtypes of AML
- Low WT1 expression in normal tissues for improved safety
- TCR sourced from healthy donor T cells intended to minimize immune toxicity

NTLA-5001: Potential Best-in-Class Engineered T Cell Therapy For AML



Inserts a **natural, high-avidity TCR** to replace native TCR for upgraded safety profile

- Activates both cytotoxic and helper T cells

Specifically **targets Wilms' Tumor 1 (WT1)**, an antigen overexpressed in >90% of AML blasts¹

- Recognizes an epitope (VLD²) presented broadly by AML blasts with the HLA-A*02:01 allele³

Modified by **proprietary cell engineering** technology for optimized cell health and function

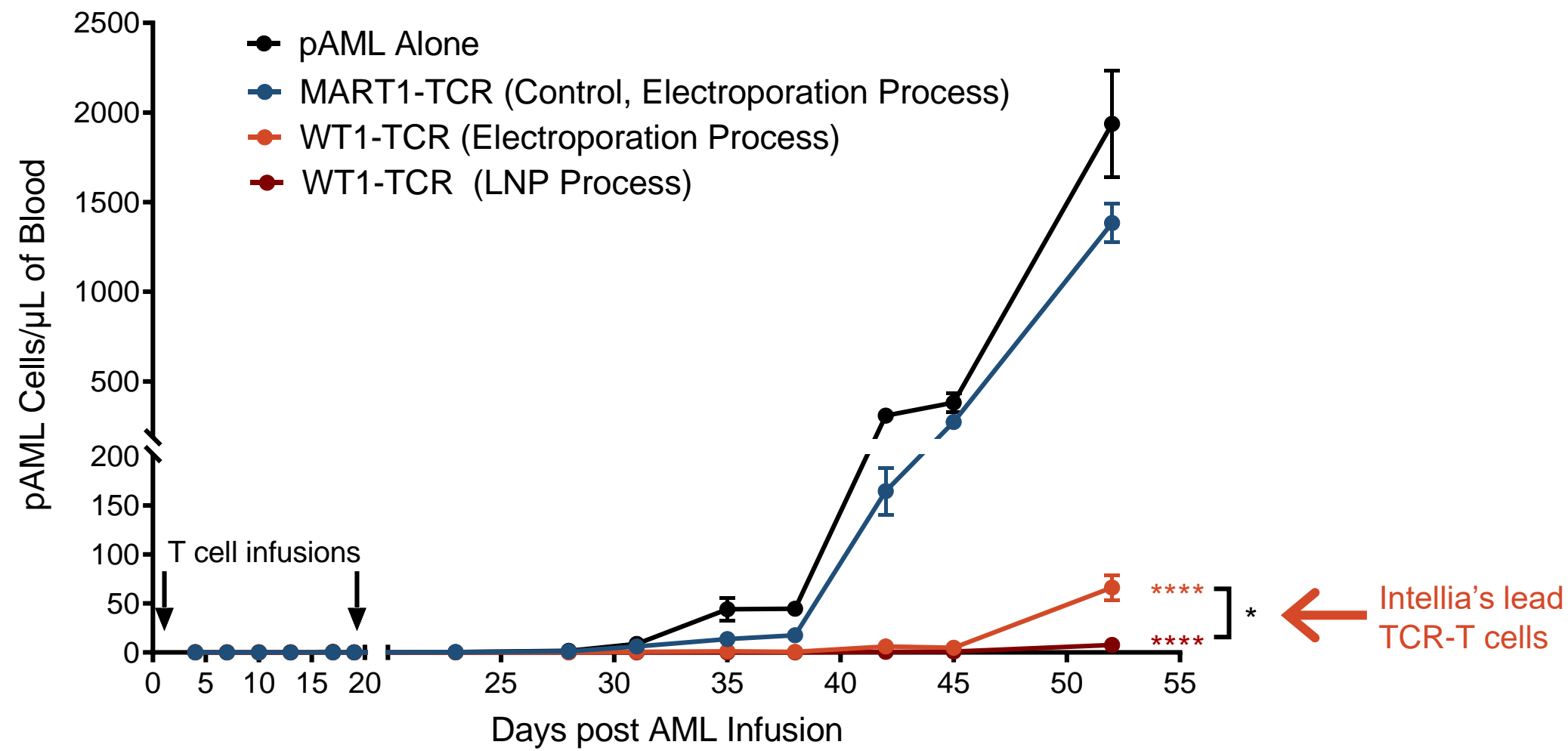
¹Cilloni et al., *J Clin Oncol*, 2009

²VLD is the WT1₍₃₇₋₄₅₎ epitope VLDFAPPGA

³Refer to <http://www.allelefrequencies.net> for HLA frequency data

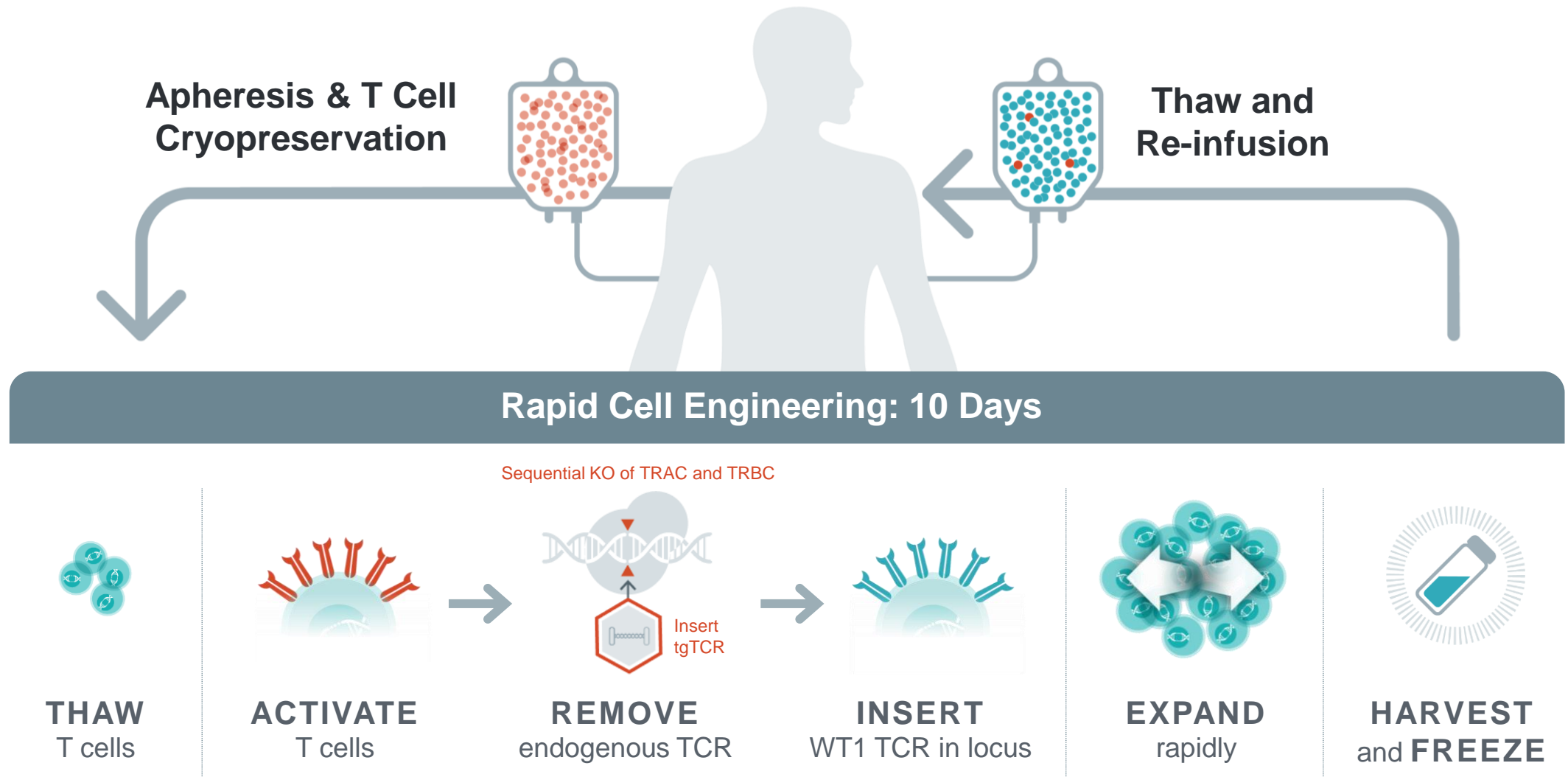
In collaboration with IRCCS Ospedale San Raffaele

NTLA-5001: Robust Anti-Tumor Efficacy Observed Against Patient-Derived AML Blasts in Mouse Model



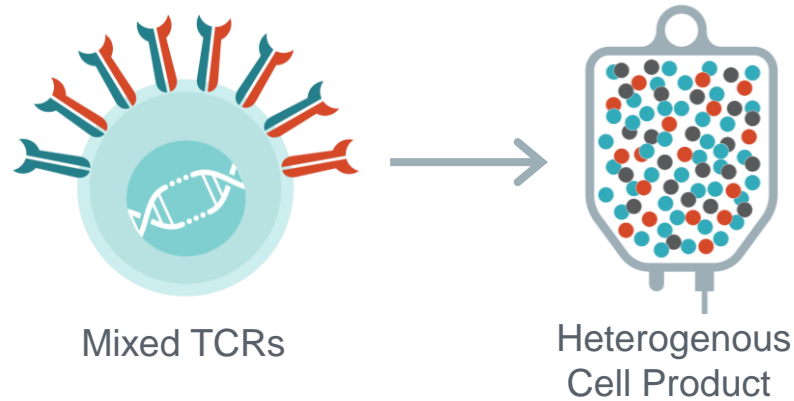
****p<0.0001, WT1 TCR vs. MART1 Control TCR (2-way ANOVA)
*p<0.05 or p<0.01, EP vs. LNP Process (2-way ANOVA)

NTLA-5001: Uniform Expression of Therapeutic TCR for Potent Tumor Targeting

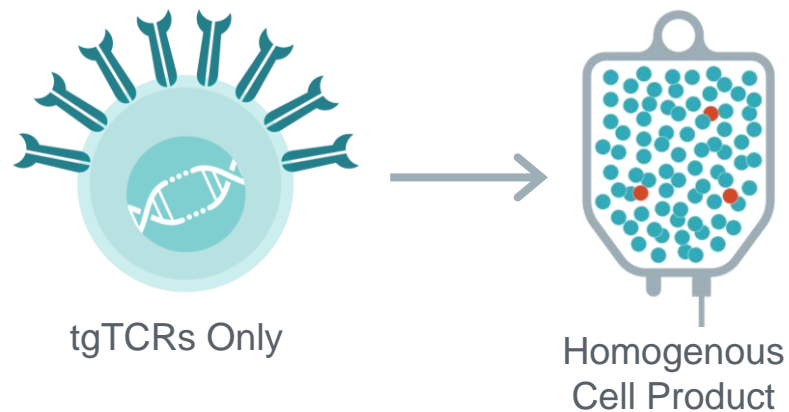


CRISPR Engineering Overcomes Key Challenges of Traditional TCR Approaches

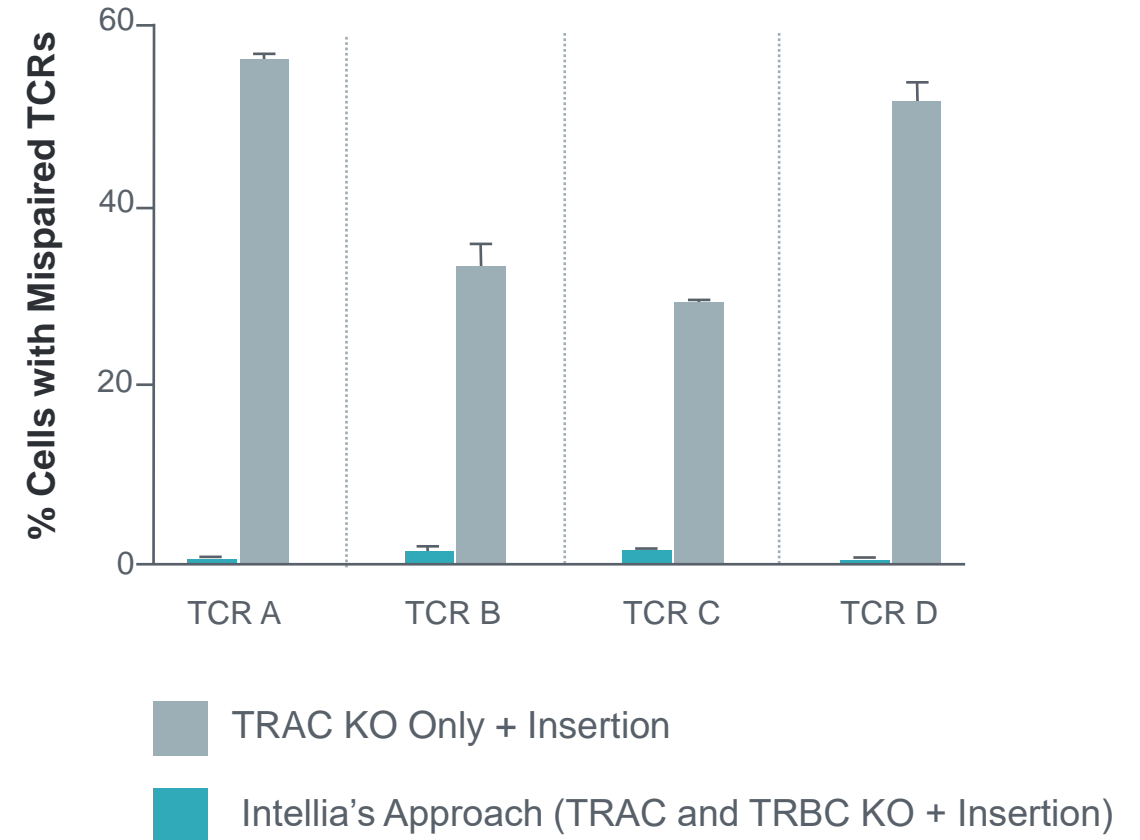
Traditional tgTCR Addition



CRISPR/Cas9 tgTCR Replacement



Removal of Endogenous TCR Prevents Mispairing



NTLA-5001 Phase 1/2a Trial Design

Open-label, multi-center study of NTLA-5001, a WT1-directed TCR immunotherapy, in adults with AML

Total Enrollment:

Up to 54 patients,
age ≥ 18 years

Key Inclusion Criteria:

- Relapsed/refractory AML after one or more therapies
- Post transplant patients are eligible
- HLA-A*02:01 positive



Intervention:

Single dose administered via intravenous (IV) infusion

PHASE 1

Dose Escalation

Two-ascending arms: Up to 3 cohorts*

ARM 1: Lower Disease Burden

ARM 2: Higher Disease Burden

PHASE 2

Expansion Cohorts

To confirm recommended dose from each arm of Phase 1

Dose 1 (N=9)

Dose 2 (N=9)

KEY ENDPOINTS

- Evaluate safety and tolerability
- Characterize cell kinetics of NTLA-5001
- Determine anti-tumor activity

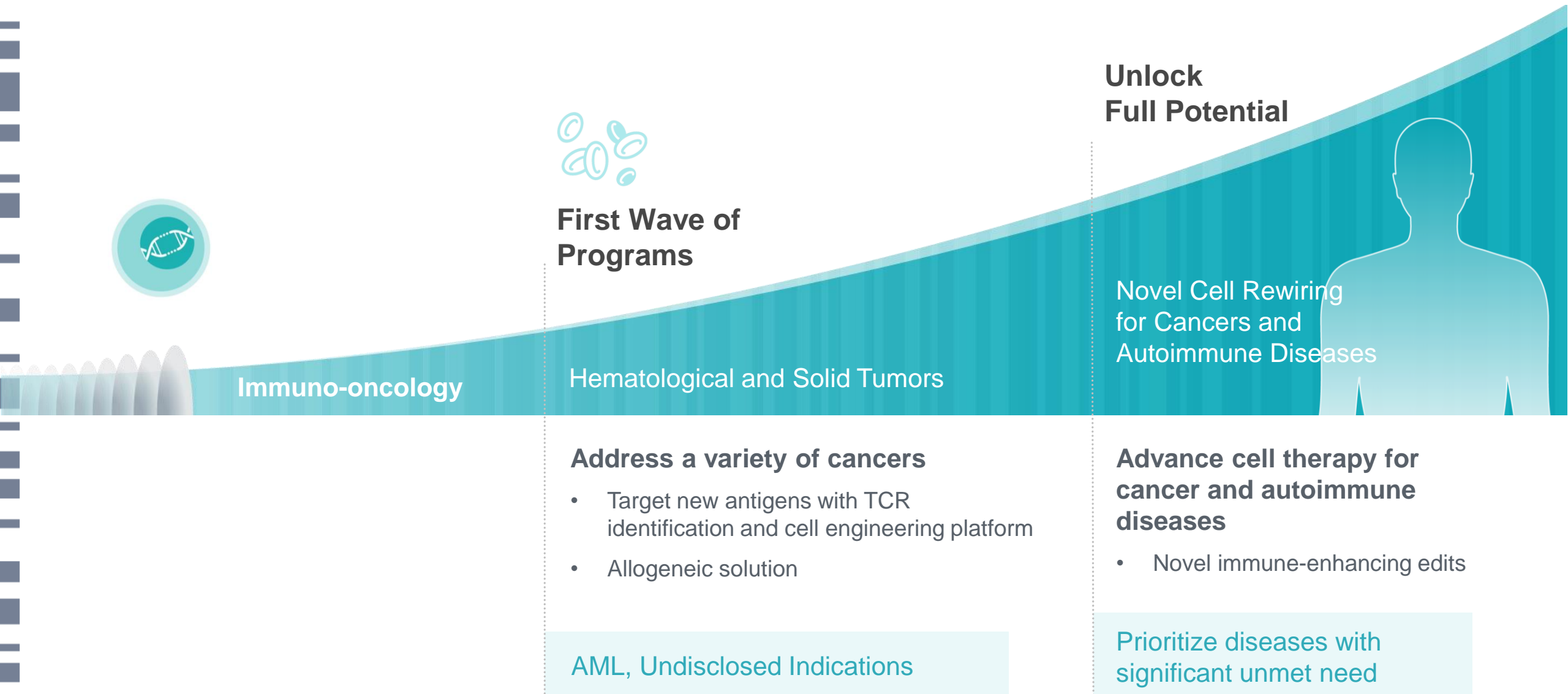
*3-6 subjects per cohort

Clinicaltrials.gov ID: NCT05066165

Lower disease burden: Patients with less than 5% blasts in bone marrow

Higher disease burden: Patients with relapsed/refractory disease with greater than or equal to 5% blasts in bone marrow

Ex Vivo Pipeline Expansion Strategy



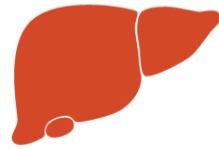
Unlocking the Full Potential of CRISPR

Solving *in vivo* delivery supports rapid expansion of pipeline to broad patient population

in vivo

Genetic diseases

CRISPR is the therapy



NTLA-2001

Unlock the liver for ATTR, NTLA-2002 for HAE and beyond

NTLA-3001 and Factor IX

Restore a functional protein via insertion for AATD and Hem B

Target bone marrow and other tissues

Modular platform

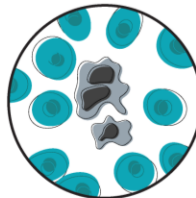
ex vivo

Immuno-oncology, autoimmune diseases

CRISPR creates the therapy

NTLA-5001

Rewire T cells to target Acute Myeloid Leukemia



Engineer allogeneic therapies



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Intellia Investment Overview

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Ex Vivo Portfolio

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In Vivo Editing of Hematopoietic Stem Cells

LNP-Based Editing of T Cells

Intellia's Allogeneic Solution

Intellia's Proprietary Base Editor

Platform: Identifying Potent and Highly Specific Guide RNAs

Strategic Collaborations

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Intellia's Genome Editing Toolbox

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World-Class Genome Editing Platform Allows for Unsurpassed Capabilities

Proprietary CRISPR-based Modular Platform

Editing Tools

CRISPR/Cas9

Base editor

Additional
enzymes

Delivery Tools

LNPs

AAVs

Additional
modalities

ENABLES SELECTING THE BEST TOOLS FOR EACH THERAPEUTIC APPLICATION:

Applies to *in vivo* or *ex vivo* application

Capable of achieving any editing strategy

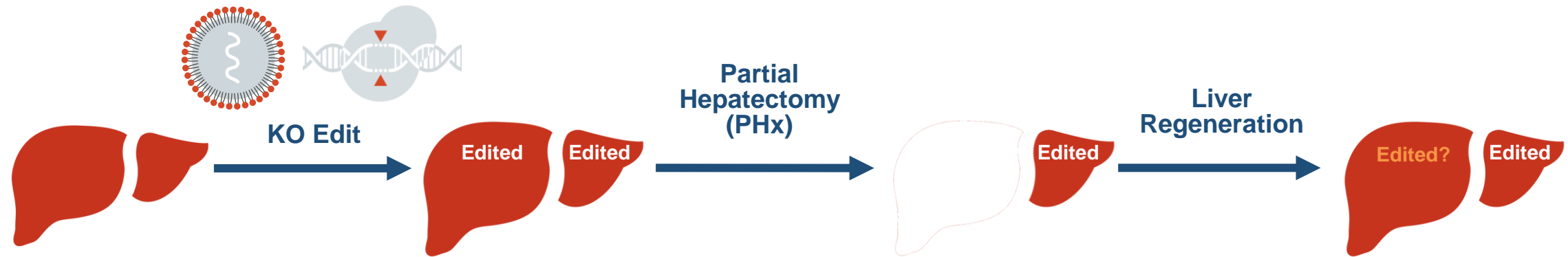
- Precise knockout and targeted insertions
- Multiplicity of edits
- Single nucleotide modifications

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Persistence of *In Vivo* Edits

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Partial Hepatectomy Model for Investigating Persistence of Knockout Genome Editing

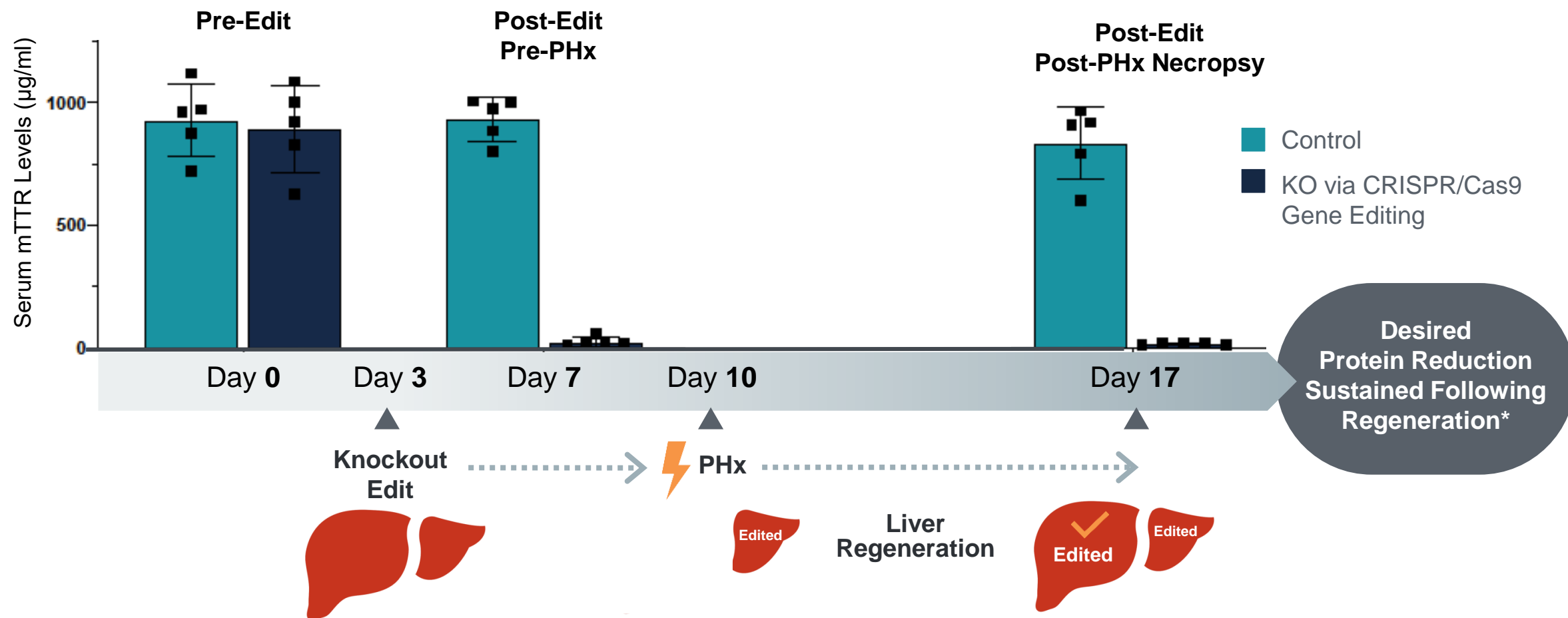


NHP studies demonstrate sustained KO editing and target TTR protein reduction carried through regular cell turnover for 12 months



Key Question: Can editing be carried through tissue regeneration following partial hepatectomy and accelerated cell division?

Protein Reduction Remains Unchanged Following Murine Liver Regeneration



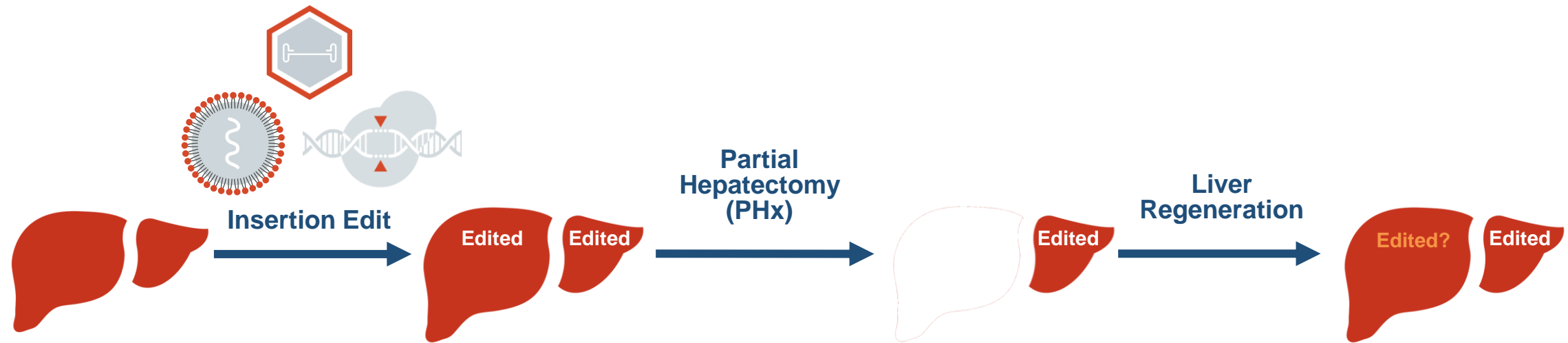
***TTR* gene editing rate similarly remains unchanged post-PHx by NGS analysis¹**

PHx: Partial Hepatectomy

*Similar results obtained for control and LNP when sham surgery was performed

¹Next generation sequencing (NGS) analysis to evaluate the frequency of insertion and deletion events (edits).

Partial Hepatectomy Model for Investigating Persistence of Insertion Genome Editing



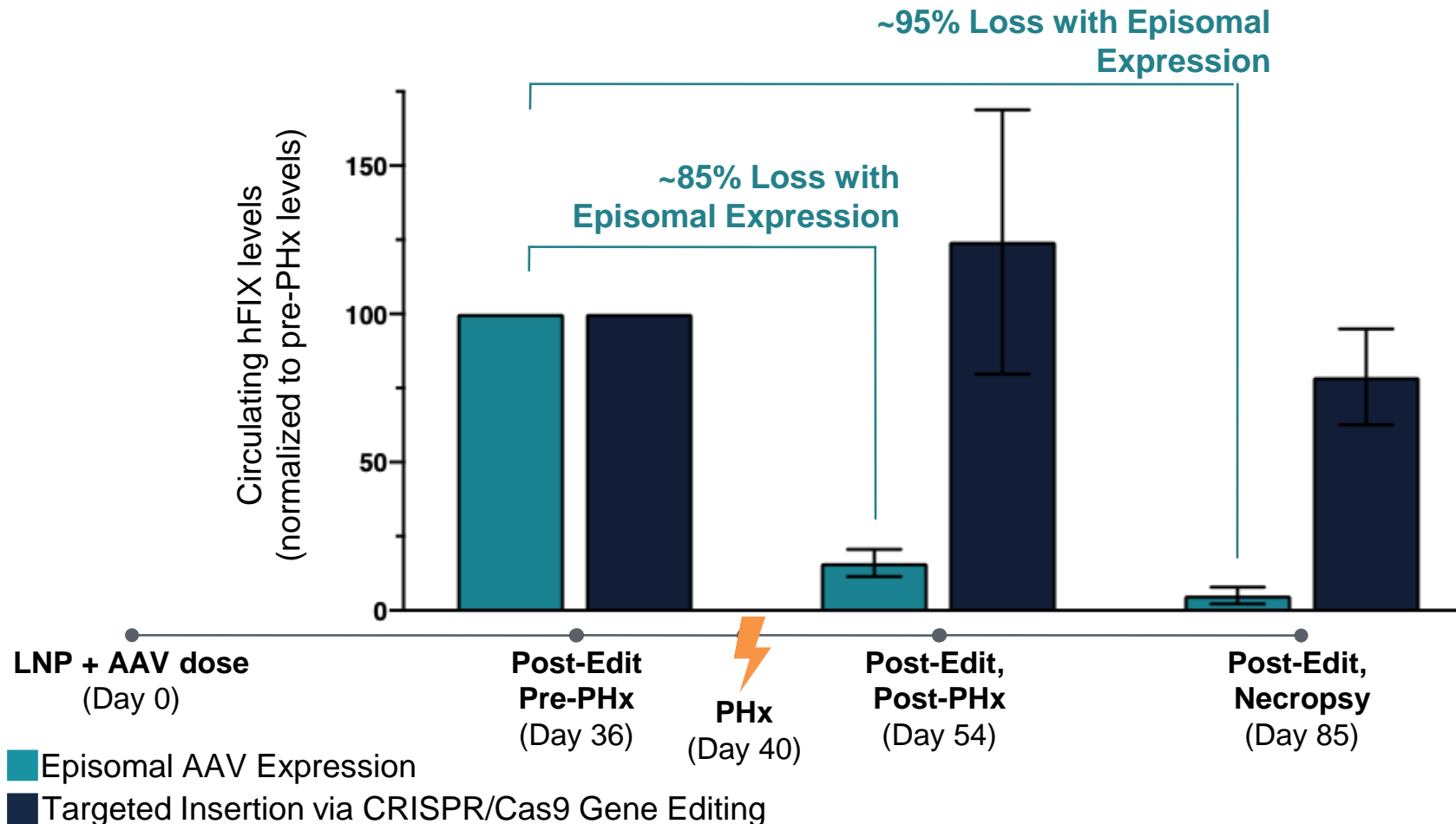
Rodent studies show sustained FIX insertion editing through 12 months, demonstrating that editing is carried through normal cell turnover



Key Question: Can insertion editing be carried through tissue regeneration following partial hepatectomy?

Persistent Protein Levels Post-PHx from Targeted Gene Insertion in Murine Model, in Comparison to Significant Loss of Protein Expression with Gene Therapy

Correlating editing rate similarly remains unchanged post-PHx by NGS analysis¹



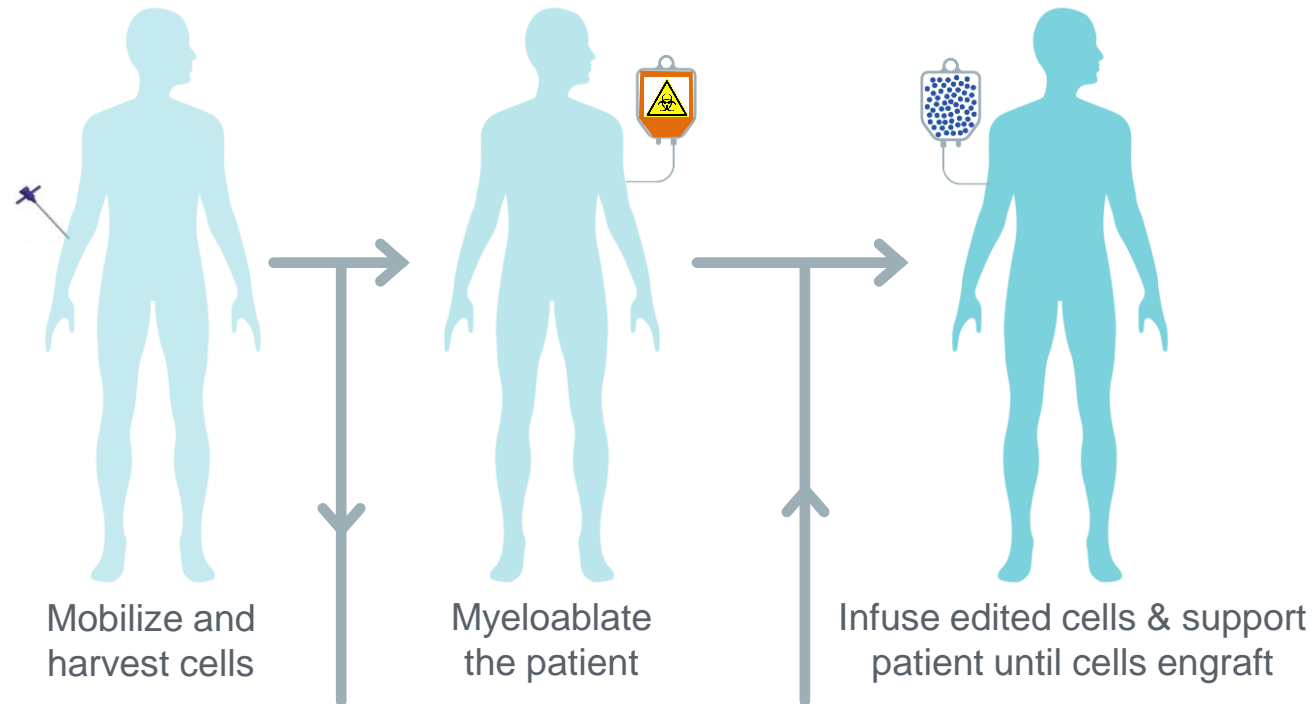
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In Vivo Editing of Hematopoietic Stem Cells

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Ex vivo SCD gene editing still has significant limitations

Complex cell manufacturing process



Conditioning regimen toxicity

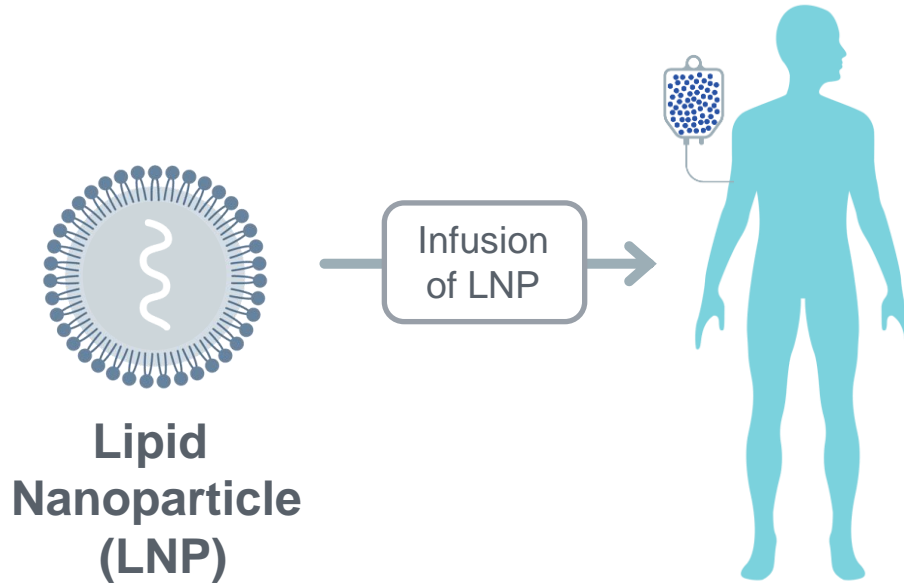
- Immunosuppression for > 1 month, predisposing to infection
- Risk of malignancy from chemotherapy drugs, especially leukemia
- Risk of infertility

Implications

- *Ex vivo* gene editing will be limited to highly selected SCD patients with severe disease
- Treatment complexity will limit access for patients in resource-poor settings

In vivo non-viral SCD gene editing could overcome these limitations

Simplified process



Potential improved safety and accessibility

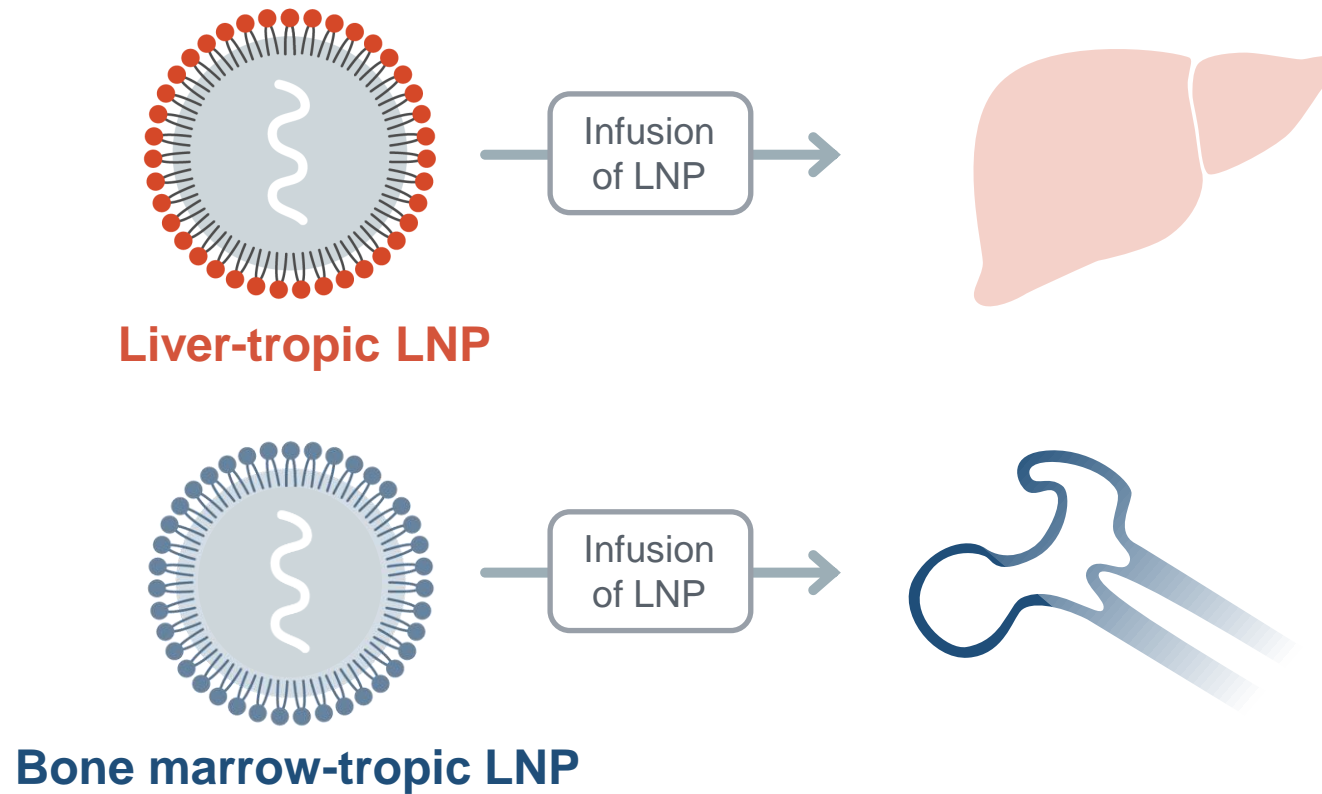
- Avoids myeloablation and associated risks of immunosuppression, malignancy and infertility
 - Approach could become mainstream therapy for SCD
- Avoids need for complex cell manufacturing or extensive supportive care post-treatment
 - Treatment simplicity could expand access to patients in resource-poor settings

Desired features of *in vivo* approach

- Provides clinically meaningful, durable HSC editing
- Allows for multidosing to reach therapeutic target
- Preserves regenerative potential of edited cells
- Translatable to human HSC population

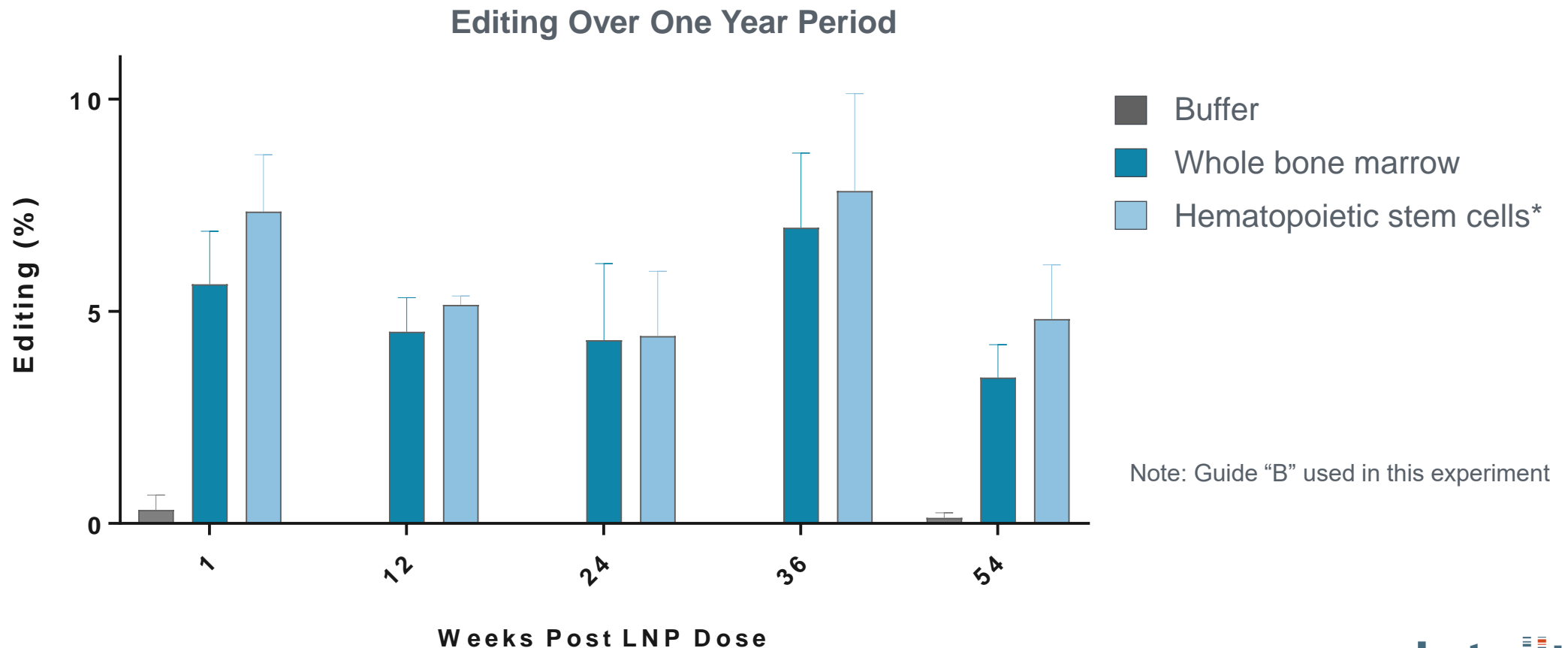
Editing HSCs *in vivo* requires LNPs with bone marrow tropism

- LNPs designed, formulated and tested *in vivo* to identify compositions with enhanced delivery to bone marrow and HSCs



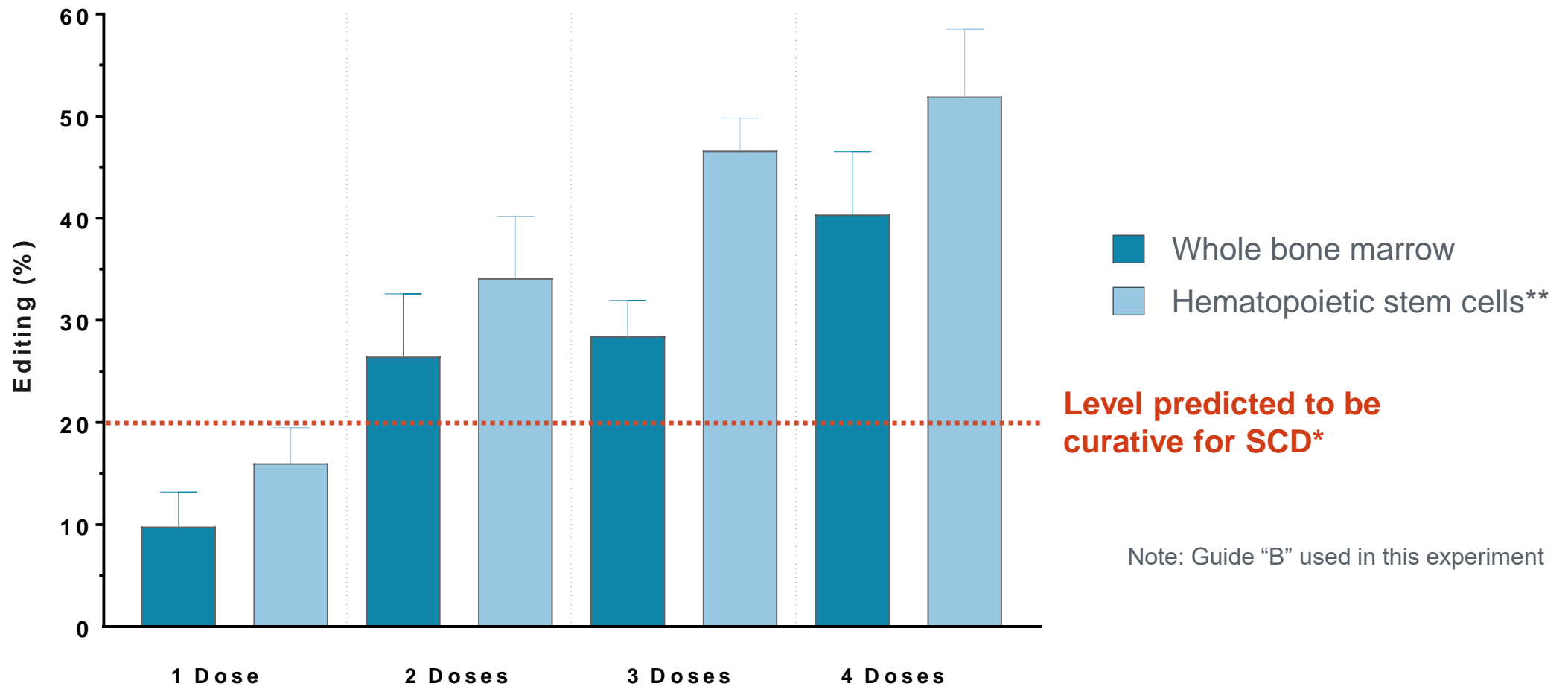
Editing of mouse bone marrow and HSCs is durable through at least one year

- Editing was similar across all time points assessed, in both whole bone marrow and HSCs populations
- Results highlight the potential for a single-course, long-lasting therapy



Editing of mouse bone marrow and HSCs increases with multidosing

- Non-immunogenic LNP delivery platform may enable stepwise “treat-to-target” approach



**Blood*. 2017;130(17):1946-1948.

**Lin-Sca-1⁺c-Kit⁺CD34⁺Flk2⁻ cell population

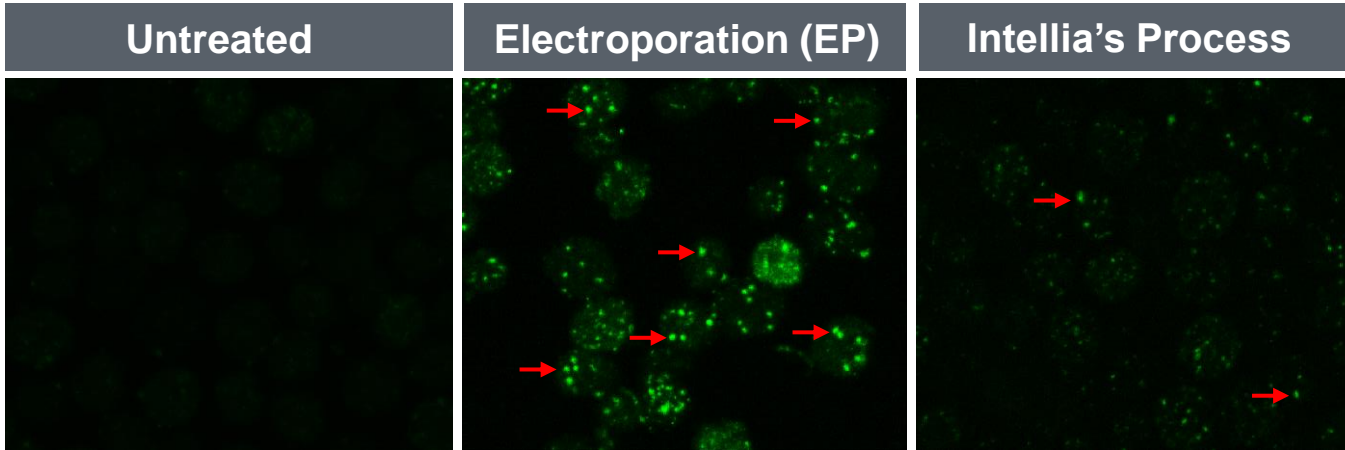
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LNP-Based Editing of T Cells

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LNP-Based Cell Engineering Technology Optimizes Cell Health and Function

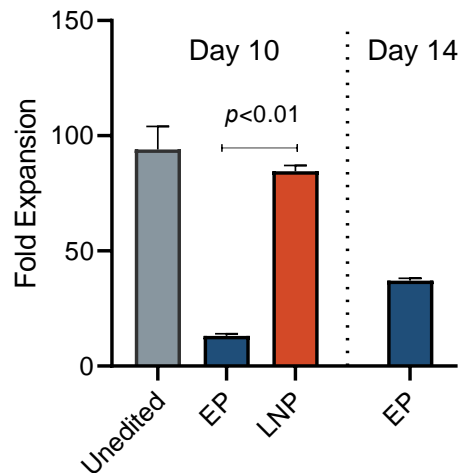
DNA Damage
γ-H2AX marker



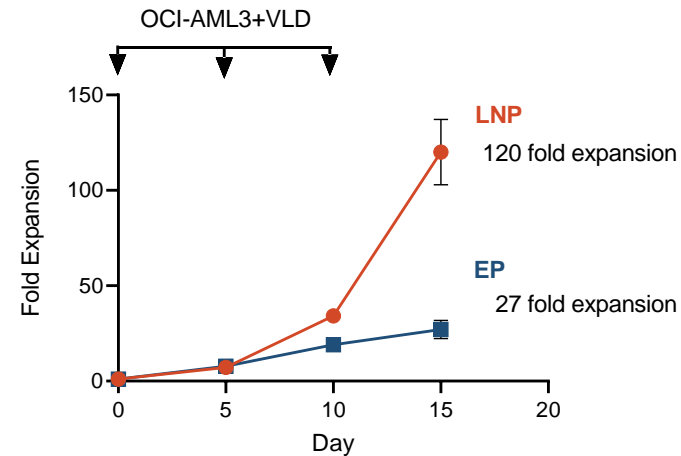
LNP approach to editing T cells

- Enables sequential editing
- Reduces safety risks from unwanted breaks caused by EP
- Produces cells with high expansion and performance

Cell Expansion

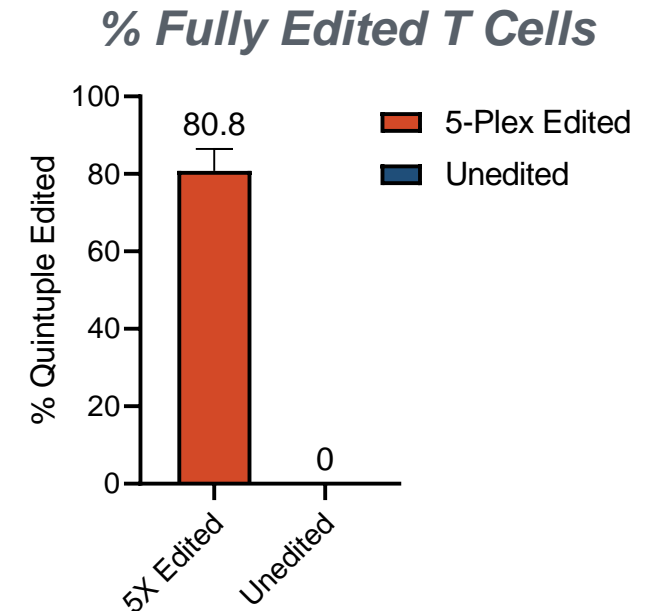
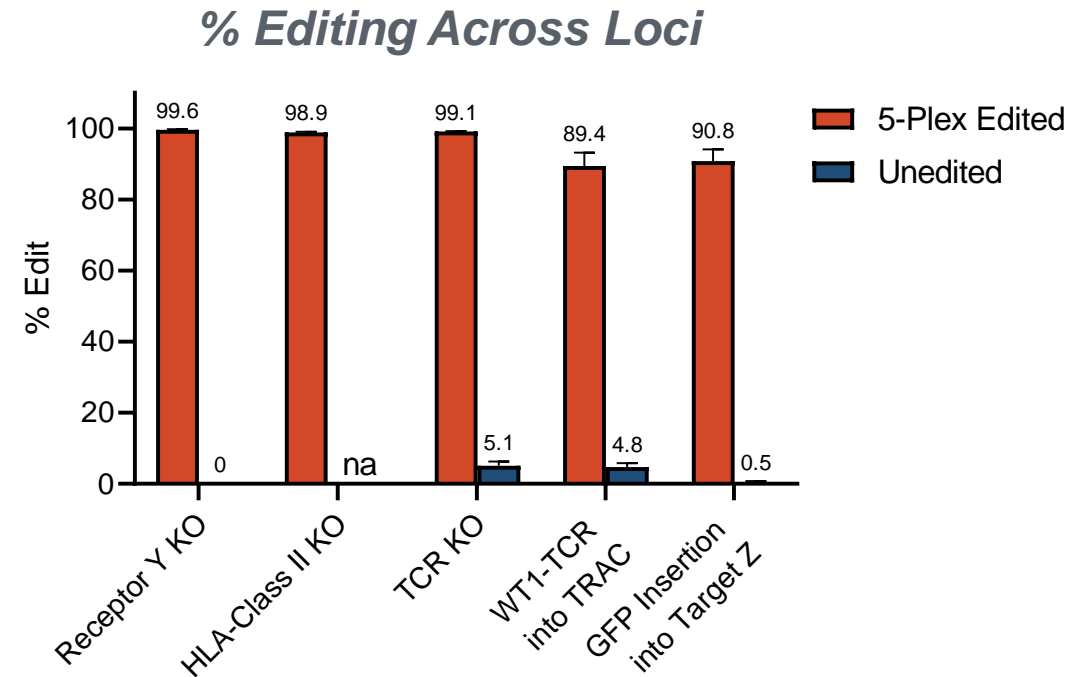
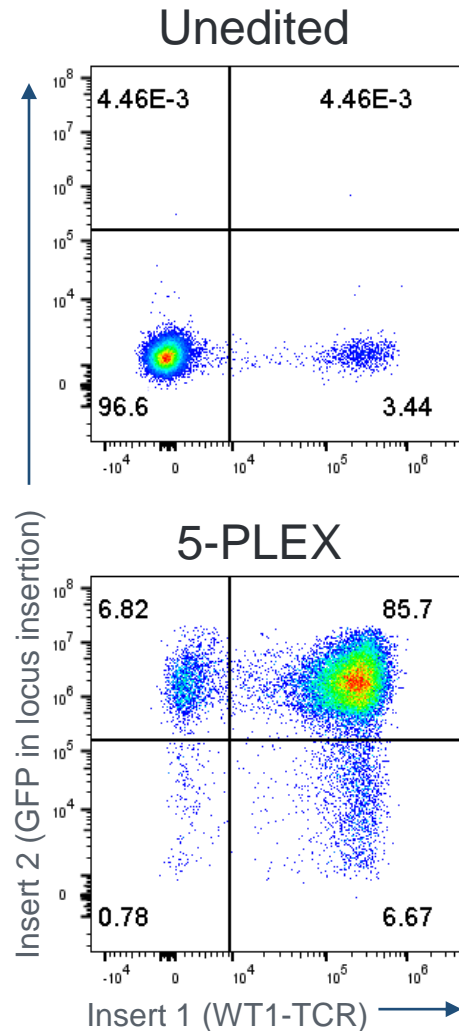


Re-stimulation Stress Test



Multiplex CRISPR/Cas9 T Cell Editing: 5 Sequential Edits with 2 Insertions

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



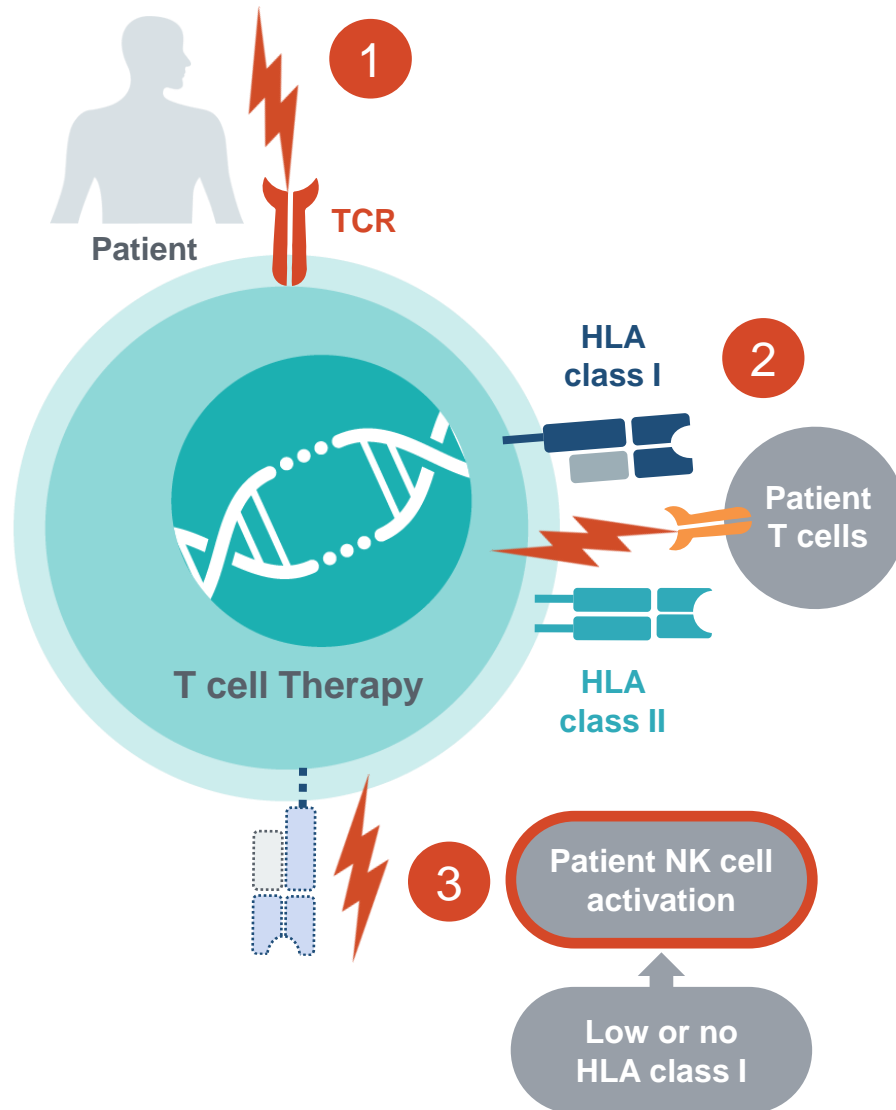
- >80% of cells have insertion of both the TCR and GFP transgene
- Cells retained high viability and complete editing of 3 other KO targets
- Modular platform for insertion of T cell supporting transgenes

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Intellia's Allogeneic Solution

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Three Immune Concerns Must Be Addressed by Allogeneic Cell Therapies



1 Graft versus host disease (GvHD)

T cell receptor (TCR) from allogeneic T cells recognizes and kills recipient (host) cells.

Largely solved with knockout (KO) of endogenous TCR

2 Rejection via host T cells

Human Leukocyte Antigen (HLA) molecules must match between donor and recipient to prevent rejection from:

- Host CD8 (HLA class I) T cells
- Host CD4 (HLA class II) T cells

3 Rejection via host natural killer (NK) cells

NK cells will attack cells that lack HLA-I expression or have low HLA-I

No validated solution yet

Immune Concerns Unaddressed by Current Allogeneic Solutions

Approach	Employ intense lymphodepletion regimen	Knockout (KO) HLA-I (B2M)	KO HLA-I & express NK inhibitor (HLA-E)	Intellia's Approach
				KO HLA-II & Receptor X*
Avoid rejection of cell therapy by host CD8 T cells	✓	✓	✓	✓
Avoid rejection of cell therapy by host CD4 T cells	✓	✗	✗	✓
Avoid rejection of cell therapy by host NK cells	✓	✗	✗	✓
Avoid profound immunosuppression	✗	✓	✓	✓

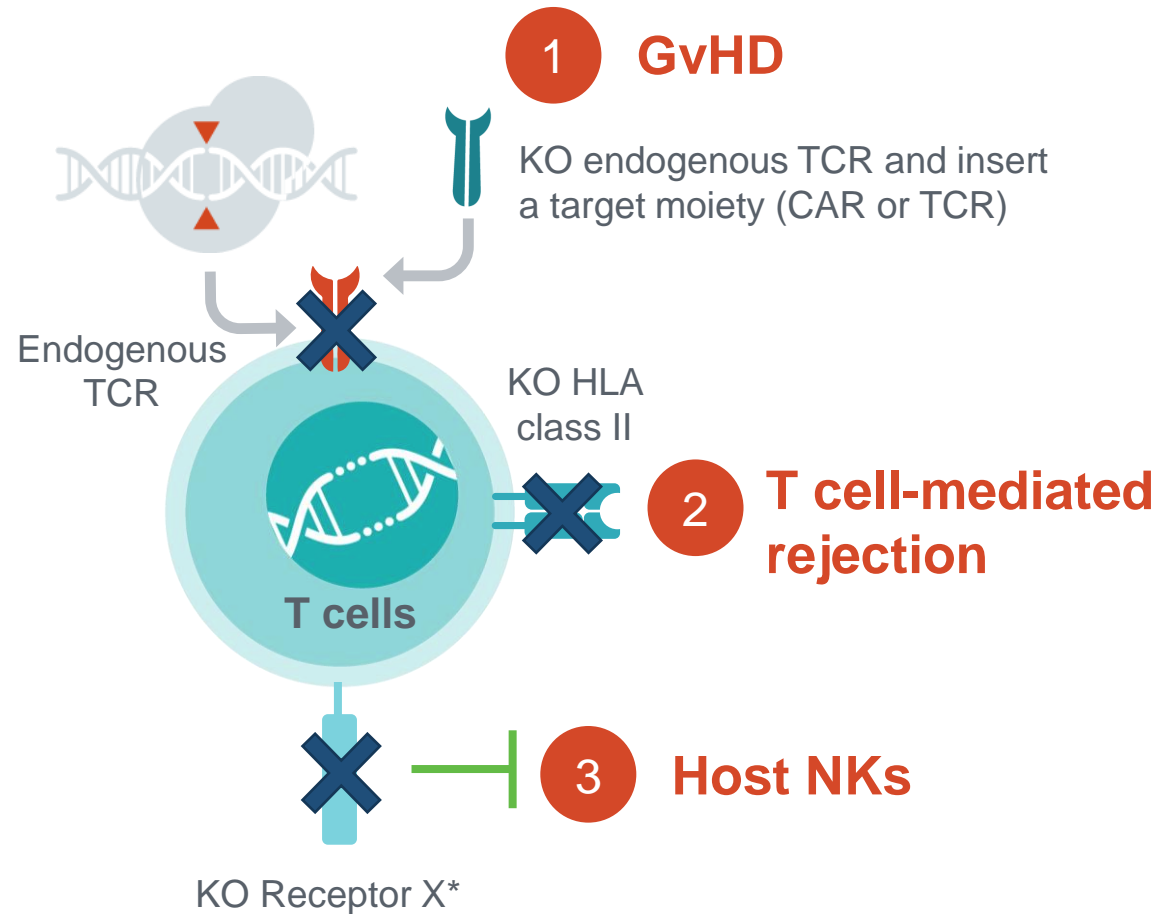
*Receptor X: Undisclosed target

B2M: Beta-2-microglobulin

HLA-E: Human leukocyte antigen class E

Slide based on preclinical data disclosed by Intellia. Intellia's cell product to be further explored in additional preclinical and clinical studies.

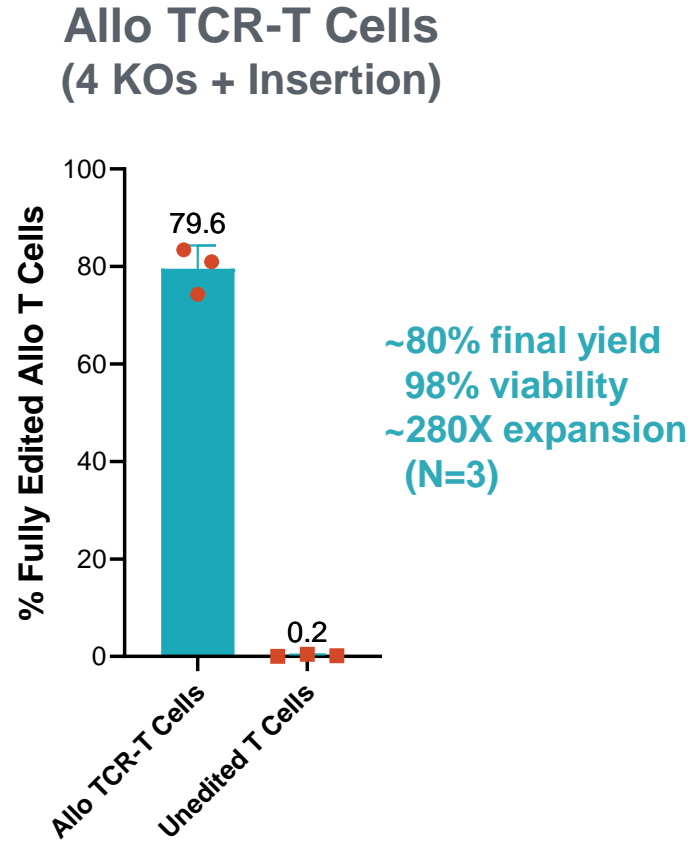
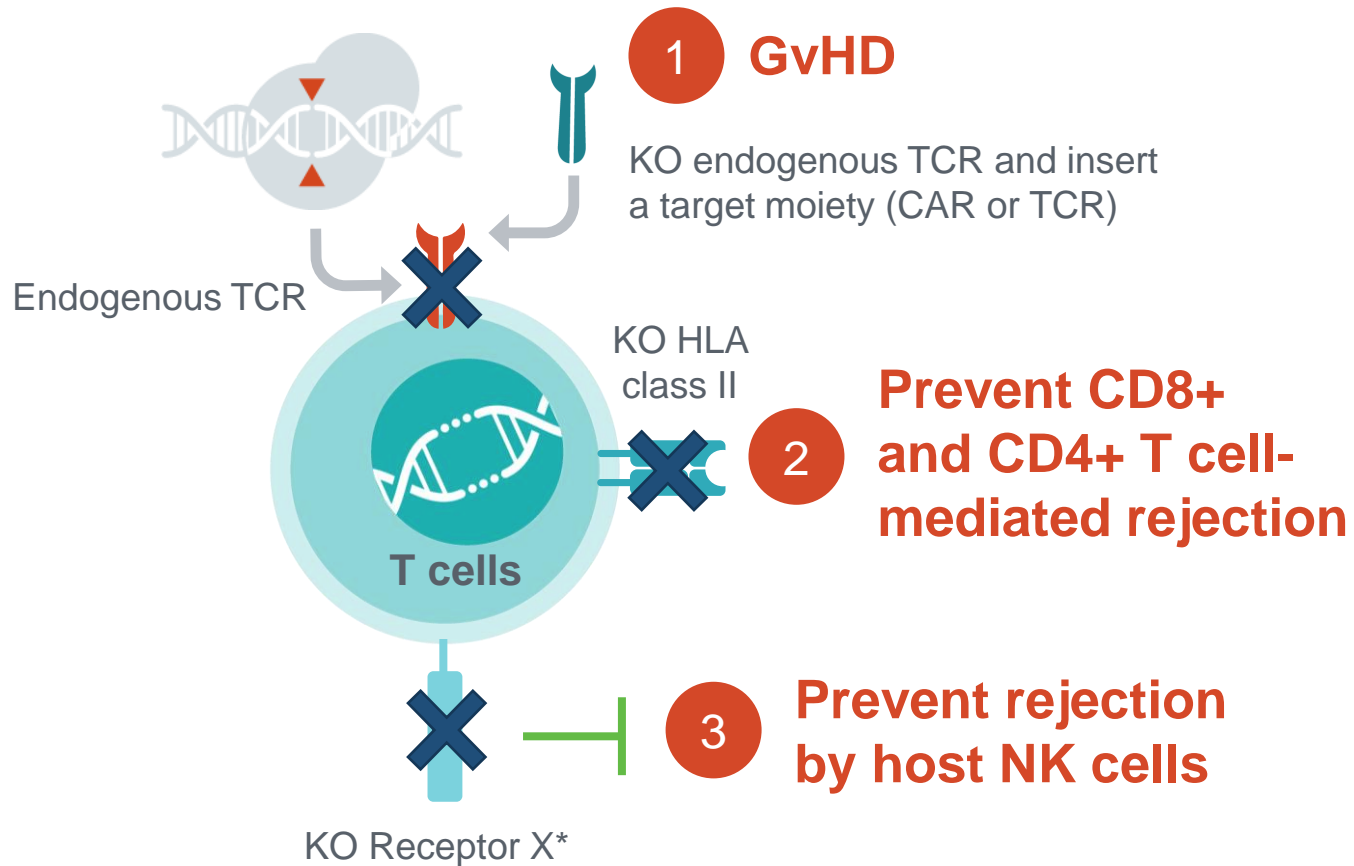
Intellia's Differentiated Allogeneic Approach Aims to Address All Three Immune Concerns



Potential Key Advantages

- Approach is applicable to CAR and TCR
- Solve for host NK and T cell rejection
- Avoid long-term immunosuppression

Intellia's Differentiated Allogeneic Approach Aims to Address Immune Requirements



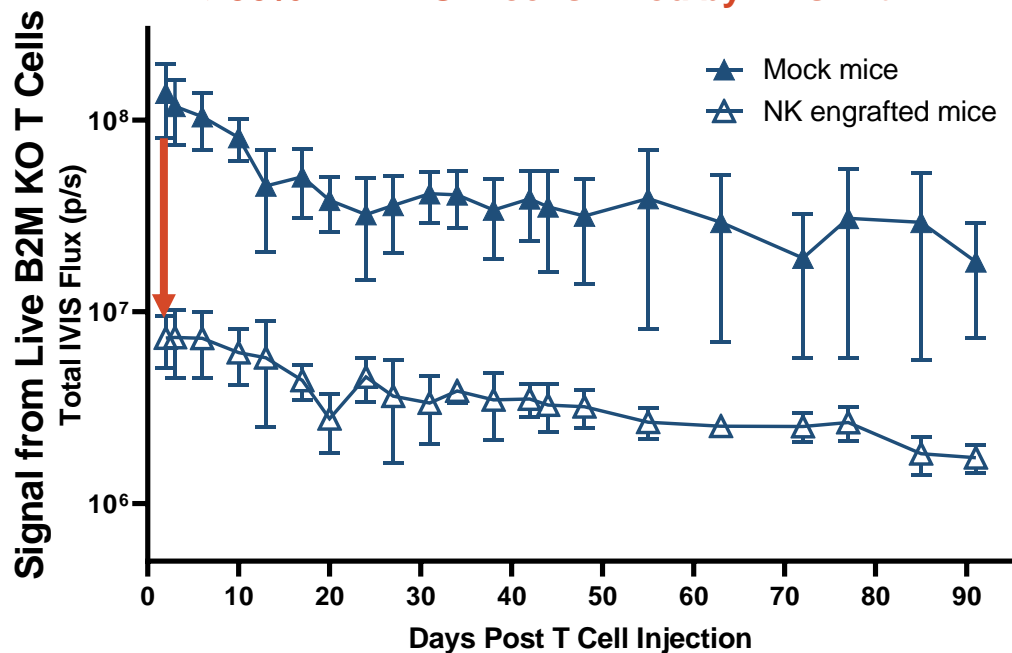
Applicable to CAR and TCR • Solve for host NK and T cell rejection • Avoid long-term immunosuppression

Allo TCR-T Cells Resist NK Cell Killing for at Least 90 Days *In Vivo*



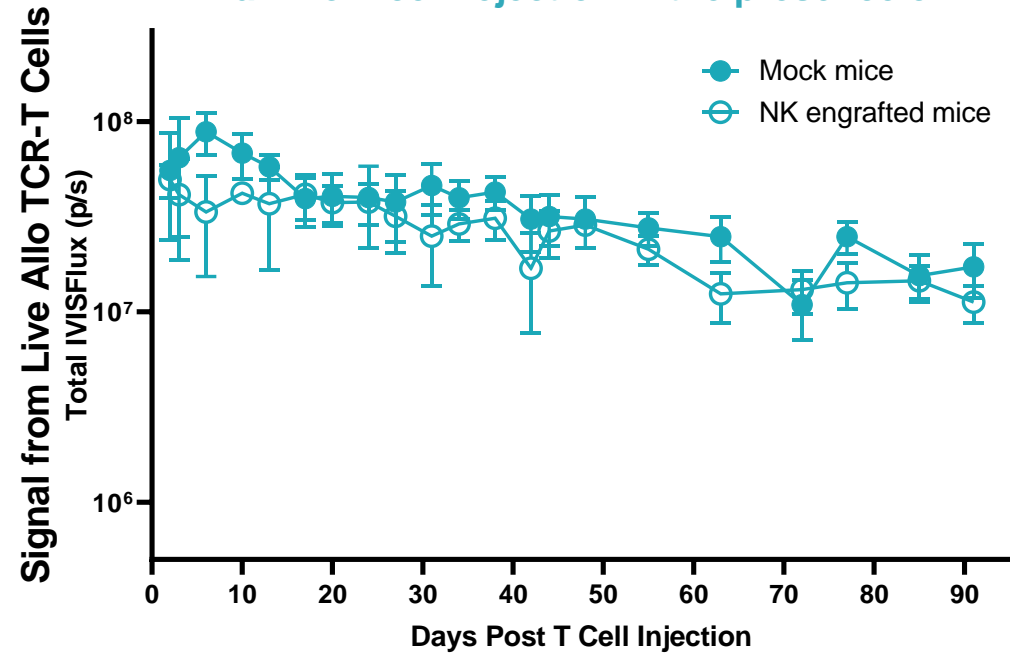
B2M Knockout T cells

>90% B2M KO T cells killed by NKs within 24h



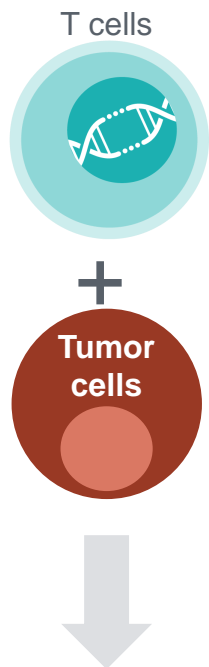
Allo TCR-T Cells

Minimal Allo T cell rejection in the presence of NK cells

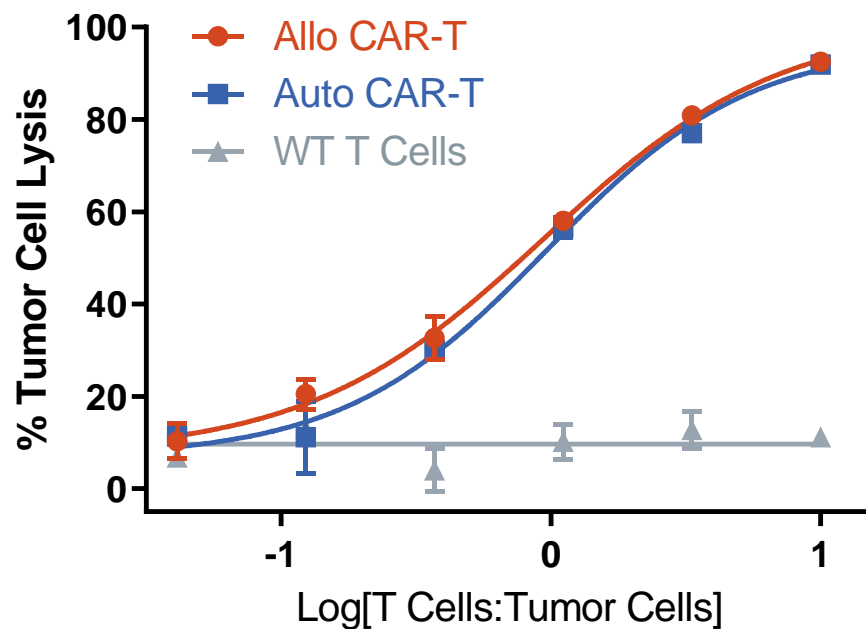


Allo T Cells Have Comparable Tumor Cell Killing Activity to Autologous T Cells

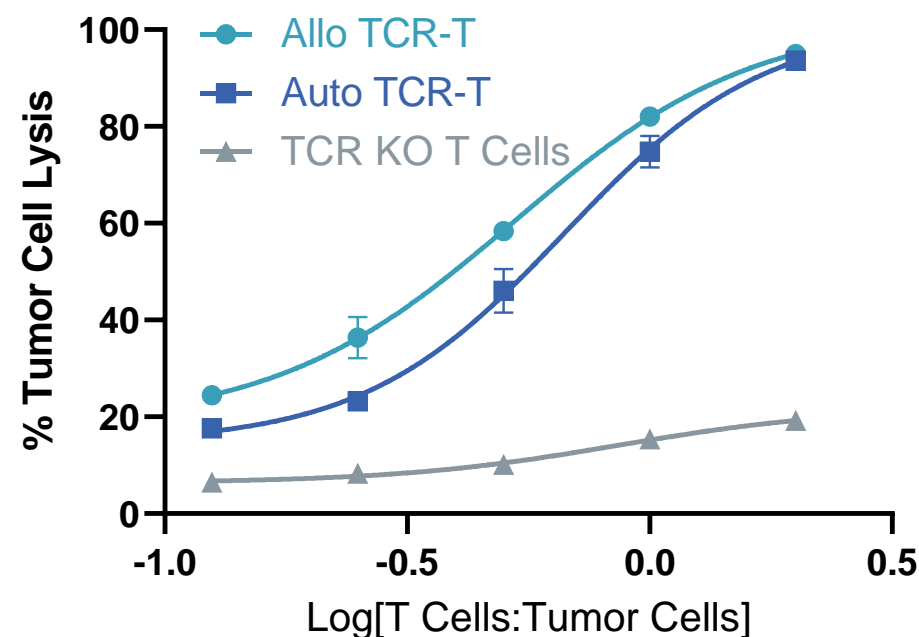
In vitro T cell
cytotoxicity assay



Allo CAR-T Cells



Allo TCR-T Cells



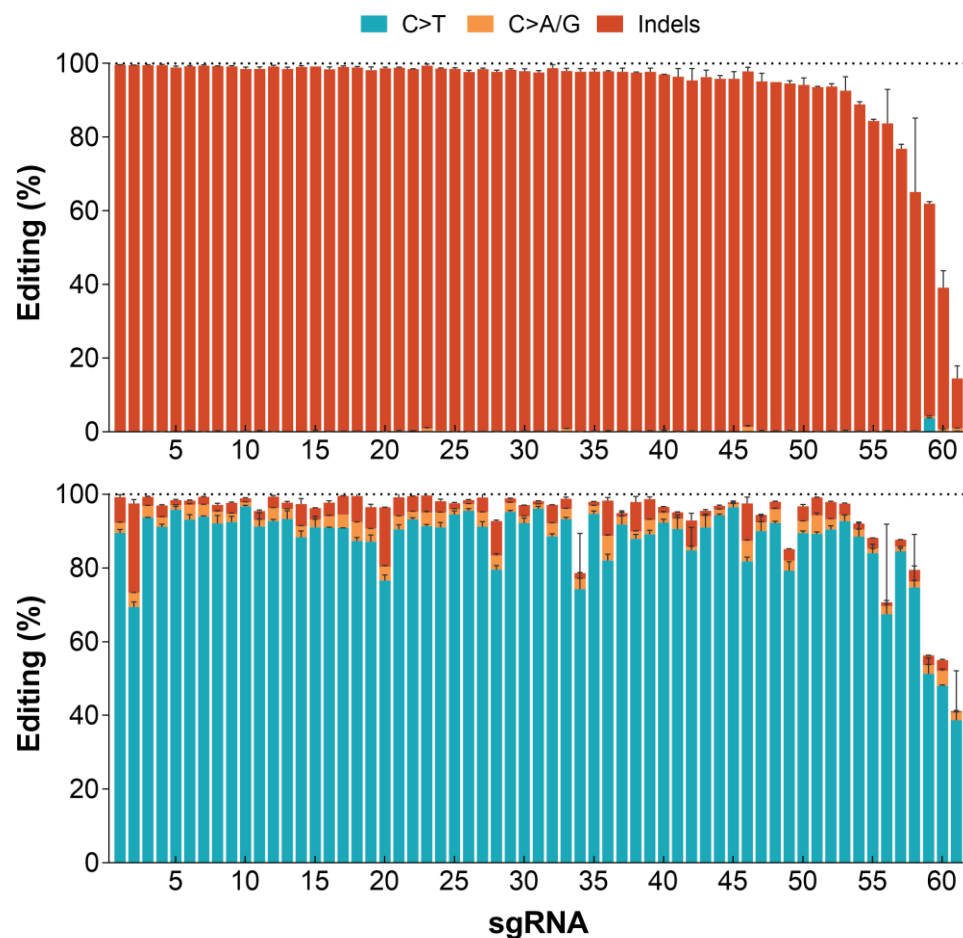
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Intellia's Proprietary Base Editor

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Intellia's Base Editor is Equipotent to Cas9 for *Ex Vivo* Editing

Intellia's base editor is highly active with similar activity to Cas9 cleavase

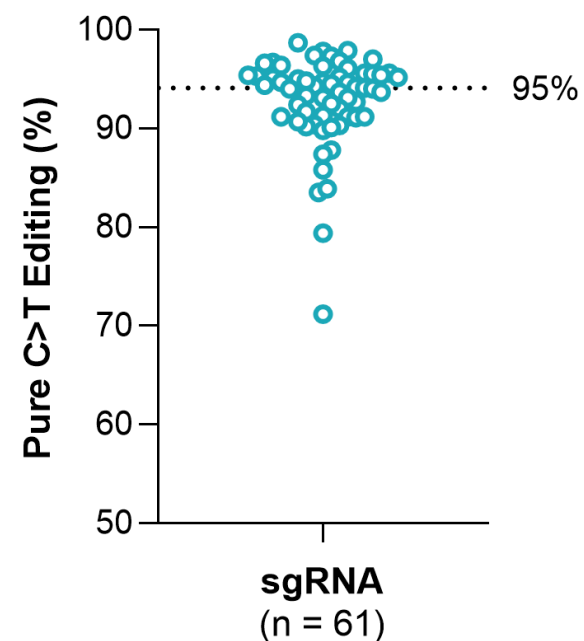


Cas9

Base Editor

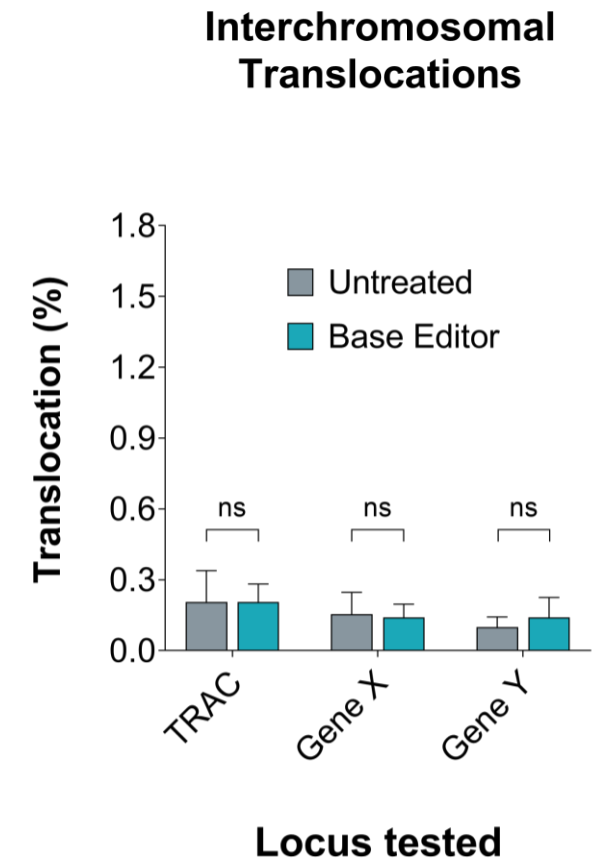
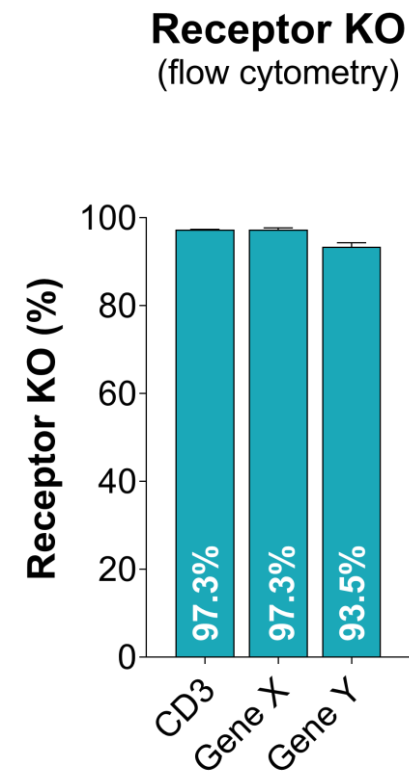
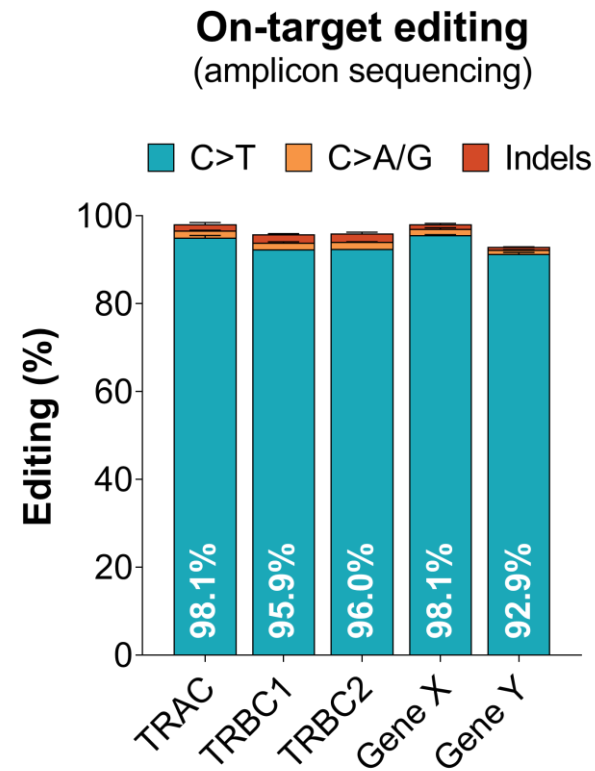
>700 constructs screened for potency
85% of guides gave >90% editing
36% of guides gave >95% C to T purity

Pure C>T edits
(without indels)



Simultaneous Knockout with Base Editing **Does Not** Lead to Translocations

- 1 Isolate primary T cells
- 2 Deliver base editor + 4 sgRNAs
- 3 Evaluate editing, receptor KO and translocations

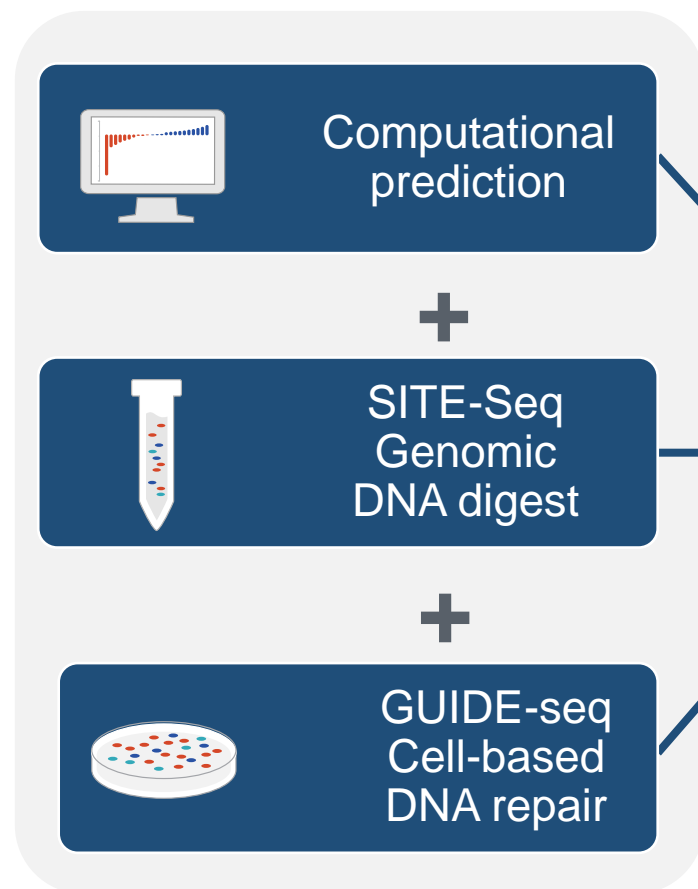




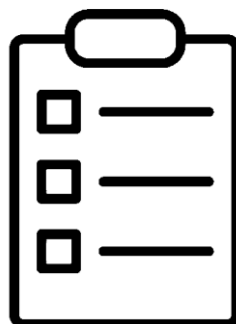
Platform: Identifying Potent and Highly Specific Guide RNAs

Comprehensive gRNA Specificity Assessment: Off-Target Workflow

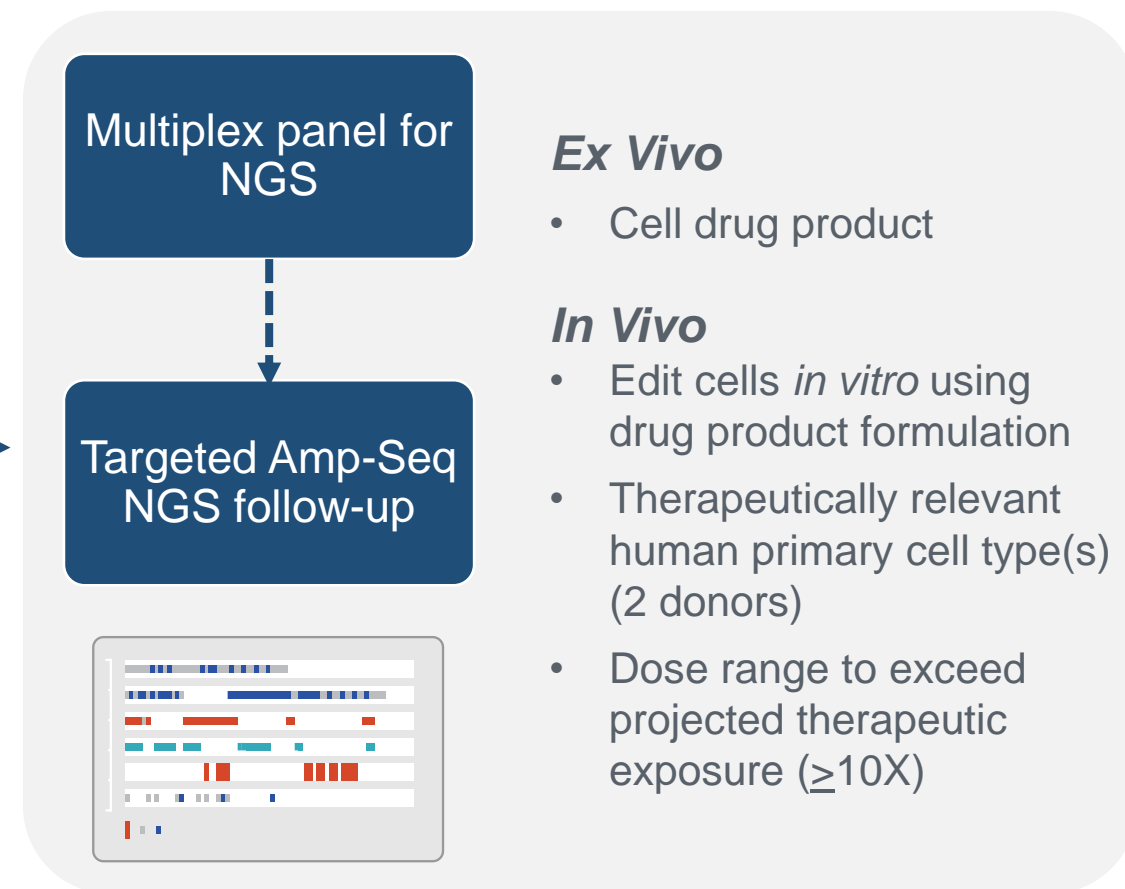
1: Discovery of Potential Off-Target Edits



Aggregate
ALL
potential
off-target
genomic loci

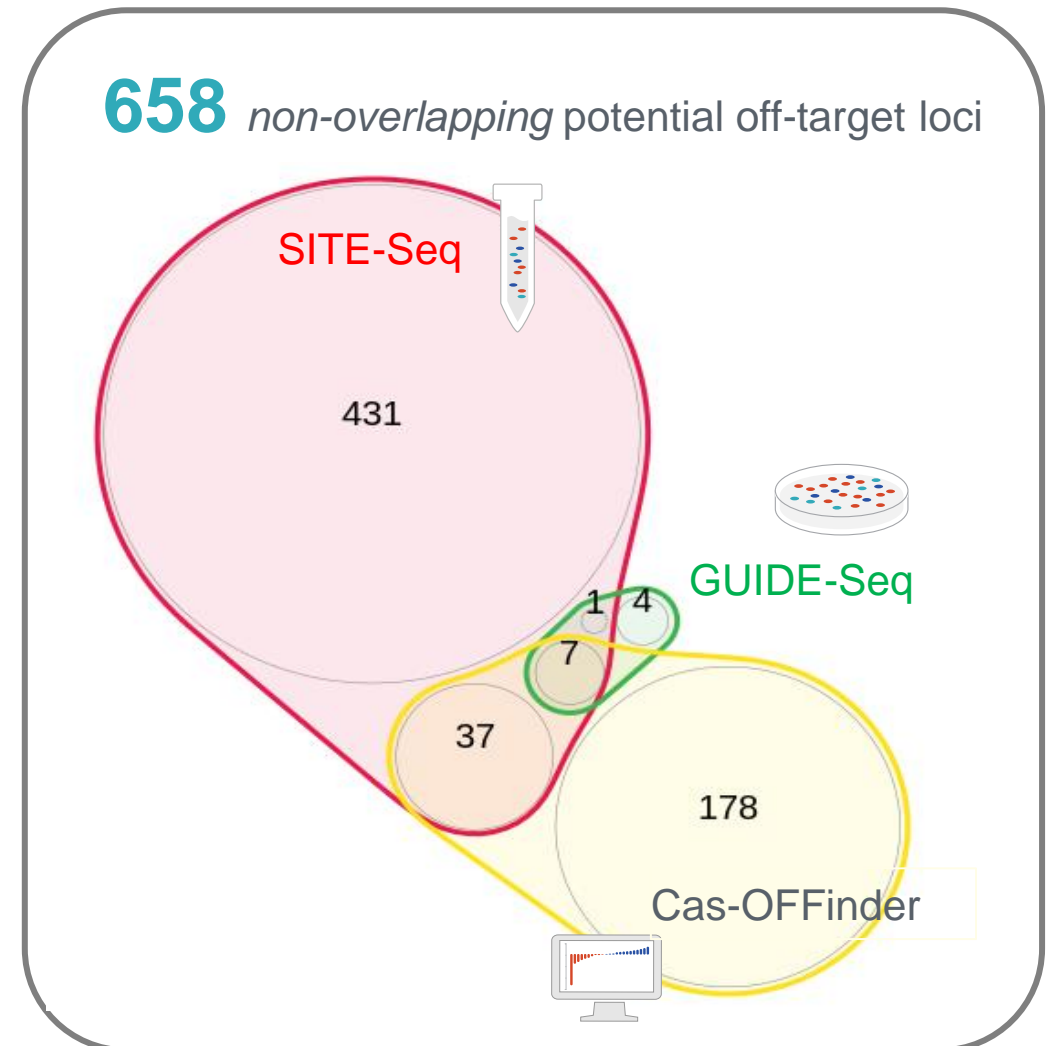
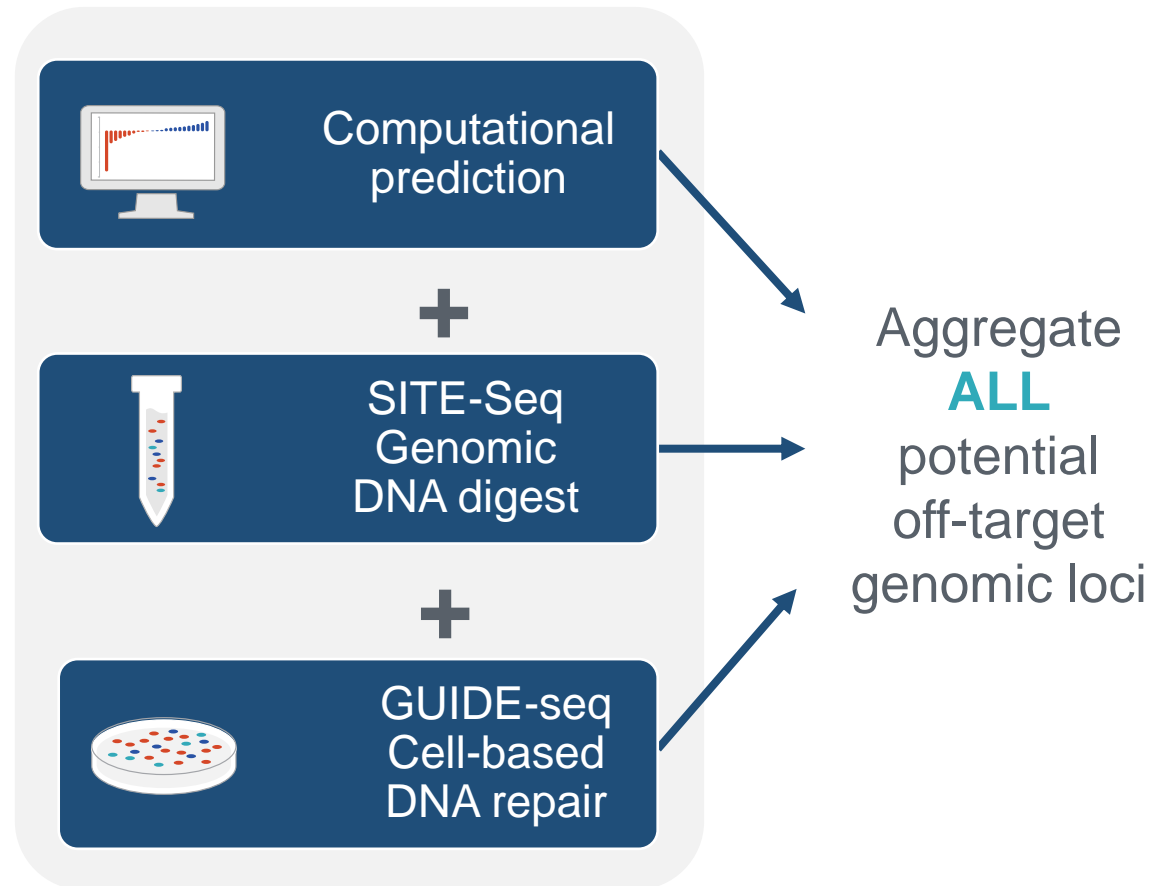


2: Validation of Off-Target Edits in Cells



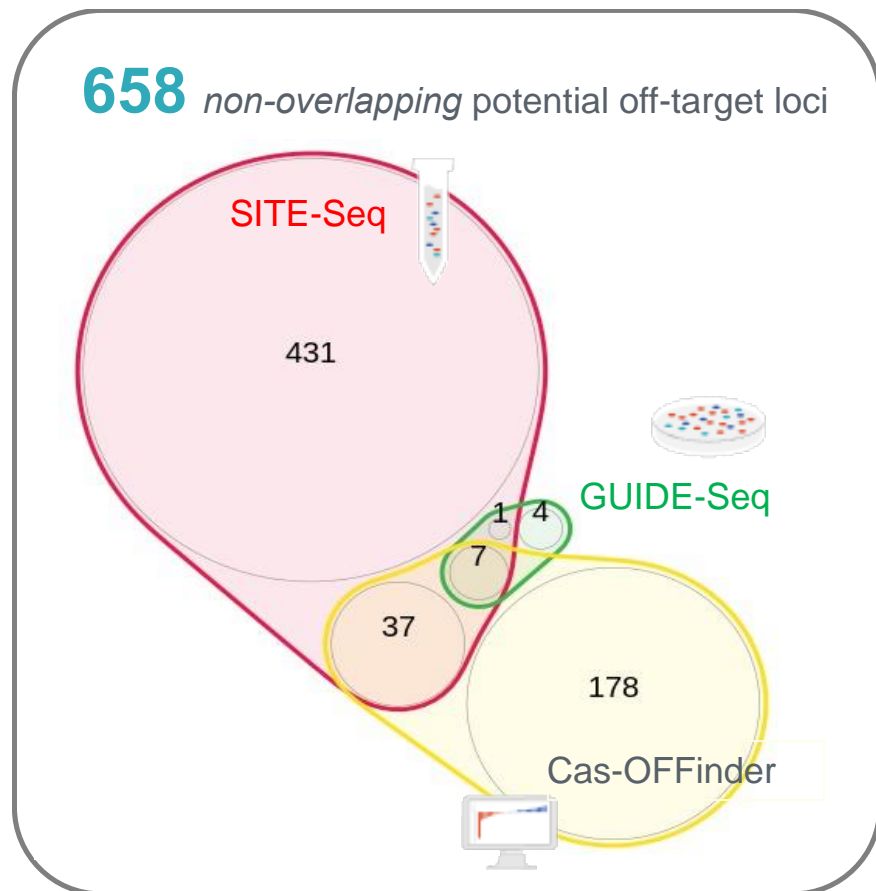
Limited Overlap in Discovered Off-Target Loci by Three Leading Methods

1: Discovery of Potential Off-Target Edits

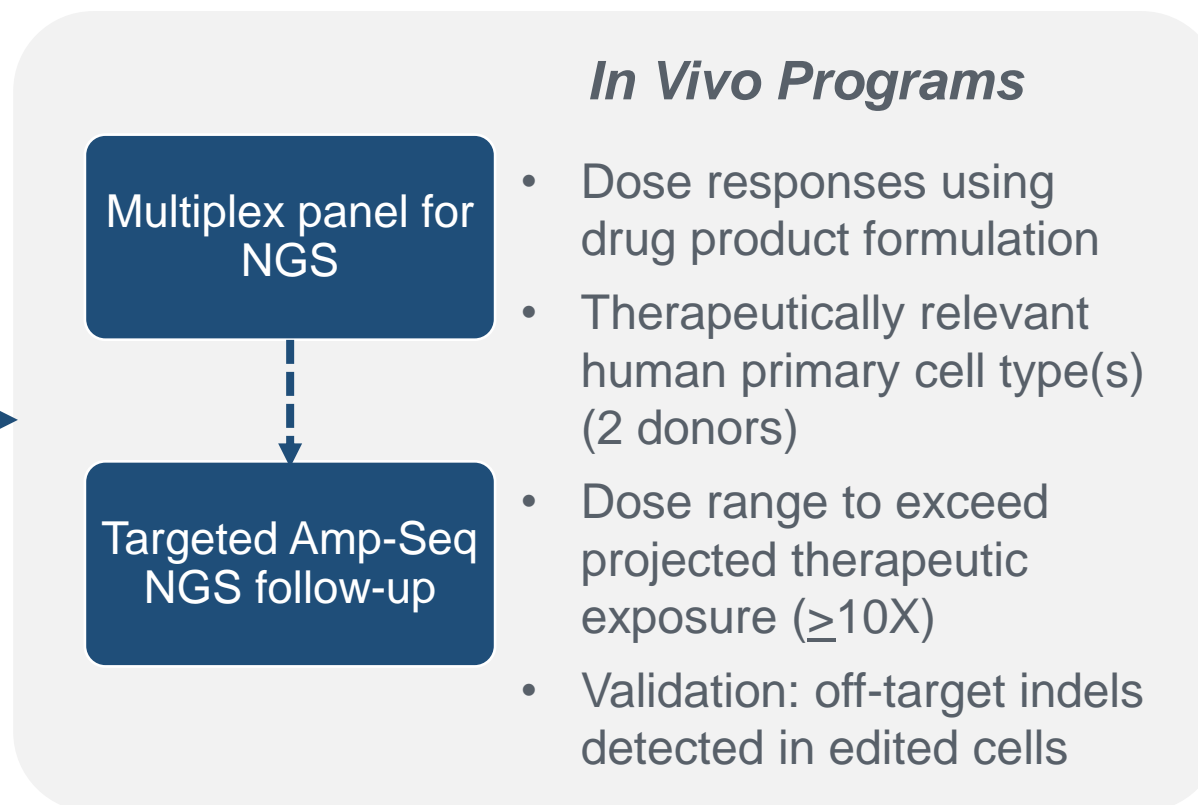


Off-Target Workflow In Practice: Representative Example

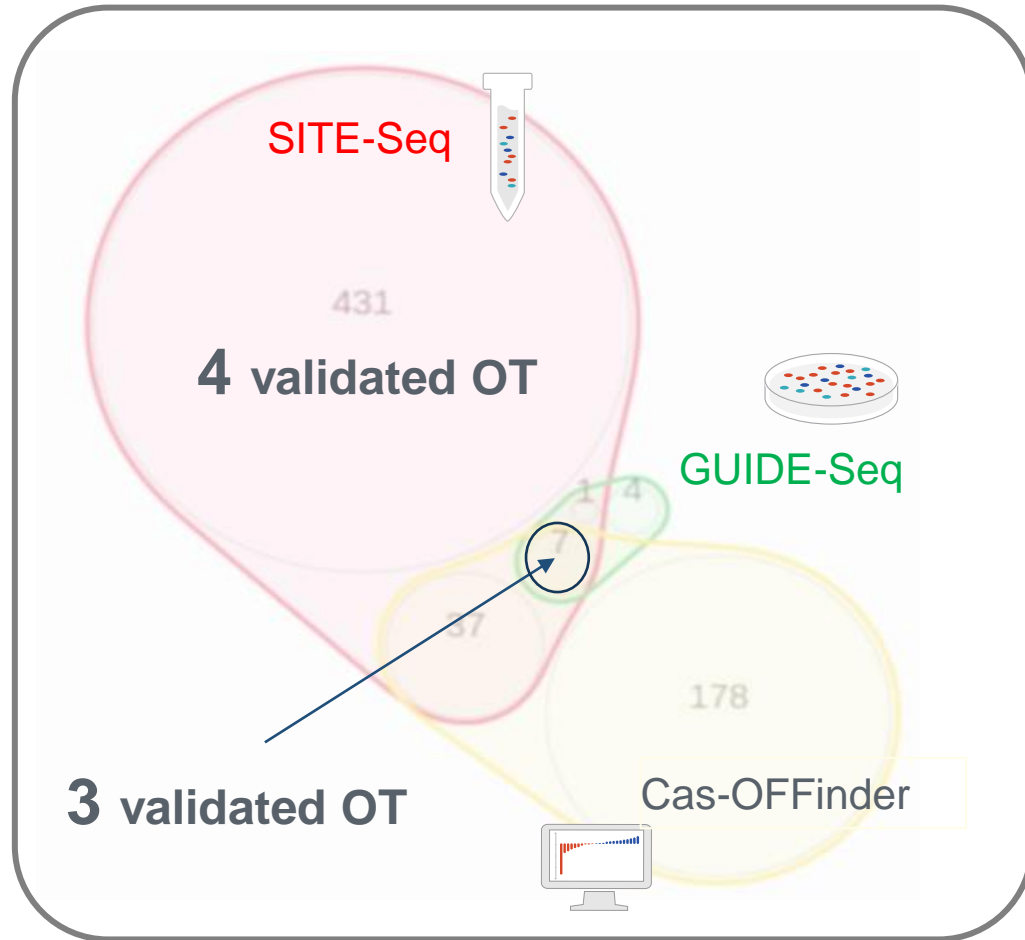
1: Discovery of Potential Off-Target Edits



2: Validation of Off-Target Edits in Cells



Validation of Off-Target Editing in Primary Human Hepatocytes at Supersaturating LNP CRISPR Concentrations to Maximize Sensitivity



658 potential off-target loci

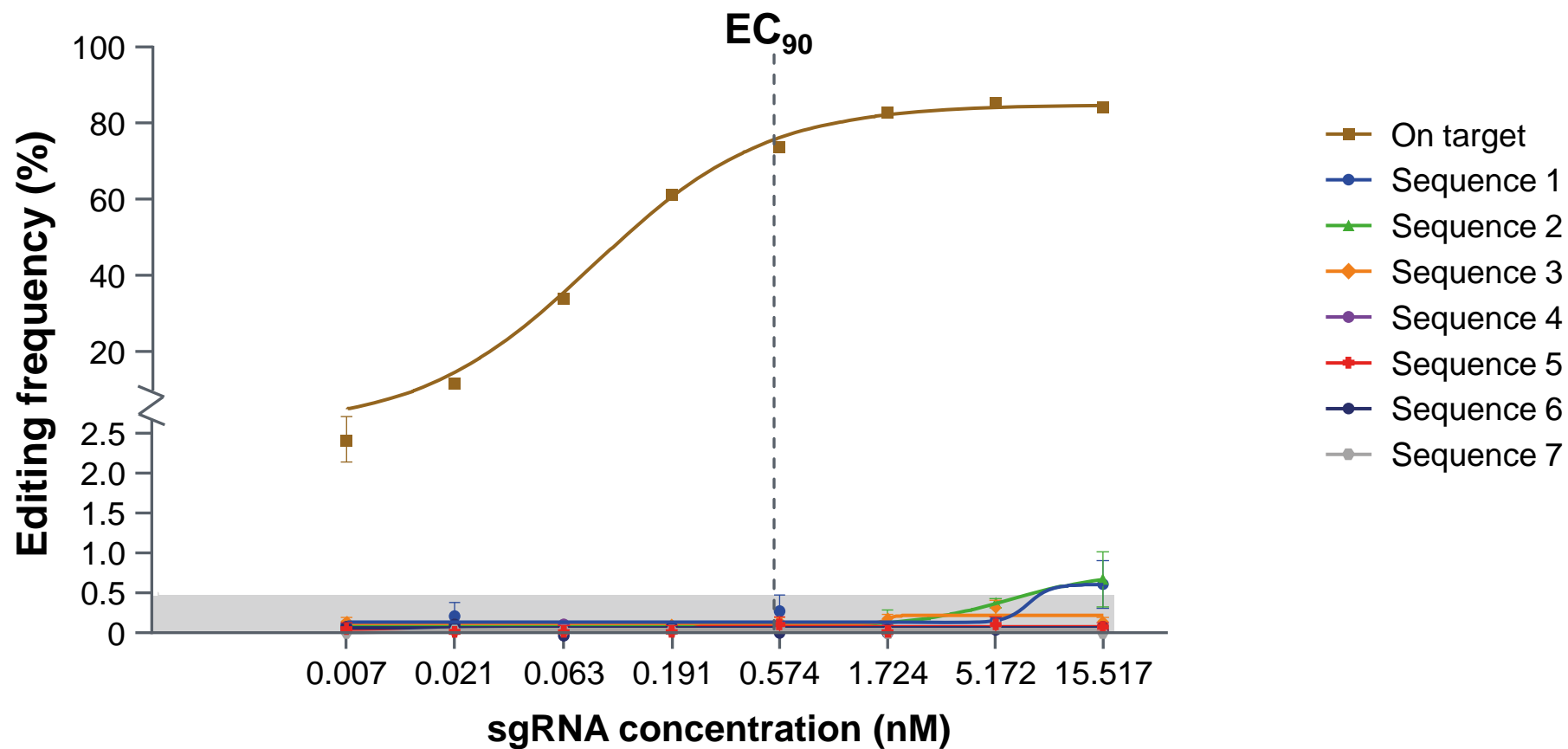


7 validated off-target (OT) loci

2 in introns and 5 in intergenic regions

- SITE-Seq discovered **100%**
- GUIDE-Seq and Cas-OFFinder discovered the same 3 out of 7 validated off-target loci **43%**
- Eliminate gRNA with validated off-target indels in regions of the genome associated with cancer

In Vitro: No detectable off-target editing with pharmacologic concentration of sgRNA



EC_{90} , concentration inducing 90% of maximal effect; sgRNA, single guide RNA



Strategic Collaborations

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Growing Intellia's Impact on Patients Through Strategic Collaborations

Increasing shareholder value:

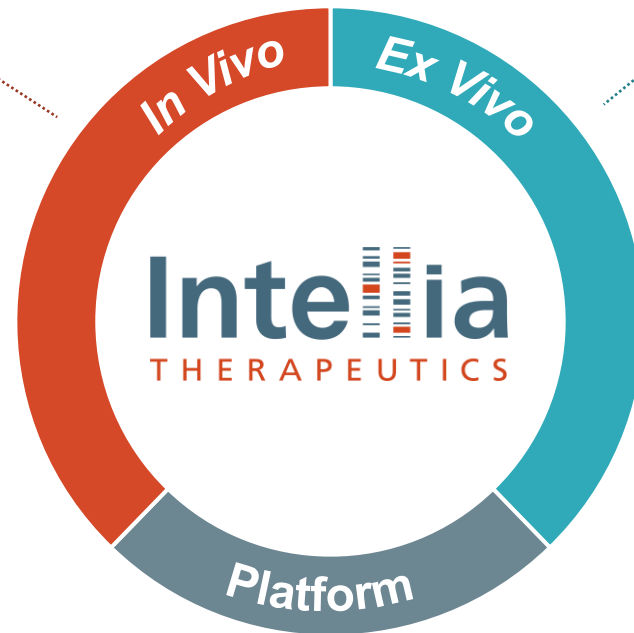
- Leveraging our technology while retaining rights to key areas of focus
- Accelerate development of programs outside key areas of focus
- Expand our pipeline with valuable rights in future commercial success
- Access external expertise to enhance our platform

Genetic diseases

REGENERON

Ophthalmology

SPARINGVISION
GENOMIC MEDICINES FOR OCULAR DISEASES



Immuno-oncology (IO)



IO (NK Cells)



Autoimmune diseases (CD19)



Sickle cell disease and IO



Foundational Partnerships Provided Access to R&D Capabilities

REGENERON

- Up to 15 *in vivo* targets with a mix of co-developed and licensed programs
 - Liver-centric product development
- **ATTR:** First selected Co/Co program
 - Intellia is lead party; Regeneron will share 25% of costs and profits
- **Hemophilia A and B:** Co/Co agreements based on targeted insertion capabilities
 - Regeneron is lead party; Regeneron will share 65% of costs and profits
- *In vivo* targets exclusively developed by Regeneron:
 - Up to \$320M in milestones per target
 - High single-to-low-double-digit royalties
- Non-exclusive license to certain platform IP on up to 10 *ex vivo* CRISPR products in defined cell types



- Advancing Phase 1/2 study for sickle cell disease based on CRISPR/Cas9-edited HSCs
- Research collaboration term concluded in December 2019
- Novartis selected various CAR-T, HSC and OSC targets for development
 - Up to \$230M in milestone payments per product
 - Mid single-digit royalties
 - All non-selected targets revert to Intellia

Intellia, Cellex and Blackstone Launch AvenCell to Develop Allogeneic Universal CAR-T Cell Therapies, With \$250M Committed Funding

Concurrent Cellex deal enables expansion and acceleration of Intellia's ex vivo pipeline with expanded manufacturing capabilities



- Rights to **two Co/Co options** in U.S. and key European countries on allogeneic universal CAR-T products
 - Intellia leads U.S. commercialization
- Additional **validation** of Intellia's proprietary allogeneic platform
- Hold substantial **equity stake** in NewCo
- **Access to Cellex cell therapy manufacturing site and allogeneic cell donations** via a preferred relationship
 - Supports Intellia's wholly owned *ex vivo* pipeline
 - Expanded capacity to handle additional pipeline growth



- Expansion of **existing Intellia-GEMoAB collaboration**
- Combines **GEMoAB's switchable universal CAR-T cell technology** with **Intellia's allogeneic platform** enabled by advanced CRISPR engineering
- Addition of **validating partner** Blackstone and **infusion of \$250M capital** to prosecute pipeline
- Clinical-stage autologous products from GEMoAB with **near-term milestones**
- **Seasoned management team**
- Access to Cellex cell therapy **manufacturing site**

Intellia

THERAPEUTICS