

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 about Intellia's beliefs and expectations regarding; our ability to build a world-class gene editing toolbox to develop an unsurpassed gene editing pipeline; the safety, efficacy and advancement of our clinical programs for NTLA-2001, also known as nexiguran ziclumeran or "nex-z", for the treatment of transthyretin ("ATTR") amyloidosis, NTLA-2002 for the treatment of hereditary angioedema ("HAE") and NTLA-3001 for the treatment of alpha-1 antitrypsin deficiency ("AATD")-associated lung disease pursuant to our clinical trial applications ("CTA") and investigational new drug ("IND") submissions, including the expected timing of data releases, regulatory filings, and the initiation and completion of clinical trials, such as initiating the Phase 3 study for the treatment of ATTR amyloidosis with polyneuropathy in 2024, presenting updated data from the ongoing Phase 1 study of NTLA-2001 in 2024, and dosing the first patient in the Phase 1 study of NTLA-3001 in 2024; the execution of its strategic priorities for 2024-2026, including the completion of patient enrollment for pivotal studies of NTLA-2001 and NTLA-2002, the planned BLA submission for NTLA-2002 for HAE in 2026, demonstrating human proof-of-concept for targeted in vivo gene insertion, initiating clinical development for its allogeneic ex vivo program, demonstrating preclinical proof-of-concept of editing in tissues outside the liver, and advancing DNA writing technology; the ability to generate data to initiate clinical trials and the timing of CTA and IND submissions; the advancement, expansion and acceleration of our CRISPR/Cas9 technology and related technologies, including DNA writing, base editing, manufacturing and delivery technologies, to advance and develop additional candidates and treatments; our ability to demonstrate our platform's modularity and replicate or apply results achieved in preclinical studies, including those in its NTLA-2001, NTLA-2002 and NTLA-3001 programs, in any future studies, including human clinical trials; our ability to optimize the impact of our collaborations on our development programs, including, but not limited to, collaborations with Regeneron Pharmaceuticals, Inc. ("Regeneron"), including our co-development programs for ATTR amyloidosis and hemophilia A, 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 ReCode Therapeutics, Inc. ("ReCode") for the development of novel genomic medicines for the treatment of cystic fibrosis, 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; the potential commercial opportunities, including value and market, for our product candidates, including the potential of NTLA-2001, NTLA-2002 and NTLA-3001 to be a single-dose treatment, the potential of NTLA-2001 to halt and reverse disease and result in lifelong, stable TTR reduction, the potential of NTLA-2002 to eliminate significant treatment burden; and the potential of NTLA-3001 to achieve normal human levels of alpha-1 antitrypsin protein and halt progression of lung disease; and our use of capital and other financial results.

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 flings 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, including uncertainties related to regulatory approvals to conduct clinical trials; 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 development of novel platform capabilities, including technologies related to editing in tissues outside the liver, base editing and DNA writing; risks related to Intellia's reliance on collaborations, including that its collaborations with Regeneron, AvenCell, SparingVision, ReCode, 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 entit



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Intellia is Leading a New Era of Medicine

Turning Nobel-Prize-Winning Science into Medicine

- Poised to bring first-ever in vivo CRISPR therapy to market
- Initiated first-ever, pivotal Phase 3 program for an in vivo CRISPR therapy
- Three Phase 3 studies actively enrolling patients

150+ patients
dosed with Intellia's
investigational *in vivo*CRISPR-based therapies

Robust pipeline of in vivo and ex vivo programs

Comprehensive gene editing toolbox



Advancing a Full-Spectrum Genome Editing Company

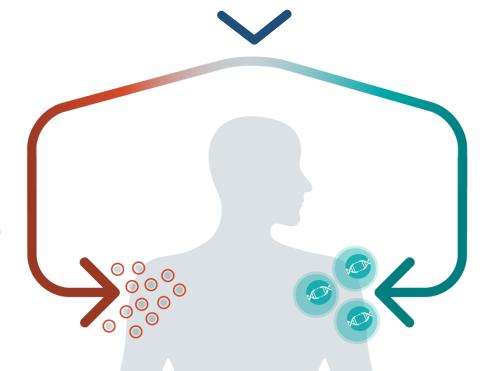
CRISPR-Based Modular Platform

EMPLOY NOVEL EDITING AND DELIVERY TOOLS

In Vivo CRISPR <u>is</u> the therapy

FIX THE TARGET GENE

Genetic diseases



Ex Vivo
CRISPR creates

the therapy

REWIRE & REDIRECT CELLS

Immuno-oncology
Autoimmune diseases



Intellia is Developing Potentially Curative Gene Editing Treatments to Transform the Lives of Patients

Full-Spectrum Strategy

Pipeline of *in vivo* and *ex vivo* CRISPR-based therapies for life-threatening diseases with high unmet need

Clinically Validated Modular Platform

Modular technology enables a reproducible path to drug discovery and development

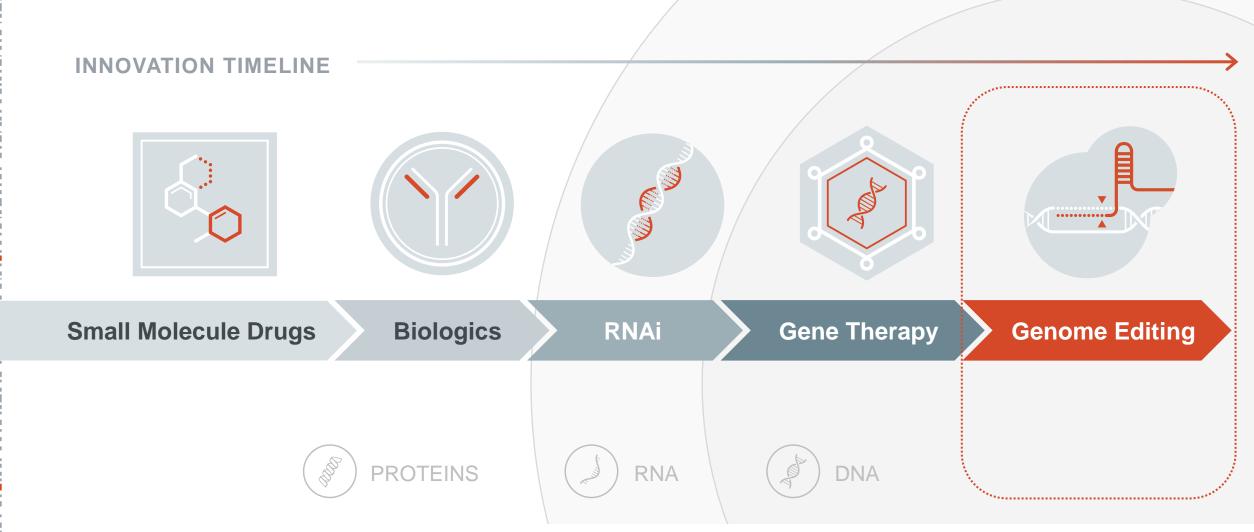
Deploying Novel Tools

Continued innovation across editing and delivery modalities for future therapeutic applications





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





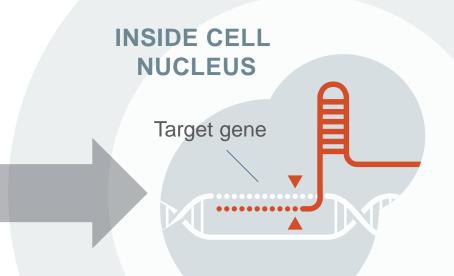
Gene Editing Starts with CRISPR/Cas9, a Two-Part, Programmable System

FOUNDATIONAL CRISPR MACHINERY



Guide RNA (gRNA) Identifies genetic target

Cas Protein Responsible for the targeted DNA editing and provides platform for other enzymatic activities



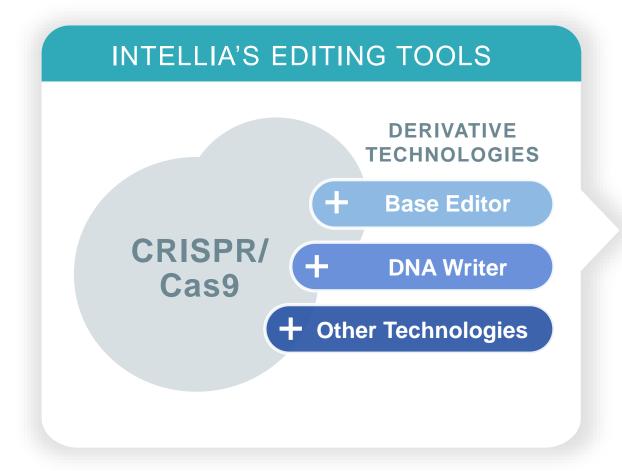
KEY FEATURES OF CRISPR/CAS9 SYSTEM

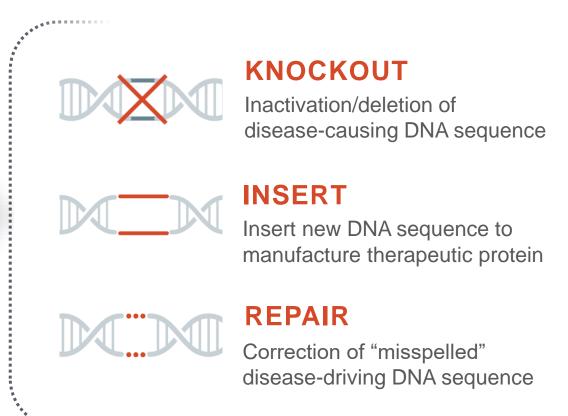
- Selectivity

- ✓ High potency ✓ Address any site ✓ Target multiple DNA sites.



CRISPR/Cas9 and Derivative Gene Editing Technologies Can Be Used to Make Any Type of Edit





INTELLIA SELECTS THE BEST TOOL FOR EACH THERAPEUTIC APPLICATION



A Tailored Approach to Maximize the Reach of Gene Editing Across Multiple Tissues

INTELLIA'S DELIVERY TOOLS



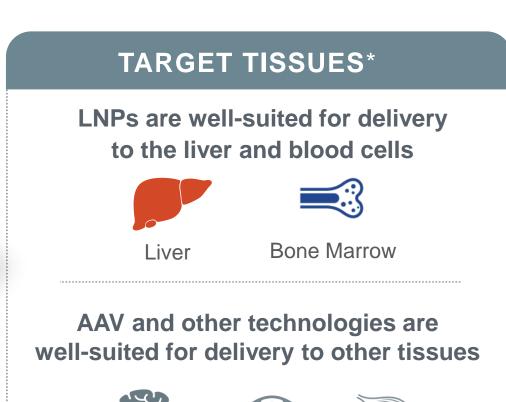
LNP: Livertargeted



LNP:
Bone marrow-targeted







Eve

CNS/PNS



Muscle

CRISPR-Based Editing Technologies are a Promising New Therapeutic Modality

Potential of CRISPR-Based Editing Technologies



Treat patients at the root cause of their disease



Single dose
treatment with potential
lifelong benefit



Reduce burden to the healthcare system over a patient's lifetime



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



The NEW ENGLAND JOURNAL of MEDICINE

October 24, 2024

CRISPR-Based Therapy for Hereditary Angioedema



The NEW ENGLAND
JOURNAL of MEDICINE

January 31, 2024

CRISPR-Cas9 In Vivo Gene Editing of KLKB1 for Hereditary Angioedema



The NEW ENGLAND
JOURNAL of MEDICINE

November 16, 2024

CRISPR-Cas9 Gene Editing with Nexiguran Ziclumeran for ATTR Cardiomyopathy



Intellia's Strategic Priorities for 2024 – 2026

- 1 Execute pivotal trials for first two *in vivo* CRISPR-based therapies
- 2 Launch next wave of *in vivo* and *ex vivo* clinical programs
- 3 Deploy new gene editing and delivery modalities

- Complete patient enrollment for pivotal studies of NTLA-2001, also known as nexiguran ziclumeran (nex-z), and NTLA-2002
- Planned BLA submission for NTLA-2002 for HAE in 2026

- Demonstrate human proof-of-concept for targeted in vivo gene insertion
- Initiate clinical development for first allogeneic ex vivo program

- Demonstrate preclinical proof-of-concept of editing in tissues outside the liver
- Advance DNA writing technology



Upcoming 2024 Key Clinical Program Milestones

NTLA-2001 Dose first patient in pivotal Phase 3 MAGNITUDE trial for ATTR-CM in Q1 2024 (nex-z) **ATTR** Continue to open new sites and enroll patients Present updated data from the ongoing Phase 1 study in 2H 2024 Initiate a pivotal Phase 3 study for ATTRv-PN by year-end NTLA-2002 Present updated data from the Phase 1 portion in 2024 HAE Initiate the Phase 3 study in 2H 2024 Present data from the Phase 2 portion in 2H 2024 NTLA-3001 Dose first patient in Phase 1 study of NTLA-3001 by year-end 2024 **AATD**



Broad Development Pipeline Fueled by Robust Research Engine

PROGRAM	APPROACH	Research and Preclinical	Early-Stage Clinical	Late-Stage Clinical	PARTNERS		
In Vivo: CRISPR is the therapy							
NTLA-2001 (nexiguran ziclumeran*): Transthyretin Amyloidosis	Knockout				Intelia REGENERON THERAPEUTICS		
NTLA-2002: Hereditary Angioedema	Knockout				Intelia THERAPEUTICS		
NTLA-3001: AATD-Lung Disease	Insertion				Intelia THERAPEUTICS		
Hemophilia A / B***	Insertion				Intelia REGENERON LEAD		
Research Programs	Knockout, insertion or repair				Intelia THERAPEUTICS		
Research Programs	Tissues outside the liver				THERAPEUTICS ReCode REGENERON SPARINGVISION		
Ex Vivo: CRISPR creates the therapy							
Research Programs	Allogeneic and other				** kyverna. THERAPEUTICS ** KERAPEUTICS		



^{*} NTLA-2001 is also known as nexiguran ziclumeran (nex-z)



^{**} Intellia is advancing both wholly owned and partnered programs.

^{***} Hemophilia A program is in the research stage; Hemophilia B is being advanced by Regeneron – Intellia is eligible for milestones and royalties.

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In Vivo

CRISPR is the therapy

GENETIC DISEASES

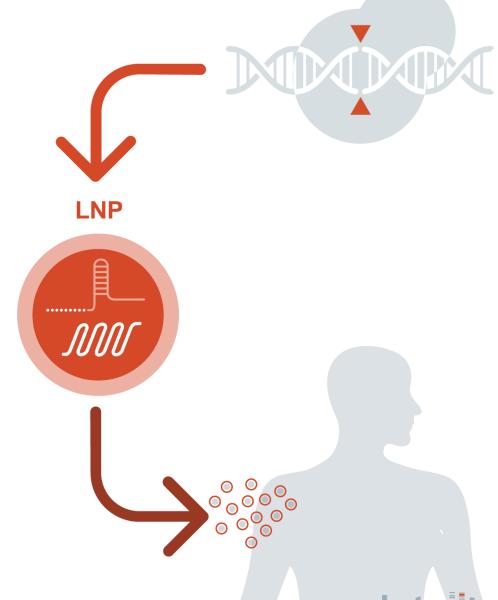
Strategic Advantages:

Potential curative therapy from a single dose

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

Potential for permanent gene knockout or gain of function by targeted 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: gRNA identifies genetic target **TTR** gRNA KLKB1 **qRNA Target**specific **mRNA**

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 (nex-z*) for Transthyretin (ATTR) Amyloidosis

About ATTR Amyloidosis

- Caused by accumulation of misfolded TTR protein
- Primarily affects the nerves and/or the heart
- Chronic dosing is required with current treatment options

Our Approach

Knock out *TTR* gene with a single-dose CRISPR-based treatment

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

Key Advantages Includes Potential to:

- Halt and reverse disease with deep and consistent TTR reduction
- Be a single-dose treatment
- Expect lifelong, stable TTR reduction



ATTR Amyloidosis: Large Commercial Opportunity with Significant Unmet Need

NTLA-2001 (nex-z)

Potential to be the best-in-class TTR reduction agent and only single-dose treatment

Prevalence^{1,2}

50,000

ATTRv patients worldwide

~200-500K

ATTRwt patients worldwide

Life Expectancy³

2-7 years after diagnosis for ATTR-CM patients

10+ years

after diagnosis for ATTRv-PN patients

Disease Burden⁴

Patients experience **highly burdensome symptoms**, including heart failure, shortness of breath, muscle weakness and sensory deficits

Commercial Opportunity^{5,6}

\$11B+
global market size
expected by 2029

\$450K+

average annual cost of TTR reduction treatment in the U.S.



¹ Hawkins et al. *Ann Med.* 2-15; 47(8): 625–638

² Compiled from various sources.

³ Luigetti et al. Ther Clin Risk Manag. 2020; 16:109-123

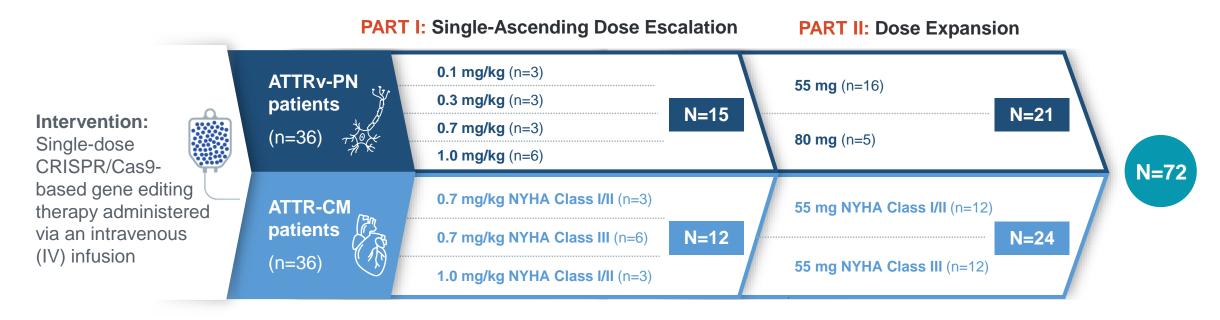
⁴ Griffin et al. JACC 2021; Intellia Patient Survey 2022

GlobalData 2023

⁶ Redbook 2023

NTLA-2001 (nexiguran ziclumeran) Phase 1 Study in ATTR Amyloidosis

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



PRIMARY OBJECTIVES

Evaluate safety, tolerability, PK and PD

Measure serum TTR levels

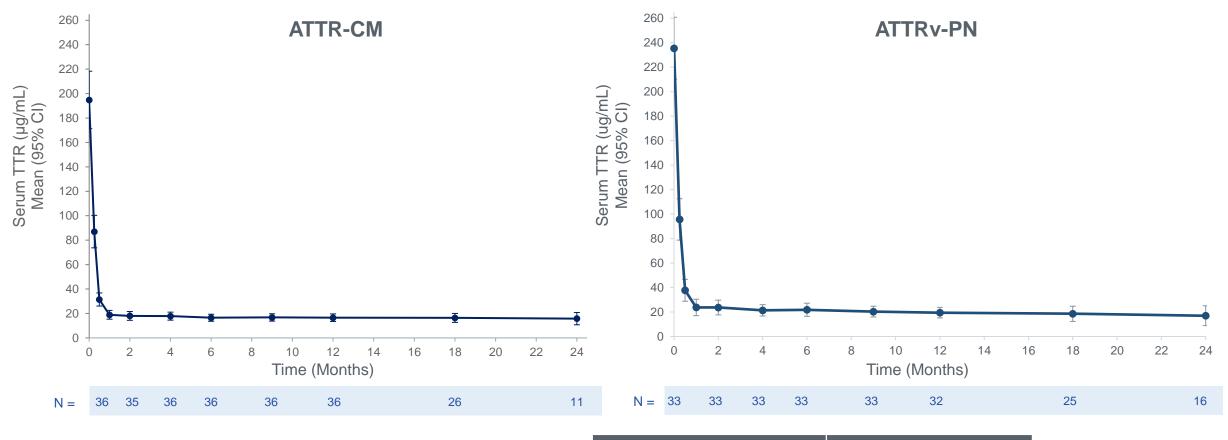
SECONDARY OBJECTIVES

Evaluate efficacy on clinical measures of:

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



Nex-z Led to Deep, Rapid and Durable Reductions in Absolute Serum TTR in Every Patient Following a Single Dose

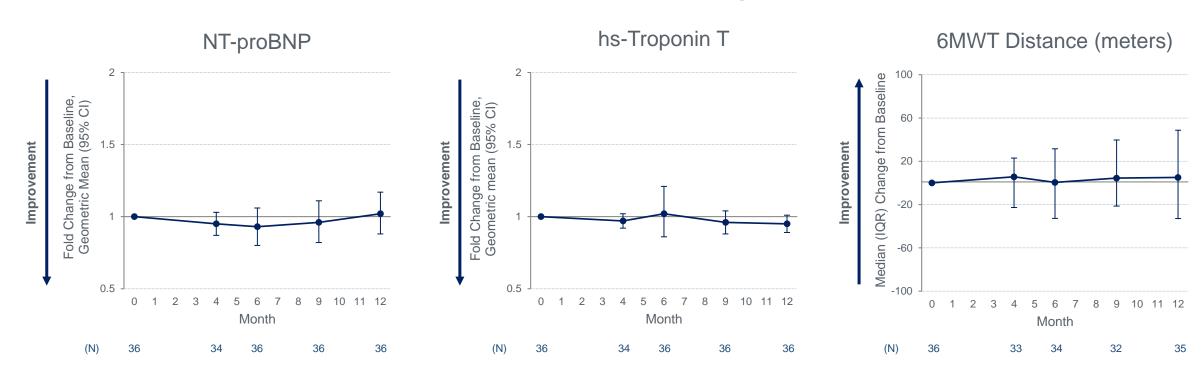


		ATTR-CM (n=36)	ATTRv-PN (n=33)
Mean (95% CI) Serum TTR at	Residual absolute TTR concentration	18.9 µg/mL (15.4 to 22.5)	23.8 µg/mL (17.0 to 30.5)
Day 28	% Change from baseline in serum TTR	-89% (-92 to -87)	-90% (-92 to -87)



Nex-z Treatment Led to Stability of NT-proBNP, hs-Troponin T, and 6MWT Over 12 Months

Markers of Disease Progression¹

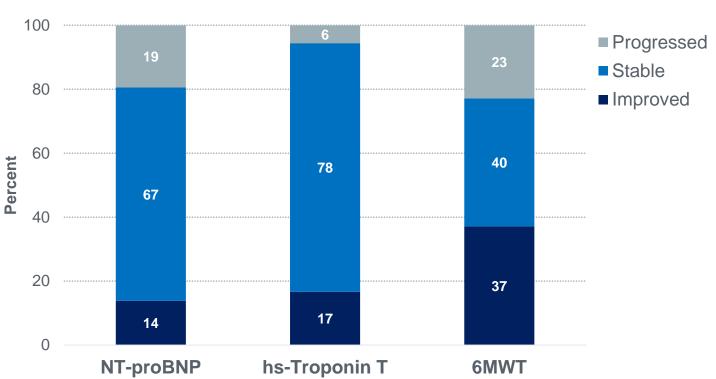


Consistent with cardiac disease marker data, similar pattern of stability or improvement was observed in symptoms, QOL and assessments of cardiac structure



Nearly 80% of Patients Demonstrated Stability or Improvement in Markers of Disease Progression

Changes in Markers of Disease Progression



Disease Progression Criteria^{1,2}:

- NT-proBNP: an increase of >700 ng/L and >30%
- hs-Troponin T: an increase of >10 ng/mL and >20%
- 6MWT: an absolute reduction of >35 m in 6MWT distance

Improvement was defined as the equivalent counter criteria

83% of NYHA class I/II patients and 47% of NYHA class III patients had no worsening in any marker at 12 months



First Clinical Evidence from Ongoing Phase 1 Study that nex-z May Favorably Impact Disease Progression in ATTR Amyloidosis

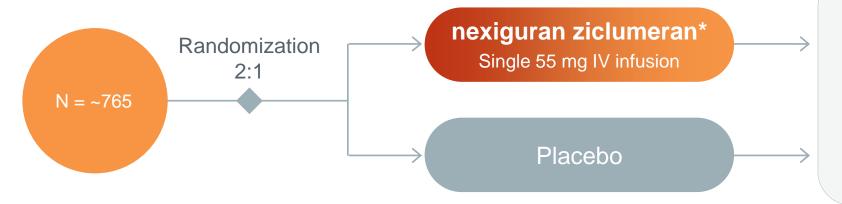
Key Takeaways:

- Single dose of nex-z resulted in deep, rapid and durable reductions in serum TTR with very low variability in patients
- Reductions in TTR were accompanied by stability or improvement based on several disease markers for ATTR-CM and ATTRv-PN
 - Including an ATTR-CM population with advanced disease who are expected to have rapid disease progression and high mortality rates
 - Including ATTRv-PN patients with early-stage disease and a subset of patients with severe neurological impairment who were previously treated with patisiran
- Favorable safety and tolerability observed





A Phase 3, Randomized, Double-blind, Placebo-controlled Study to Evaluate NTLA-2001 (nexiguran ziclumeran*) in Patients with ATTR Amyloidosis with Cardiomyopathy (ATTR-CM)



Primary Endpoint

 Composite endpoint of CV-related mortality and CV-related events

Key Secondary Endpoints

- Serum TTR
- KCCQ-OS score

Key Eligibility Criteria:

- Adult patients with diagnosis of either hereditary or wild-type ATTR-CM
- NYHA Class I III
- NT-proBNP baseline ≥ 1000 pg/mL

Stratification:

- NAC stage
- TTR genotype: wild-type vs. mutant
- Concomitant tafamidis use vs. no tafamidis

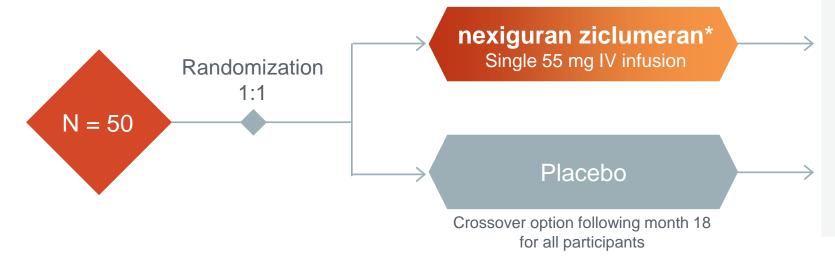
Study Duration:

- Dependent on occurrence of prespecified number of CV events and a minimum of 18 months follow-up
- Majority of patients are expected to have ≥ 30 months of follow-up for the primary analysis





A Phase 3, Randomized, Double-blind, Placebo-controlled Study to Evaluate NTLA-2001 (nexiguran ziclumeran*) in Patients with Hereditary ATTR Amyloidosis with Polyneuropathy (ATTRv-PN)



Primary Endpoints

 Change from baseline in mNIS+7 at month 18 and serum TTR at day 29

Key Secondary Endpoints

 Change from baseline in Norfolk QOL-DN, mBMI and serum TTR and absolute serum TTR at month 18

Key Eligibility Criteria:

- Adult patients with diagnosis of ATTRv-PN
- NIS 10 130
- PND score of ≤ 3B
- Naïve to silencers; washout of stabilizers

Stratification:

- NIS score <50 vs. ≥ 50
- *TTR* genotype: early onset V30M vs others

Study Duration:

 All patients have completed the month 18 visit





NTLA-2002 for Hereditary Angioedema (HAE)

About HAE

- Genetic disease characterized by recurring, severe and unpredictable swelling in various parts of the body
- Despite availability of existing therapies, significant unmet need persists
- Chronic dosing is required with current treatment options

Our Approach

Knock out *KLKB1* gene with a single-dose CRISPR-based treatment

Reduce kallikrein activity to prevent attacks

Key Advantages Includes Potential to:

- Be a single-dose treatment
- Provide extensive and continuous reduction in kallikrein activity
 - Intended to minimize the risk of breakthrough attacks
- Eliminate significant treatment burden



HAE: Large Commercial Opportunity with Significant Unmet Need

NTLA-2002

Potential to be the best-in-class HAE prophylaxis agent and only single-dose treatment

Prevalence¹

150,000+

HAE patients worldwide

Diagnosis²

20 years old

average age of diagnosis

Symptom onset typically occurs by 12 years old

Disease Burden³

50-60%

patients continue to have HAE attacks despite existing therapies

- Attacks can result in hospitalizations
- Patients subject to lifetime of attack risk and chronic treatment

Commercial Opportunity^{4,5}

\$6B+

global market size expected by 2029

\$500K+

annual U.S. cost of leading prophylactic treatment



¹ Zuraw BL. Clinical practice. Hereditary angioedema. N Engl J Med. 2008 Sep 4;359(10):1027-36. doi: 10.1056/NEJMcp0803977. PMID: 18768946.

² Farkas et al. Allergy. 2017. 72;300-313

³ Banjerii et al. Ann Allergy Asthma Immunol. 2020. 124;600-607

⁴ GlobalData 2023

⁵ Redbook 2023

NTLA-2002 Phase 1/2 Trial Design

International, multicenter 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

75 mg (n=3)

50 mg (n=4)

25 mg (n=3)

PHASE 2

Expansion study to confirm recommended dose

Randomized

50 mg (n=10)

25 mg (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)



Phase 2 Study: NTLA-2002 Continues to Be Well Tolerated Across All Dose Levels

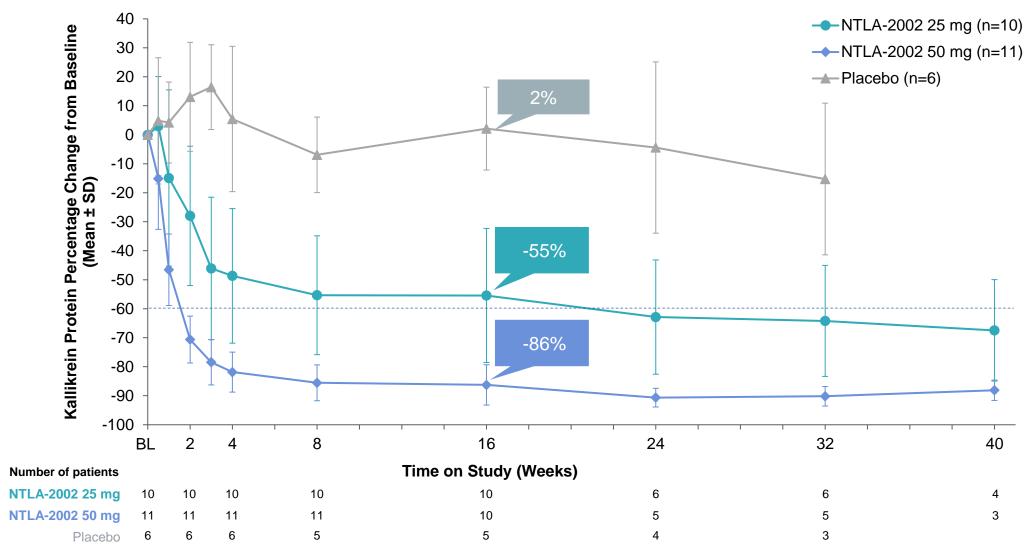
TEAEs in ≥2 Patients After NTLA-2002 Administration (pooled), n (%)	NTLA-2002 25 mg (n=10)	NTLA-2002 50 mg (n=11)	Placebo (n=6)
Any TEAE	10 (100)	11 (100)	6 (100)
Headache	4 (40)	4 (36)	1 (17)
Fatigue	3 (30)	3 (27)	2 (33)
Nasopharyngitis	3 (30)	3 (27)	2 (33)
Back pain	3 (30)	2 (18)	0
Upper respiratory tract infection	3 (30)	2 (18)	1 (17)
Cough	3 (30)	1 (9)	0
Infusion-related reaction	1 (10)	3 (27)	1 (17)
COVID-19	2 (20)	1 (9)	1 (17)
Ear infection	2 (20)	0 (0.0)	0
Epistaxis	0	2 (18)	1 (17)
Influenza-like illness	1 (10)	1 (9)	0
Oropharyngeal pain	1 (10)	1 (9)	1 (17)
Pyrexia	0	2 (18.2)	0
Sinusitis	1 (10)	1 (9)	0

- All TEAEs were Grade 1 or 2*
- No SAEs in patients treated with NTLA-2002
- 4 IRRs with NTLA-2002; 2 led to temporary interruption of study drug
 - Each instance resolved without sequelae and both patients received the full dose
- No clinically significant laboratory abnormalities
 - 1 patient had transient Grade 2 increase in ALT on Day 22



ALT, alanine aminotransferase; IRR, infusion-related reaction; SAE, serious adverse event; TEAE, treatment-emergent adverse event. *Common Terminology Criteria for Adverse Events (CTCAE) Grading

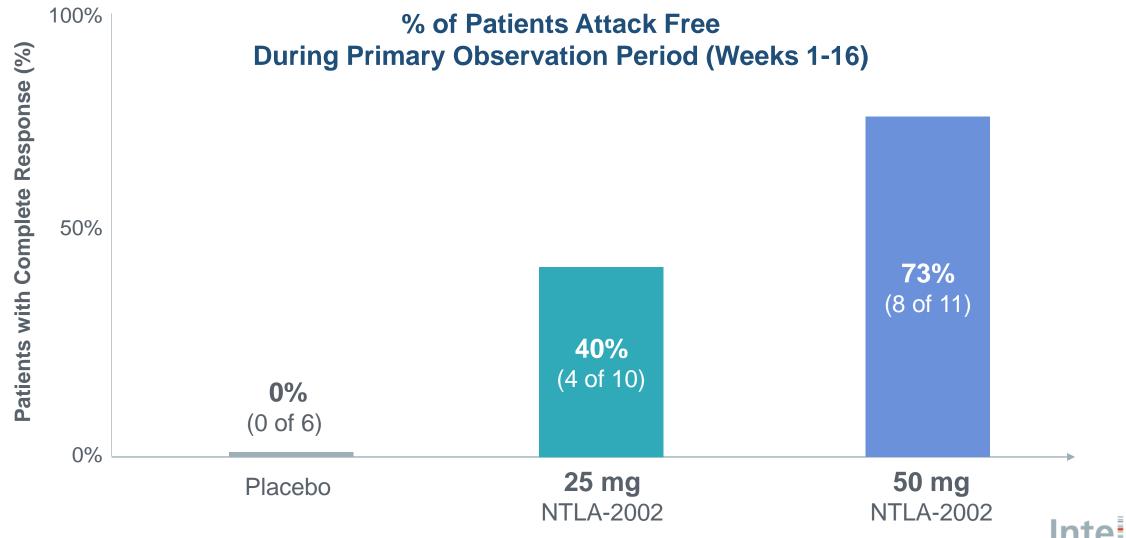
Phase 2 Study: A Single Dose of NTLA-2002 Showed Dose-Dependent and Durable Reductions in Plasma Kallikrein Protein Over Time



For post-baseline assessments, only scheduled visits completed by at least 3 patients in each arm are presented. Dashed line represents targeted minimum reduction. BL, baseline; SD, standard deviation.



Phase 2 Study: Eight of 11 Patients Receiving a Single 50 mg Dose Experienced a Complete Response – Attack-Free and No Subsequent Treatment Required



Phase 1/2 Data Demonstrated the Potential of a Single Dose of NTLA-2002 to Be a Functional Cure for Patients With HAE

Key Takeaways

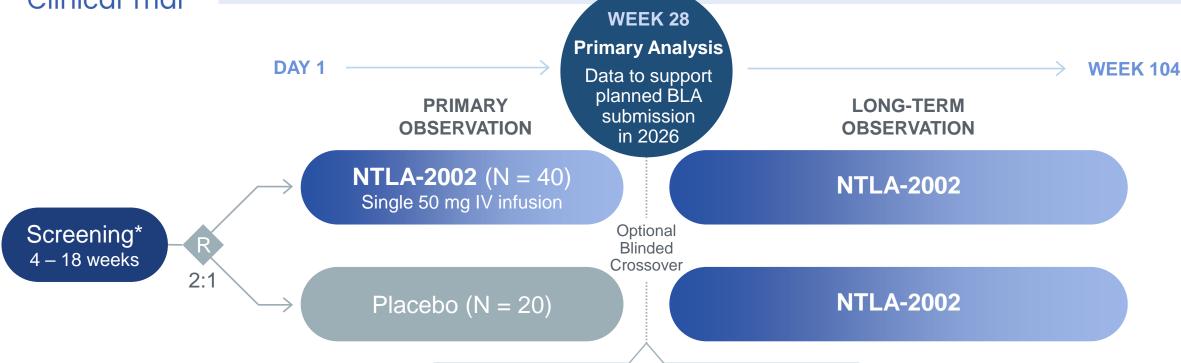
- Single 50 mg dose led to majority of patients (12/15) achieving complete elimination of attacks through the latest follow-up
- Robust and durable attack reductions observed in all patients
- Dose-dependent and durable reductions in plasma kallikrein protein achieved
- Highly encouraging safety and tolerability profile observed

50 mg dose selected for Phase 3 study





A Phase 3, Randomized, Double-blind, Placebo-Controlled Study to Evaluate the Efficacy and Safety of NTLA-2002 in Patients with Hereditary Angioedema (HAE)



PRIMARY ENDPOINT:

Time-normalized number of investigator-confirmed HAE attacks over

weeks 5 through 28

Clinicaltrials.gov ID: NCT06634420

^{*} Patients on long-term prophylaxis are required to wash out of therapy prior to the run-in period of screening.



NTLA-3001 for Alpha-1 Antitrypsin Deficiency (AATD)-Associated Lung Disease

About AATD

- Genetic disorder leading to progressive lung and/or liver disease¹
- >60K AATD patients in the U.S.^{2*}
- ~250K AATD patients globally^{3*}

Our Approach

Targeted insertion of a functional SERPINA1 gene into the albumin locus

Continuous expression of functional AAT protein at normal levels

Key Advantages

- Designed to be a single-dose treatment
- Aims to achieve normal human levels of AAT protein and halt progression of lung disease



¹ Remih et al. Curr Opin Pharmaco 2021; 59:149-156

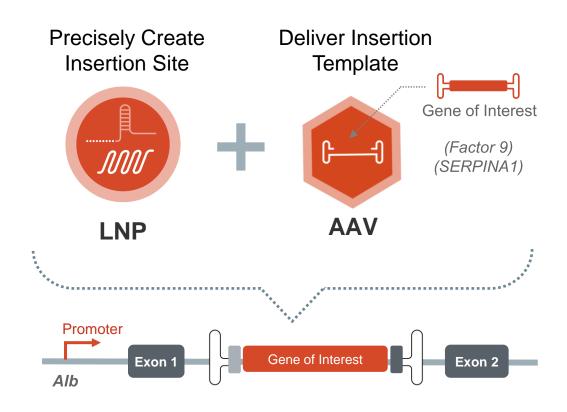
² Brantly M. Clin Chem. 2006; 52:2180-2181

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

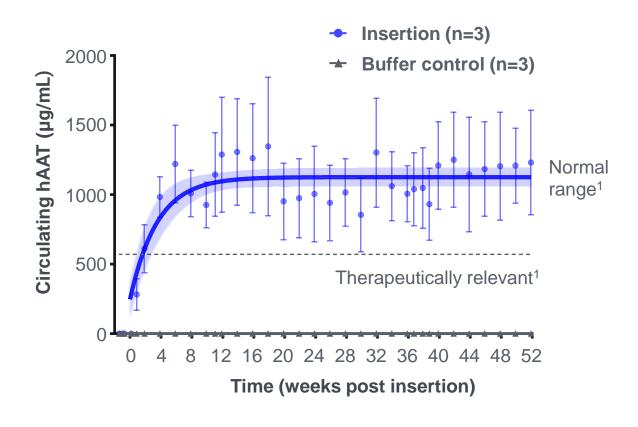
^{*} In severe AATD patients defined as individuals with Pi*ZZ genotype.

Durable Production of Physiologic Levels of hAAT Through One Year in NHP

Insertion Platform Enables Targeted, Stable Gene Insertion in the Albumin Locus



Human AAT (hAAT) Expression





NTLA-3001 Phase 1/2 Study Design

International, multi-center, open-label study to evaluate NTLA-3001 in adults with alpha-1 antitrypsin deficiency (AATD)-associated lung disease

Total Enrollment: Up to 30 patients, age 18-75 years with AATD-associated lung disease

Key Eligibility:

- ZZ or Z/null genotype
- < 11 µM AAT serum concentration
- FEV1 > 35% to < 65%

Intervention:

Single administration of intravenous (IV) AAV and LNP infusions

PHASE 1 Single-Ascending Dose N = Up to 18 patients* AAV LNP Dose 1 Up to 3 doseescalation cohorts Dose 3

PHASE 2 Expansion study

N = ~12 patients

Administer dose selected from Phase I

KEY ENDPOINTS

- Evaluate safety, tolerability
- Measure change in serum AAT levels



Significant Opportunities to Unlock Full Potential of In Vivo Platform

CRITERIA USED TO SELECT POTENTIAL FUTURE CANDIDATES:

Unmet need • Population size • Technical feasibility

Potential Liver Development Programs*

RARE DISEASES**

- Blood disorders
- Lysosomal storage diseases
- Metabolic diseases

PREVALENT DISEASES**

- Chronic viral diseases
- Dyslipidemia
- Hypertension
- NASH

Unlocking Full Potential of Genome Editing

TARGET TISSUES













Expansion into tissue-specific diseases



^{*} This is a selection of potential liver targets and does not represent all future opportunities.

^{**} Individual targets could be developed by Intellia, Regeneron or through collaborations.

^{***} In collaboration with SparingVision

^{****} In collaboration with ReCode

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Ex Vivo

CRISPR creates the therapy

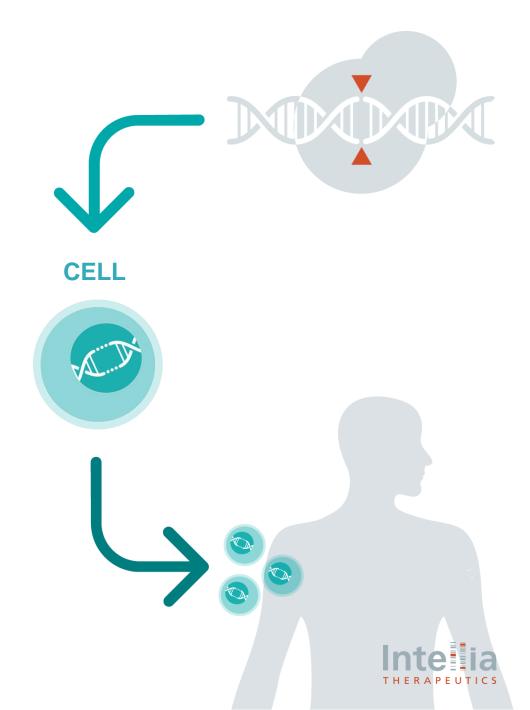
IMMUNO-ONCOLOGY / AUTOIMMUNE DISEASES

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	Other Approaches		
Gene Editing Approach	Delivery	LNP	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?		×	×	



LNP-based, sequential process



Precise CRISPR KOs & insertion(s)

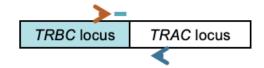


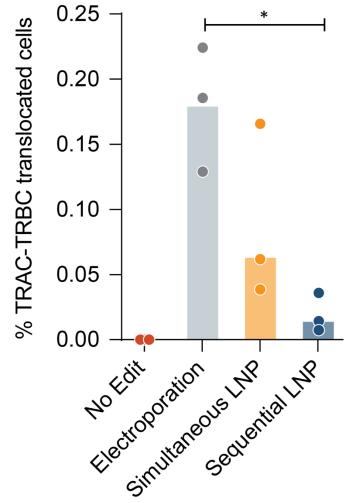
Quality cell product

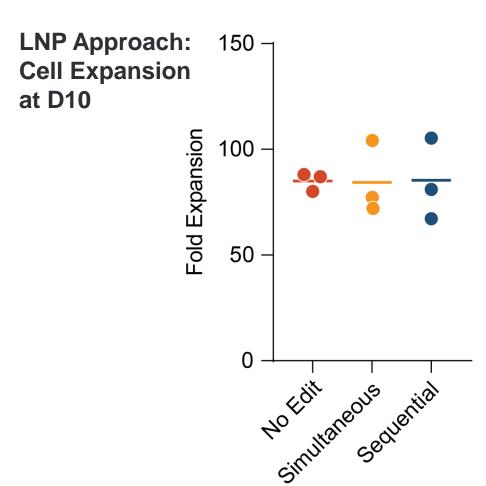


Sequential Editing with LNP Approach Minimizes Translocations While Retaining Robust Cell Viability and Expansion

ddPCR assay to detect *TRAC-TRBC* translocations

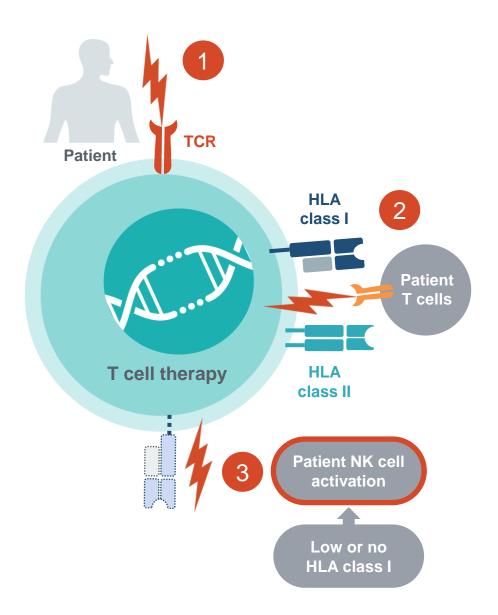








Three Immune Concerns Must Be Addressed by Allogeneic Cell Therapies



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

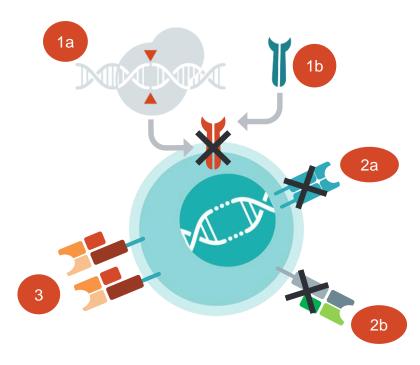
 Human leukocyte antigen (HLA) molecules must match
 between donor and recipient to prevent rejection from:
 - Host CD8 (HLA class I) T cells
 - Host CD4 (HLA class II) T cells
- Rejection via host natural killer (NK) cells

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

No validated solution yet



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



Key Potential Advantages

- ✓ Approach is applicable to CAR and TCR
- ✓ Solves for host NK and T cell rejection
- ✓ Avoids long-term immunosuppression

Intellia's Editing Strategy Main Objective of Edit Prevent Graft-versus-Host Knockout endogenous TCR Disease (GvHD) Insert target CAR or TCR **Direct T cell for tumor killing** Knockout HLA Class II **Prevent CD4-mediated rejection** Knockout HLA-A only **Prevent CD8-mediated rejection** Block NK cell activation and Partial HLA Class I match avoid NK-mediated rejection





Realizing the Promise of Gene Editing

At Intellia, we innovate every day to make CRISPR-based medicines a reality for patients.

This is just the beginning of the gene editing revolution.



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- Persistence of In Vivo Edits
- In Vivo Editing of Hematopoietic Stem Cells
- Intellia's Allogeneic Solution
- Platform: Identifying Potent and Highly Specific Guide RNAs
- Strategic Collaborations
- Abbreviations

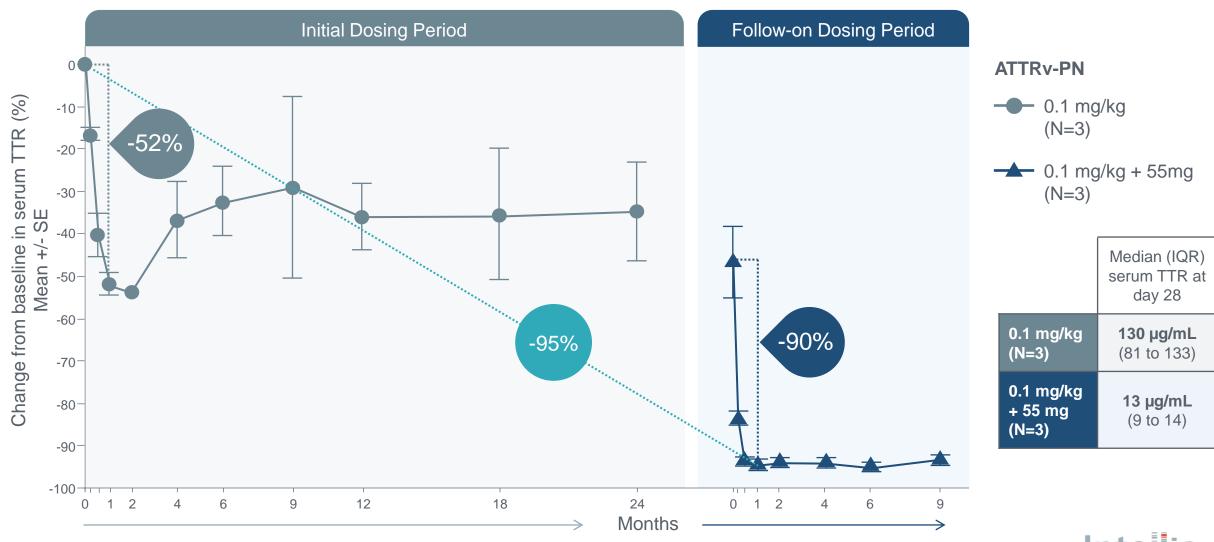


Re-dosing with Intellia's LNP Delivery Platform

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Clinical Proof-of-Concept that Redosing with Intellia's LNP-Delivered Gene Editing Technology Led to a Targeted Additive Pharmacodynamic Effect



Follow-on Dosing Was Well-Tolerated and Did Not Lead to Any Safety Findings

TEAEs by Maximum Toxicity Grade and Preferred Term Reported in Patients After Receipt of a Follow-On Dose (n=3)

Preferred Term ^a , n (%)	0.1 mg/kg + 55 mg (n=3)	Maximum CTCAE Toxicity Grade	
Any TEAE	2 (67%)		
COVID-19	1 (33%)	1	
Fatigue	1 (33%)	1	
Hand fracture	1 (33%)	1	
Headache	1 (33%)	1	
Infusion-related reaction	1 (33%)	1	
Nausea	1 (33%)	1	
Vulvovaginal candidiasis	1 (33%)	2	

- 8 12 months of follow-up for patients who received a follow-on dose
- No clinically significant changes in liver enzymes, platelets or coagulation parameters



This presentation includes data for an investigational product not yet approved by regulatory authorities.

Data cutoff April 12, 2024. Note: For each preferred term, patients reporting more than one adverse event are counted only once.

A Grade 1 AE of "Hereditary neuropathic amyloidosis" was reported for one patient at the time of the data cutoff and confirmed to be entered in error.

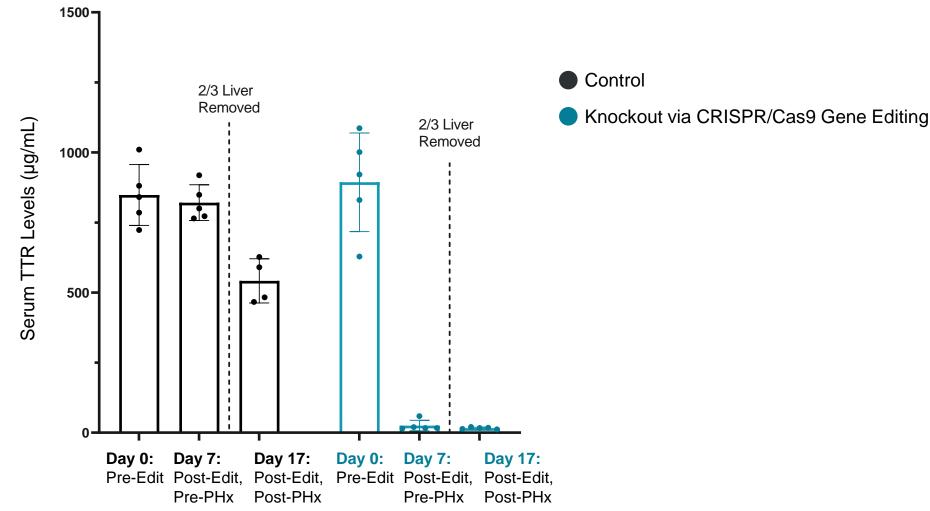
Adverse events are coded to preferred term using MedDRA, version 26.0.

Persistence of In Vivo Edits

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Protein Reduction Remains Unchanged Following PHx Murine Model of Liver Regeneration



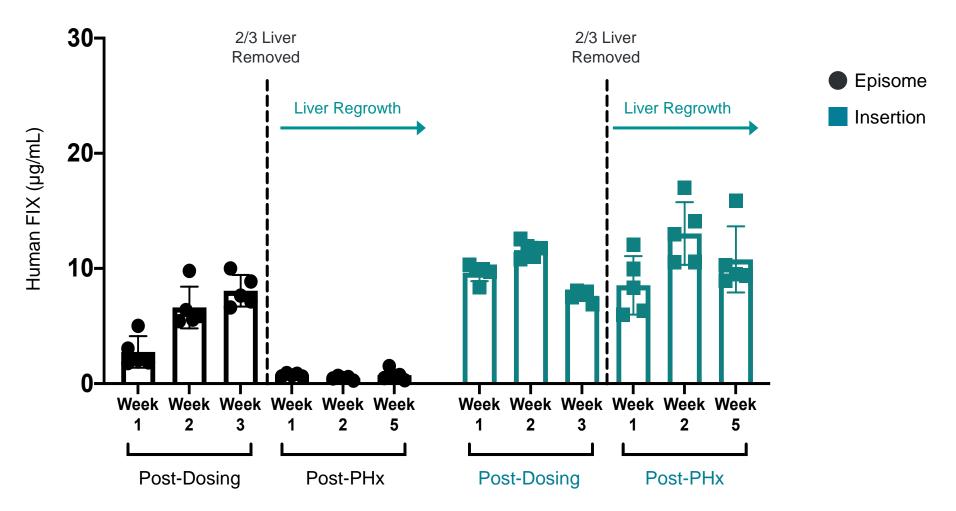
^{*} Similar results obtained for control and LNP when sham surgery was performed.

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



REGENERON

Gene Insertion Provides a Durability Advantage Over Conventional AAV Episomes in a PHx Murine Model of Rapid Liver Growth





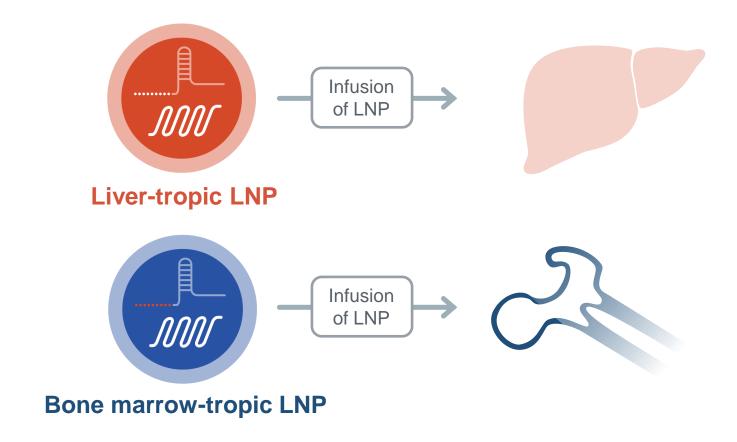
In Vivo Editing of Hematopoietic Stem Cells

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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

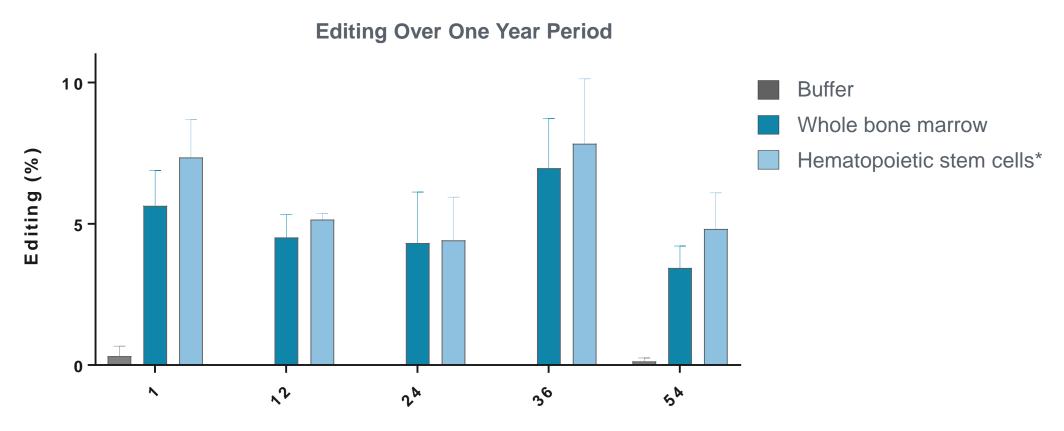




Editing of Mouse Bone Marrow and HSCs is Durable Through At Least One Year

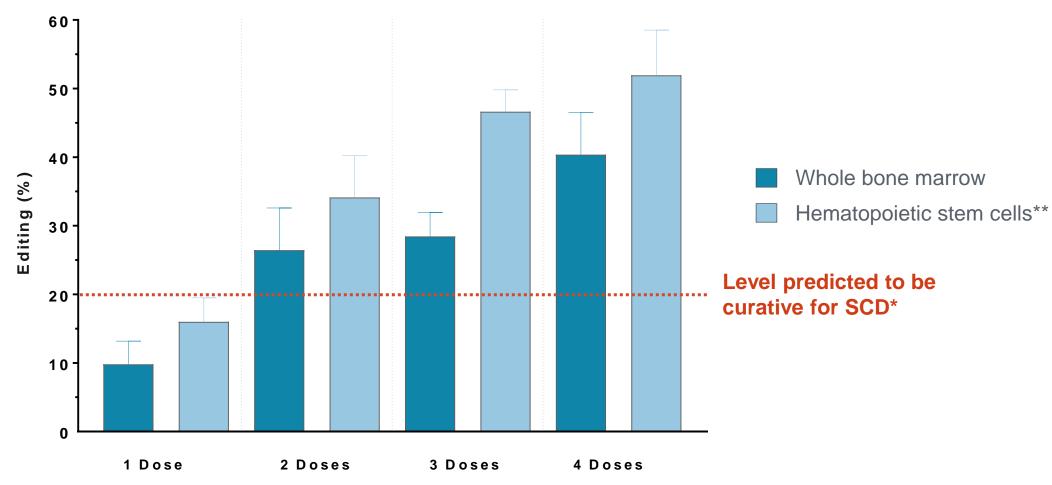
Weeks Post LNP Dose

- 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

Intellia's Allogeneic Solution

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Immune Concerns Unaddressed by Current Allogeneic Solutions

				Intellia's Approach
Approach	Employ intense lymphodepletion regimen	Knockout (KO) HLA-I (B2M)	KO HLA-I & express NK inhibitor (HLA-E)	KO HLA-II & partial HLA Class I match
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	×			

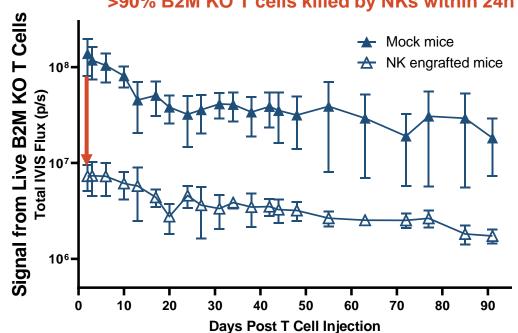


Allo TCR-T Cells Resist NK Cell Killing for at Least 90 Days In Vivo



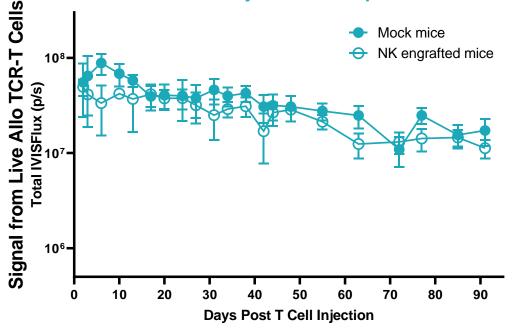
B2M Knockout T cells

>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



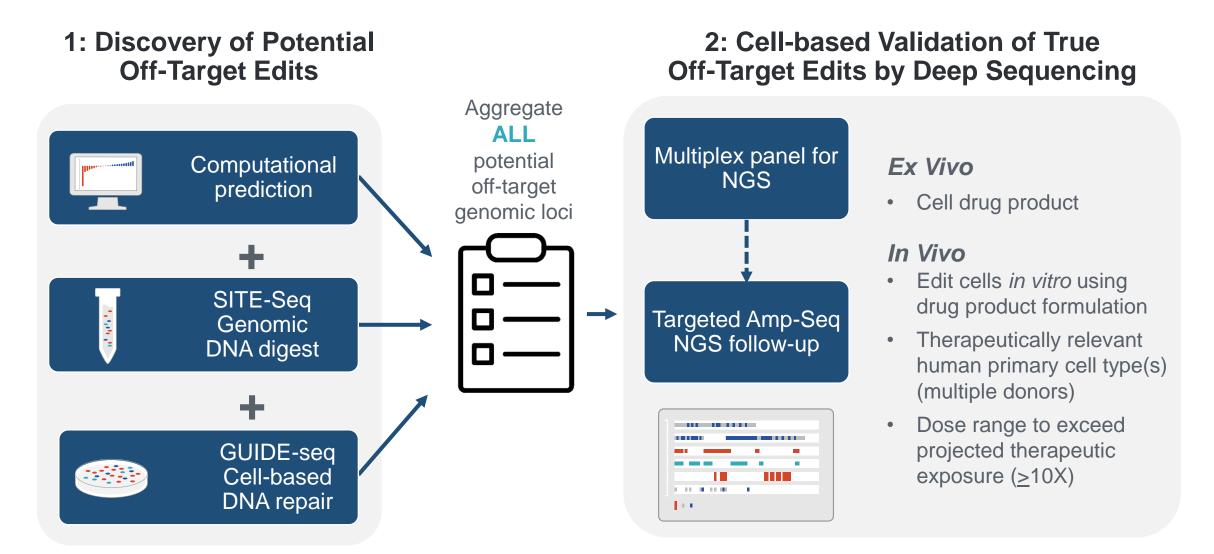


Platform: Identifying Potent and Highly Specific Guide RNAs

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Comprehensive gRNA Specificity Assessment: An Off-Target Workflow

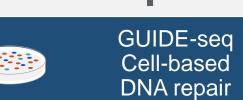




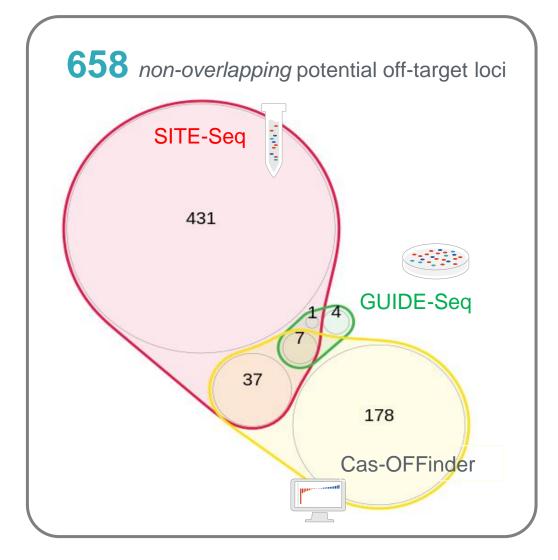
Limited Overlap in Discovered Off-Target Loci by Three Leading Methods

1: Discovery of Potential Off-Target Edits Computational prediction

SITE-Seq Genomic DNA digest



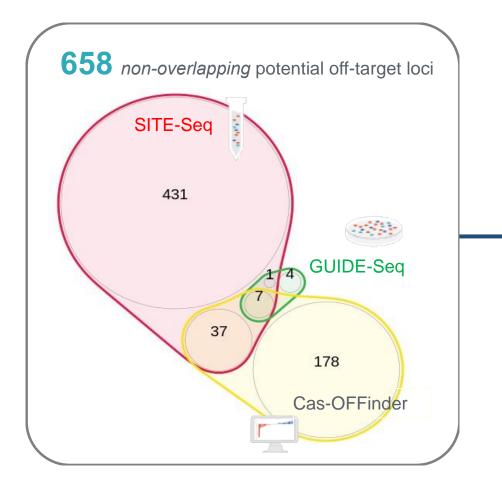
Aggregate
ALL
potential
off-target
genomic loci





Off-Target Workflow In Practice: Representative Example

1: Discovery of Potential Off-Target Edits



2: Validation of Off-Target Edits in Cells

Multiplex panel for

NGS

Targeted Amp-Seq

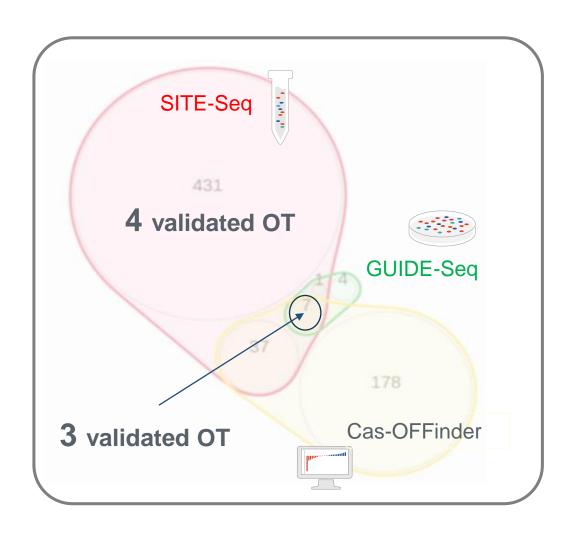
NGS follow-up

In Vivo Programs

- Dose responses using drug product formulation
- Therapeutically relevant human primary cell type(s) (2 donors)
- Dose range to exceed projected therapeutic exposure (>10X)
- Validation: off-target indels detected in edited cells



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



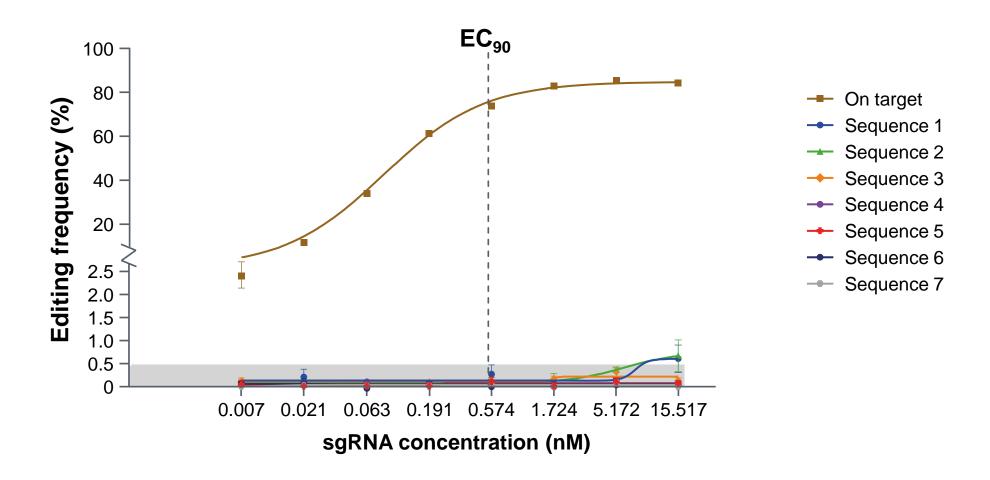
658 potential off-target loci

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 offtarget indels in regions of the genome associated with cancer



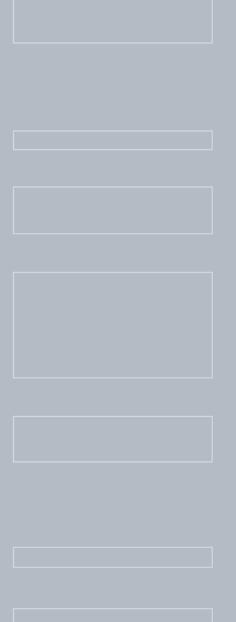
In Vitro: No Detectable Off-Target Editing with Pharmacologic Concentration of sgRNA





Strategic Collaborations

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Growing Intellia's Impact on Patients Through Strategic Collaborations





Collaborations Helping to Accelerate the Development of CRISPR-Based Therapies

REGENERON

Collaboration Overview:

- Up to 15 in vivo targets with a mix of co-developed and licensed programs
 - Liver-centric product development
- ATTR (in vivo knockout): Intellia is lead party;
 Regeneron will share 25% of costs and profits
- Hemophilia A (in vivo insertion): 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 for up to 10 ex vivo CRISPR products in defined cell types
- New research collaboration as of September 2023 to develop treatments for neurological and muscular diseases

Click below to learn more about our other collaborations







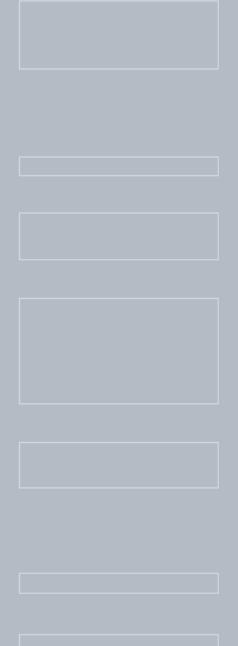






Abbreviations

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Abbreviations

AATD: alpha-antitrypsin deficiency

AAV: adeno-associated virus

AE: adverse event

AESI: adverse event of special interest

AI: autoimmune disease

ALT: alanine aminotransferase

AST: aspartate transaminase

ATTR amyloidosis: transthyretin amyloidosis

ATTRv: hereditary ATTR amyloidosis

ATTRwt: wild-type ATTR amyloidosis

ATTR-CM: ATTR amyloidosis with cardiomyopathy

ATTRv-PN: hereditary ATTR amyloidosis with

polyneuropathy

B2M: beta-2-microglobulin

BL: baseline

BLA: biologics license application

CAR-T: chimeric antigen receptor T cells

CNS: central nervous system

CTCAE: Common Terminology Criteria for Adverse Events

CV: cardiovascular

ddPCR: digital droplet polymerase chain reaction

DSB: double strand break

GvHD: graft-versus-host disease

EC₉₀: concentration inducing 90% of maximal effect

FEV1: Forced expiratory volume in 1 second

FOD: follow-on dose

Gr: Grade

gRNA: guide RNA

HAE: hereditary angioedema

Hem A/B: hemophilia A/B

HLA-I / II: human leukocyte antigen class I / II

HLA-E: human leukocyte antigen class E

HSC: hematopoietic stem cells

IO: immuno-oncology

IQR: interquartile range

IRR: infusion-related reaction

KCCQ-OS: Kansas City Cardiomyopathy Questionnaire-

Overall Summary

KLKB1: kallikrein B1

LNP: lipid nanoparticle

MedDRA: Medical Dictionary for Regulatory Authorities

mRNA: messenger RNA

NAC: National Amyloidosis Centre

NASH: nonalcoholic steatohepatitis

nex-z: nexiguran ziclumeran

NHP: non-human primate

NK: natural killer

NT-proBNP: N-terminal-pro-B-type natriuretic peptide

NYHA: New York Heart Association

PD: pharmacodynamics

PHx: partial hepatectomy

PK: pharmacokinetics

PNS: peripheral nervous system

Pt: patient

SAE: serious adverse event

SE: serious event

SCD: sickle cell disease

SD: standard deviation

sgRNA: single-guide RNA

TCR: T cell receptor

TEAE: treatment-emergent adverse event

TTR: transthyretin



