

Precision gene editing designed to deliver durable, curative medicines

Hemophilia A lead program | IND 2026

Corporate Presentation
May 2026



Forward-looking statements

This presentation includes forward-looking statements, including forward-looking statements within the meaning of the Private Securities Litigation Reform Act of 1995. All statements other than statements of historical facts contained in this presentation are forward looking statements, including statements regarding our cash runway, strategy and plans, industry environment, potential growth opportunities, and the therapeutic potential of our programs. The words “believe,” “may,” “will,” “estimate,” “continue,” “anticipate,” “design,” “expect,” “could,” “plan,” “potential,” “predict,” “seek,” “should,” “would,” or the negative version of these words and similar expressions are intended to identify forward-looking statements.

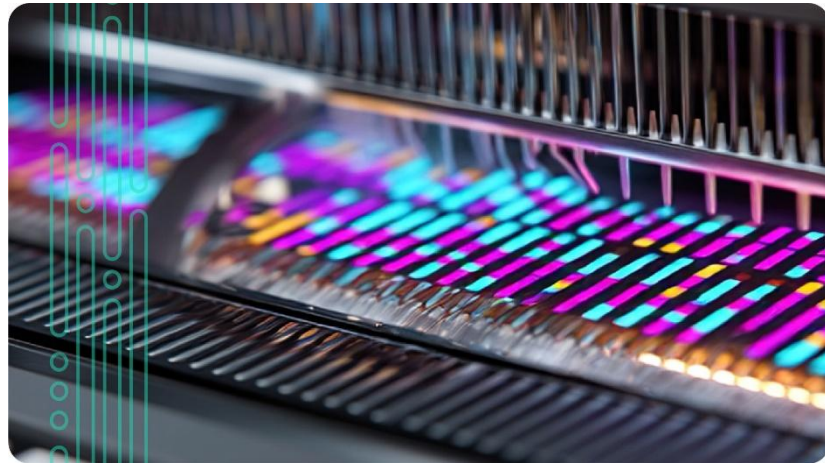
We have based these forward-looking statements on our current expectations and projections about future events and trends that we believe may affect our financial condition, results of operations, strategy, short- and long-term business operations and objectives, and financial needs. These forward-looking statements are subject to a number of risks, uncertainties and assumptions, including but not limited to, our ability to develop and advance our programs and product candidates, our ability to maintain and establish collaborations or strategic partnerships, our regulatory approvals and filings, and other risks, uncertainties and assumptions identified in our filings with the Securities and Exchange Commission (the “SEC”), including our most recent Form 10-K and Form 10-Q filed with the SEC, and any subsequent filings with the SEC.

Moreover, we operate in a very competitive and rapidly changing environment, and it is not possible for our management to predict all risks, nor can we assess the impact of all factors on our business or the extent to which any factor, or combination of factors, may cause actual results to differ materially from those contained in any forward-looking statements we may make. In light of these risks, uncertainties and assumptions, the forward-looking statements and circumstances discussed in this presentation may not occur and actual results could differ materially and adversely from those anticipated or implied in the forward-looking statements. You should not rely upon forward-looking statements as predictions of future events. Although we believe that the expectations reflected in the forward-looking statements are reasonable, we cannot guarantee that the future results, levels of activity, performance or events and circumstances reflected in the forward-looking statements will be achieved or occur. Moreover, except as required by law, neither we nor any other person assumes responsibility for the accuracy and completeness of the forward-looking statements. We undertake no obligation to update publicly any

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A differentiated *in vivo* gene editing company advancing curative genetic medicines



An *in vivo* CRISPR gene editing company capitalizing on its proprietary technologies to create curative genetic medicines

Focusing on lead program MGX-001 in hemophilia A advancing to the clinic

Expanding indications leveraging site-specific gene integration system and partnered assets targeting cardiometabolic indications

Differentiated gene editing beyond CRISPR/Cas9

20,000 +

Signature editing systems from Metagenomi's database

Proprietary CRISPR genome editing



High specificity

Precise editing



Durable integration

Large gene insertion



Broad targeting

Expanded genome access

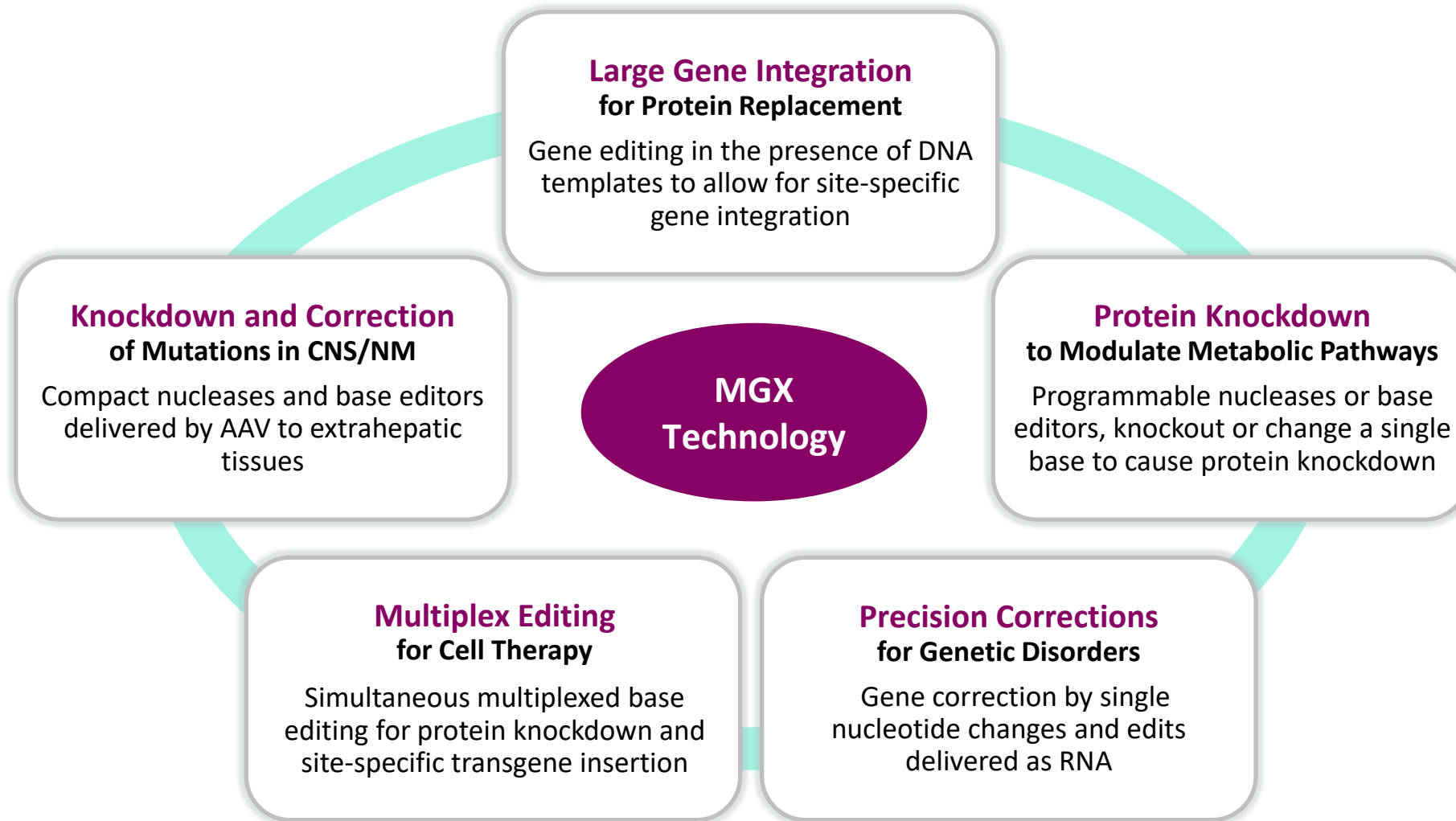


Multiplexed editing

Multi-gene capability

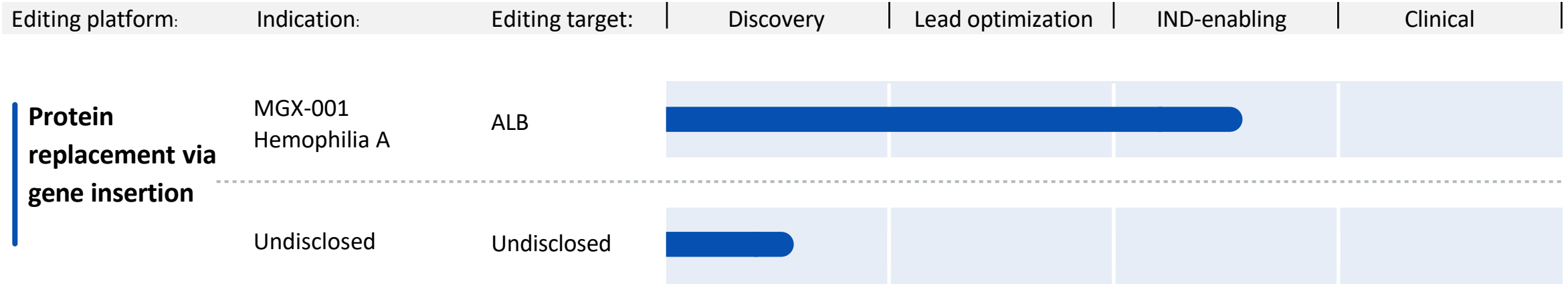
We improve editing precision and expand genome targeting and editing functionality beyond CRISPR/Cas9 to effectively address genetically-driven diseases.

A versatile platform designed to address a wide range of genetic diseases

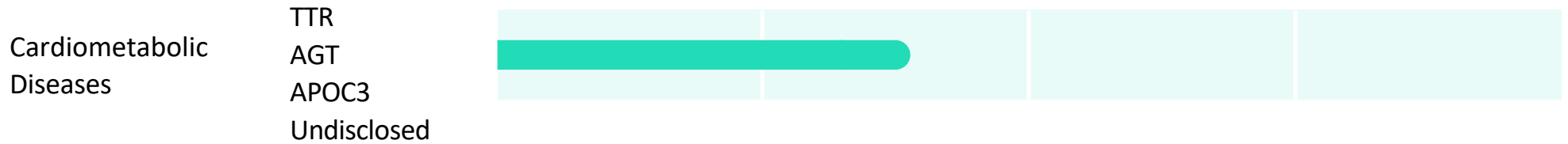


Potentially capable of correcting any type of genetic mutation found anywhere in the human genome

Focused pipeline anchored by differentiated lead program

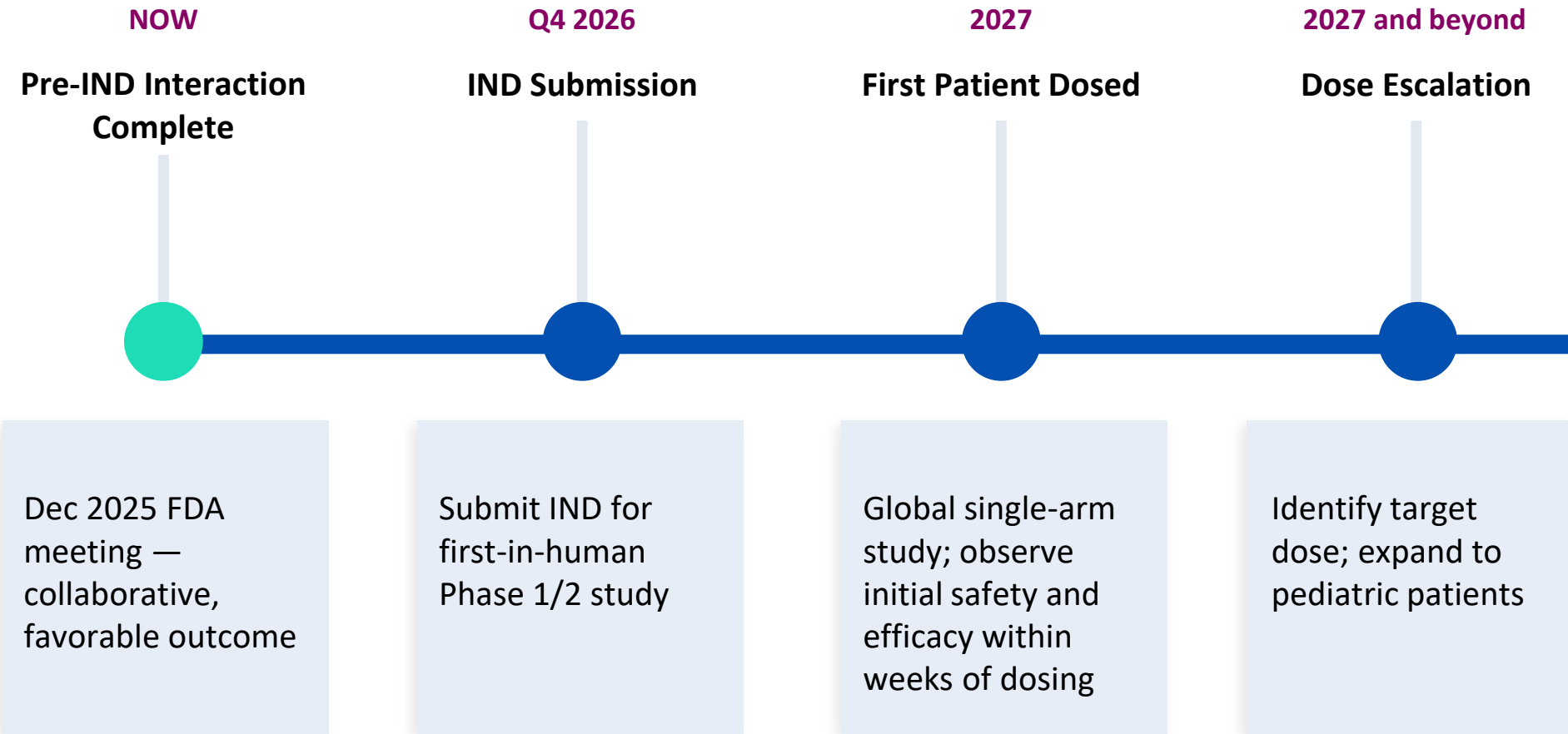


Protein reduction via gene knockout



Metagenomi is exploring opportunities to pursue neuromuscular disease targets & liver disease targets such as A1AT and Wilson Disease, as well as business development to expand therapeutic applications including cell therapy.

MGX-001 development roadmap to clinic



Goal: Enable a new standard of care for hemophilia A – a one-time cure freeing patients from lifelong treatment burden and bleeding risk

\$140.2 million in cash, cash equivalents, and available-for-sale marketable securities at end of Q1 2026
Runway anticipated to support operations through Q4 2027

MGX-001 - designed to deliver a durable, one-time treatment for Hemophilia A

Hemophilia A: large, validated market still lacking a durable cure



~26,500

patients in the U.S.¹

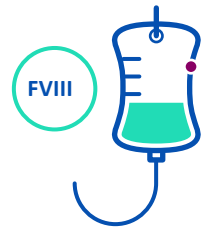
~500,000

worldwide²

Hemophilia A is the most common X-linked inherited and de novo bleeding disorder, largely affecting males.

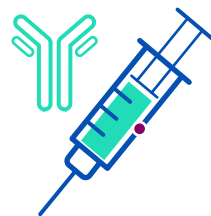
Caused by variety of mutations in the Factor VIII (FVIII) gene leading to loss of functional FVIII protein.

Current Standard of Care:



Factor VIII replacement therapy

- IV typically dosed 1 - 3 times/week
- Significant adherence challenges
- Risk of breakthrough bleeding
- Chronic treatment, non-curative



Bi-specific antibody "mimetic"

- SQ dosed 1, 2 or 4 weeks post loading
- Risk of breakthrough bleeding
- Treatment burden, non-curative

Recent Curative Gene Therapy Attempted:



- Variable initial efficacy
- Decline in FVIII levels over time
- High risk of prolonged corticosteroid use
- Not suitable for pediatric patients

Annual treatment cost³:

~\$565K - \$750K

Lifetime treatment cost:

~\$18M - \$24M⁴

1 - Soucie, J.M., et al, 2020. Haemophilia. Vol. 26, no. 3, pp. 487-493.

2 - Stonebraker, J. S., et al, 2010. Haemophilia. Vol. 16, pp. 20-32.

3 - ICER. Gene Therapy for Hemophilia B and A: Final Evidence Report. Dec 22, 2022.

4 - Curtis R et al. Poster presented at: 65th ASH Annual Meeting & Exposition; December 11, 2023; San Diego, CA.

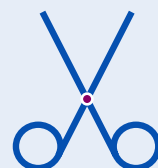
Genome editing offers a potentially ideal curative approach for Hemophilia A

Hemophilia A is an ideal indication for genome editing approach:

- Monogenic and well-characterized biology with clear biomarker
- Clearly defined target threshold of curative FVIII level & wide safety range
- Robust preclinical models and regulatory familiarity
- Strong advocacy and infrastructure
- Clear opportunity for a durable cure

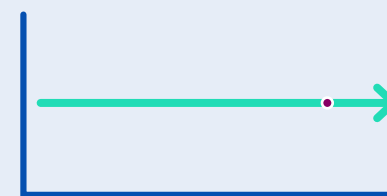
MGX-001 is uniquely suited for patients of all ages:

Technology:



proprietary
Type V
nuclease

Durability:

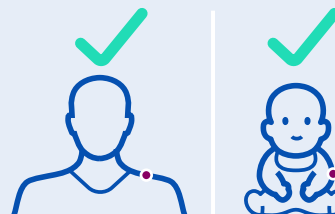


Regulatory status:



IND-enabling
stage

Pediatric potential:



MGX-001 is a potentially durable, curative approach for adults and children – the population with the most to gain

Compelling preclinical profile achieved across efficacy, durability, and safety

Extensive and supportive preclinical data set

- FVIII activity achieved in **curative range** with clear dose response in NHPs
- **Durable** FVIII activity over approximately 19-month study in NHPs
- **Encouraging safety profile** with single doses of steroids, and no genotoxicity observed

Novel mechanism of action

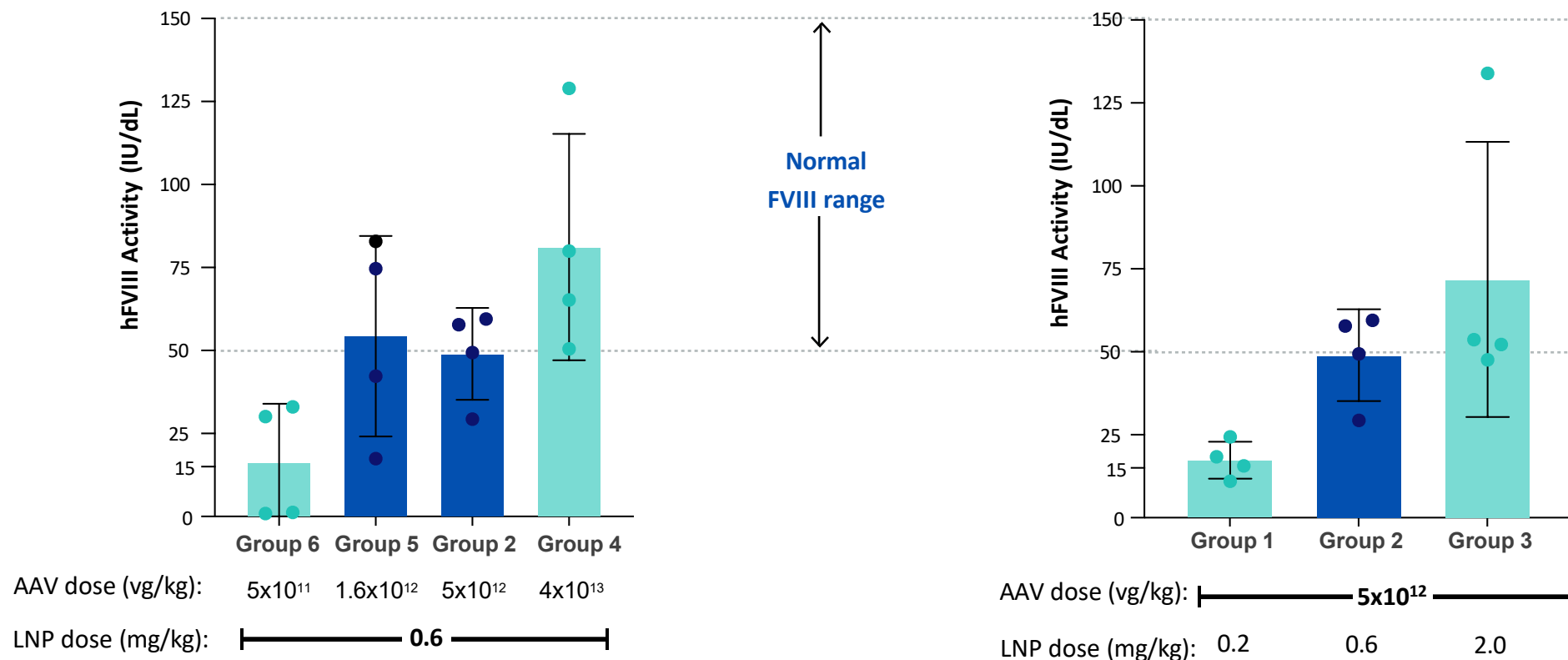
- FVIII integration leveraging **albumin promoter** to achieve normalized activity level
- **Promoterless FVIII gene** delivered by AAV effective at **lower dose** than approved gene therapies
- Precise FVIII integration facilitated by proprietary CRISPR nuclease MG29-1 achieving **no detectable off-target** editing

Compelling potential clinical profile

- Enables **endogenous production** of FVIII supporting hemostatic regulation
- Potential to normalize FVIII levels and deliver meaningful clinical benefit for both **adults and pediatric** patients
- Goal to be one-time durable cure allowing patients the freedom of a **hemophilia free mind**

Curative FVIII range achieved

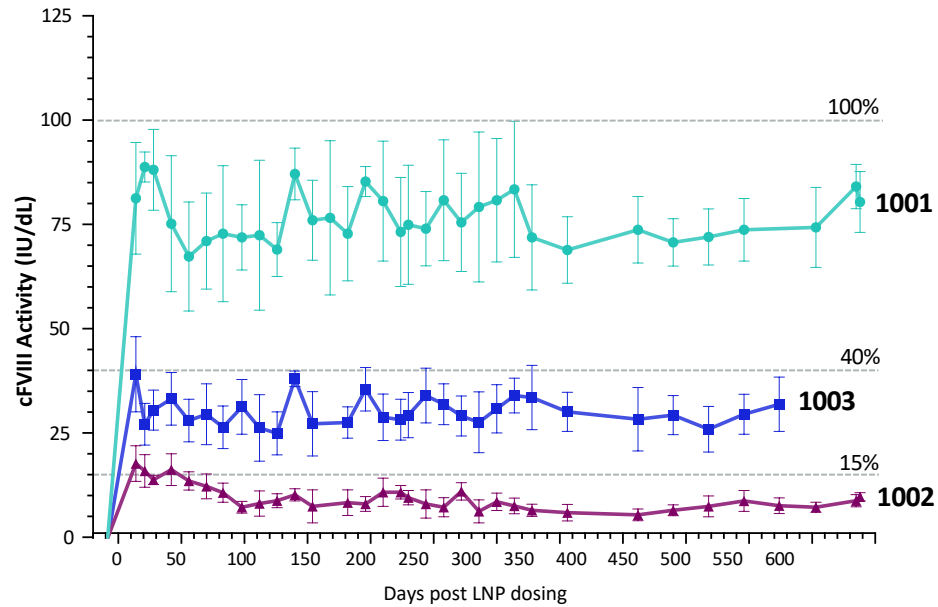
Dose range finding study in NHP identified minimum and optimal efficacious doses



FVIII activity for each animal is the mean of values on d5, d8, d11 post LNP dosing measured with a capture-chromogenic assay. hFVIII activity was stable from d5 to d11 post LNP. Xyntha was used for the standard curve.

Durable FVIII expression demonstrated ~19 months

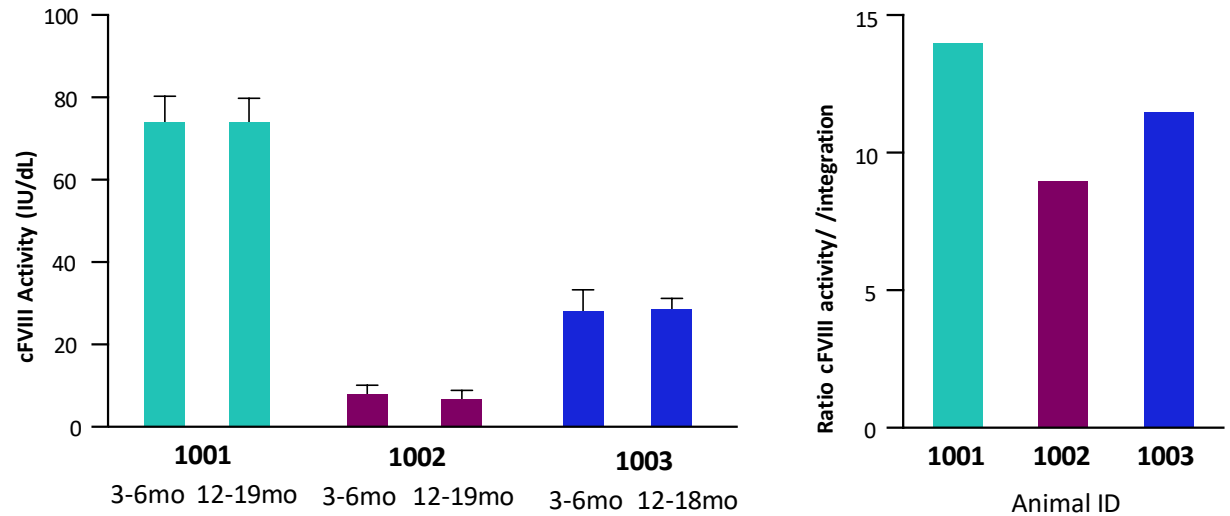
NHP durability study: Stable expression over ~19 months



FVIII activity values are the mean and standard deviation of at least 3 independent assay runs with each sample run in at least duplicate in each assay.

Animal 1003 died on day 540 (17.8 mo) post LNP, assessed as unrelated to the treatment.

Plasma FVIII activity levels unchanged between 3-6 months and 12-19 months and correlate to integration:

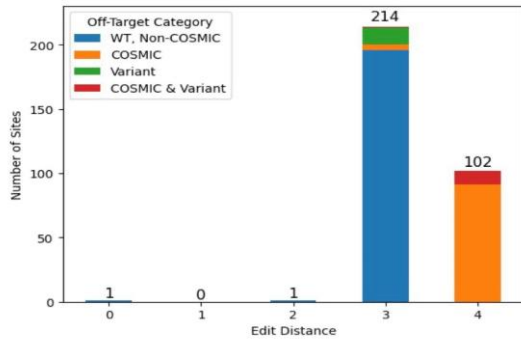


1 - Integration in forward orientation (copies per 100 haploid genomes, average of 5 liver lobes).

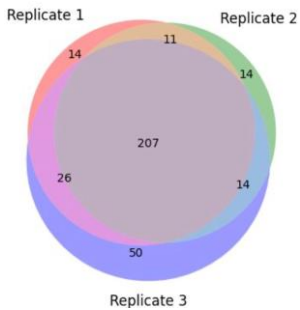
No genotoxicity observed

Discovery of potential off-target sites

1. In silico off-target discovery:



2. Biochemical off-target discovery:

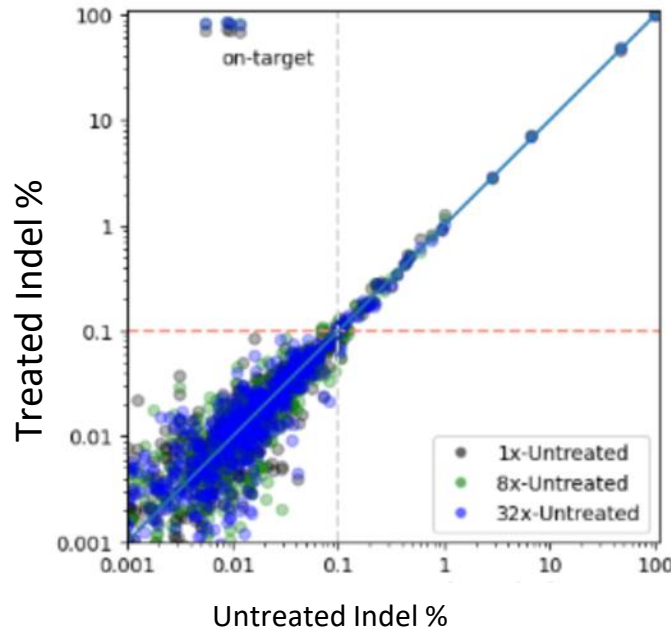


3. In cell off-target discovery:

No potential off-targets were discovered in cell-based assays.

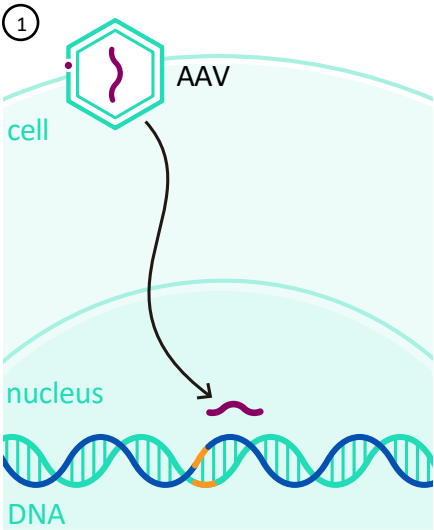
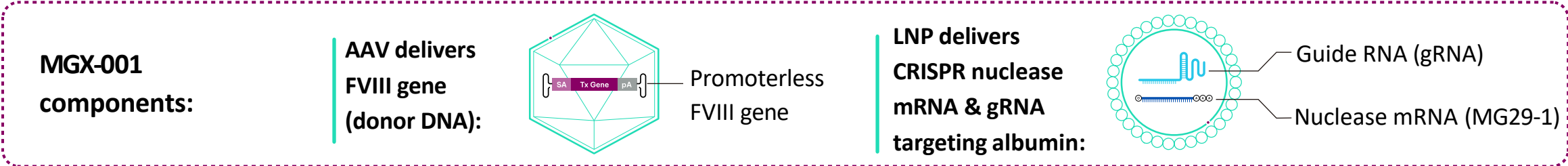
No validated off-target sites observed

Three independent primary human hepatocyte donors:

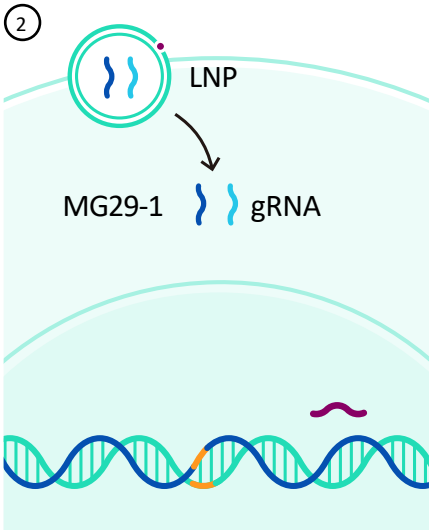


Genome integrity maintained as observed via off-target editing and AAV integration assays.

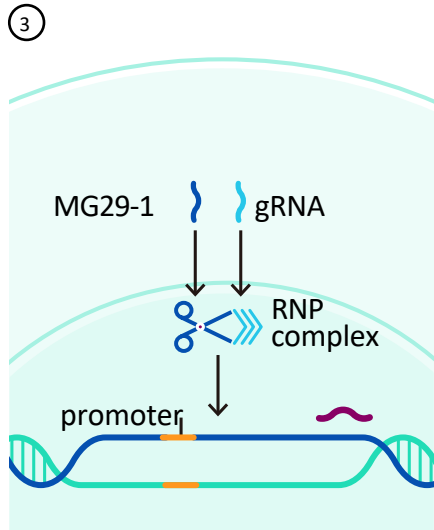
MGX-001: leveraging natural promoter through FVIII gene integration



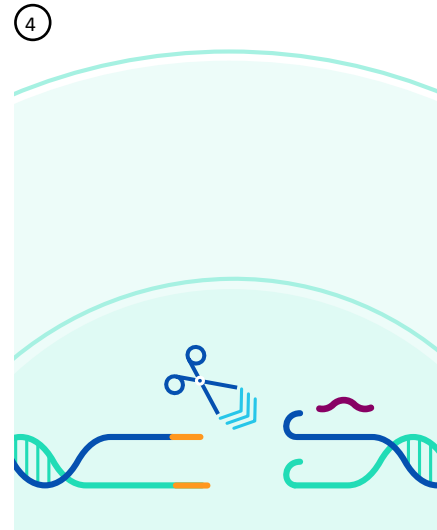
AAV delivers FVIII gene



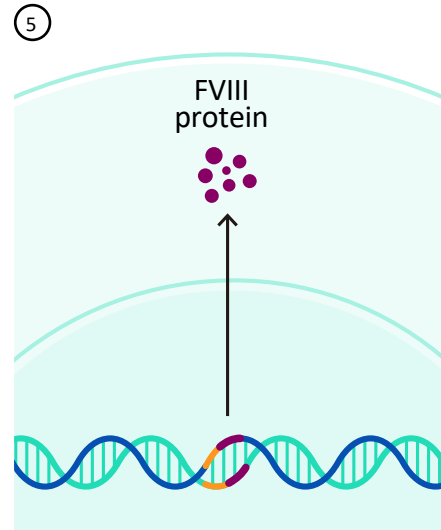
LNP delivers CRISPR cargo



CRISPR cargo forms complex



Cut at albumin locus and FVIII gene integrates



Albumin promoter drives sustained FVIII production

Potential to deliver a durable cure for both adult and pediatric patients with hemophilia A

Designed to enable endogenous FVIII expression for hemostatic regulation

Compelling pre-clinical data

- Curative FVIII activity
- Durable FVIII expression

Encouraging safety profile

- Minimal steroid use
- Promoterless AAV application
- No off-target editing observed

Established regulatory framework and defined clinical endpoints

- pre-IND interaction completed
- IND submission on track for Q4 2026
- First-in human in 2027

Our goal: To enable a new standard of care for hemophilia A



Expanding applications of site-specific large gene integration system

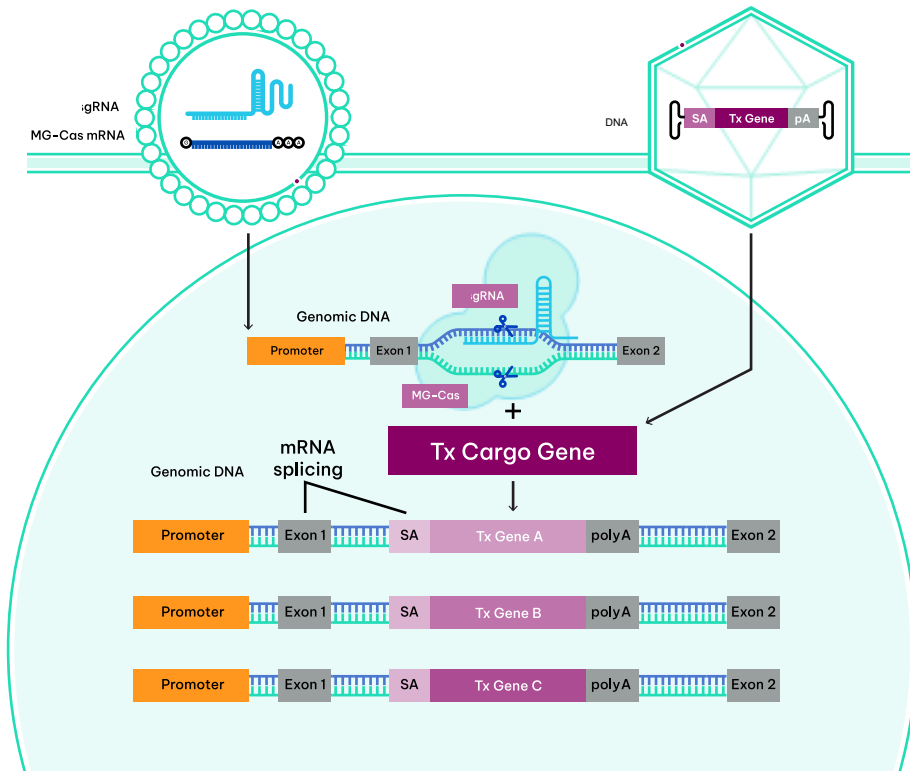
Leverage site-specific large gene integration across additional indications



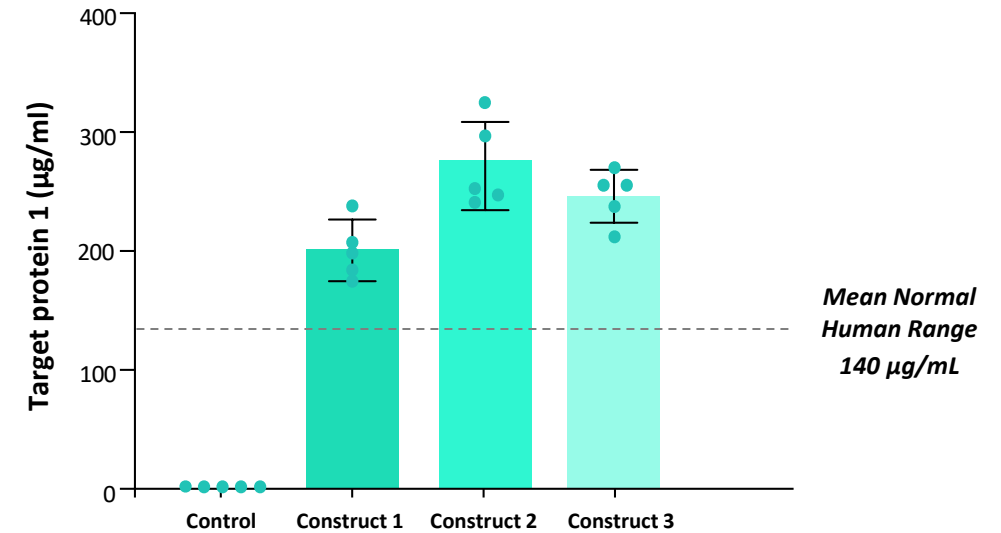
Expanding MGX-001 site-specific gene integration system into additional therapeutic targets

LNP delivers nuclease mRNA and guide targeting albumin site

AAV delivers Transgene (donor DNA)

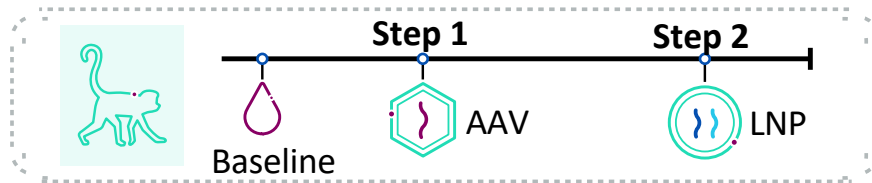


Normal circulating levels of target protein achieved in secreted protein disorder mice with multiple construct designs

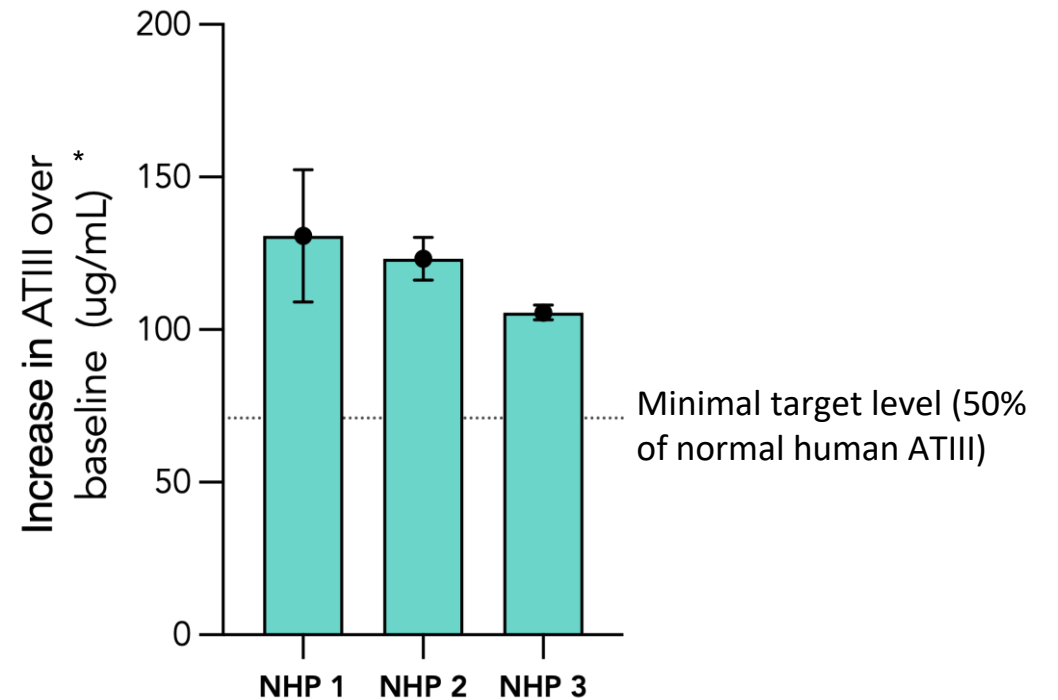


- Above normal human protein expression achieved in mouse plasma
- Insertion assessed with multiple DNA template constructs
- LNP and AAV dose titration can be used to fine tune therapeutic window

Proof-of-concept demonstrated in additional disease models



Achieved circulating AT-III protein exceeding curative target of 50% of normal human levels



*Data are the mean of day 8 and 11 post-dosing minus the mean of days 0, 4 and 7 pre-dose

N=3
In-life, study
Ongoing at 46 days

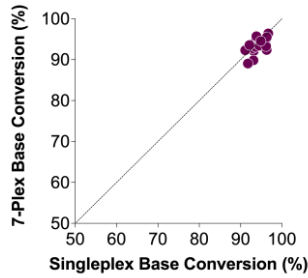
AAV dose (vg/kg) + LNP dose (mg/kg)

Animal	AAV dose (vg/kg)	LNP dose (mg/kg)
Animal 1001		
Animal 1002	1.0 x 10 ¹³	1.0
Animal 1003		

- Severe antithrombin (AT-III) deficiency increases risk of venous thromboembolism (VTE)
- On average patients with severe disease have 50% of the normal amount of AT-III in their blood (70 ug/ml) ^{1,2}
- Replacing the missing AT-III with at least 50% of normal is expected to be a functional cure

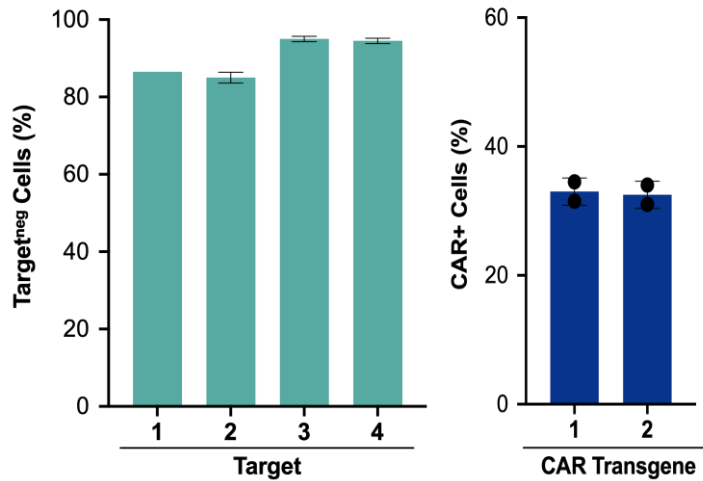
Broaden therapeutic
potential with
transformative gene editing

Multiplex editing and compact editors enable next-generation therapies

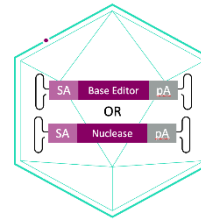


Multiplex editing in T-cells achieved across seven unique gene targets at as high efficiency as single plex editing

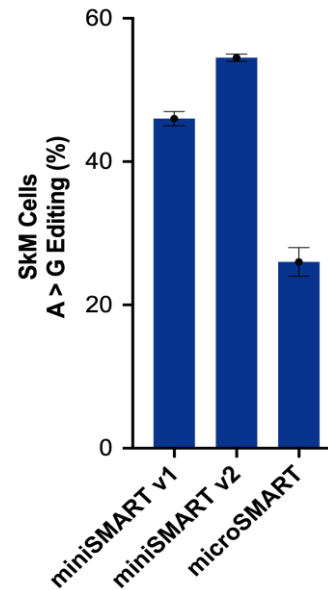
Combined protein knockdown and CAR knock-in in a single-step for cell therapy applications



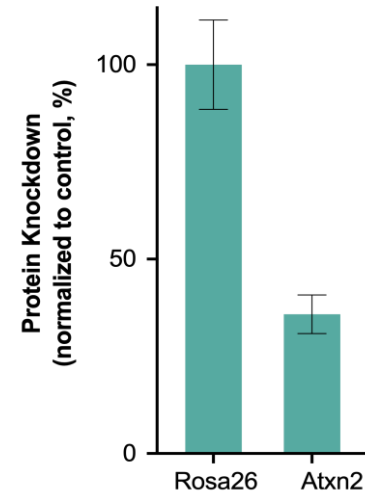
Four-plex knockdown and dual CAR knock-in in T cell with MGABE100 + 5 gRNA + DNA CAR template¹



Compact nucleases and base editors demonstrated high editing efficiency and compatibility with AAV delivery^{2,3}



Multiple compact base editors² demonstrated effective neuromuscular targeting



Compact nuclease MG119-28 achieved 64% knockdown of Atxn2 protein in mice⁴

Strategic partnership expands reach into large cardiometabolic markets

Current indications:

TTR

AGT

APOC3

Undisclosed

IONIS

- MGX's in vivo genome editing complements Ionis leadership in cardiometabolic space
- 4 targets: two co-development and co-commercialization options
- Multibillion dollar TAM

Building a leading gene editing company focused on cures

- Broad and differentiated library of **proprietary gene editing** technologies representing a significant long-term value driver
- Advancing **MGX-001 for hemophilia A** with clear development path and well-defined clinical and regulatory endpoints
- Extending **beyond hemophilia A**, large gene integration system opens potential to address other protein deficiencies
- Pairing our gene-editing capabilities with complementary expertise to **accelerate development through collaboration**



Thank you