Wednesday, October 31 | 8:25 – 8:50 am ET

ASO Design, Discovery and Research innovation in ASO technology

PRESENTER

Nano-rare

Konstantina Skourti-Stathaki, PhD

Hosted by:

Biogen

Director, ASO Design and Discovery





Integrated Processes & Building a Cohesive Team

Providing an optimal ASOs to nano-rare patients is the product of high-quality processes and years of expertise and experience



Konstantina Laurence (Laury) Sarah Glass **Julie Douville** (Nadina) Skourti-Mignon Stathaki Molecular Geneticist **Toxicologist RNA** Expert Neuroscientist Clinical trial expertise ASO expertise ASO discovery and **ASO** expertise Non-clinical Operational **Clinical development** design Management development Establishing Systems Clinical & Safety Supporting Patient Creating Preclinical to Processes Patient Mgmt & **Optimal ASOs** Journey Regulatory **Creating Unified** Treatment Communication ATTC - RMC **RMC - IND Cohesive Team** STAR - DSMB /Education

Amy Williford

Communications Educator Fundraising expertise



n-Lorem Benefits From More Than 35 Years of ASO Expertise and Builds on continuous Research Innovation to Assure Each Patient is Treated With the Best ASO Possible





Drugs and Their Targets

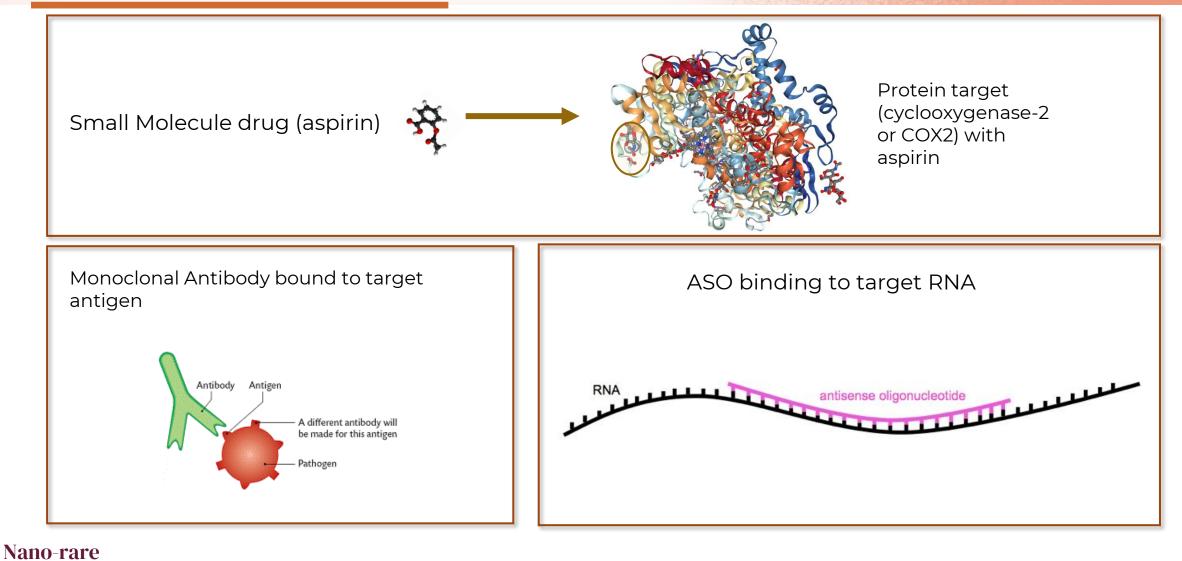
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ASO Technology Makes it Possible to Do What We Do at n-Lorem

Attribute	Small Molecule	Monoclonal Antibodies	RNA-Targeted (ASO)
Information Content		++	+++++
Rules of Engagement	Still largely unknown	Partially understood	Well understood, easy to use
Learning from Previous Drugs	"change a methyl change the drug"	Limited transferability	Broad transferability
Cost of Drug Discovery	Extremely expensive	Less expensive, but costly	Modest
Rate of Drug Discovery	Slow – Decades	Years	Months
Selectivity for Target		++	++++
Versatility	++++	+	++++
Cost of Goods	++++		++++
Advancing Technology	No	Minimal	Extensive, rapid advancement



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What Do We Do in the n-Lorem Lab and How Do We Do it?





The n-Lorem Lab Does ASO Discovery, Directed and Basic Research

- ASO Discovery
- Directed research to better understand targets and mutations
- Basic research to advance ASO technology

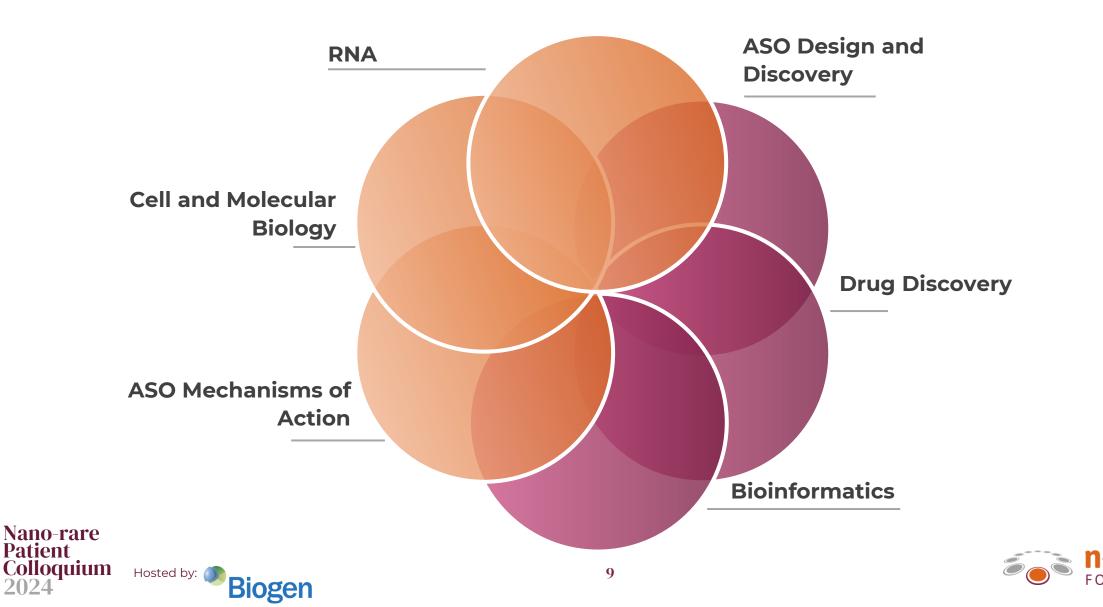


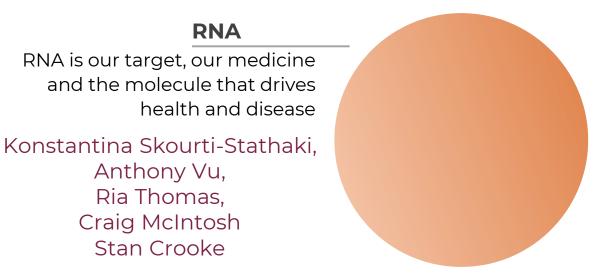
The n-Lorem Lab Does ASO Discovery, Directed and Basic Research

- ASO Discovery
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- Basic research to advance ASO technology

We purposefully created a lab and a team with the right skills, right experience to meet the demand in the highest quality possible

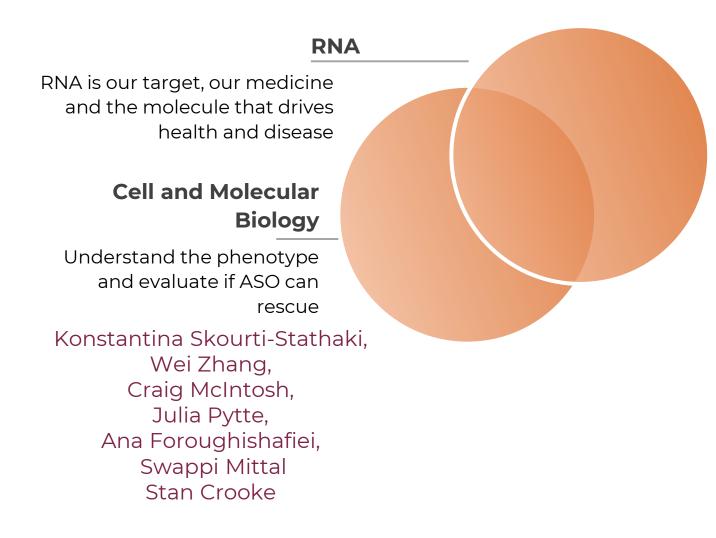


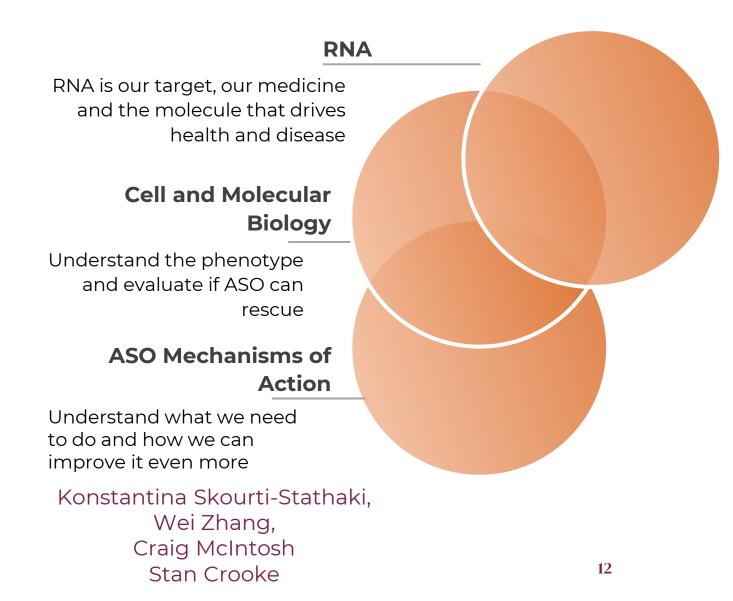


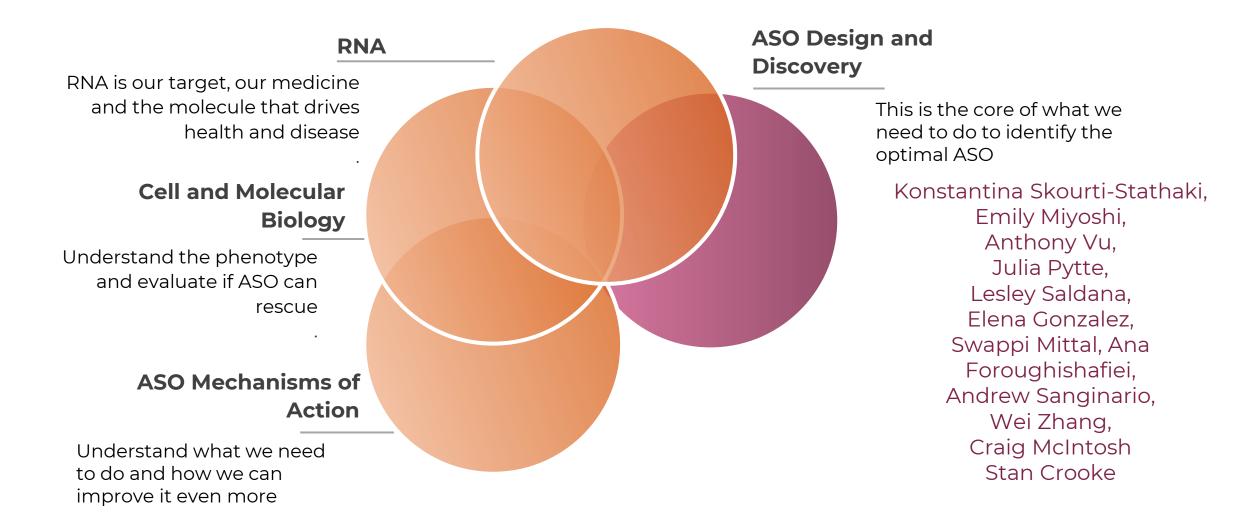


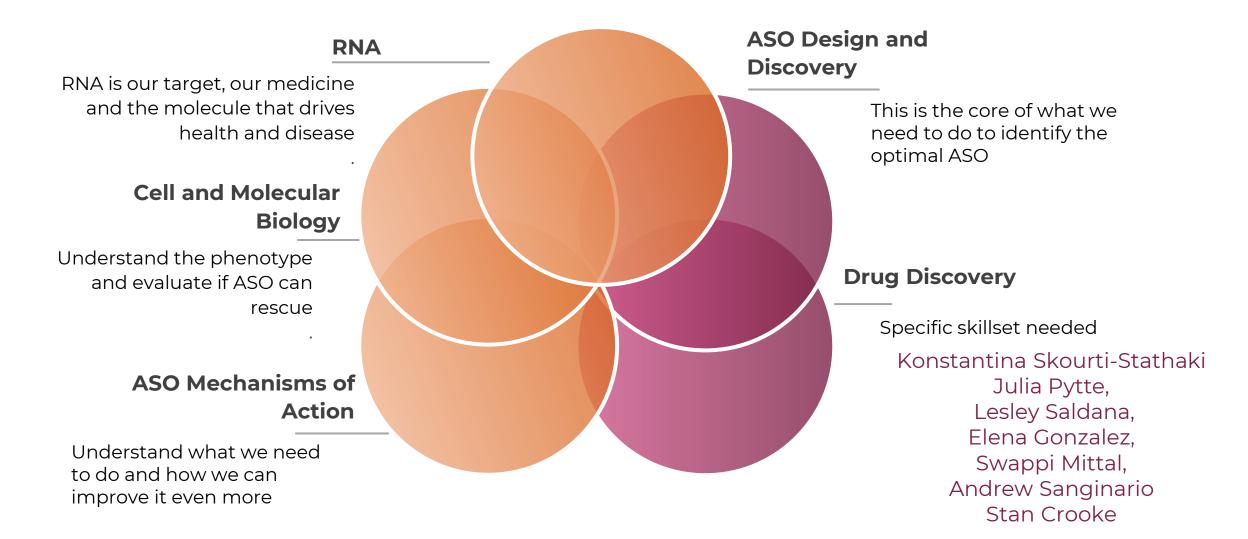


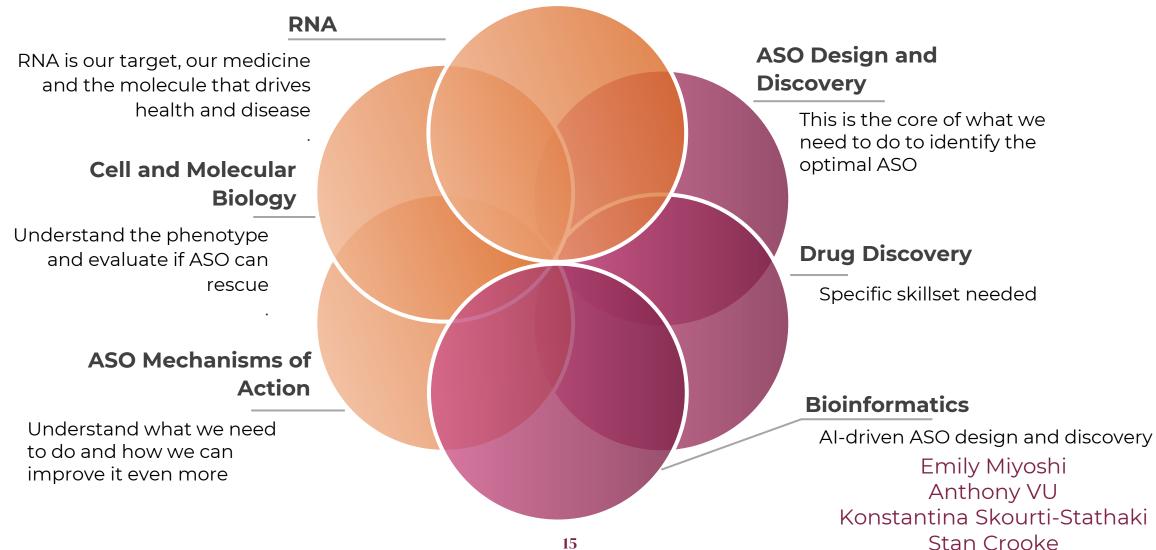




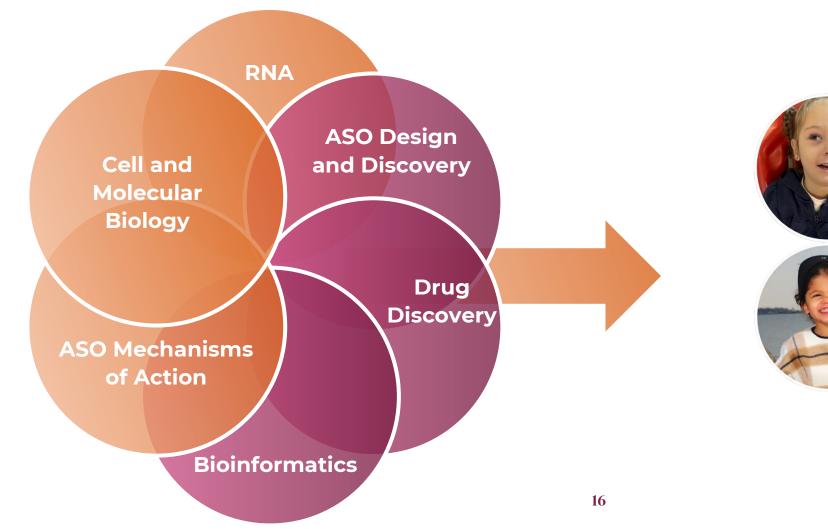








These skills and experience mean that we provide the best possible decisions for our patients



We started as a very small lab that we paid for at lonis but given the demand, we needed a more robust capacity





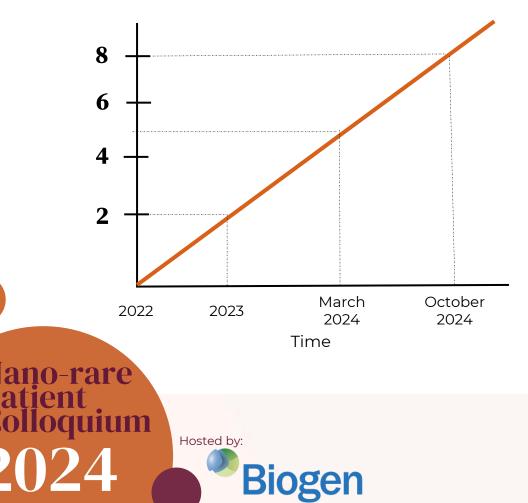
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Here is what we have accomplished.



Rapid Progress with a Newly Established Lab

ASO discovery programs per quarter







Conclusions

- In less than 2 years, we now have ~27 programs that have completed discovery and are progressing into preclinical development
- We are at a rate of completing ~8 programs per quarter and continuing to grow
- We have the capacity to double or triple next year, if we raise the funds





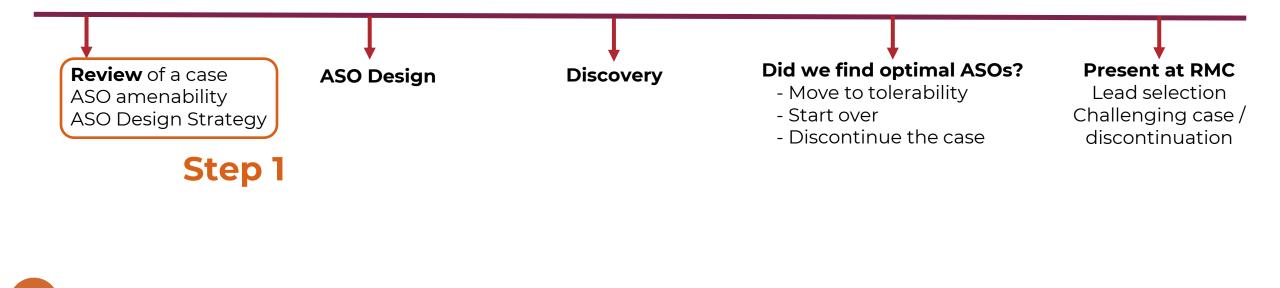
ASO Design and Discovery Process







Checkpoints of ASO Design and Discovery Process







Can ASO Technology Meet the Needs of the Patient?

- Our task is to determine if an ASO strategy may be devised:
 - Is the mutation amenable to an ASO?
 - Impact: Are we confident that we know the nature of the mutation.
 - What ASO design is optimal? Do we know/ understand enough about the proximal pathological molecular mechanisms?
 - Impact: Will the ASO be able to correct these pathological mechanisms?
 - By understanding the cellular phenotype, we can ask if we can correct relevant phenotype
 - We often add new insights into the pathology of the disease.
 - What is the ASO design strategy?

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- Impact: Do we need an allele-selective ASO, a non-allele selective ASO, or a steric-blocking ASO?
- What mutations can we not fix? True null mutations



Can ASO Technology Meet the Needs of the Patient?

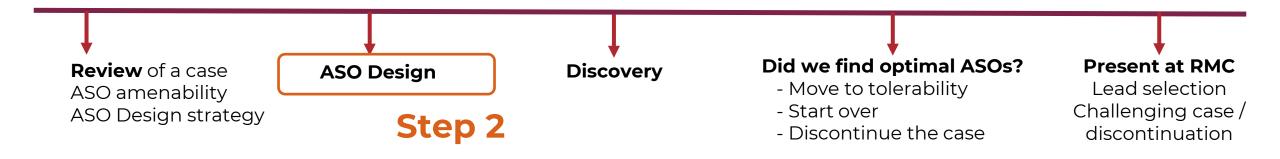
Areas of consideration for ATTC

- Clinical Genetics : Favorable Benefit risk ratio
- ASO Design & Discovery: Amenable to H1 ASO; allele-selectivity required for now but if additional experiments are done, we could consider non-AS approach
- Clinical Development: Straightforward Treatment Goals
- Clinical Operations: Logistically straightforward with clear support from institution.





Checkpoints of ASO Design and Discovery Process







Different ASOs for Different Strategies

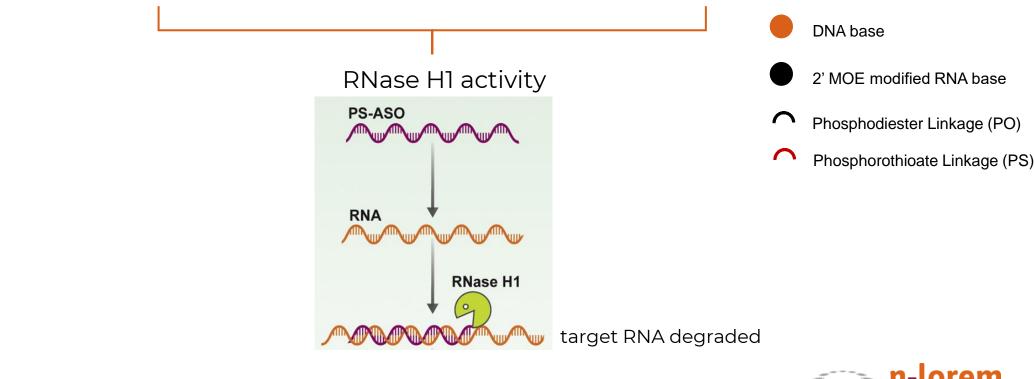
Gapmer (5-10-5; RNA-DNA-RNA)

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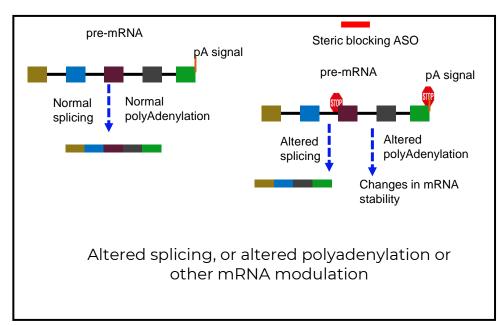




Different ASOs for Different Strategies

Splicing/ steric blocking ASO (18mer, Full MOE)

5' **TACGCTGAGTTGCCGTA** 3'



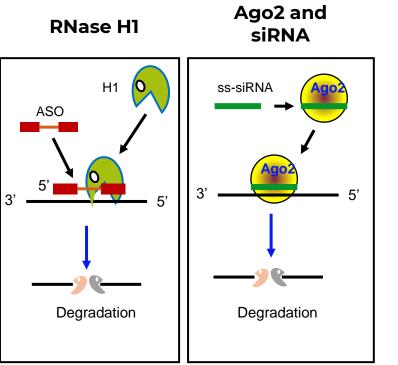
2' MOE modified RNA base
 Phosphorothioate Linkage (PS)





2 ASO Mechanisms: RNA Degradation

 We have led the way in understanding all kinds of mechanisms to be used with ASOs

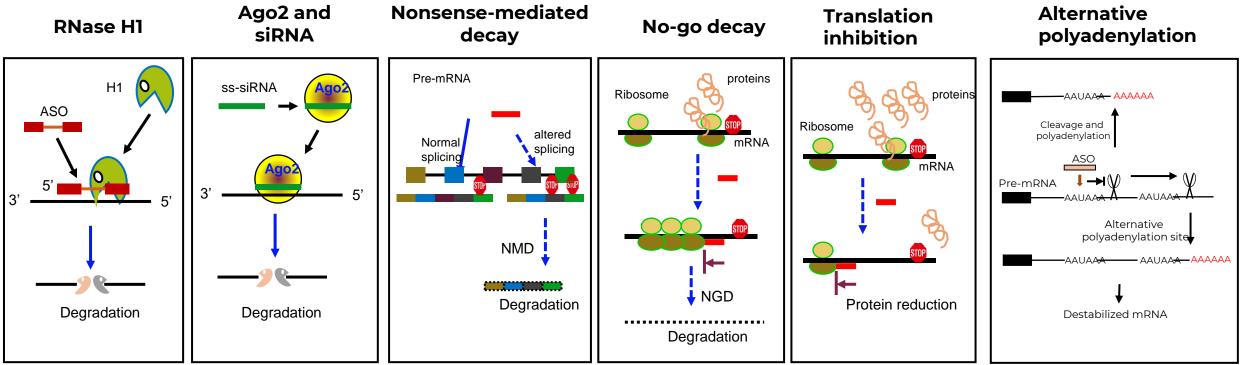


Crooke ST. Curr Mol Med. 2004;4(5):465-87; Crooke ST. NAT, 2017;27(2):70-77; Crooke et al., Nat. Biotech. 2017, 35(3):230-237; Lima WF et al., Cell. 2012;150(5):883-94; Yu D. et al., Cell. 2012 150(5):895-908; Hanecak R et al., J Virol. 1996, 70(8):5203-12; Ward AJ., et al., NAR 2014;42(9):5871-9; Rigo F et al., NCB, 2012;8(6):555-61; Liang XH et al., NAR, 2019;47(13):6900-6916; Crooke ST et al., NAR, 2020, 48(10):5235-5253; Crooke ST et al., (2020) JACS 142(35):14754-14771, Crooke ST et al., Nature Review Drug Discovery, 2021, 1-27, Crooke ST et al., JBC, 2021. 296:1-39; Crooke ST et al., Biochem Pharm, 2021 Jul;189:114196.



2 ASO Mechanisms: RNA Degradation and mRNA Modulation

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AI-informed ASO Design Makes ASO Discovery Highly Efficient

- 35 years of ASO expertise
- Millions of ASOs

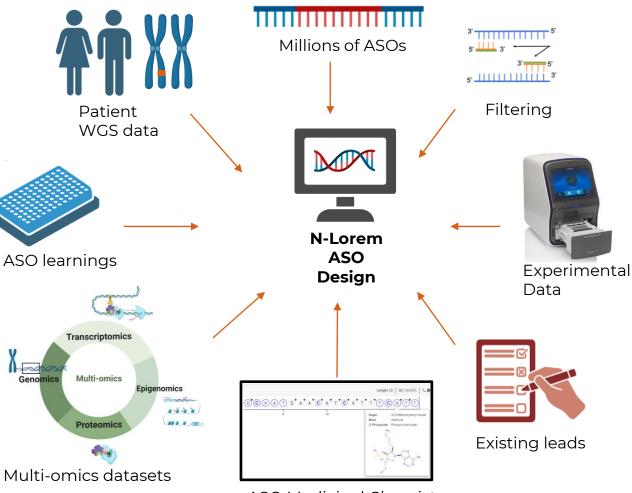
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- All relevant cell and animal models
- Hundreds of thousands of patients
- Continuous learnings from WGS data
- Design ASOs to reach more patients
- State-of-the-art integration of multi-omics data

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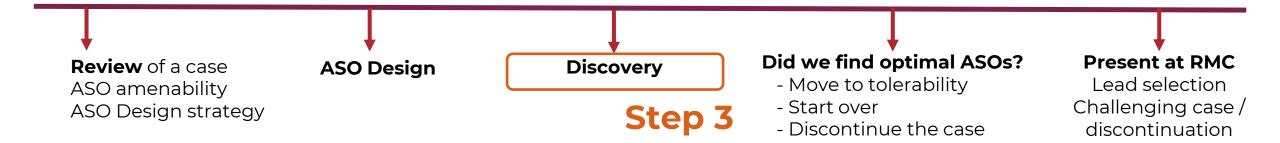


ASO Medicinal Chemistry



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Checkpoints of ASO Design and Discovery Process

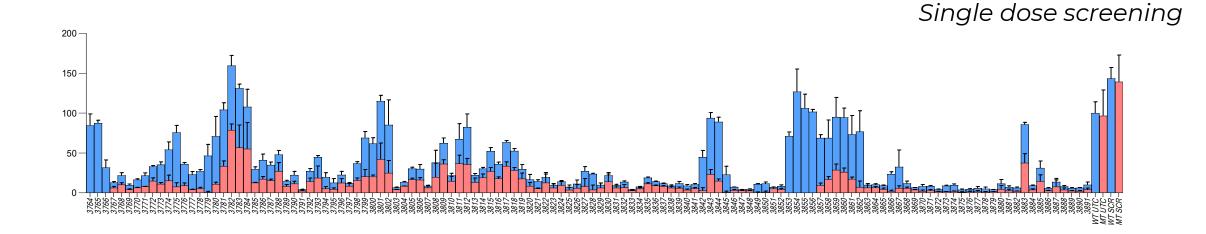






A Rigorous, High-throughput, High-quality ASO Discovery Process: Discovery of an Allele-selective ASO

Screening Step	Purpose	Approximate Minimum Numbers of ASOs Typically Evaluated	Minimum Criteria
Primary ASO screen	To identify optimal sites in target RNA for ASO and H-1 binding	~500	>80% target reduction

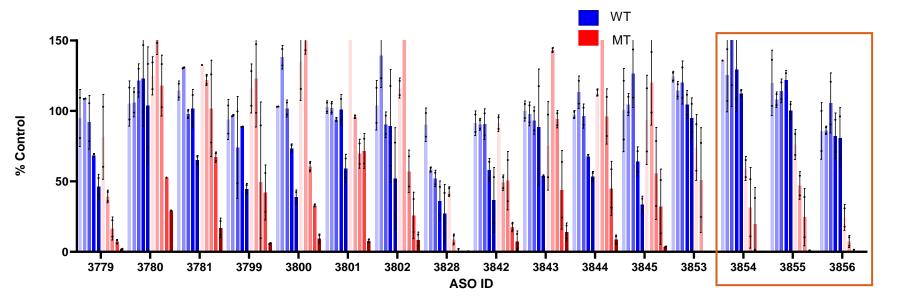






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Dose response screening



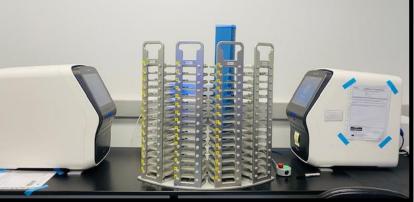
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In vitro off-target analysis	To confirm selectivity of ASO for target RNA vs. any worrisome off-target	As many as necessary	~10-fold difference in IC50s for target RNA vs. off target
BJAB Assay	To exclude activators of innate immunity	~50-75	Less than 2-fold increase in TNF-alpha at high ASO concentrations



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Critical Decisions and Challenges We Address

Decisions:

- Discontinue a case
- Do we have the best ASO possible, or do we need to redesign?

Challenges:

- Encounter differences with published literature/ data
- We don't know enough about the nature of the mutation
- Highly structured RNA
- Small number of SNPs for allele-selective ASOs





We Do Everything We Can Before We Discontinue a Case

- Are the ASOs we have sufficient, given the function of the gene?
- If we do not know, we go ahead and perform research experiments
 - How does the cell tolerate the specific ASO?
 - Does it still do its normal function?
 - Can the ASO rescue the pathological phenotypes?
- If allele-selectivity is not sufficient, we bring the case to RMC and recommend to discontinue





A Minor Error in Sequencing Proved to be a Major Problem for ASO Discovery

 Discrepancy in indel sequence was resolved enabling the treatment of all four patients with the same ASO

Genetic Report		Long-read WGS	
Indel Variant	Sequencing Method	Variants	Sequencing Method
c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>G</mark> ACAGGT	Sanger seq	c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>T</mark> ACAGGT	Long-read WGS
c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>T</mark> ACAGGT	NGS (likely short read)	c.2053-3358_2053-3350 delinsTGTTTTTTACAT <mark>T</mark> ACAGGT	Long-read WGS
c.2053-3358_2053-3350 delins (sequence not specified)	Sanger/capillary seq	c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>T</mark> ACAGGT	Long-read WGS
c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>G</mark> ACAGGT	Sanger seq and NGS (likely short read)	c.2053-3358_2053-3350 delinsTGTTTTTACAT <mark>T</mark> ACAGGT	Long-read WGS



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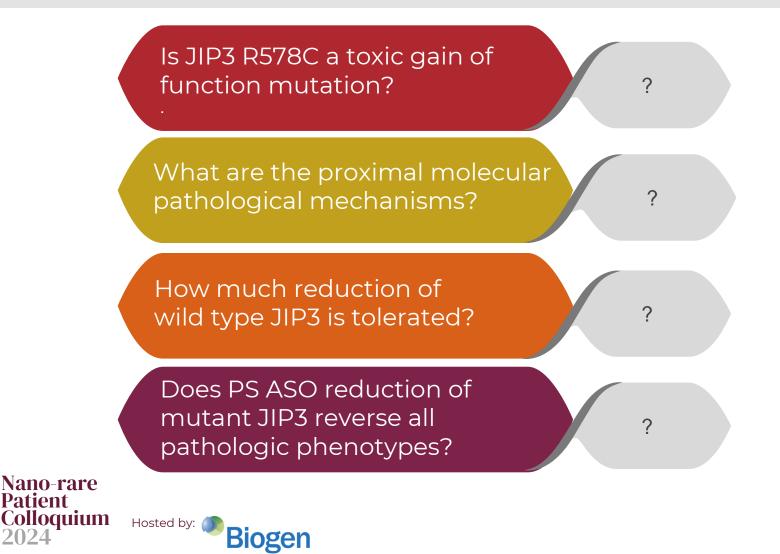
Directed Research to Better Understand Targets and Mutations





What is the Nature of JIP3 R578C Mutation and is it **ASO Amenable?**

Open questions that needed answering before moving to ASO discovery



Patient



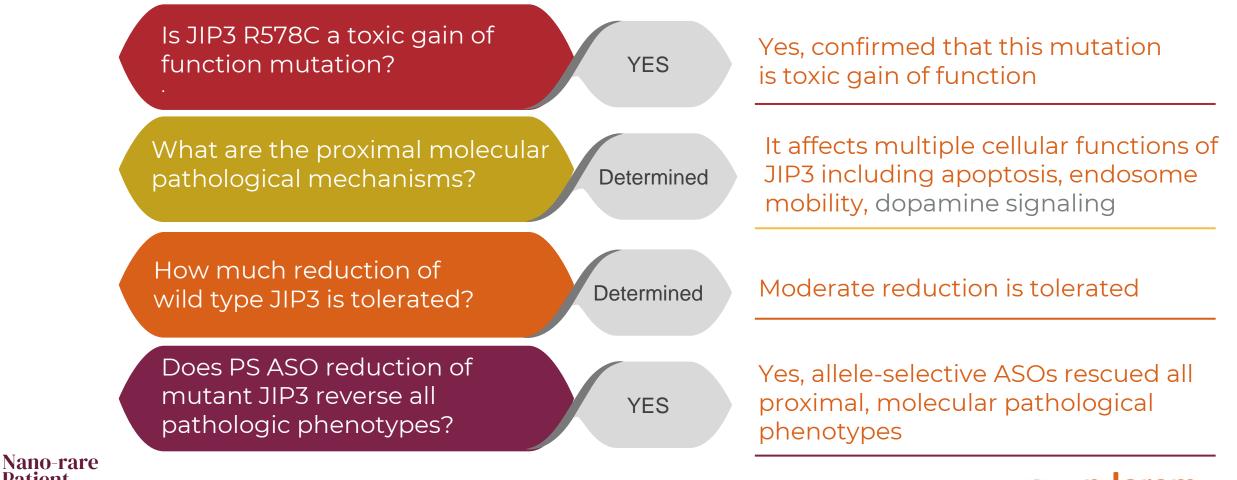
What is the Nature of JIP3 R578C Mutation and is it ASO Amenable?

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We answered these questions to design and discover the most optimal ASOs





Basic Research to Advance ASO Technology and Reach More Patients





How Can We Expand to Reach More Patients?

- Mechanisms to avoid adverse events and immune activation
 Why? We could make even better ASOs and be even more efficient
- Enhance allele-selectivity
 - Why? ~50% of our programs are allele-selective. We can make better ASOs and reach our patients faster and help even more patients
- Upregulation strategies with ASOs
 - Why? ~40% of our declined cases require an ASO-mediated protein upregulation strategy. We want to be able to accept and treat these patients

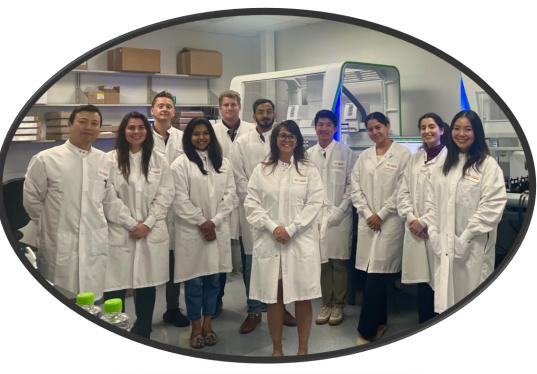
We have the knowledge and expertise to address these issues today





Conclusions

- We have an outstanding team in place to discover optimal ASOs
- We have the capabilities to do this rapidly, in a costeffective and high-throughput manner







Conclusions

- We have an outstanding team in place to discover optimal ASOs
- We have the capabilities to do this rapidly, in a costeffective and high-throughput manner
- We apply and share our learnings for each patient to maximize our opportunities
- 100% commitment to the patient: The most important decision that we are making in every step of the way is –

Is the ASO sufficient and optimal for this patient?







Join Us As We Make The World A Better Place, One Patient, One Family at a Time