



# Chemoenzymatic Ligation for siRNA and sgRNA Manufacturing: Recent Advances and Development Strategies

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TIDES USA: Bridging Research to Reality. San Diego – May 22<sup>nd</sup> 2025

## Disclosures

- Dr. Butler is an employee of Hongene Biotech Corporation and serves on the board of directors of Akte Therapeutics



## Presentation overview

- Recap of OTS conference presentation
- Chemoenzymatic ligation for siRNA synthesis
  - Inclisiran case study
  - Divalent siRNA case study
  - siRNA summary
- Chemoenzymatic ligation for sgRNA synthesis
  - 100mer G211 sgRNA case study
  - Hongene/ReciBioPharm partnership
- Engineered thermostable T4 RNA ligase
- Control of stereochemistry
- Conclusions



## Recap: Presentation at OTS conference, Montreal, October 2024

**Title:** The Evolving Landscape of Oligonucleotide Manufacturing: Meeting Future Demand with Ligation Technologies

### Topics covered:

- The evolving market for sgRNA and siRNA therapeutics
- Three generations of oligonucleotide synthesis technology
- Mechanism, properties, and immediate applications of ligation technology
- Examples of products synthesized by ligation at Hongene
- CMC features of ligation
- Current and future oligonucleotide fragment manufacturing processes

Email me - [david.butler@hongene.com](mailto:david.butler@hongene.com) - for a copy of the OTS deck, which can also be [viewed here](#)



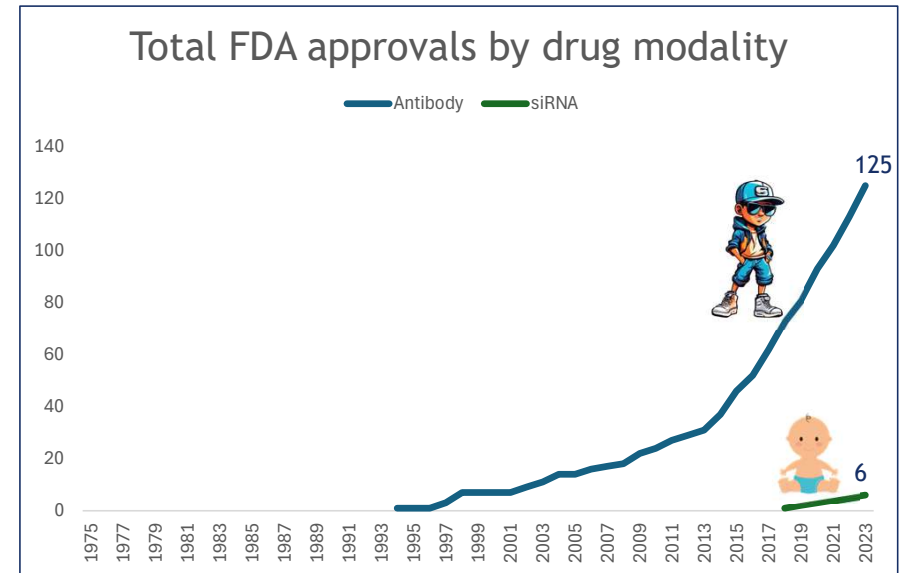
## Market potential of siRNA drugs

Market demand for siRNA drugs and CMC challenges expected to grow significantly over the next decade

Cardiovascular disease area	Lead siRNA drug	Target	Phase	Possible approval
Hypercholesterolemia	Inclisiran	PCSK9	4	Approved
Hypertriglyceridemia	Plozasiran	APOC3	3	~2025
Elevated Lp(a)	Olpasiran	LPA	3	~2027
Hypertension	Zilebesiran	AGT	2	~2030
NASH/MASH	Rapirosiran	HSD	2	~2030
Obesity	<i>Multiple</i>	INHBE	1	2030s

- Several disease areas have potential to reach >10M US patients
- inclisiran + 4 could be approved by 2030
- Research suggests >10 t API required by early 2030s

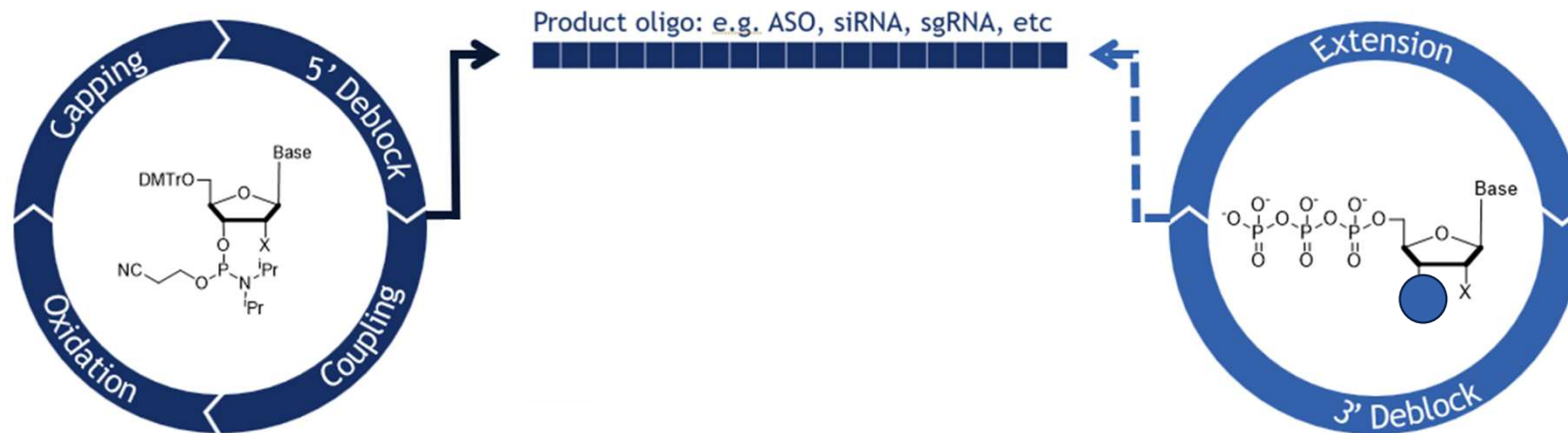
siRNA is in its infancy in its life trajectory as a therapeutic drug modality



The Antibody Society. Therapeutic monoclonal antibodies approved or in regulatory review. (Aug 2024); [Antibody therapeutics product data - The Antibody Society](#)



# Three generations of oligonucleotide synthesis technology



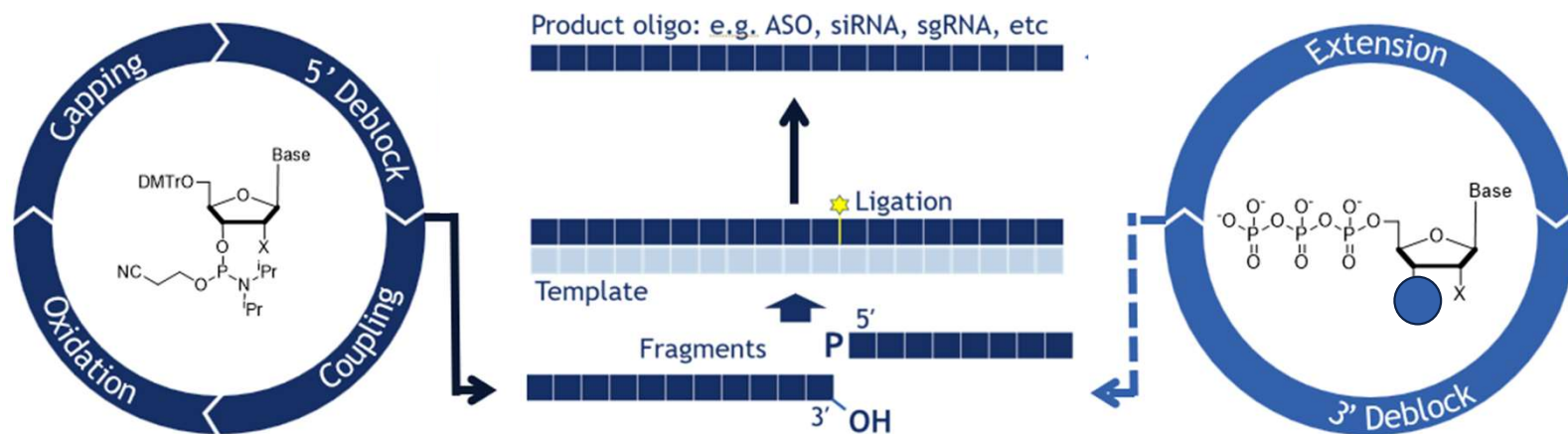
GENERATION 1	
Raw materials	Phosphoramidites
Development status	>40 years, current paradigm
Product purity	Lower
Product yield	Lower
Sustainability	>3,000 kg RM/kg API <sup>1</sup>

GENERATION 3	
Raw materials	NTPs, 3'-protected NTPs, enzymes
Development status	Very early <sup>2,3,4,5</sup>
Product purity	Enzymes need to be engineered
Product yield	Enzymes need to be engineered
Sustainability	Best (aqueous)

1. B. Andrews et al, *JOC*, 2021; 2. E.R. Moody et al, *Science*, 2023; 3. N. Sabat et al, *Front Chem*, 2023; 4. D. Wiegand et al, *Nat Biotechnol*, 2024; 5. S. Forget et al, *NAR*, 2025.



# Three generations of oligonucleotide synthesis technology



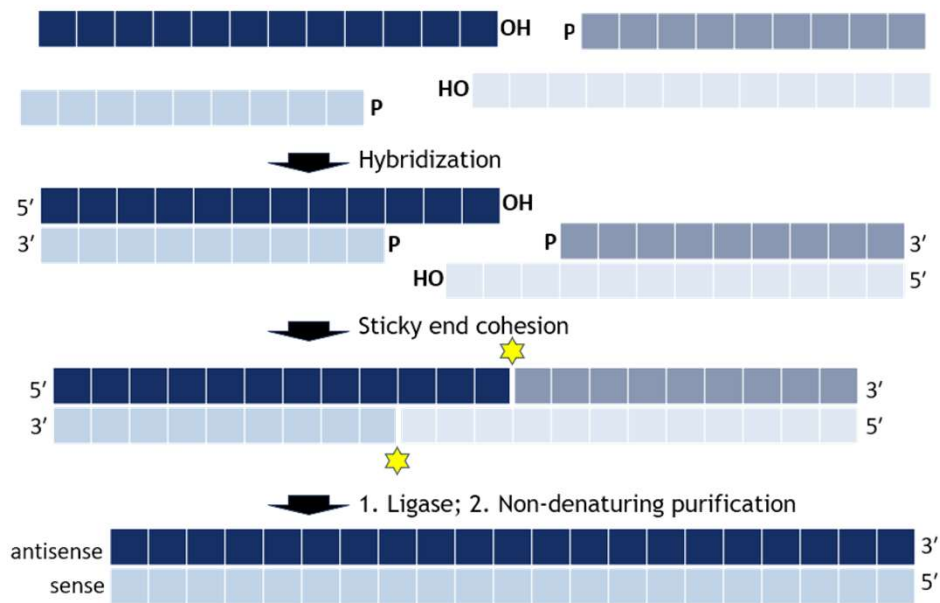
	GENERATION 1	GENERATION 2	GENERATION 3
Raw materials	Phosphoramidites	Oligonucleotide fragments	NTPs, 3'-protected NTPs, enzymes
Development status	>40 years, current paradigm	Ready for manufacturing	Very early <sup>2,3,4,5</sup>
Product purity	Lower	Higher	Enzymes need to be engineered
Product yield	Lower	Higher	Enzymes need to be engineered
Sustainability	>3,000 kg RM/kg API <sup>1</sup>	Better (partly aqueous)	Best (aqueous)

1. B. Andrews et al, *JOC*, 2021; 2. E.R. Moody et al, *Science*, 2023; 3. N. Sabat et al, *Front Chem*, 2023; 4. D. Wiegand et al, *Nat Biotechnol*, 2024; 5. S. Forget et al, *NAR*, 2025.

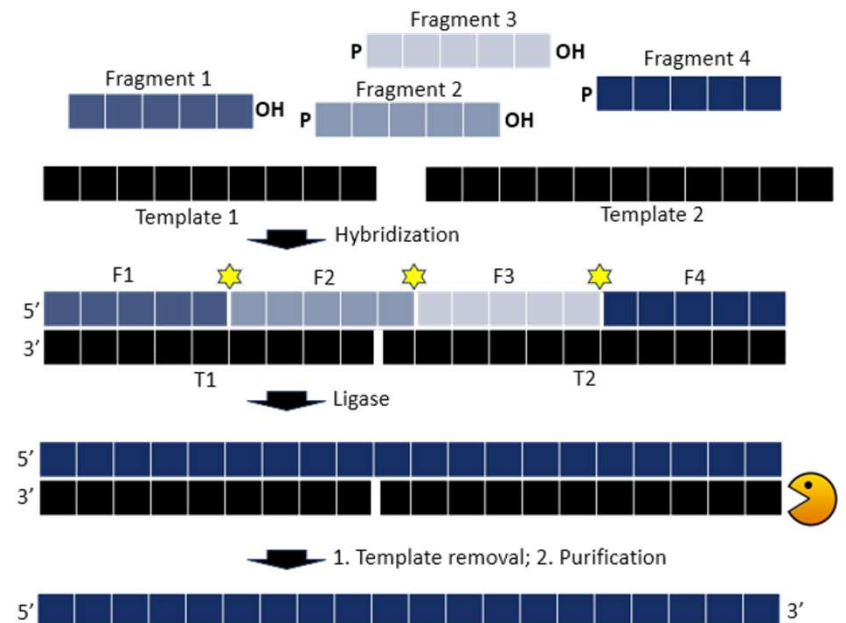


# Hongene's chemoenzymatic ligation processes for siRNA and sgRNA

## 1. siRNA Sticky end ligation<sup>1,2,3,4</sup>



## 2. sgRNA Splinted (template) ligation<sup>5,6</sup>



1. H.G. Khorana, *Pure Appl Chem*, 1968; 2. A. Susic et al, *Bioconj Chem*, 2014; 3. S. Paul et al, *ACS Chem Biol*, 2023; 4. S. Kajimoto et al, *AEM*, 2022; 5. M. Moore et al, *Methods Enzymol*, 2000; 6. N.Sabat et al, *Nat Commun*, 2024.



# Chemoenzymatic ligation for siRNA synthesis



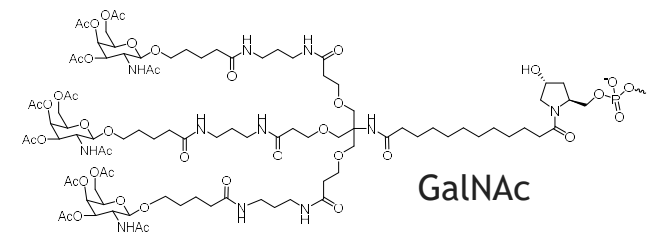
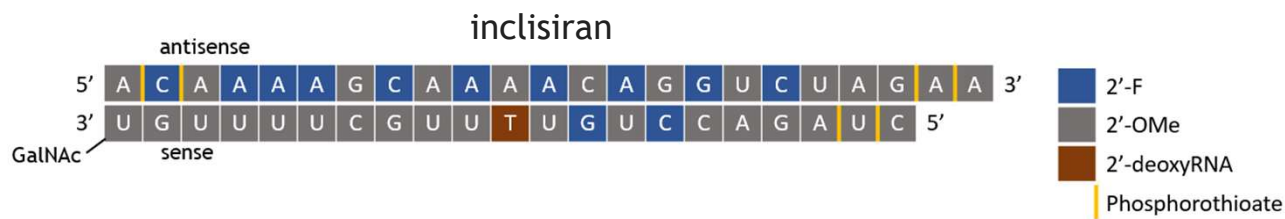
# siRNA ligation case study inclisiran



# siRNA ligation case study - inclisiran

## Background

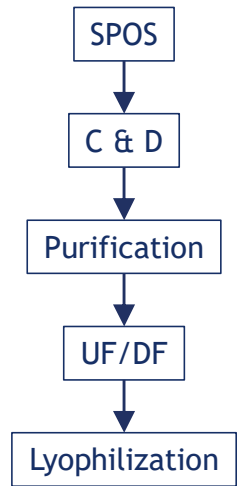
- inclisiran (LEQVIO®)
  - siRNA targeting PCSK9
  - Adjunct to diet and statin therapy for adults with primary hyperlipidemia, including HeFH, to reduce LDL-C
- Contains chemistries typically found in siRNA therapeutics
  - Ribose: 2'-F, 2'-OMe, 2'-deoxy
  - Backbone: PO with PS on ends
  - Targeting: Triantennary GalNAc
- Excellent model siRNA for chemoenzymatic ligation PoC studies



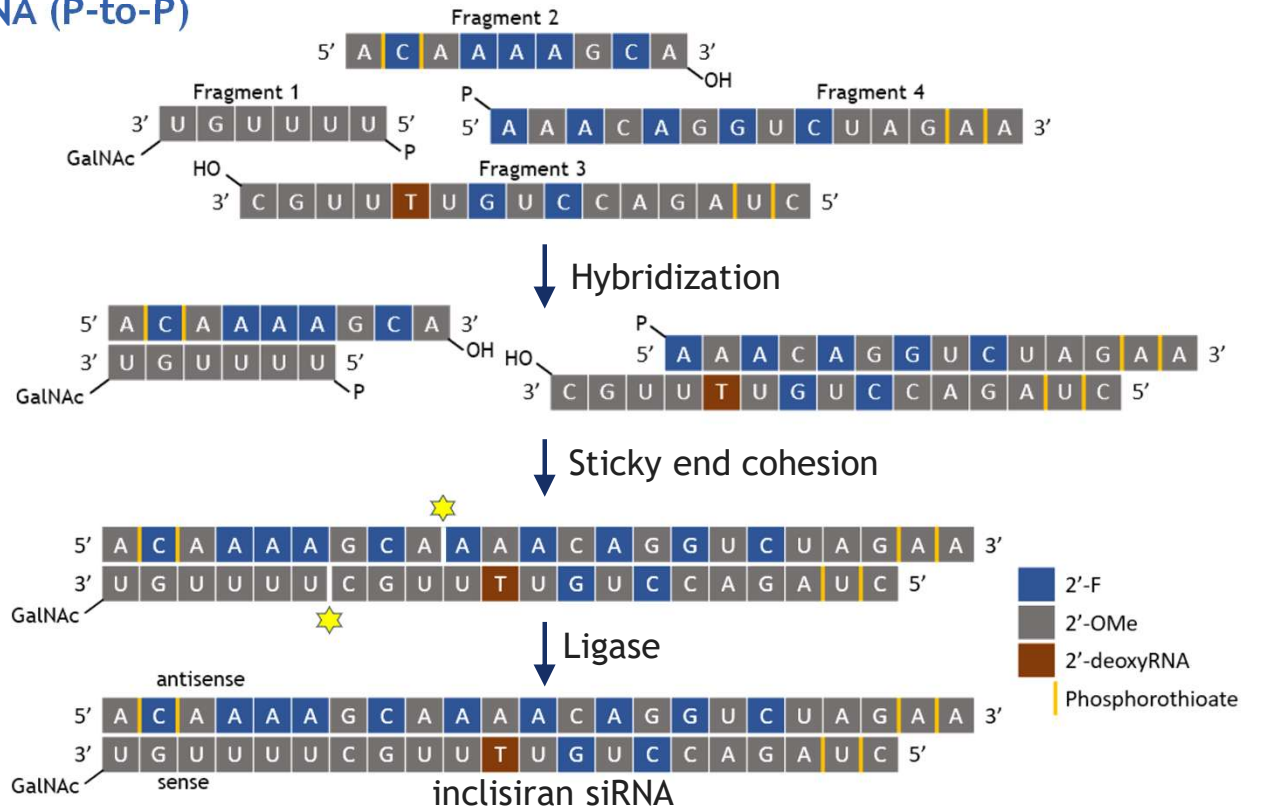
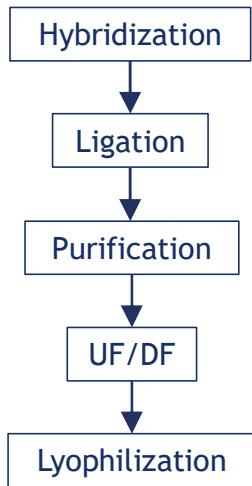
# Inclisiran: Chemoenzymatic process design

## Purified fragments and purified siRNA (P-to-P)

### Oligonucleotide fragments



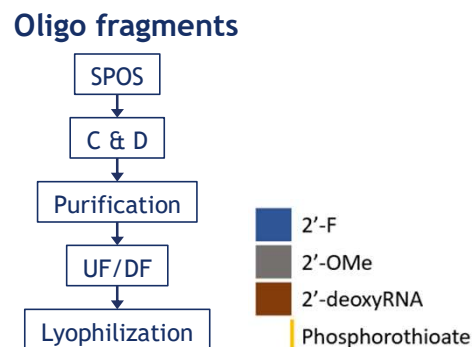
### Chemoenzymatic ligation



# Inclisiran: Control of oligonucleotide fragment purity

## IPRP-HPLC/MS

- Oligo fragment and siRNA purified by HPLC (P-to-P strategy)
- Most conservative approach - Clean oligo fragments for ligation
- Oligo fragment purity and impurities are well-controlled



Fragment	Sequence and chemistry	Length	Purity	Yield
1		6	98.9%	39.2%
2		9	98.4%	45.2%
3		15	97.9%	55.5%
4		14	96.3%	35.8%

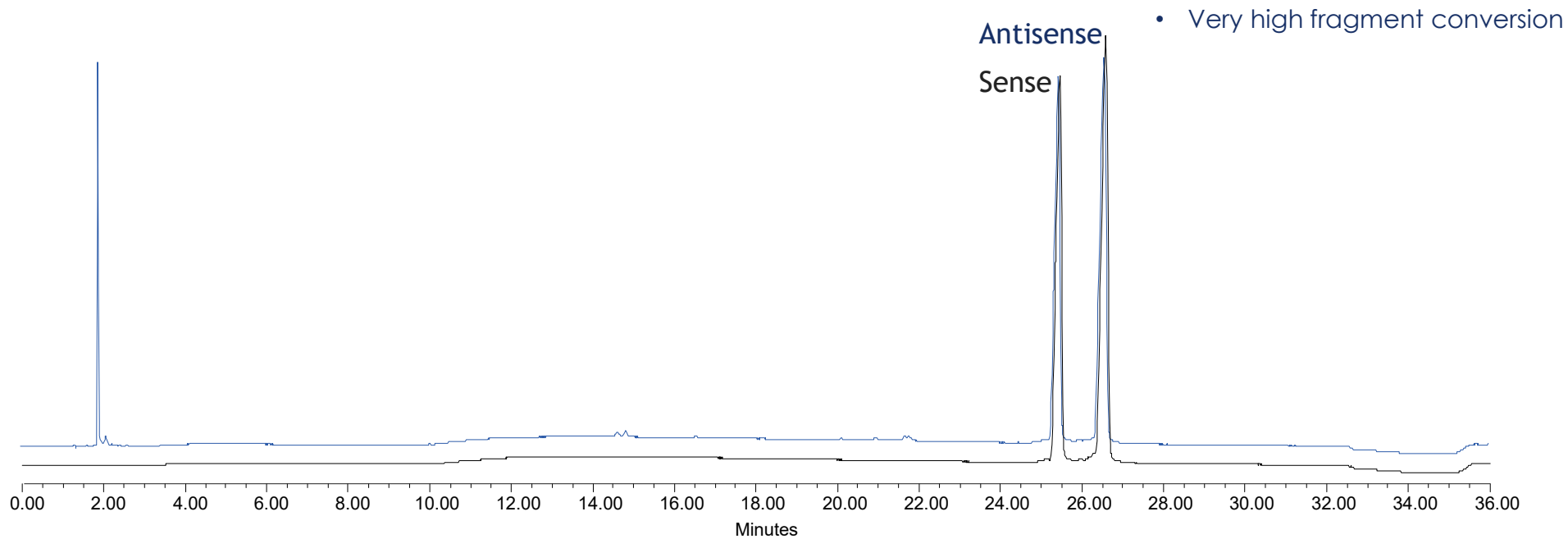
Unoptimized yields based on calculated molar extinction coefficients. Purity measured by IPRP-HPLC



# Inclisiran: Control of product purity and impurities

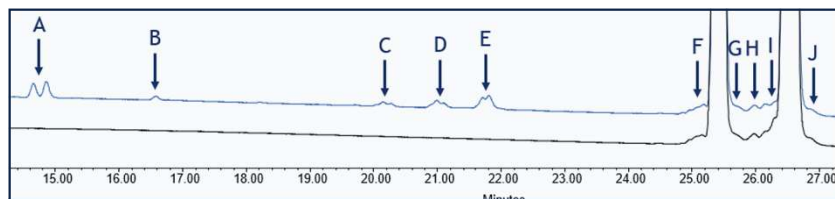
IPRP-HPLC/MS

Crude: 95.1%  
Purified: 96.8%



# Inclisiran: Control of product purity and impurities

IPRP-HPLC/MS



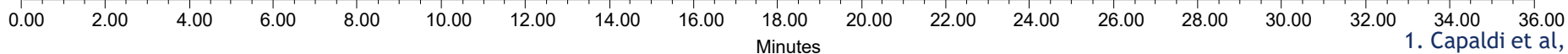
Crude: 95.1%  
Purified: 96.8%

Impurity peak	Assignment by MS	Crude area %	Purified area %
A	Fragment 2	0.6%	N.D.
B	Fragment 1	0.1%	N.D.
C	Fragment 4 minus 5'-P-1nt	0.4%	N.D.
D	Fragment 4 minus 5'-P	0.4%	N.D.
E	Fragment 4 + 328Da	0.9%	N.D.
-	<b>Sense</b>	<b>42.6%</b>	<b>43.1%</b>
F	SS-16Da, SS-1nt	0.3%	0.6%
G	SS-204Da, SS+12Da	0.2%	0.4%
H	AS-20Da	0.2%	0.5%
I	AS-16Da, AS-20Da, AS-2Da	0.3%	1.1%
-	<b>Antisense</b>	<b>52.5%</b>	<b>53.7%</b>
J	SS+1nt	0.2%	0.3%

Antisense

Sense

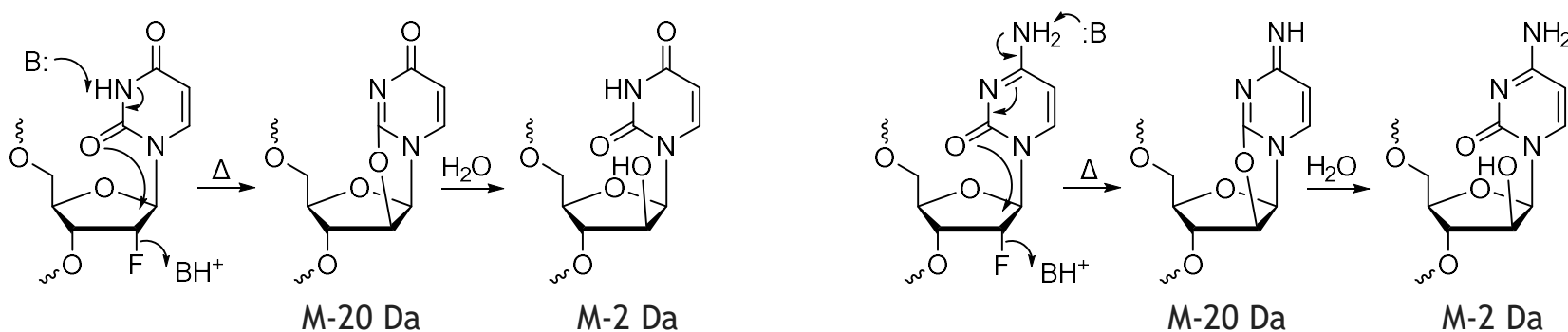
- Very high fragment conversion
- Residual fragments easily removed during purification
- No new ligation-related impurities
- Impurities all SPOS related
- All impurities below suggested 1.0% ID threshold (Capaldi et al<sup>1</sup>)
- M-20 Da and M-2 Da lyophilization-related and elevated in purified product



1. Capaldi et al, NAT, 2017.



## Degradation of siRNA containing 2'-F pyrimidines



- Pathway previously reported for siRNA duplexes containing 2'-F pyrimidines on storage at elevated temperatures
  - 2,2'-anhydropyrimidines (M-20 Da) were observed predominantly during storage as solid<sup>1</sup>
  - 2,2'-anhydropyrimidines hydrolysis products (arabinosyl pyrimidines, M-2 Da) observed predominantly during storage in solution<sup>1,2</sup>
  - Conversion of 2,2'-anhydro to arabinosyl pyrimidines increases with neutralization of liberated  $HF^3$
- Aligned with our observations of elevation of these impurities following lyophilization

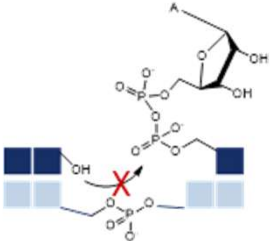
1. C.J. Calvitt et al, *Oligonucleotides*, 2010; 2. S. Seiffert et al, *Anal Biochem*, 2011; I.L. Doerr et al, *JOC*, 1967.



# Oligonucleotide fragments are a focus of impurity control

## Two reasons for improved purity for Gen 2 vs Gen 1

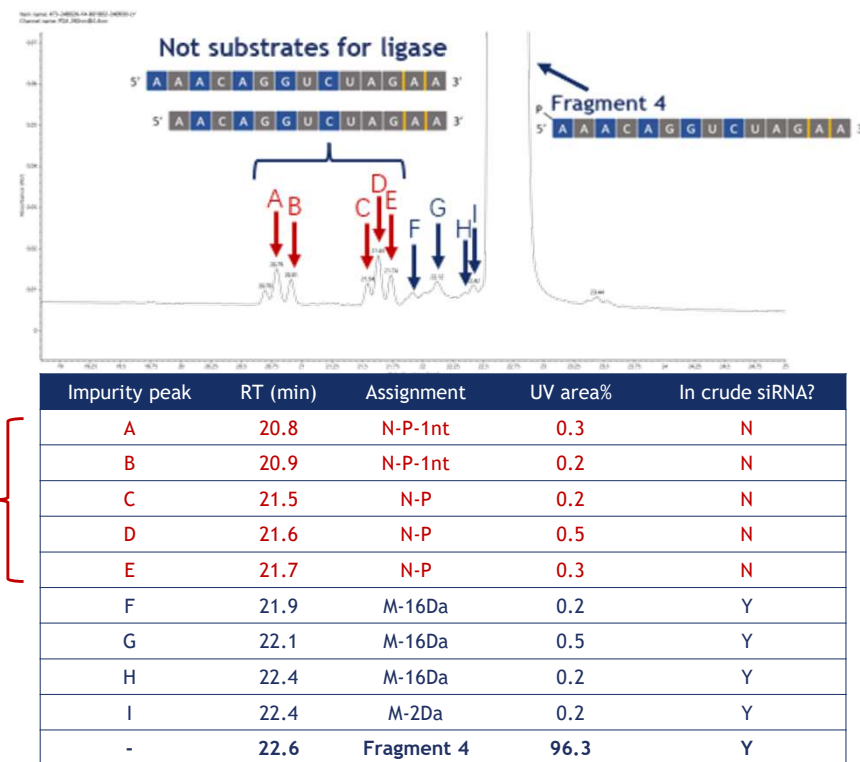
1. SPOS-related impurities such as N-1nt, N+1nt, PO, M-20Da, M-2Da, CNET, iBu, and GalNAc degradants reduced compared to full-length synthesis
  - Less opportunity to compound and accumulate
2. Some oligonucleotide fragment impurities aren't good substrates for ligase
  - These don't persist in drug because they are purged during siRNA column chromatography



Many impurities in oligonucleotide fragments are purged

- Not ligase substrates (e.g. 5'(N-1nt) or
- Inefficient hybridization

Fragment 4 impurities - IPRP-HPLC/MS



# Ligation case study

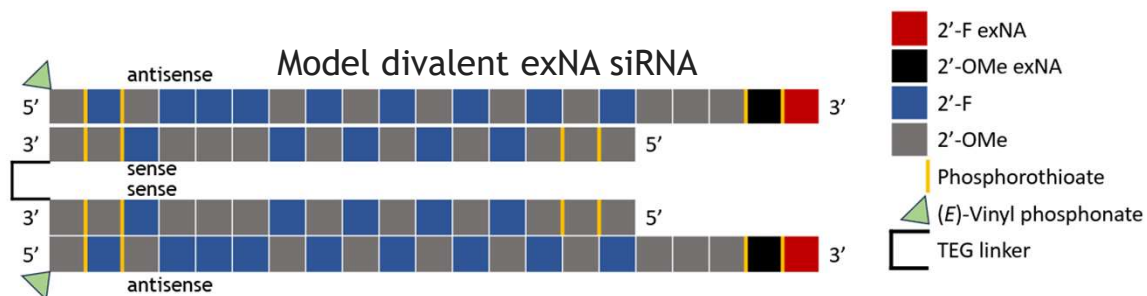
## Divalent exNA siRNA



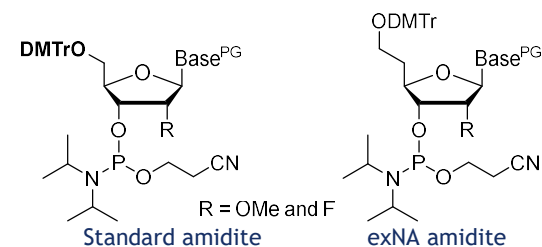
# Ligation case study: Divalent exNA siRNA

## Background

- UMass has invented divalent siRNAs with enhanced efficacy. These contain exNA chemistry<sup>1</sup>
- Scientists at Broad, UMass and Harvard recently identified a divalent siRNA clinical candidate for Prion disease<sup>2</sup>
- Hongene manufactured this clinical candidate using SPOS for the clinical trial
- We were curious as to whether divalent siRNA chemistry was amenable to chemoenzymatic ligation
- A model divalent siRNA construct was selected for PoC studies



## Novel exNA phosphoramidite Starting Materials scaled for GMP oligonucleotide CDMO services

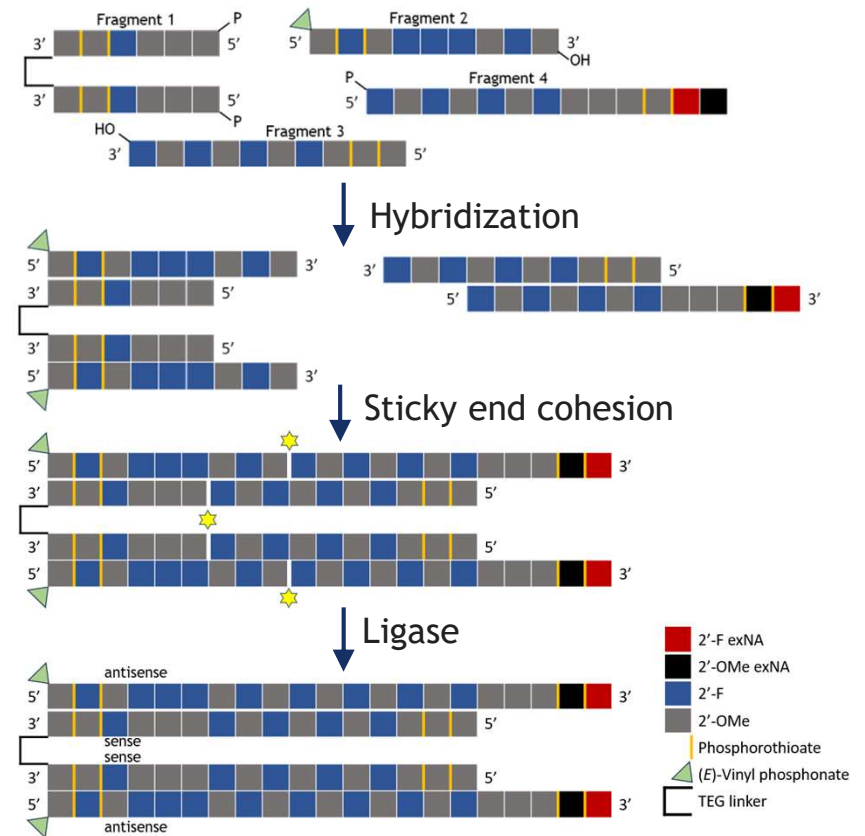
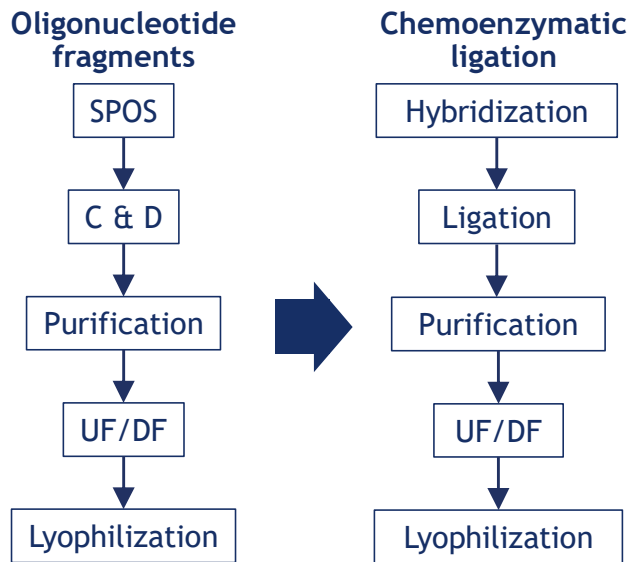


1. K. Yamada et al, *Nat Biotech*, 2024; J.E. Gentile et al, *bioRxiv*, 2024.



# Divalent siRNA: Chemoenzymatic process design

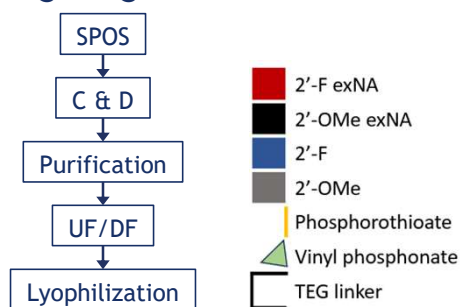
## Purified fragments and purified siRNA (P-to-P)



# Divalent siRNA: Control of oligonucleotide fragment purity

## IPRP-HPLC/MS

### Oligo fragments



Fragment		Length	Purity	Yield
1		12	96.2%	26.3%
2		9	95.4%	48.2%
3		10	96.0%	42.5%
4		12	93.6%	30.2%

Unoptimized yields based on theoretical molar extinction coefficients. Purity measured by IPRP-HPLC



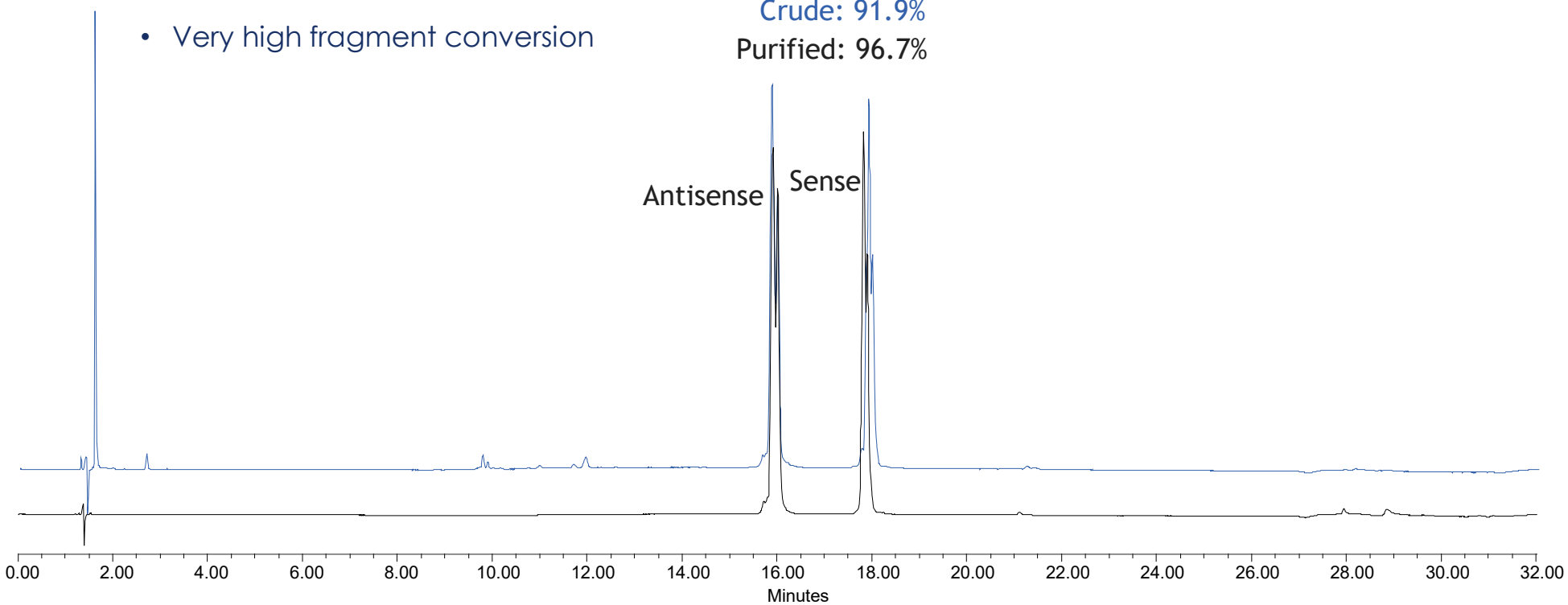
# Divalent siRNA: Control of purity and impurities

## IPRP-HPLC/MS

- Very high fragment conversion

Crude: 91.9%  
Purified: 96.7%

Antisense      Sense



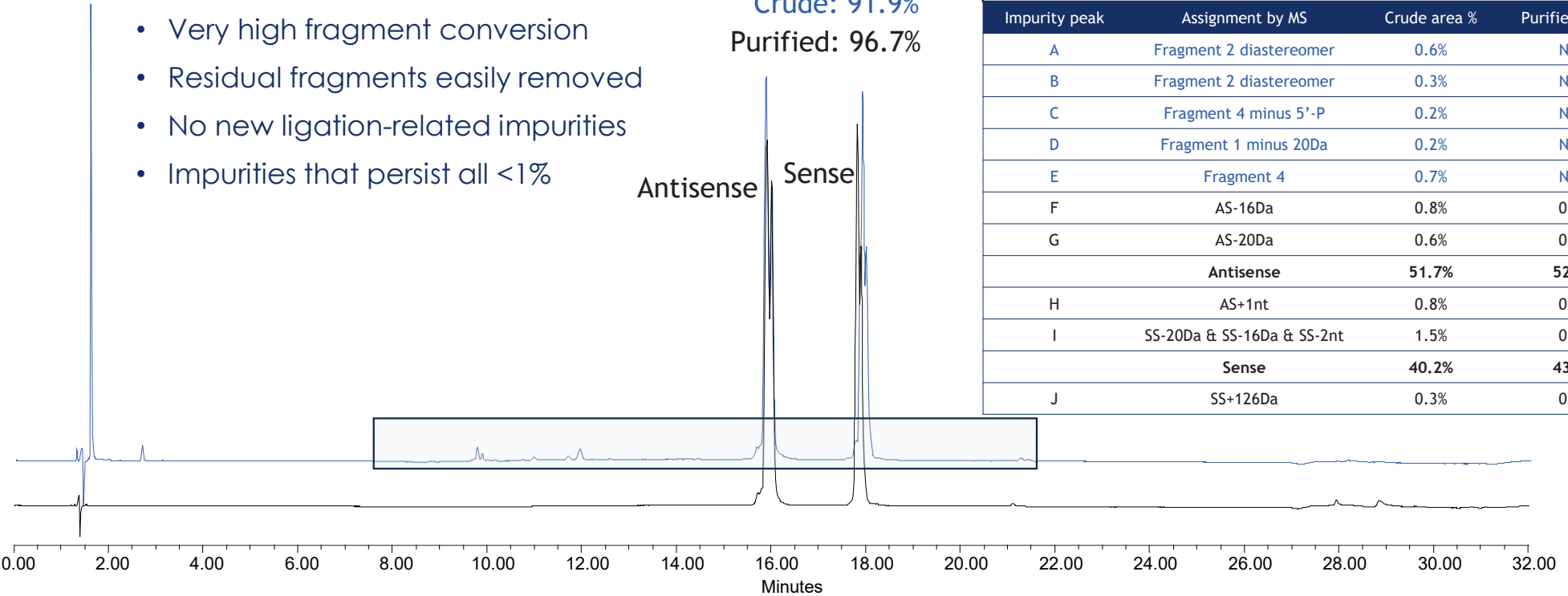
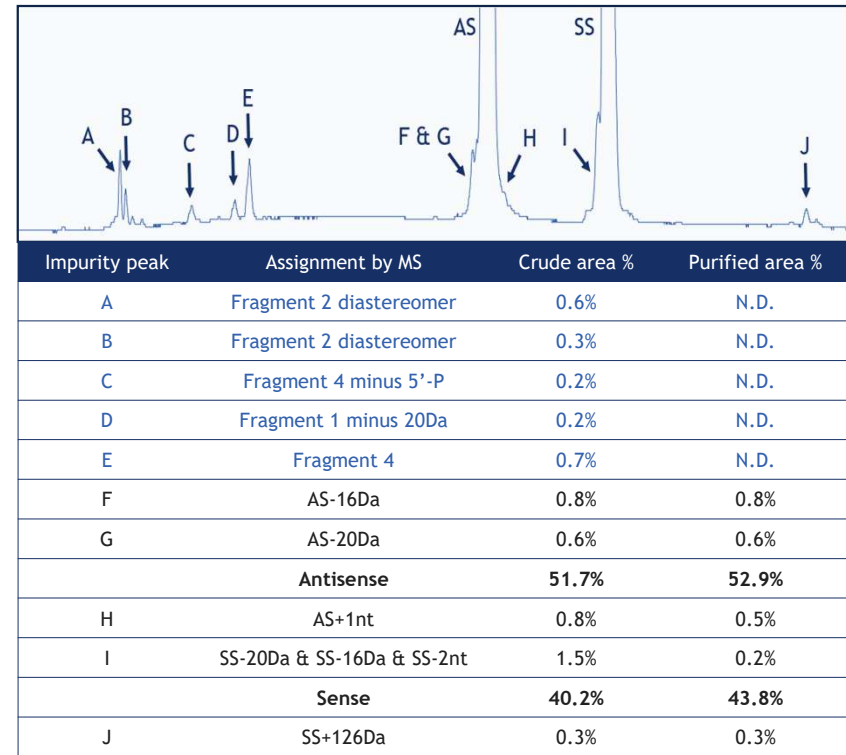
# Divalent siRNA: Control of purity and impurities

## IPRP-HPLC/MS

- Very high fragment conversion
- Residual fragments easily removed
- No new ligation-related impurities
- Impurities that persist all <1%

Crude: 91.9%  
Purified: 96.7%

Antisense Sense



# siRNA summary



## Summary of exemplary siRNAs synthesized by ligation

Molecule	GMP	Oligonucleotide chemistry				Synthesis strategy <sup>1</sup>	Yield <sup>2</sup>	Purity <sup>3</sup>
		SS/AS	2'-Ribose mods	Backbone	GalNAc			
Inclisiran siRNA	-	21/23	2'-OMe, 2'-F, 2'-deoxy	PS/PO	✓	P-to-P	26%	97%
Divalent siRNA	-	16/21	2'-OMe, 2'-F, exNA, (E)-VP	PS/PO/TEG	x	P-to-P	19%	97%
C-to-P siRNA	-	21/23	2'-OMe, 2'-F	PS/PO	✓	C-to-P	43%	96%
CDMO siRNA	No	ND	2'-OMe, 2'-F	PS/PO	✓	C-to-P	~960 g	95%
CDMO siRNA	Yes	ND	2'-OMe, 2'-F	PS/PO	✓	P-to-P	~1,020 g	97%

1. P-to-P = HPLC purified fragments and purified siRNA; C-to-P = UF/DF processed fragments and HPLC purified siRNA; 2. % Yields based on theoretical MEC, calculated from lowest yielding fragment; 3. Denaturing IPRP-UPLC method

- Managing quality and cost of production
  - P-to-P generally adopted as the most conservative for PoC studies and FIH GMP batch
  - C-to-P gives similar purity as P-to-P, but with higher yield
  - C-to-C will be most effective strategy to manage costs for high volume CVD indications
  - Batch mode and single use bioreactors (scalable); at least 100 g siRNA per L is feasible
  - Plenty of room to optimize oligonucleotide fragments processes for scalability and yield



# Chemoenzymatic ligation for sgRNA synthesis



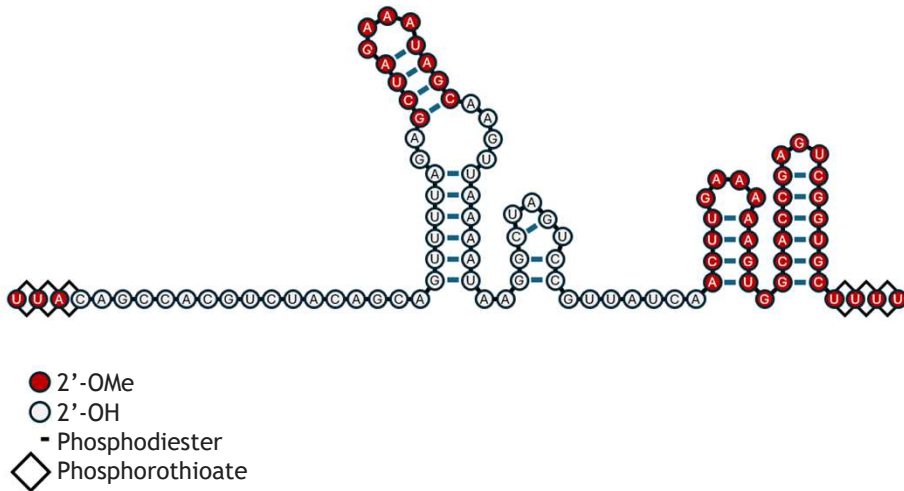
# Ligation case study

## G211 sgRNA



# Ligation case study: G211 100mer sgRNA

Highly modified 100mer G211 sequence



## Background

- We sought out a relevant sgRNA sequence for PoC ligation study
- *In vivo* TTR gene editing with LNP delivery paper published by Intellia<sup>1</sup>
- Highly modified 100mer G211 sequence selected
- Highest editing of all guides tested
- Contains sgRNA chemistry typically found in non-virally delivered gene editing drugs
  - Ribose: 2'-OH, 2'-OMe
  - Backbone: PO, PS on ends

1. Finn et al, *Cell Reports*, 2018.

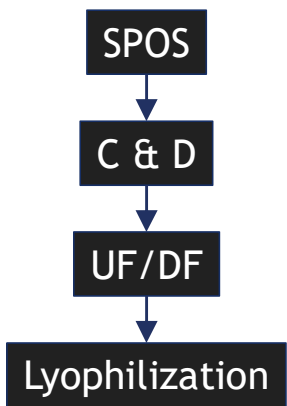




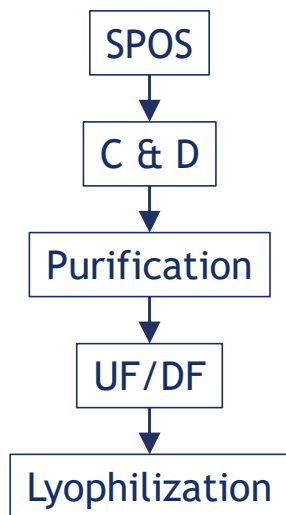
# Chemoenzymatic process flow for step 1 and step 2

Purified oligonucleotide fragments and crude splints

## DNA splints



## Oligo fragments

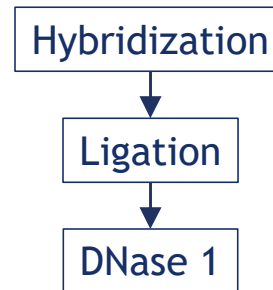


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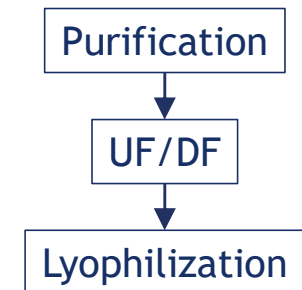


## Chemoenzymatic ligation

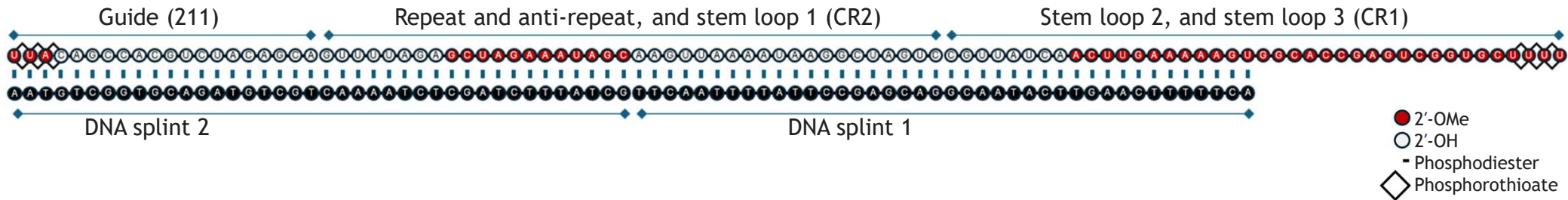
### Upstream



### Downstream



# Components for assembly of ultra-high purity sgRNA



Step	Sequence	Length	5'-P	Purif <sup>n</sup>	Purity <sup>1</sup>	Sequence information (5' to 3' fragments, 3' to 5' splints)
1	G211-1	9	N	HPLC	98.4%	UUA C A G C C A
	G211-2	11	Y	HPLC	98.9%	C G U C U A C A G C A
	G211 splint	20	-	UF/DF	82.6%	A A T G T C G G T G C A G A T G T C G T
2	G211	20	N	HPLC	98.1%	UUA C A G C C A G G U C U A C A G C A
	CR1	40	Y	HPLC	90.4%	C G U U A U C A A C U U G A A A A G U G G C A C C G A G U C G G U G C U U U U
	CR2	40	Y	HPLC	81.4%	G U U U U A G A G C U A G A A A U A G C A A G U U A A A A U A A G G C U A G U C
	DNA splint1	40	-	UF/DF	70.4%	T T C A A T T T T A T T C C G A G C A G G C A A T A C T T G A A C T T T T T C A
	DNA splint2	40	-	UF/DF	77.5%	A A T G T C G G T G C A G A T G T C G T C A A A A T C T C G A T C T T T A T C G

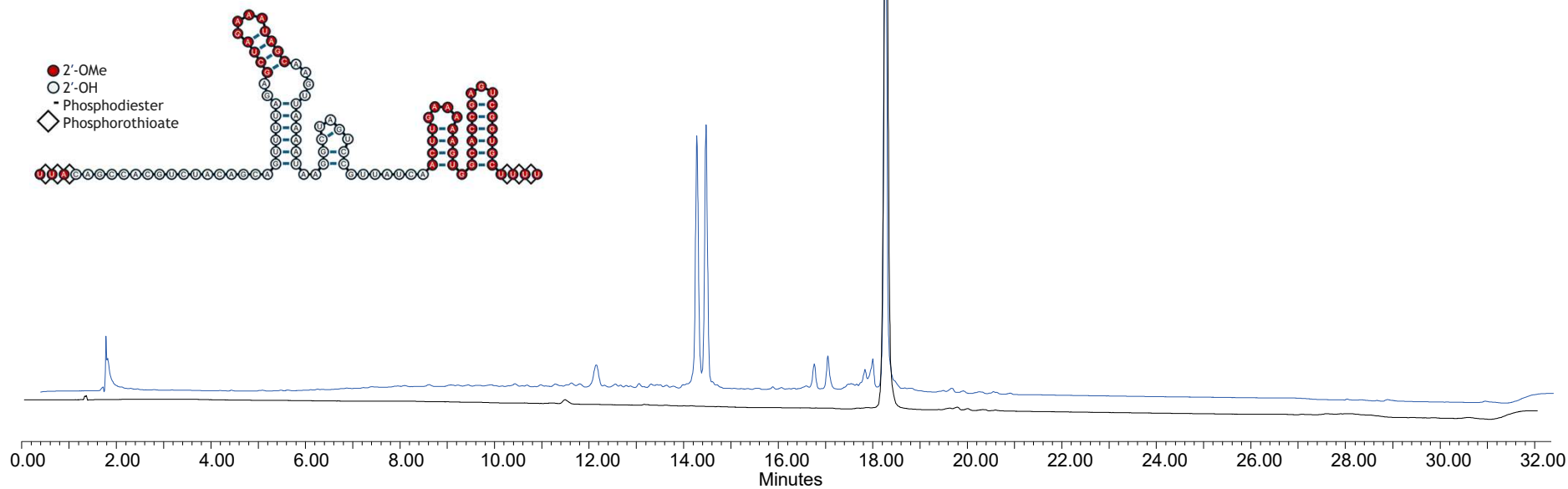
1. Purity measured by IPRP-HPLC/MS



# sgRNA before and after HPLC purification

## IPRP-HPLC/MS

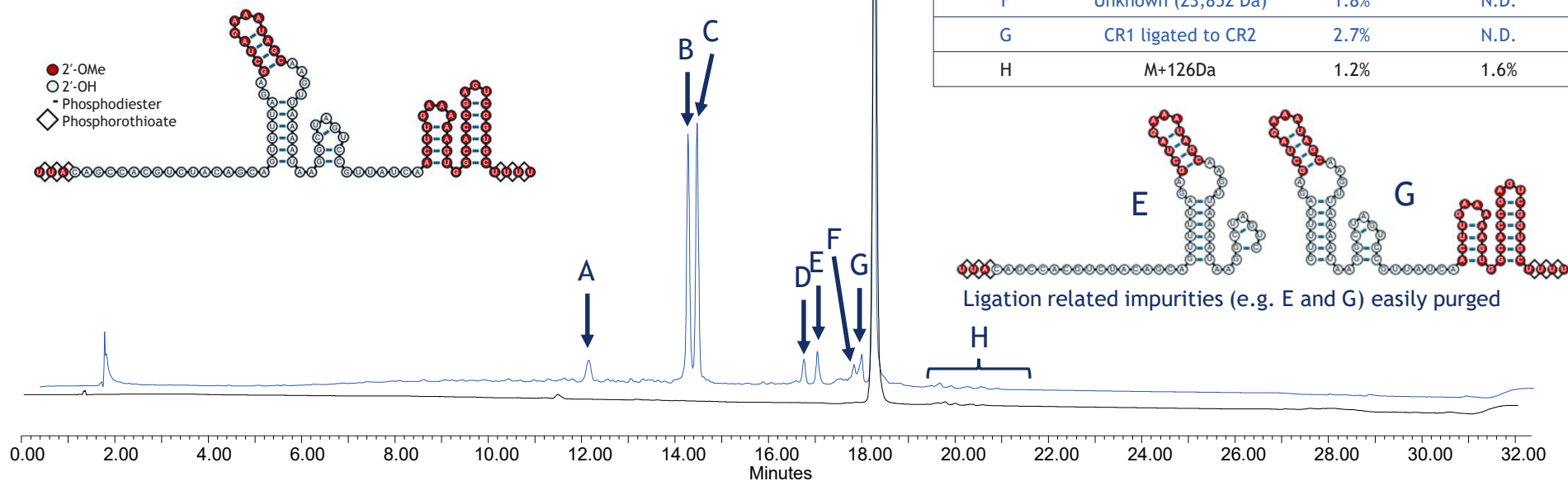
Crude after ligation, before DNase 1: 39.0%  
HPLC purified: 96.8%



# sgRNA before and after HPLC purification

## IPRP-HPLC/MS

Crude after ligation, before DNase 1: 39.0%  
HPLC purified: 96.8%



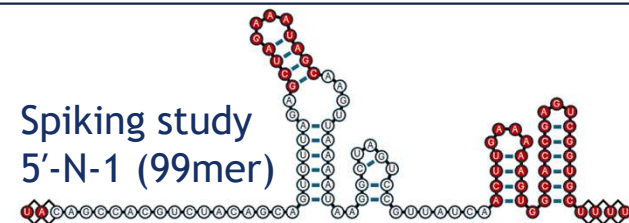
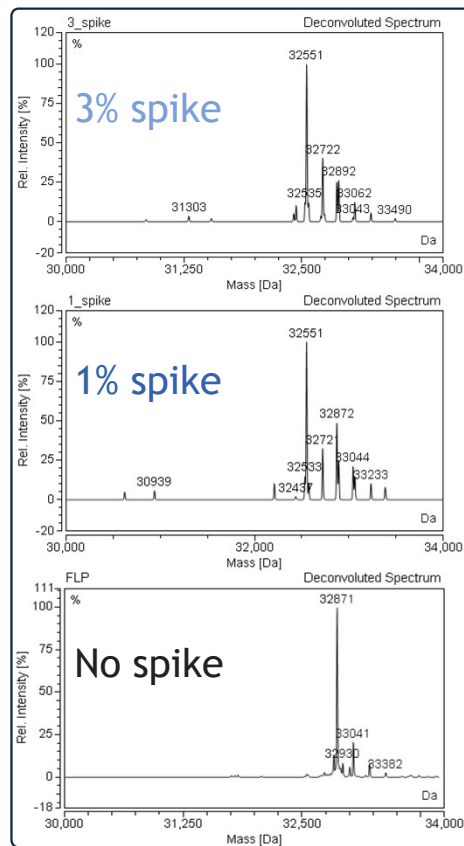
Impurity peak	Assignment by MS	Crude area %	Purified area %
A	G211	2.5%	N.D.
B	DNA splint 2	16.5%	N.D.
C	DNA splint 1	16.1%	N.D.
D	CR2	1.8%	N.D.
E	G211 ligated to CR2	2.2%	N.D.
F	Unknown (23,852 Da)	1.8%	N.D.
G	CR1 ligated to CR2	2.7%	N.D.
H	M+126Da	1.2%	1.6%



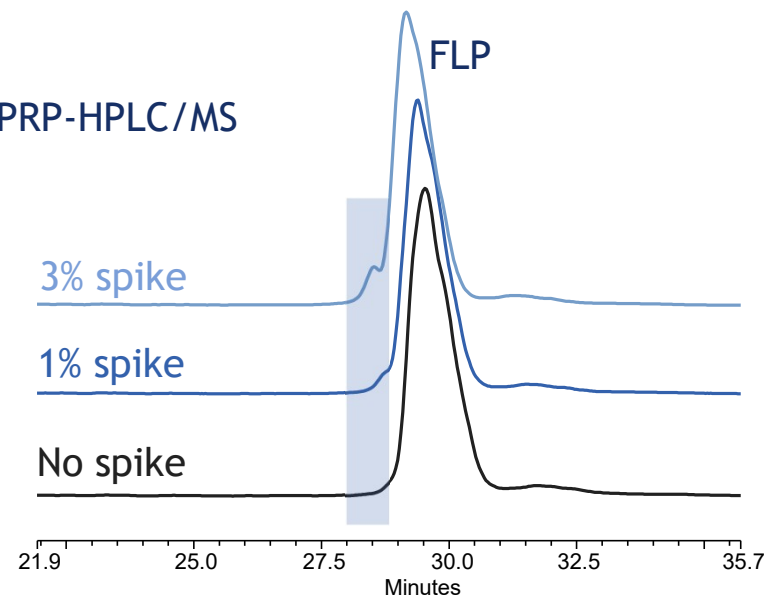
# sgRNA purity and impurity control

## Heavy reliance on HPLC and MS

- HPLC methods for separation of closely-related impurities (e.g. N-1nt)
- Required MS capabilities
  - Intact mass identification
  - Impurity profiling
    - 1Da resolution...?
  - Sequence confirmation



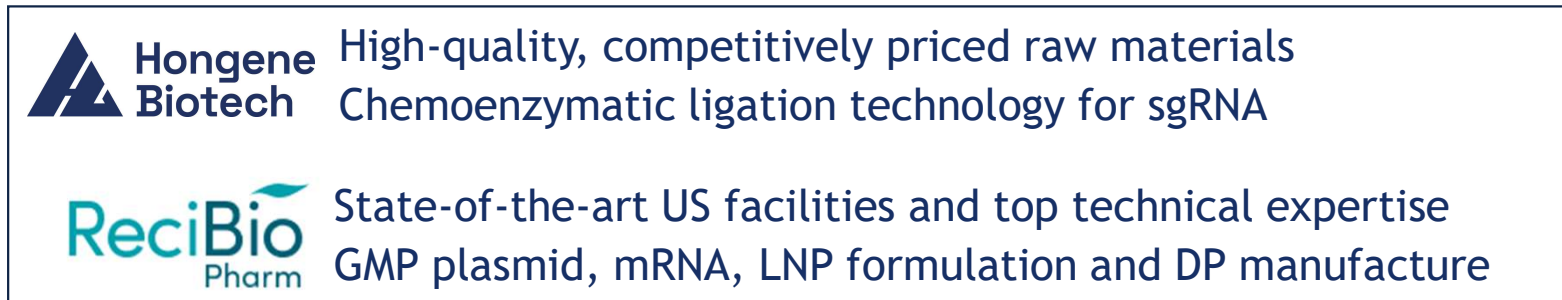
IPRP-HPLC/MS



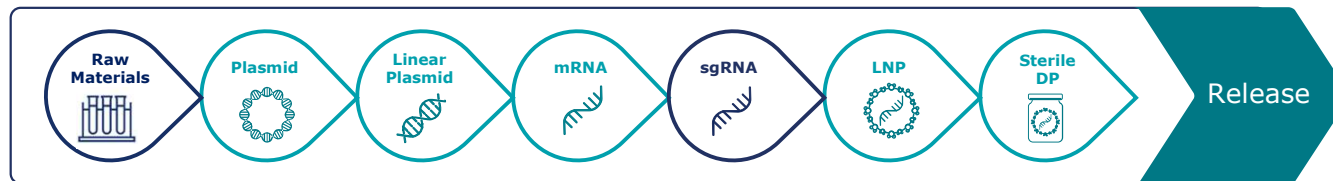
# Gene editing drug CDMO services under one roof in MA, USA

ReciBioPharm Hongene Biotech partnership

## The value proposition



One facility, one point of contact, one master services agreement, one quality agreement



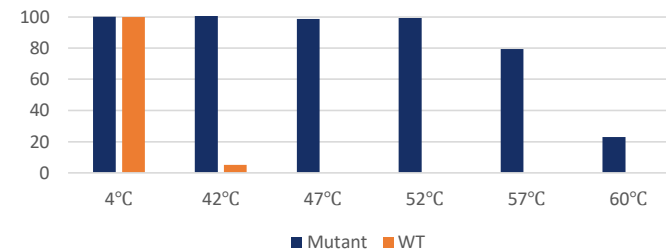
# Engineered thermostable T4 RNA ligase



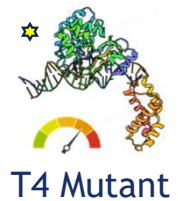
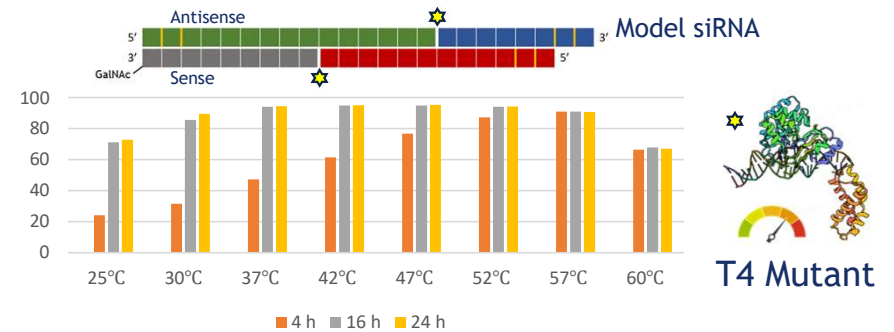
# Engineered thermostable T4 RNA ligase is highly active at 52°C

- Rationale for thermostable enzyme
  - Facilitates ligation at higher temperature
  - Reduced secondary structures
  - Improved selectivity for fragments with optimal hybridization
  - Mitigation of incorrectly ligated byproducts
  - Leverage C-to-C strategy further driving down cost

T4 mutant retains activity after incubation at 52°C



T4 mutant is maximally active between 52-57°C



Activity study: 20ul enzyme solution (0.1mg/mL in 50mM tris, 10mM Mg(Ac)<sub>2</sub>, 5mM ATP) was incubated at the indicated temperature for 8h. This solution was then added to 20ul volume of 2.5 mM each of four oligo fragments and ligation was allowed to proceed at 33°C for 2h. Ligation activity calculation: Percentage of siRNA product (Temp)/siRNA product (4°C).



# Control of stereochemistry



# Collaboration with Bruker: Control of stereochemistry using NMR



Hongene  
Biotech



## Structural Analysis by NMR:

- Nondestructive
- In formulation
- High-resolution (atomic)
- Sensitive to changes
- Selective

Booth #32  
Harbor Foyer



Current  
Phase

- NMR method development to distinguish stereochemistry of oligonucleotide fragments, single strands, & duplex

Phase 2

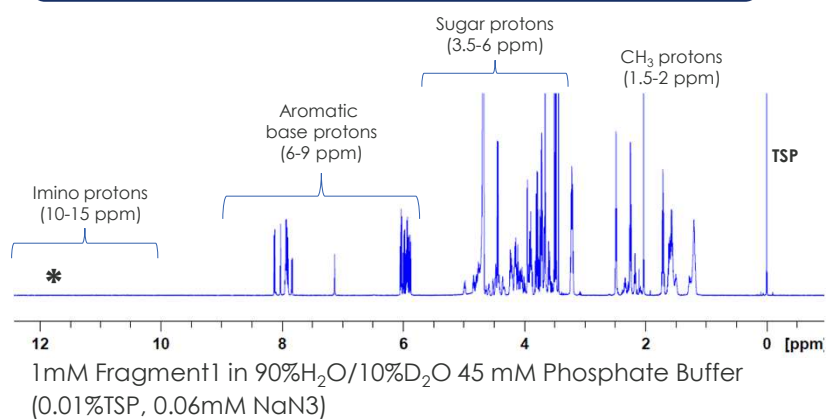
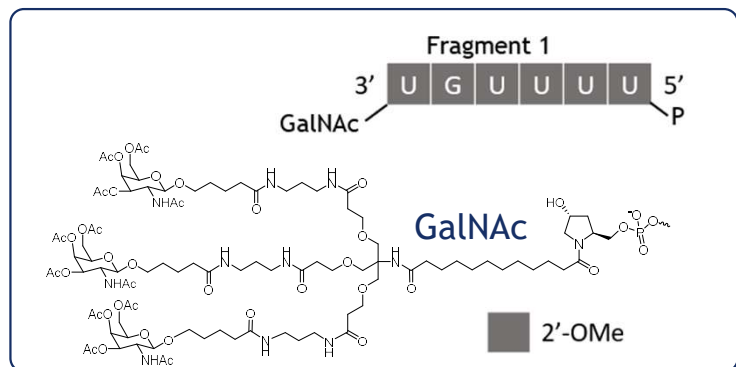
- Compare the effects of synthesis conditions (e.g. activators) on the stereochemistry by NMR

Phase 3

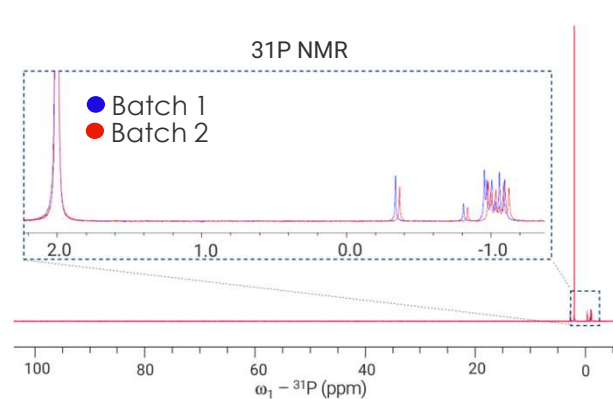
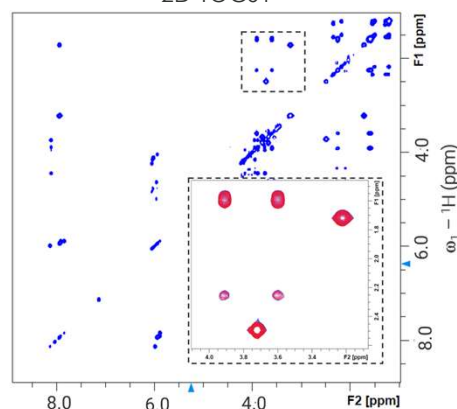
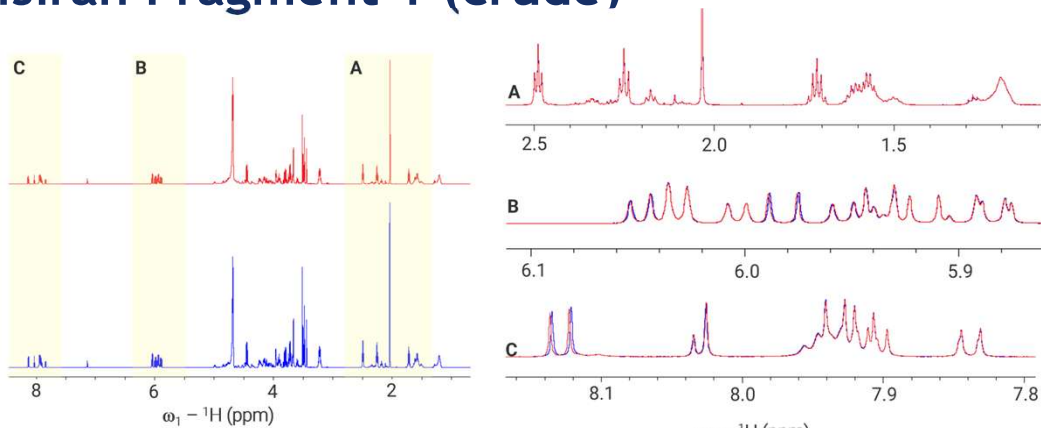
- Study stereochemical control of stereo-random oligos by NMR, compare with stereo-pure reference materials



# Batch to batch comparison of inclisiran Fragment 1 (crude)



\*imino protons not observed in ss oligo fragment



# Conclusions



## Conclusions

- Market demand for siRNA drugs is anticipated to grow significantly over the next decade
- This demand will be met by adoption of oligonucleotide fragment ligation technology
- Advantages are improved quality, yield, environmental sustainability and scalability
- We have manufactured 1 kg of siRNA GMP drug substance using ligation technology
- We envision no significant regulatory hurdles to wider adoption of the technology
- C-to-P and C-to-C purification strategies can be leveraged for large scale manufacturing
- Engineered T4 RNA ligases will provide future efficiencies and cost improvements
- Oligonucleotide fragments are manufactured today using scale-limited SPOS
  - Future demand will require scalable processes adapted to batch reactors
- High purity sgRNA for CRISPR based gene editing drugs is accessible by splinted ligation



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Thank you!

