

ASOs

(and a bit about other oligonucleotide based medicines)

April 27, 2023

Chris Hart, Ph.D.
Co-Founder / CEO
Creyon Bio, Inc.

Outline

- What are ASOs, how do they relate to other Oligonucleotide based medicines (OBMs)
- How do they work
- When ASOs (or other OBMs) may be appropriate for developing a treatment, or for basic research
- Overview of developing ASO medicines
- Questions

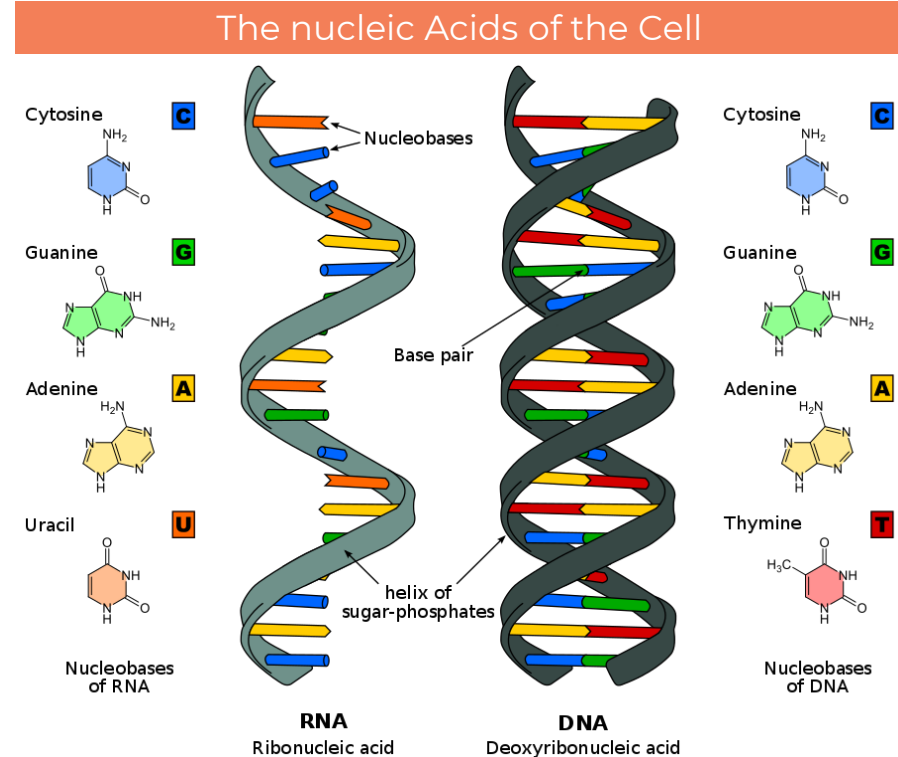
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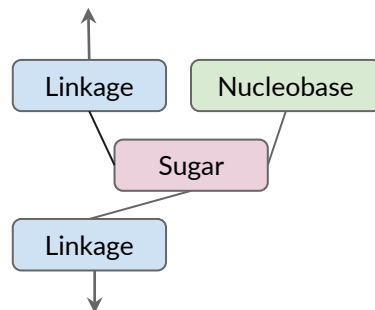
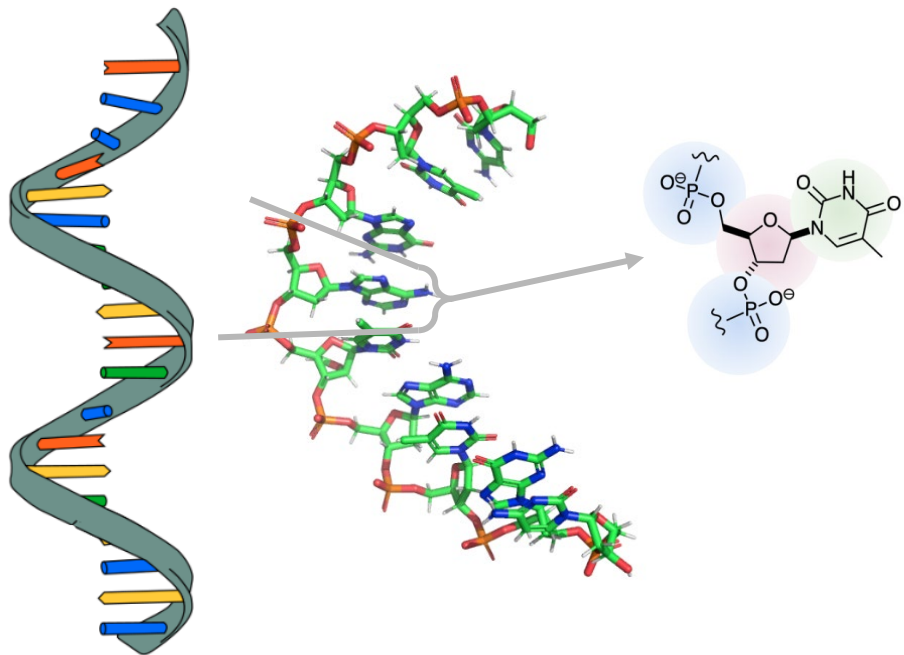
What are OBMs: Chemically synthesized, short, chemically modified, nucleotide polymers.

OBMs:

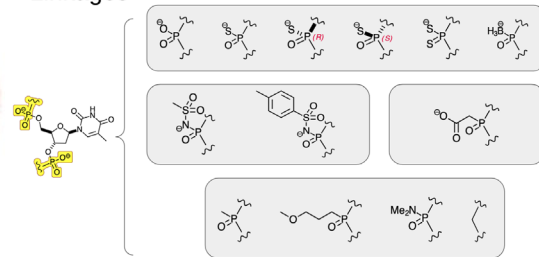
- Typically 15-22 nt, almost always <100 nt
- DNA/RNA-*like* molecules that are synthetically created (e.g., not created in cells or through enzymology)
- Chemically modified to impart better pharmacological properties (e.g., make them good drugs)



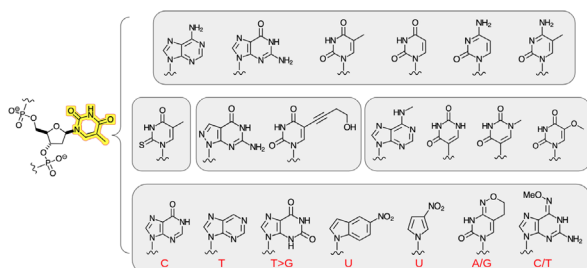
OBM design requires optimization of both sequence and chemistry



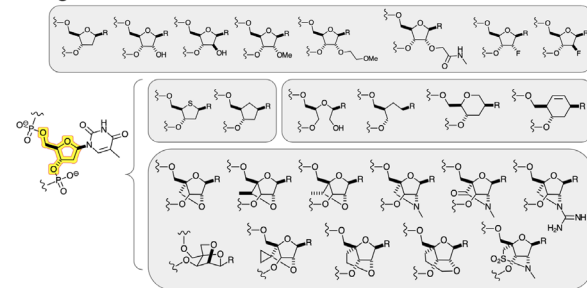
Linkages



Nucleobase

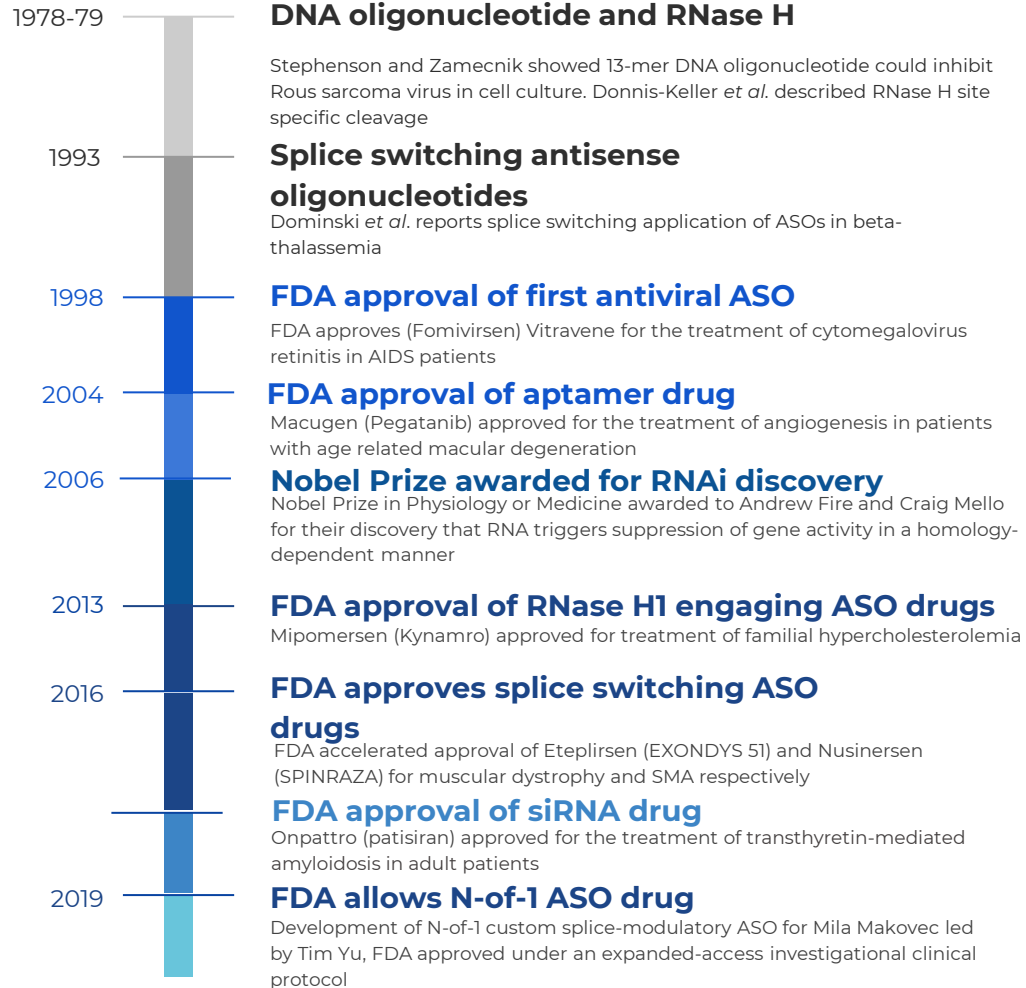


Sugar



Oligonucleotide Based Medicines (OBMs) have been researched for decades

- There is a rich understanding of the mechanism-of-action and molecular biology/pharmacology of OBMs
- Multiple types of OBMs have been approved - ASOs, siRNAs, aptamers
- Many pharma and biotech companies are actively advancing ASOs and siRNAs for rare and common diseases



Different types of OBM's use different chemistries tuned to optimize different molecular mechanisms; have different pharmacology

Antisense Oligonucleotides (ASOs)

- Single stranded (14-21 nt)
- Can be absorbed into cells with or without targeting
- Work through multiple molecular mechanisms:
 - Splice altering
 - RNAseH1 Knock Down
 - Other steric blocking

Short interfering Ribonucleic Acids (siRNAs)

- Double Stranded (20-25 nt)
- Requires delivery formulation or conjugation
- Works by loading into RISC and directing it to degrade specific mRNAs

Guides, Aptamers, and more

- Typically longer (30-80nt)
- Variable cellular uptake
- Work through multiple molecular mechanisms:
 - aptamers - Ab like binding
 - recruit endogenous enzymes (e.g., ADAR)
 - recruit exogenous enzymes (e.g., CRISPR)

Oligonucleotide Based Medicines (OBMs) can do many things

AGCCGCAAAG
TCGGCGTTTC

Precise Watson-Crick-Franklin
Hybridization Defined Target Recognition

*Control Gene
Expression*

*Guides for
Editing Enzymes*

*General Steric
Blocking*

siRNA (RISC KD)

ADAR

Viral Packaging

ASO (RNaseH1 KD)

CRISPR

Block RNA binding

Splice Modulation

...

Repeat expansion

uORF blockage

Influence RNA/DNA
structures

NMD control

...

...

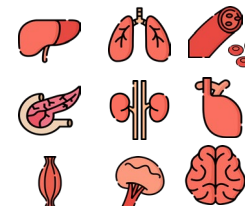
Favorable Pharmacology

Multiple routes of
administration



Active for weeks
to months

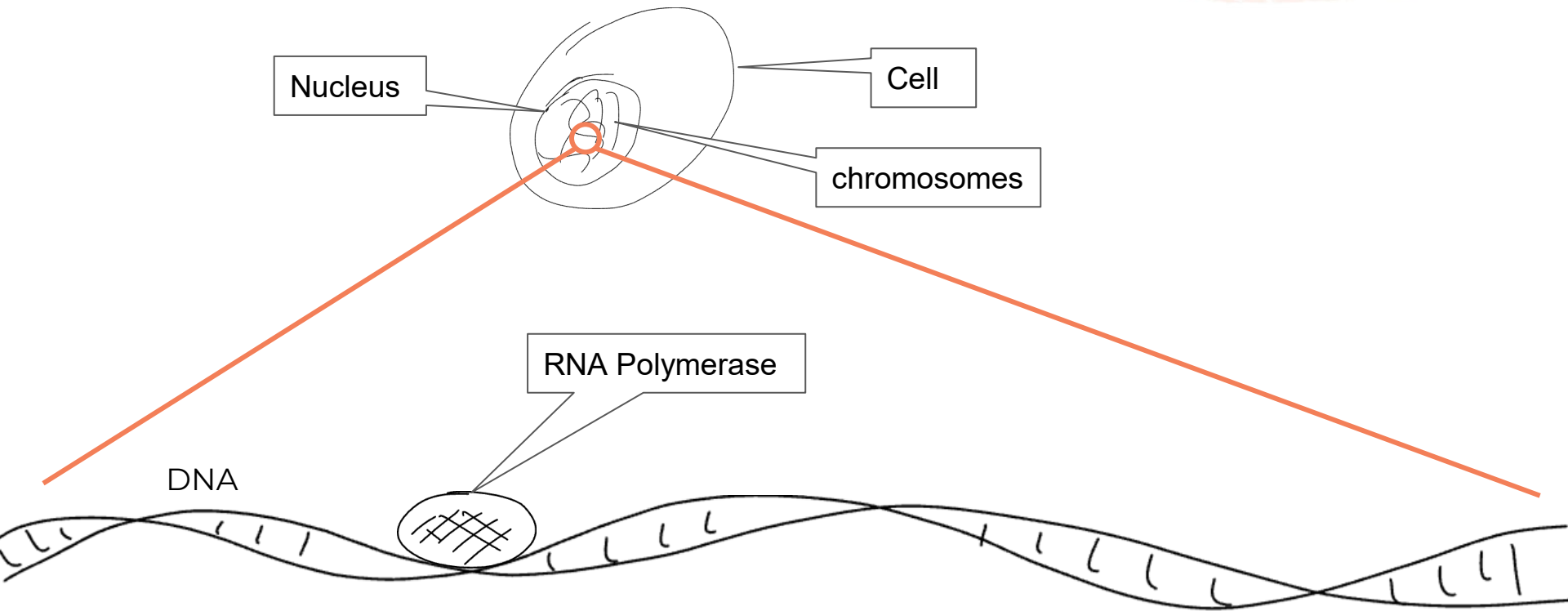
Broad
Tissue
Access



Outline

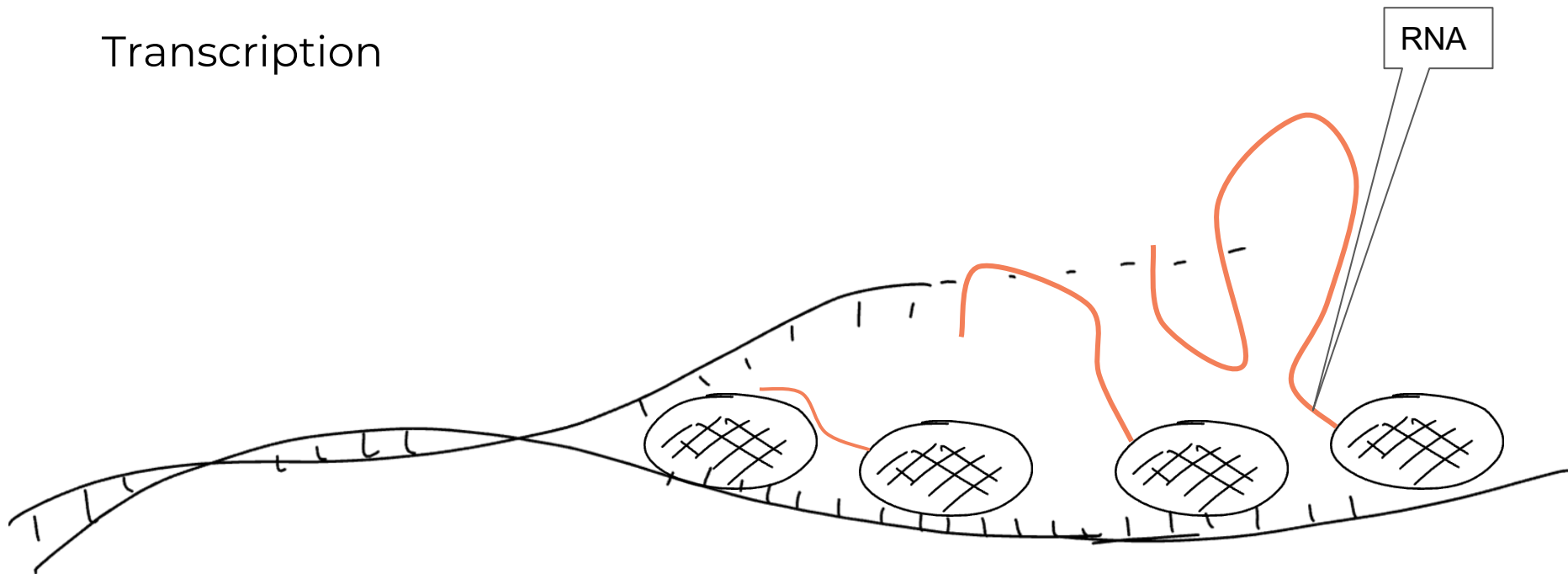
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ASOs (and other OBM) typically target the genetic basis of disease -- typically by modulating RNA



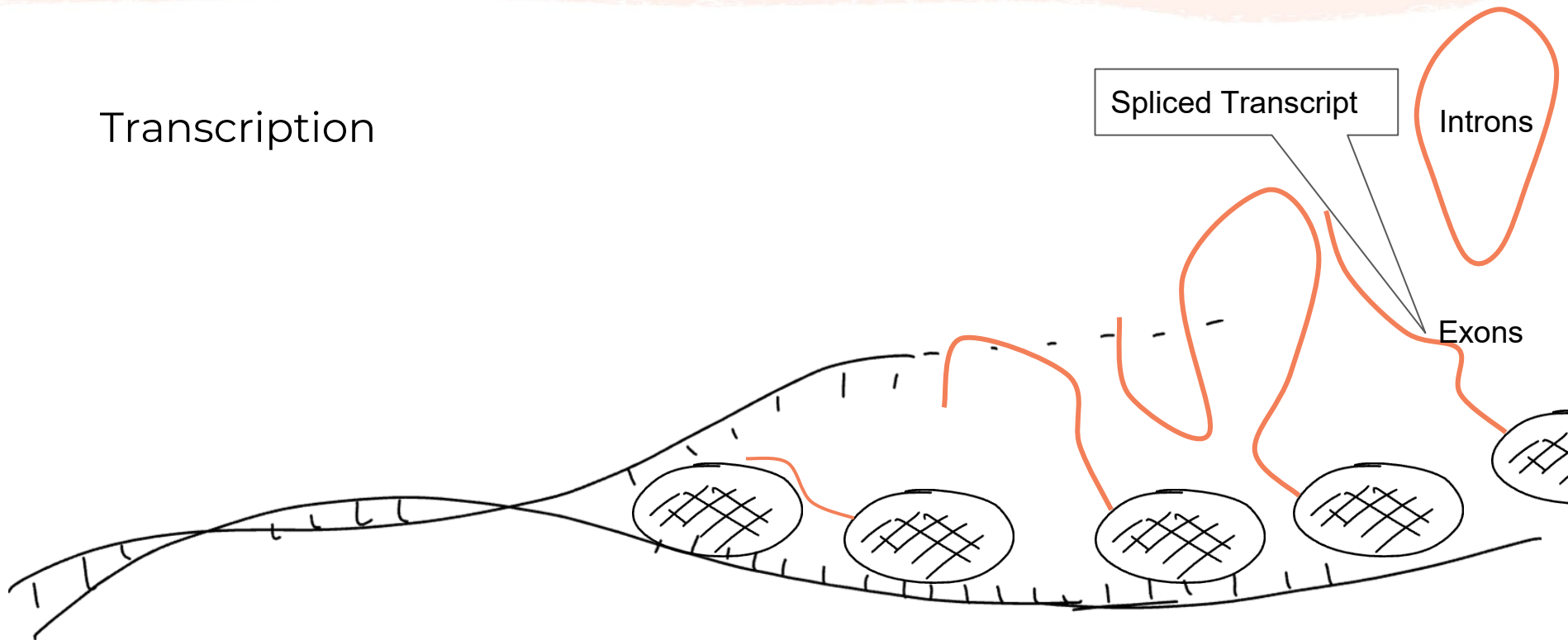
ASOs (and other OBM)s typically target the genetic basis of disease -- typically by modulating RNA

Transcription



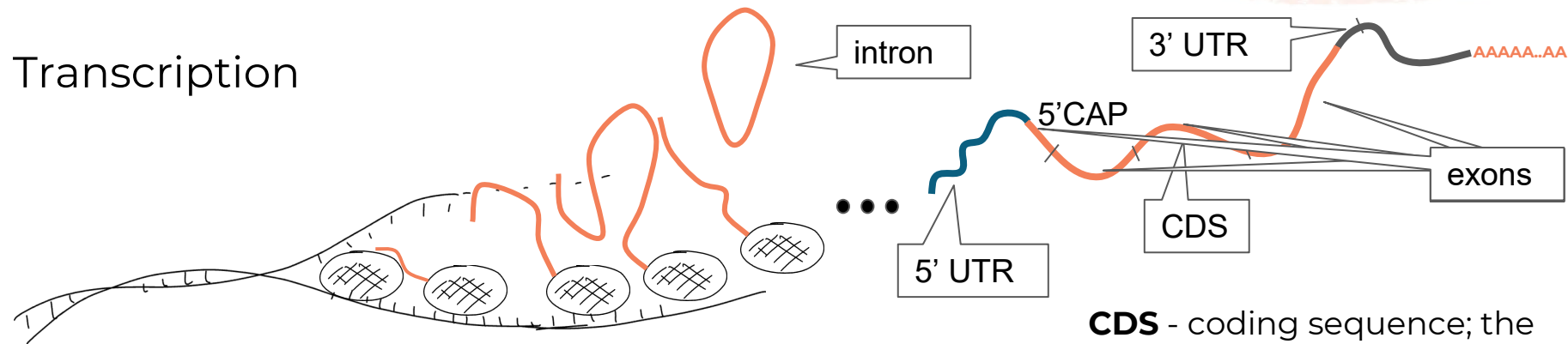
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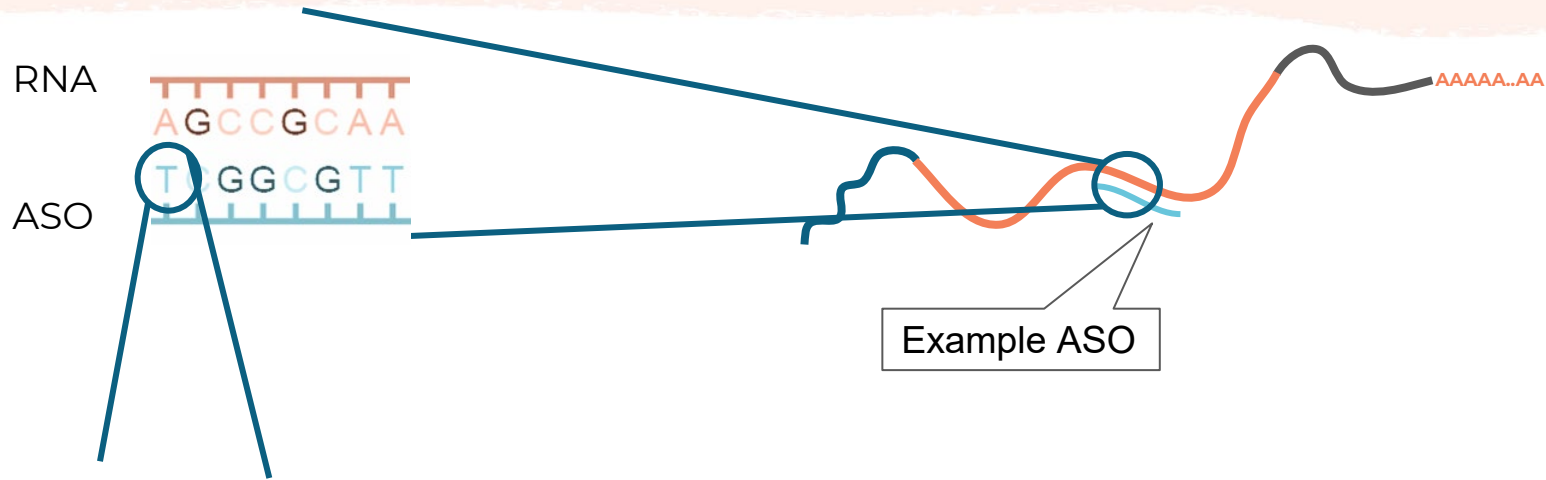
Transcription



CDS - coding sequence; the region of the RNA that codes for a protein

UTR - untranslated regions; regulatory domains of the RNA

OBNs target RNA through well defined Watson-Crick-Franklin hybridization



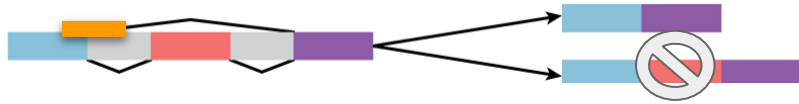
ASOs can target any RNA sequence

- Introns
- Exons
- 5' or 3' UTRs
- mRNA or pre-mRNA
- Protein coding or non-coding

Outline

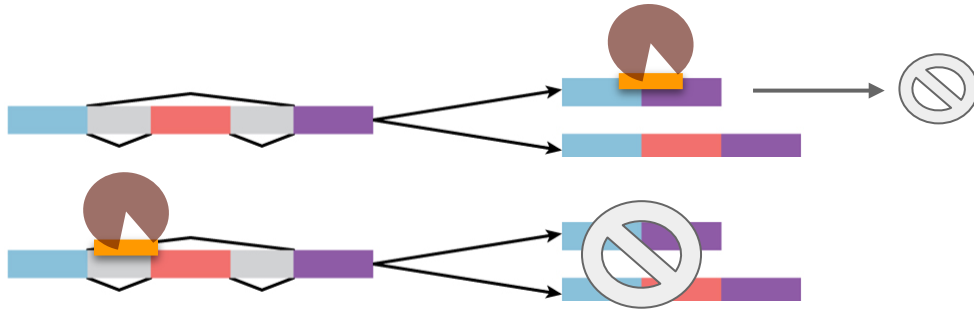
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Oligonucleotide based medicines (OBMs) modulate gene expression directly



Occupancy Only

e.g., splice modulating



Enzymatic RNA knockdown

e.g., ASO-directed RNase H1 KD

- Correct aberrant disease causing splicing
- Upregulate gene expression by skipping NMD-inducing transcripts
- Works on mRNA, pre-mRNA, ncRNA (e.g., the druggable and undruggable)
- Allele selectivity achievable allowing for correction of toxic GoF diseases
- Additional enzymatic systems can be engaged (e.g., RISC, ADAR)

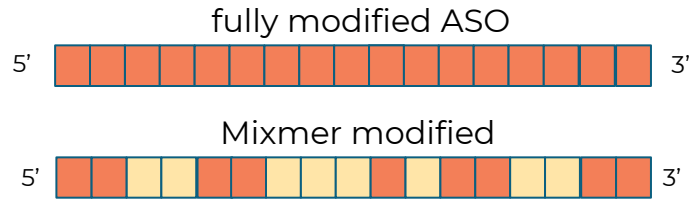
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Well established ASO architectures



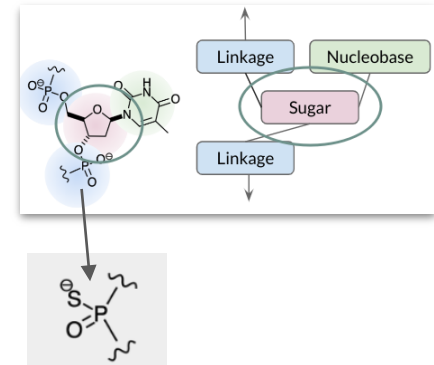
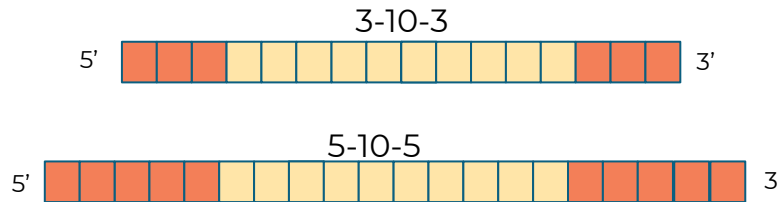
*Steric hindrance
(splice shifting, ...)*



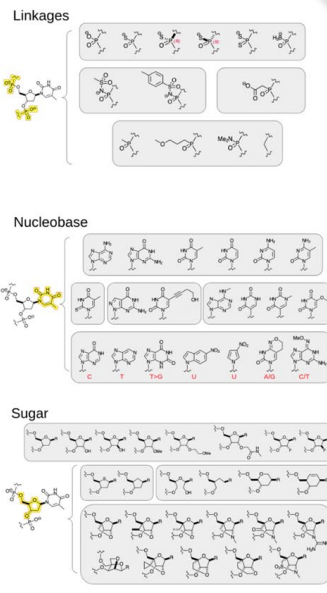
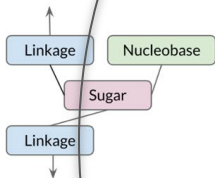
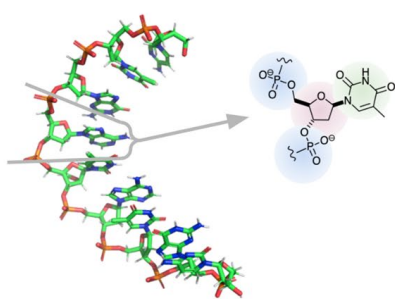
□ DNA

□ 2'OMe/2'O-MOE/2'F/LNA/PS

Gapmer modified ASOs
(RNaseH1 engaging)



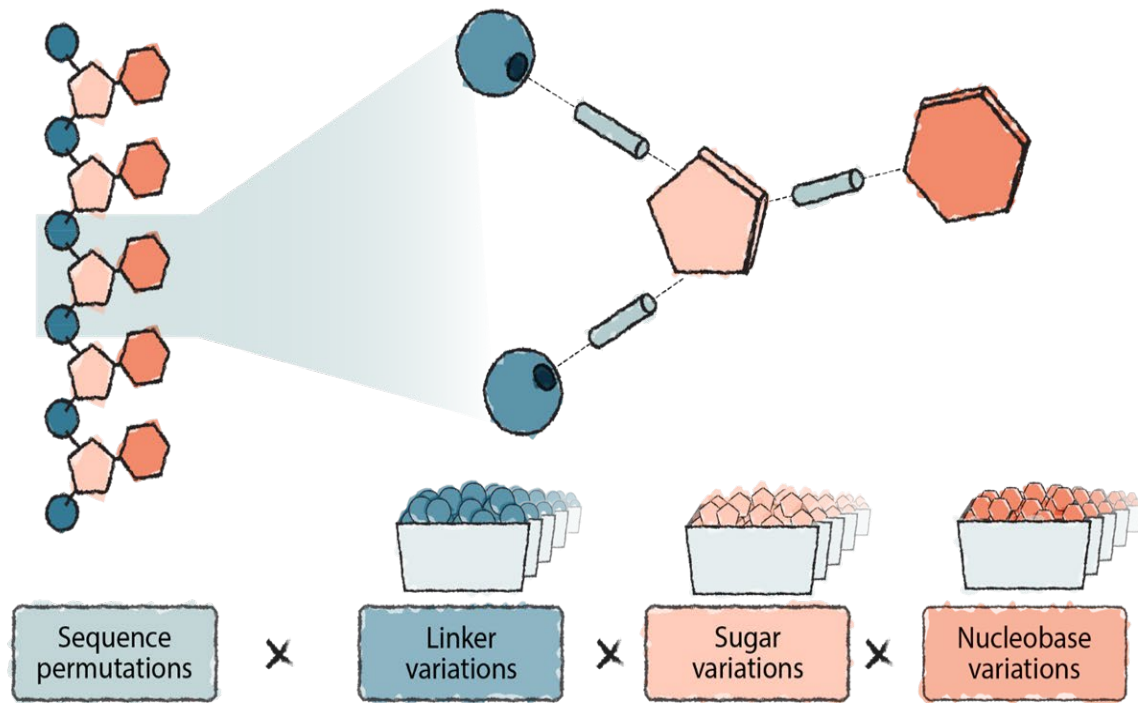
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Prepared for RARE Entrepreneur Bootcamp, Oct. 2022
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There is a growing number of interesting chemistries that can be used in ASOs and other OBMs.

Design Caveat



OBM's are challenged by exponential combinations

Sequence and Chemistry Collude...

Optimization is Hard

Potential toxicities induced by ASOs

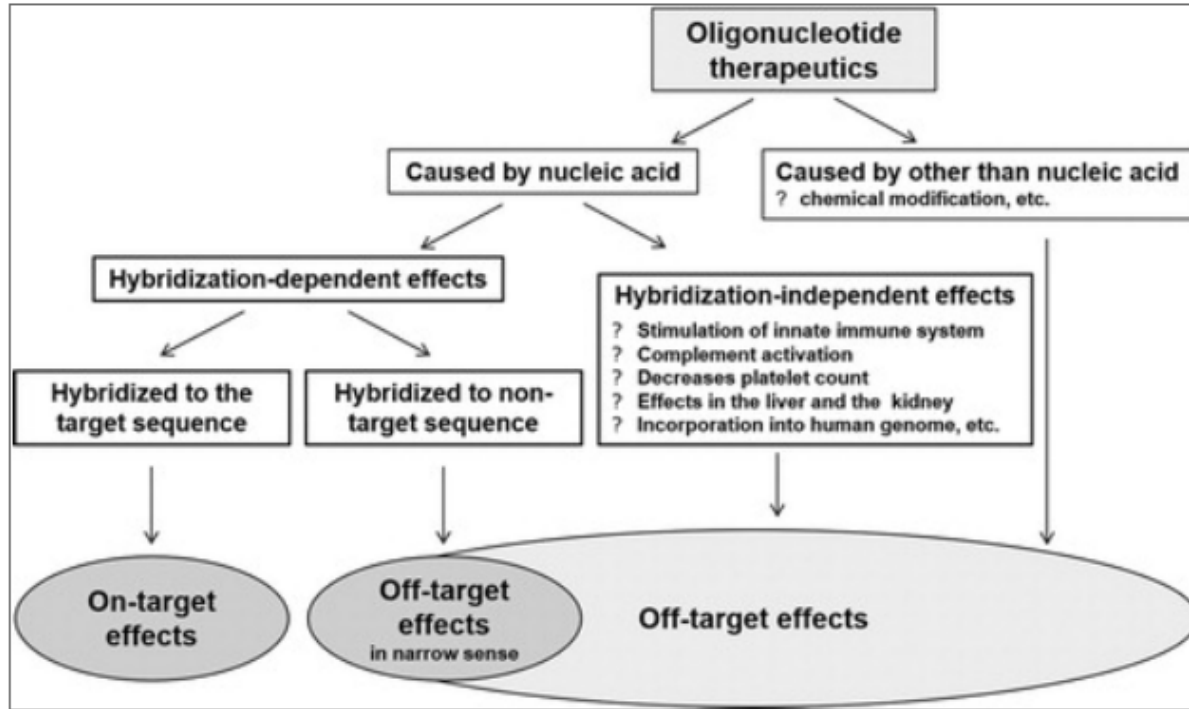


Image from: Hirabayashi et al. Considerations of the Japanese Research Working Group for the ICH S6 & Related Issues Regarding Nonclinical Safety Assessments of Oligonucleotide Therapeutics: Comparison with Those of Biopharmaceuticals. Nucleic Acid Therapeutics. Apr 2021.114-125.<http://doi.org/10.1089/nat.2020.0879>

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Other insights into on-target tolerability concerns

- Mouse or other model organism phenotypes
 - Caveats:
 - Humans ≠ Mouse ≠ fly ≠ worms ...
- Modern Biology Tools:
 - High-content imaging
 - Functional genomics
 - iPS derived cellular models (organoids, neurons, ...)

Assessing hybridization-dependent off-target effects

Search for all perfect matched and near-matched sequences in the relevant transcriptome.

- ASOs - active at all expressed RNA (e.g., pre-mRNAs, ncRNAs)

Caveats:

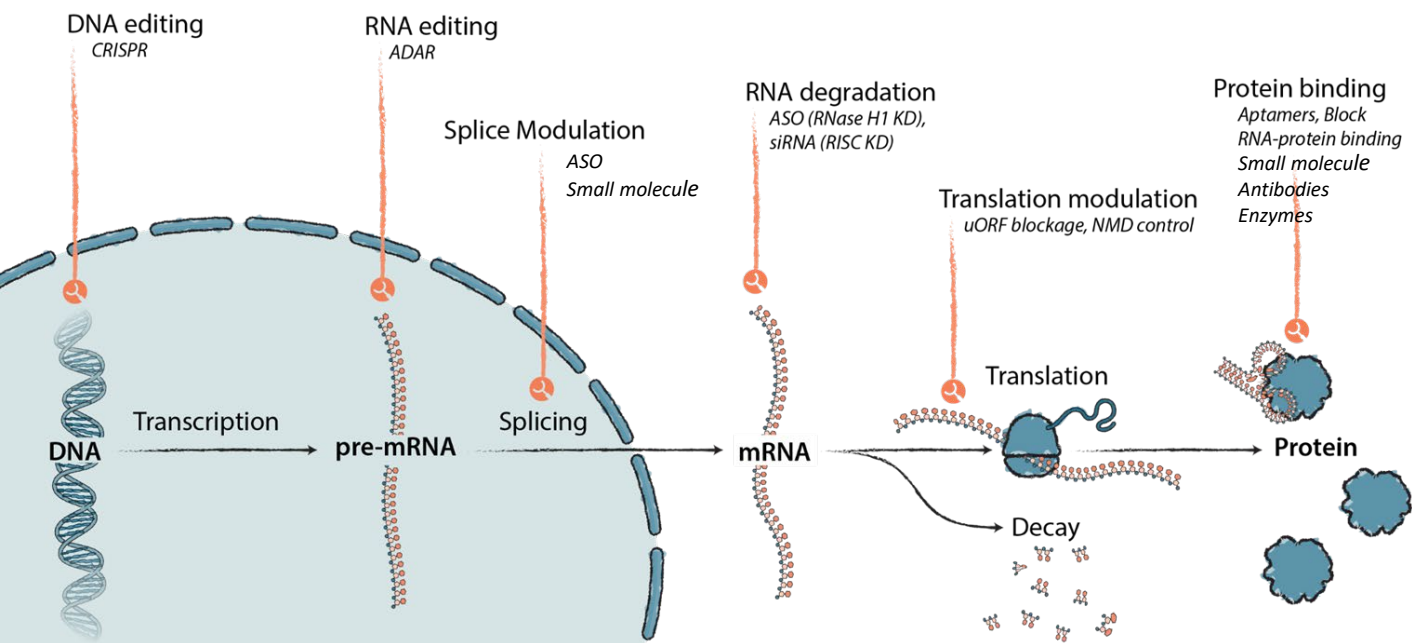
- Sequence analysis tools built for NGS analysis (e.g., bowtie, STAR) or evolutionary searches (BLAST, BLAT) have optimizations that may miss important hits



CREYON

Modalities Overview

Rare disease is often genetic ∴ interacting/correcting affected downstream processes, machines and structural elements of the cell offer possible treatments



Key consideration of different modalities is where to intervene to correct the genetic processes in the most efficient way

Different pathogenic variants within the same rare disease may be amenable to different modalities