Engineered Enzymes to Overcome Scalability and Sustainability Challenges of Nucleic Acid Therapeutics Manufacturing

Mathew Miller, PhD Associate Director, Life Sciences Technology and Applications

Stephanie Forget, PhD Senior Scientist I



Codexis core business pillars

Based on CodeEvolver[®] platform to accelerate enzyme discovery and commercialization



Enzymes from Nature Commercially Relevant Enzyme **Pharmaceutical Manufacturing** enzymes for small molecule production

Value Creating Products Life Sciences enzymes for NGS applications and DNA/RNA synthesis

Biotherapeutics

enzymes as oral drugs; engineered transgenes and capsids for gene therapy



About the Speaker



Mathew Miller, Ph.D.

Associate Director of Life Science Technology and Applications, Codexis

Mathew joined Codexis in 2009. He is currently focused on enzymes used in mRNA manufacturing, including directed evolution for T7 RNA polymerase. Matt has previously lead protein engineering projects for next generation sequencing and enzymatic DNA synthesis. He received his Ph.D. from the University of California, San Francisco.



mRNA is a highly versatile therapeutic modality with manufacturing challenges



Applications

CODEXIS

- Vaccines (SARS-CoV2, others)
- Cancer vaccines (Immuno-oncology)
- Gene editing (CRISPR/Cas, others)
- Therapeutic proteins

mRNA manufacturing challenges

- mRNA capping efficiency
- Undesired side products (dsRNA)
- mRNA yield
- Downstream processing

Eukaryotic mRNA cap: structure and function

7-Methylguanosine (7mG) Cap:



CODEXIS®

Vicens, Q., Kieft, J. S. & Rissland, O. S. Revisiting the Closed-Loop Model and the Nature of mRNA 5'-3' Communication. *Mol Cell* **72**, 805–812 (2018).

Co-transcriptional capping

Dinucleotide cap analog:



Cap analog is the most expensive reaction component

Enzyme Engineering targets:

Increased capping efficiency



• Reduce cost of the process



- Increase mRNA safety and efficacy
- Reduce cost of the process



CodeEvolver[®] technology rapidly generates enzymes with desired properties through iterative rounds of enzyme optimization



CODEXIS®

Co-transcriptional capping: ARCA cap analog



• Inefficiently incorporated by WT T7RNAP



Codex[®] HiCap RNA Polymerase

Increased dinucleotide analog capping efficiency (ARCA)



- Allows higher capping efficiency for a given concentration of ARCA, GTP
- Allows for >95% capping efficiency in batch IVTs without GTP starvation, preserving yield and lowering cost

CODEXIS®

Co-transcriptional capping: CleanCap® AG cap analog



- Higher incorporation by WT T7RNAP
- Native Cap-1 structure
- More expensive to produce and use

CODEXIS

Codex[®] HiCap RNA Polymerase

CODEXIS®

Increased trinucleotide (CleanCap® AG) capping efficiency

Codex[®] HiCap RNA Polymerase selectively incorporates CleanCap[®] AG



Allows for higher capping efficiency or equivalent efficiency with lower cap concentrations (1.5mM), reducing cost

Capping efficiency

Double-stranded RNA side products



- dsRNA is generated by WT T7 via an RNA-dependent RNA polymerase activity
- Pattern recognition receptors (TLR3, RIG-I) recognize dsRNA and induce interferon signaling, causing inflammation
- dsRNA signaling inhibits protein translation
- dsRNA contamination impacts safety and efficacy of mRNA therapeutics
- Polishing step required to remove dsRNA adds complexity and cost

CODEXIS

Codex[®] HiCap RNA Polymerase

Reduced dsRNA formation



- 3' extensions for the HiCap RNA Polymerase sample are significantly reduced vs the WT sample
- Residual dsRNA from HiCap RNA Polymerase without purification is similar to that from purified WT polymerase sample
- dsRNA sandwich ELISA shows ~50-fold reduction in total dsRNA side products
- Lower dsRNA eliminates the need for polishing simplified workflow and lower cost

CODEXIS®

Reduced dsRNA formation correlates with reduced immune stimulation and increased protein translation



GFP reporter translation post-transfection

- Codex[®] HiCap RNA Polymerase shows significant reduction in IFN signaling compared to WT T7
- Codex[®] HiCap RNA Polymerase shows significant increase in protein translation in HeLa cells
- Lower dsRNA improves safety and efficacy, using a simplified workflow

CODEXIS

Codex[®] HiCap RNA Polymerase attributes

	Codex [®] HiCap RNA Polymerase	Wild-type T7 polymerase
Reduced dsRNA production	• • •	•
Co-transcriptional capping efficiency	• • •	••
IVT yields	• • •	• • •
Reduced capping reagent	• • •	•
Manufacturing simplification	• • •	•
Finished product yield	• • •	•

Information previously presented by Tyler Goodwin, Ph.D., Head of Non-viral Delivery Development and Vector Core, Tune Therapeutics

Higher quality mRNA and improved manufacturing cost & efficiency



Engineered Enzymes to Overcome Scalability and Sustainability Challenges of Nucleic Acid Therapeutics Manufacturing

Mathew Miller, PhD Associate Director, Life Sciences Technology and Applications

Stephanie Forget, PhD Senior Scientist I



RNAi: High Demand / Constrained Supply

Demand Drivers



RNA delivering on promise of personalized medicines (10+ FDA approved therapies have reached market in past 5yrs)

RNAi therapeutics as a modality is growing rapidly with >450 assets in pipelines

RNAi therapies are treating the previously untreatable diseases

Production & Supply Challenges



Chemical RNA synthesis produces **Millions** of liters of chemical waste... and growing!

Phosphoramidite synthesis will be challenged to meet demand of 1,000s kg of RNAi p.a. by 2030

Critical solvent supply constraints likely (Acetonitrile)



Enzymes are poised to spur innovation and disrupt RNAi manufacturing

Scalability



Purity & Yield



Sustainability



Enzyme catalytic activity enables multiple cycle use Evolution targets >99% incorporation efficiency

Aqueous waste streams & lower solvent use



About the Speaker



Stephanie Forget, PhD Senior Scientist I, Codexis

Technical lead for enzymatic RNA synthesis program. Obtained PhD in Chemistry at Dalhousie University, Canada. Studied biosynthetic enzymes involved in rare sugars found in natural products. Postdoctoral fellow at University of British Colombia, Canada. Studied oxidative enzymes through reaction profiling.









ECO Synthesis[™] Technology

(Enzyme-Catalyzed Oligonucleotide Synthesis)







Final process in development

CODEXIS

ECO Synthesis[™] Technology

- Controlled addition of modified RNA bases (TdT)
- Deblocking of 3'blocking group (phosphatase)
- Supply of 3'blocked NTP substrates (multiple enzymes)



Enzyme Performance

- High incorporation efficiency (>99%)
- No sequence bias

At-Scale Process Requirements

- Controlled addition of monomers
- Low impurity production
- High volumetric productivity

- High manufacturing yields
- Scalable supply of nucleotide triphosphates





Enzyme Performance

- High incorporation efficiency
- No sequence bias

At-Scale Process Requirements

- Controlled addition of monomers
- Low impurity production
- High volumetric productivity

- High manufacturing yields
- Scalable supply of nucleotide triphosphates





Enzyme Performance

- High incorporation efficiency
- No sequence bias

At-Scale Process Requirements

- Controlled addition of monomers
- Low impurity production
- High volumetric productivity

- High manufacturing yields
- Scalable supply of nucleotide triphosphates





Enzyme Performance

- High incorporation efficiency
- No sequence bias

At-Scale Process Requirements

- Controlled addition of monomers
- Low impurity production
- High volumetric productivity

- High manufacturing yields
- Scalable supply of nucleotide triphosphates



Enzyme Performance: A highly engineered TdT

Key Enzyme Challenges

- Does not naturally recognize modified RNA
- Poor soluble expression and stability
- Enzymes function close to physiological conditions
- Manufacturability for large scale enzyme needs





Enzyme Performance: A highly engineered TdT

% Incorporation efficiency of N+1 additions over multiple rounds of evolution

3'-Terminus		Starting	Iterative Rounds of Evolution							
Sequence	NTP	TdT -								
ft m mAmCmU m *	fATP-3P	0	0	0	0	1	54	91	94	
	fUTP-3P	0	0	0	0	1	56	93	94	
	mCTP-3P	0	0	0	0	1	54	91	89	
	mGTP-3P	0	0	0	0	1	24	87	78	
	mATP-3P	0	0	0	0	1	39	88	66	
	*mGTP-3P	0	0	0	0	0	0	2	55	
	*mUTP-3P	0	0	0	0	0	0	19	24	
	fUTP-3P	0	0	0	0	2	33	74	92	
mGmAmC	mATP-3P	0	0	0	0	1	16	60	90	
	fATP-3P	0	0	0	1	2	34	77	88	
	mCTP-3P	0	0	0	0	0	13	66	86	
	*mUTP-3P	0	0	0	0	0	0	11	79	
	*mGTP-3P	0	0	0	0	0	0	3	64	
mGTP-	mGTP-3P	0	0	0	0	0	0	2	55	
AT*mG		4	1	4	49	75	82	47	56	
AmU*mG		0	0	0	12	46	77	75	41	
mAmU*mG	mATP-3P	0	0	0	0	5	69	82	68	
mAmUfG		0	0	0	0	0	2	1	66	
mUmGmA		0	0	0	1	2	38	82	86	
mAfUCmC	mATP-3P	0	0	0	0	4	58	88	86	
mAmG(MOE)C		0	0	0	0	0	4	14	84	
mC*mG*mA		0	0	0	0	2	39	75	75	
mCmUmG		0	0	0	0	4	82	86	72	
mAmUmC		0	0	0	0	2	57	84	63	
mAmUfU		0	0	0	0	0	0	0	59	
*mA*mA*mC		0	0	0	0	1	30	54	57	
mAmUfC		0	0	0	0	0	1	1	51	
mCmGmA	fATP-3P	0	0	0	1	4	82	93	92	
*fAfGmA		4	2	4	19	50	79	85	85	
mC*mG*mA		0	0	0	1	3	47	81	82	
*fA*fAfG		0	0	0	0	0	44	56	70	
fCfGfA		0	0	0	3	26	50	55	65	
mU*fA*fA		0	0	0	0	3	11	17	34	
fGmAfU		6	5	6	3	12	42	52	30	
fC*fG*fA		0	0	0	0	4	14	30	14	

So far...

- Incorporation of relevant 2'-modifications with 3' phosphate blocking group
 - 2'-deoxyfluoro and 2'-methoxy
- Incorporation of α-PS bonds
- Recognition of modified initiator sequences
- Improving incorporation efficiency of each nucleotide (goal >99%)

Note: All incorporating nucleotides contain 3'-blocking group; "*" denotes alpha PS-bond; "m, r, f" denotes 2'-OMe, 2'-OH, or 2'-F modifications, respectively

ECO Synthesis[™] Technology: Controlled addition of monomers



Target Sequence:

5' – Biotin/FAM...mA mU mA mA mG mA fA – 3'

✓ 4 Cycles for proof of concept

 Feasibility demonstrated for modified RNA synthesis



ECO Synthesis[™] Technology: Controlled addition of monomers



Demonstrated...

✓ Nucleoside → NTP conversion → 3'Blocked-NTP



"Two-step-one-pot" synthesis of 3'-blocked nucleotides

Provides scalable, sustainable, economic supply of required ECO Synthesis[™] monomers



Accomplished to date

- Progress on critical TdT performance
- Proof of Concept for iterative nucleotide addition
- Concept for enzymatically-derived source of 3'-blocked nucleotides

Next Steps

- Increase % monomer incorporation to reduce impurities
- Sustainable & scalable supply of nucleotides
- Scale-up to process-relevant conditions
- Achieve gram-scale synthesis of modified RNA



Reimagine enzymes for mRNA manufacturing and RNAi synthesis



