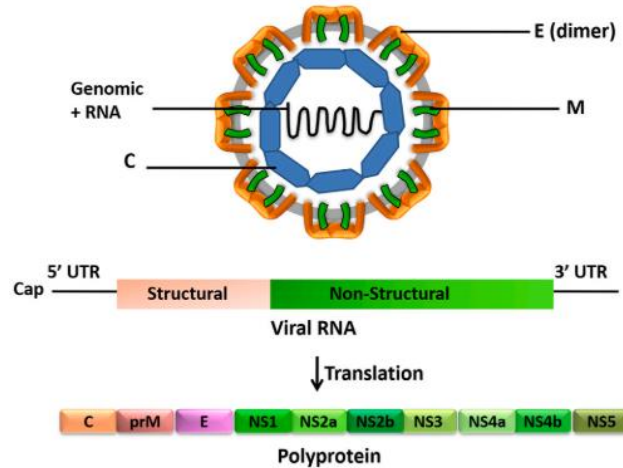


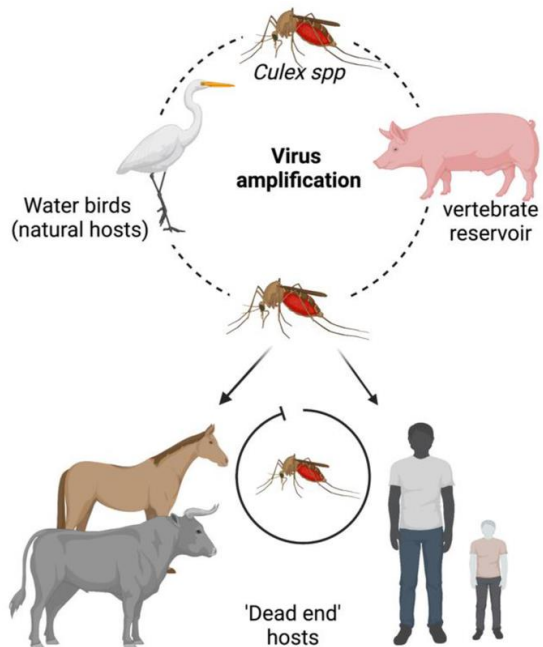
# **Development of novel mRNA vaccine against Japanese encephalitis virus (JEV)**

## Japanese encephalitis virus



### Schematic representation of the Japanese encephalitis virus genome

<https://doi.org/10.1016/j.mam.2021.100994>



### Cycle of JEV infection and amplification

<https://doi.org/10.3390/vaccines11040742>

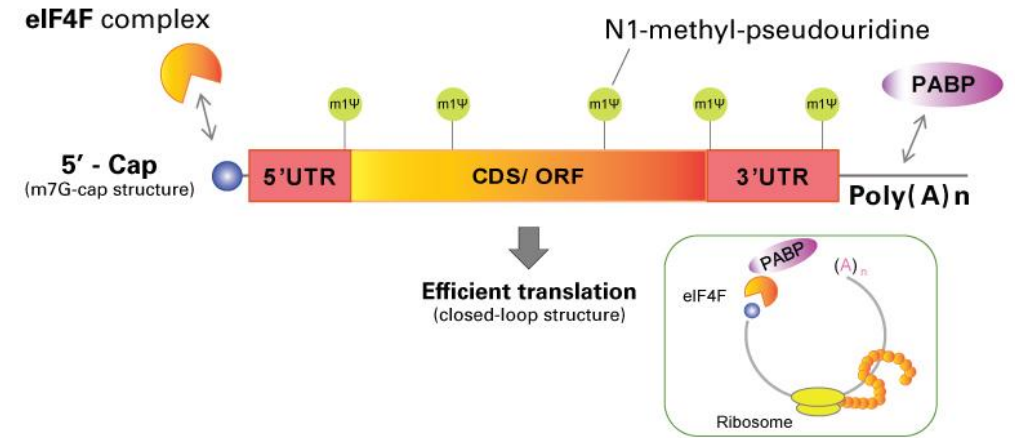
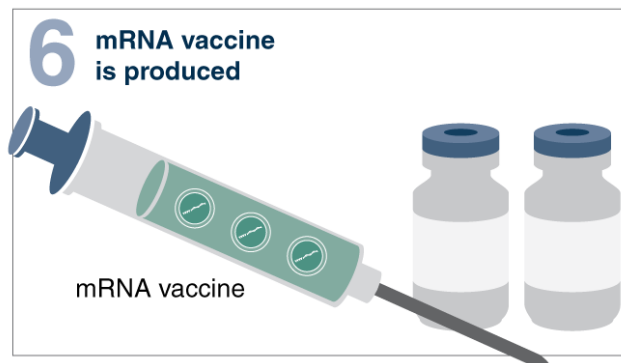
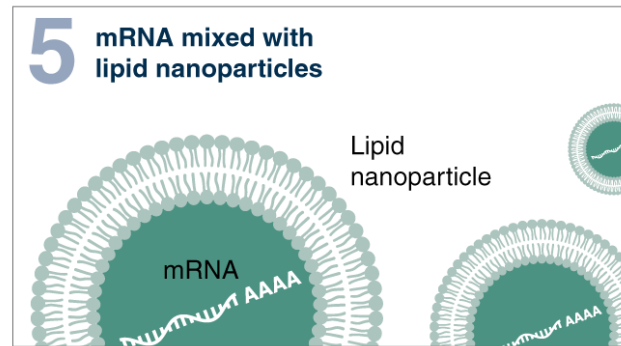
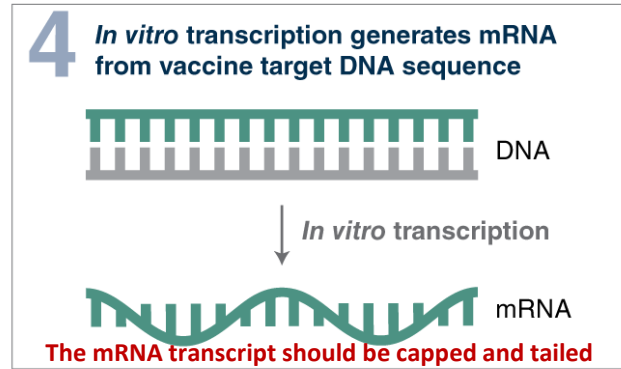
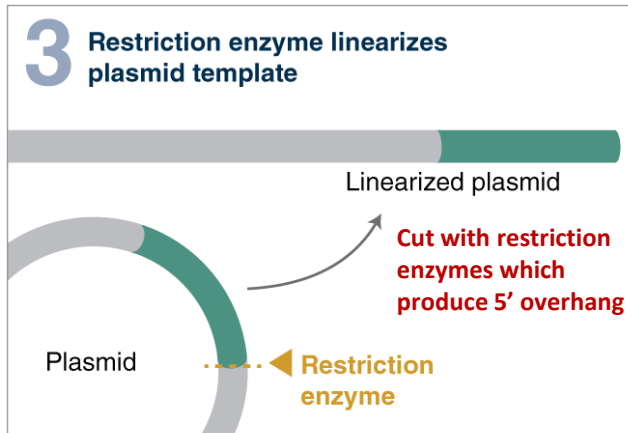
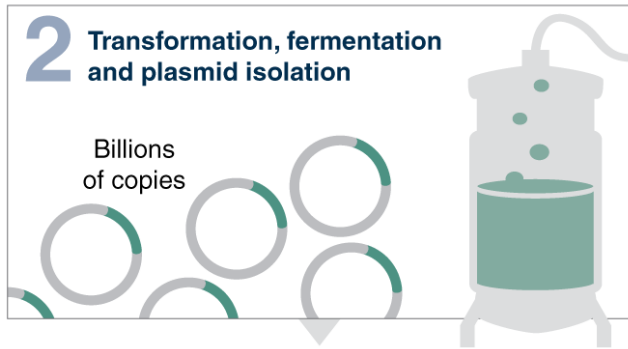
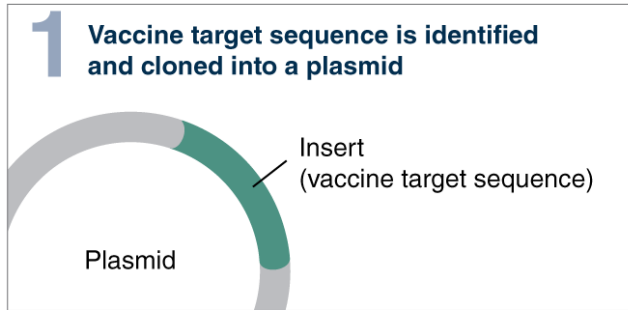
### Global geographical distribution of Japanese encephalitis

<http://wwwnc.cdc.gov/travel/yellowbook/2014/chapter-3-infectious-diseases-related-to-travel/japanese-encephalitis>

<https://doi.org/10.1007/s40121-013-0018-2>

Japanese encephalitis virus in India			
JEV Genotype	Strain name	Isolation	Accession Number; Size (bp)
I	JEV K94P05	Korea	AF045551; 10963
III	JEV P20778	Vellore	AF080251; 10977
V	JEV Muar	Malaysia	HM596272; 10988

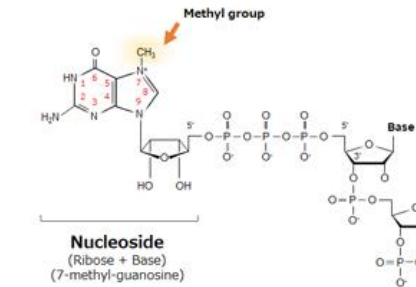
# mRNA vaccine generation



**In vitro-transcribed mRNA structure.** eIF4F = Eukaryotic initiation factor 4F, 5' UTR = 5' untranslated region, CDS/ORF = coding sequence/open reading frame, 3' UTR = 3' untranslated region, PABP = poly(A) binding protein, m1Ψ = N1-methylpseudouridine.

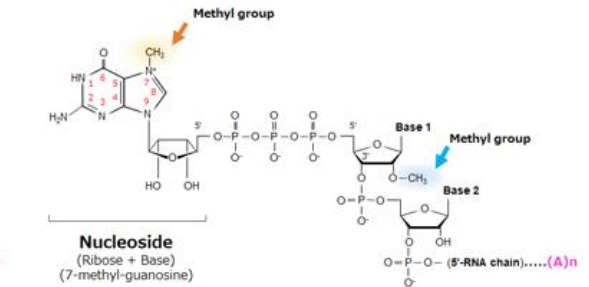
## Cap-0 structure

m<sup>7</sup>G-ppp-N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>...CDS...Poly(A)



## Cap-1 structure

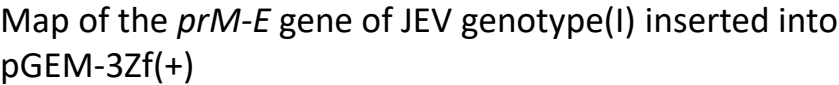
m<sup>7</sup>G-ppp-N<sub>1</sub>N<sub>2</sub>N<sub>3</sub>...CDS...Poly(A)



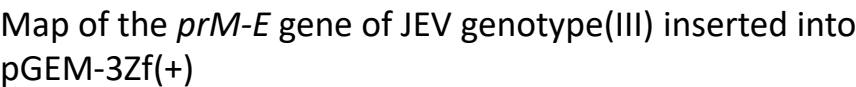
- The 5' cap is guanosine with a methyl group on the 7-position; often abbreviated as **m7G**. The cap connects via triphosphate linkage (**PPP**) to the first nucleotide at the 5' end of mRNA (the +1 position).
- In eukaryotes, the first nucleotide (+1) adjacent to the 5' cap can be further modified to create a cap-1 structure, which suppresses immunogenicity.

## Workflow for mRNA vaccine production

Restriction enzymes used for linearization: *Bam*HI



Restriction enzymes used for inserting into the plasmid: *EcoRI* and *PstI*  
Restriction enzymes used for linearization: *XhoI*





# mRNA sequence synthesis (Takara)

prM-E mRNA sequence JEV Genotype III:

AGAGCGGCCGCUUUUUCAGCAAGAUUAAGCCCAGGGCAGAGCCAUCUAUUGCUUACAUUUGCUUCUGACACAACUGUGUUCACUAGCAACCUCAAACAGACACCAUGAAGUU  
GUCAAUUUCCAGGGAAAGCUUUUGAUGACCAUCAACAACACGGACAUUUGCAGACGUUAUCGUGAUUCCCACCUCAAAAGGAGAGAACAGAUGUUGGGUCCGGGGCAAUCGAC  
GUCGGCUACAUGUGUGAGGACACUAUCACGUACGAAUGUCCUAAGCUCACCAUGGGCAAUGAUCCAGAGGAUGUGGACUGUUGGUGUGACAACCAAGAAGUCUACGUCCAAU  
AUGGACGGUGUACGCGGACCAGGCAUUCCAAGCGAAGCAAAAGAUCCGUGUCGGUCCAAACACAUGGGGAGAGUUCACUAGUGAAUAAAAAAGAGGCUUGGCUGGAUUCAAC  
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GUGGUAUUACCAUCCUCCUGCUGUUGGUCGCUCCGGCUUACAGUUUCAAACUGUCUGGGAAUGGGCAAUCGUGACUUCGUAGAAGGAGCCAGCGGAGCCACCUGGGUGGAC  
UUGGUGUUAGAAGGAGACAGCUGCUUGACAAUUAUGGCAAACGACAAACCAACAUUGGACGUCCGCAUGAUCAACAUCGAAGCUAGCCAACUUGCUGAGGUCAGAAGUUACU  
GCUAUCAUGCUUCAGUCACUGACAUCUCGACGGUGGCUCGGUGCCCCACGACUGGAGAAGCCCACAACGAGAAGGGAGCUGAUAGUAGCUAUGUGUGCAAACAAGGCUUCAC  
CGAUCGUGGGUGGGGUAACGGAUGCGGACUUUUUGGGAAGGGAAGCAUUGACACAUGUGCAAAAUUCUCCUGCACCAGUAAAGCGAUUUGGAGAAACAUAUCCAGCCAGAAAA  
CAUCAAAUACGAAGUUGGCAUUUUUGUGCAUGGAACCACCACCUCGGAAAACCAUGGGAAUUAUUCAGCGCAAGUUGGGGCGUCCCAGGCGGCAAAGUUUACAGUAACACCC  
AAUGCUCUUCGACAACCCUCAACUUGGUGACUACGGAGAAGUCACACUGGAUUUGUGAGCCAAGGAGUGGAUUAACACUGAAGCGUUUUACGUCAUGACCGUGGGGUGCAA  
AGUCAUUGUUGGUCCACAGGGAAUGGUUCCAUGAUCUCGCUCUCCCUUGGACGUCCCCUUCGAGCACAGCGUGGAGAAACAGAGAACUCCUCAUGGAAUUUGAAGAGGCGCA  
CGCCACAAAACAGUCCGUUGUUGCUCUUGGGUCACAGGAAGGAAGCCUCCAUCAGGCGUUGGCAGGAGCCAUCGUGGUGGAGUACUCAAGCUCAGUGAAGUUAACAUCAGGC  
CACCUGAAAUGCAGGCUGAAAAUGGACAAACUGGCUCUGAAAGGUACAACCUAUGGCAUGUGCACAGAAAAAUUCUCGUUCGCGAAAAACCCGGCGGACACUGGUCACGGAAC  
AGUUGUCAUCGAACUUUCCUACUCUGGGAGUGAUGGCCCUUGCAAAAUUCCGAUUGUCUCCGUUGCGAGUCUUAUUGACAUGACCCCCGUCGGGCGGCUGGUGACAGUGAA  
CCCCUUUGUCGCGACUUCAGCGCCAACUCAAGGUGCUGGUCGAGAUGGAACCCCCCUUCGGAGACUCCUACAUCGUAGUUGGAAGGGGAGACAAGCAGAUCAACCACCAUU  
GGCACAAAGCCGGAAGCACGCUGGGCAAGGCCUUUUAACGACUUUGAAGGGAGCUCAAAGACUGGCAGCGUUGGGCGACACAGCCUGGGACUUUGGCUCUAUUGGAGGGG  
UCUUAACUCCAUAAGGGAAAGCCGUUCACCAAGUGUUUGGUGGUGCCUUCAGAACACUCUUCGGGGGAUUGUCUUGGAUCACACAAGGGCUAAUUGGGGGCCCUACUACUCU  
GGAUGGGCGUCAACGCACGAGACCGAUCAAUUGCUUUGGCCUUCUUAAGCCACAGGAGGUGUGCUCGUGUUCUUAAGCGACCAUUGUGCAUGCUAGCUCGCUUUCUUGCUGUC  
CAAUUUCUAUUAAGGUUCCUUUGUUCCCUAAGUCCAACUACUAAACUGGGGGGAUAUUAUGAAGGGCCUUGAGCAUCUGGAUUCUGCCUAAUAAAAACAUUUAUUUUCAU  
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AAUAAAAACAUUUAUUUUCAUUGCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA  
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

**5'UTR (β-globin):**

AGAGCGGCCGCTTTTTTCAGCAAGATTAAGCCCAGGGCAGAGCCATCTATTGCTTACATTTGCTTCTGACACAACCTGTGTTCA  
CTAGCAACCTCAAACAGACACC

**3'UTR (β-globin):**

AGCTCGCTTTCTTGCTGTCCAATTTCTATTAAAGGTTCTTTGTTCCCTAAGTCCAACCTACTAACTGGGGGATATTATGAAGG  
GCCTTGAGCATCTGGATTCTGCCTAATAAAAAACATTTATTTTCATTGCAGCTCGCTTTCTTGCTGTCCAATTTCTATTAAAGG  
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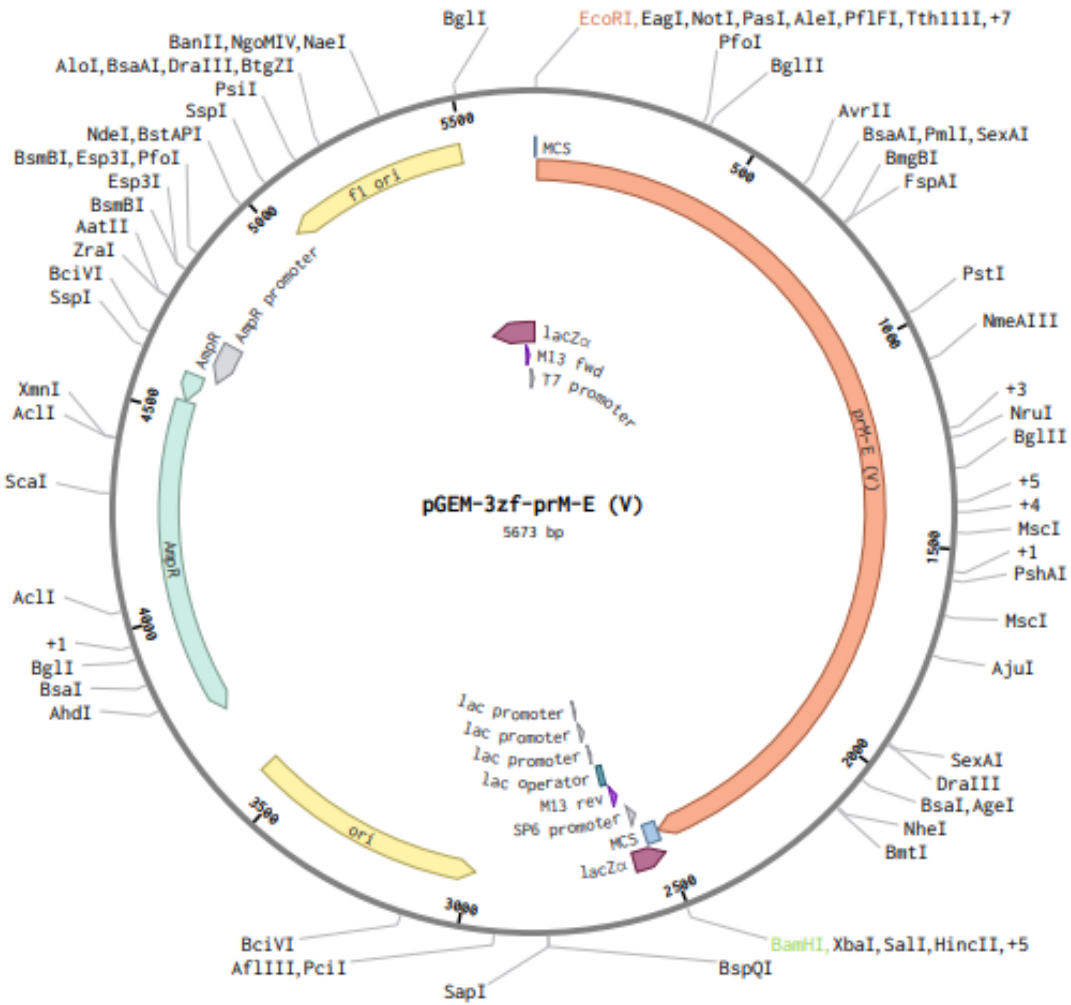
**JEV prM-E (V)**

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AACTTGATGCTGGTAATGACCCAGAGGACATTGACTGTTGGTGCGACAAACAAGCCGTGTATGTCCAGTATGGGCGTTGCA  
CGAGGACCAGGCACTCCAGGAGAAGTAGAAGATCTGTGTCAGTGCAAACCCACGGAGAAAGCTCCCTAGTGAACAAAAA  
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TCTCGTGGCAGTGGCACTCGGATGGATGCTTGGCAGCAACAACGGCCAGCGTGTGGTGTTCACAATTCTTGTGTTGGT  
CGCACCCGCATACAGCTTTAACTGCCTAGGCATGGGCAACCGCGACTTCATTGAAGGAGTCAGCGGAGCCACGTGGGTAG  
ACCTGGTGCTGGAAGGAGACAGTTGCCTCACCATCATGGCGAACGATAAACCAACACTGGACGTGCGCATGATAAACATTG  
AAGCCACGCAACTGGCTGAAGTACGAACCTATTGCTACCACGCTACAGTGGCTGACATTTCAACAGTAGCAAGATGCCCA  
CGACTGGAGAAGCCCACAACACAAGACGAGCCGATAGCAGTTATGTTTGCAAGCAAGGCTATACAGACCGTGGATGGGGA  
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AATACAGCCAGAAAATATCAAATATGAAGTTGGAGTATTTGTCCATGGAACCACAACAGCCGAGAACCATGGAACTACTCC  
GCTCAGATTGGAGCTTCCCAGGCTGCCAAGTTCACCATCACGCCCAATGCTCCTTCCATCACCTGAAGCTTGGGGACTACG  
GAGAAGTCACAATGGATTGCGAGCCTCGTAGTGGATTAACTGAAGCATTTTATGTGCTGACCGTTGGGACTAAGTCGTT  
TCTAGTCCATCGCGAATGTTTAAATGATTTGGCGCTTCCATGGCTGTCTCCATCTAGCACAACTGGAGAAACAGAGAGATC  
TTGCTGGAATTTGAAGAAGCCACGCGACGAAACAGTCTGTTGTTGCACTTGGATCACAAGAGGGAGCTCTACACCAGGC  
TCTGGCTGGCGCCATAGTGGTGGAGTATTCTAGCTCAGTGAAGTAACTTCTGGCCACCTCAAATGTAGACTAAAAATGGAC  
AAGTTGGCCTTGAAAGGAACCACTATGGCATGTGCACAGAGAAGTTCTCCTTTTCGAAAAACCCAGCTGACACTGGTCAT  
GGCACGGTCGTCATAGAATTGCAGTACACTGGCACTGATGGACCGTGCAAGATACCATCTCTTCACTGGCCAGCCTGAATG  
ATTTGACTCCAGTTGGCAGATTGGTGACAGTCAATCCTTTTGTGTCACATCCACTGCCAACTCGAAAGTTTTTGGTGGAACT  
TGAACCACCGTTTGGAGATTCATTATTGTTGTTGGGAGAGGAGACAAGCAGATTAACCACCATTTGGCACAAGGCAGGCA  
GTTTCGCTGGGAAAGGCTTTTACCCTACCTGAAAGGTGCCAGAGGTTAGCTGCCCTTGCGACACGGCCTGGGATTTT  
GGGTCCATTGGAGGAGTTTTTAATTCCATTGGCAAGGCCGTGCACCAGGTGTTTGGAGGAGCTTTTAGAACACTTTTTGGT  
GGCATGTCTTGGATAACACAAGGATTGATGGGAGCACTGCTGCTGTGGATGGGTATCAATGCGCGAGACCGGTCGATCGCA  
CTGGCCTTTCTTGCTACAGGAGGCGTGCTCTTGTCTTGGCTACCAATGTCCACGCT

**Poly (A) Tail:** 120 nt

Restriction enzymes used for inserting into the plasmid: *EcoRI* and *BamHI*

Restriction enzymes used for linearization: *BamHI*



Map of the *prM-E* gene of JEV genotype(V) inserted into pGEM-3Zf(+)

## Modified mRNA Vaccines Protect against Zika Virus Infection

Justin M. Richner,<sup>1,9</sup> Sunny Himansu,<sup>2,9</sup> Kimberly A. Dowd,<sup>3</sup> Scott L. Butler,<sup>2</sup> Vanessa Salazar,<sup>1</sup> Julie M. Fox,<sup>1</sup> Justin G. Julander,<sup>4</sup> William W. Tang,<sup>5</sup> Sujun Shrestha,<sup>5</sup> Theodore C. Pierson,<sup>3</sup> Giuseppe Ciarrella,<sup>2,\*</sup> and Michael S. Diamond<sup>1,6,7,8,10,\*</sup>

<sup>1</sup>Department of Medicine, Washington University School of Medicine, St. Louis, MO 63110, USA

<sup>2</sup>Valera LLC, a Moderna Venture, 500 Technology Square, Cambridge, MA, 02139, USA

<sup>3</sup>Viral Pathogenesis Section, National Institutes of Health, Bethesda, MD 20892 USA

<sup>4</sup>Institute for Antiviral Research, Utah State University, Logan, UT, 84335 USA

<sup>5</sup>Division of Inflammation Biology, La Jolla Institute for Allergy and Immunology, La Jolla, CA 92037, USA

<sup>6</sup>Department of Pathology and Immunology, Washington University School of Medicine, St. Louis, MO 63110, USA

<sup>7</sup>Department of Molecular Microbiology, Washington University School of Medicine, St. Louis, MO 63110, USA

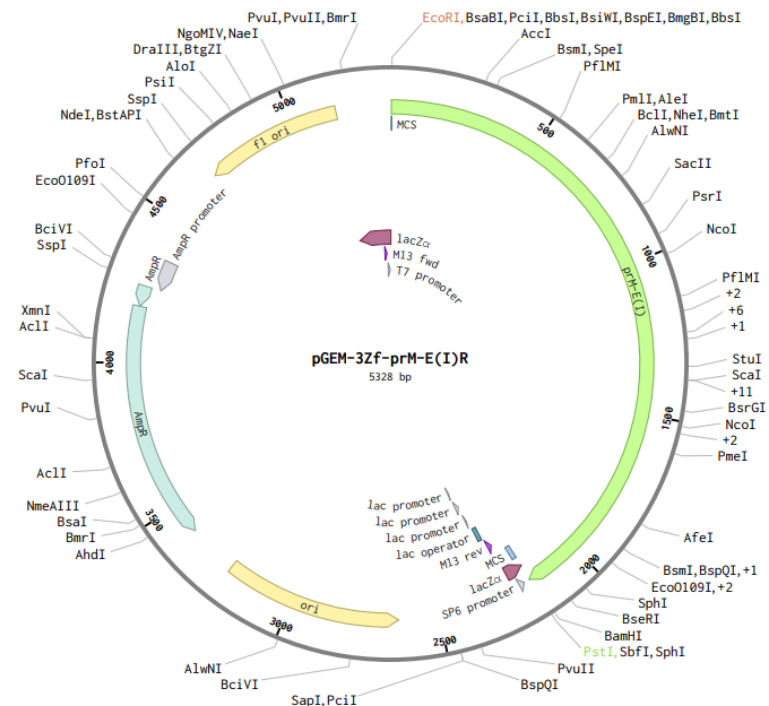
<sup>8</sup>The Andrew M. and Jane M. Bursky Center for Human Immunology and Immunotherapy Programs, Washington University School of Medicine, St. Louis, MO 63110, USA

<sup>9</sup>Co-first authors

<sup>10</sup>Lead Contact: Michael S. Diamond

\*Correspondence: [diamond@wustl.edu](mailto:diamond@wustl.edu) (M.S.D.), [Giuseppe.Ciarrella@Valeratx.com](mailto:Giuseppe.Ciarrella@Valeratx.com) (G.C.)

<http://dx.doi.org/10.1016/j.cell.2017.02.017>



Map of the *prM-E* gene of JEV genotype(I) inserted into pGEM-3Zf(+)

GAATTCGGGAAATAAGAGAGAGAAAAGAAGAGTAAGAAGAAATATAAGAGCCACCATGAAGCTATCAAACCTTTCAAGGAAAGCTTCTGATGACCATCAACAACACGGACATTGCGGACGTCATCGTGATCCCCACCTCAAAGGTGAAAACAGATGTTGGGTCCGAGCAATCGACGTTGGTTACATGTGTGAAGACACCATCACGTACGAATGTCCGAAGCTTGCCGTGGGCAACGATCCGGAAGACGTGGACTGCTGGTGCGACAATCAAGAAGTCTACGTGCAGTATGGTCGCTGCACACGGACCAGGCATTCCAAACGAAGCAGAAGATCCGTTTCAGTCCAAACGCATGGGGAAAGCTCACTAGTGAACAAAAAAGAGGCTTGCTGGATTGACGAAGGCCACGCGATACCTCATGAAAACGGAGAAGTGGATCATAAGGAACCTGGTTATGCTTTCCTGGCGGCGGCACCTTGATGGATGCTTGGCAGCAACAGTGGCCAACGTGTGGTGTCTACTATTCTCTTGCTGTTGGTCTCGGCTTACAGTTTTAACTGTCTGGGAATGGGGAATCGGGATTTTCATAGAAGGAGCCAGTGGAGCCACGTGGGTGGATCTGGTGTAGAAGGAGATAGCTGTTTGA CAATTATGGCAAACGACAAACCAACTAGATGTCCGCATGATCAACATTGAAGCTAGCCAACCTGCTGAAGTCAGGAGTTA CTGTTATCACGCTTCAGTCACTGACATTTCAACGGTGGCTCGATGCCCCATGACTGGAGAAGCCCACAACGAAAAACGTGCT GACAGCAGCTACGTGTGCAACAAGGCTTTACTGACCGCGGATGGGGAAATGGATGTGGACTTTTCGGGAAAGGAAGCAT TGACACATGCGCAAAATTTCTGTACCAGTAAGGCCATTGGAAGAATGATCCAACCAGAGAACATCAAGTACGAGGTTGG CATATTCGTGCACGGAACCACCACCTCGGAAAACCATGGGAATTAAGTACGCGCAAGTAGGAGCGTCTCAAGCAGCAAAGTT TACTGTAAGTCCAAATGCTCCCTCAATAACCCTCAAGCTTGGTGATTATGGAGAAGTCACACTGGATTGTGAACCAAGGAGT GGACTGAACACTGAAGCGTTCTATGTCATGACCGTGGGTTCGAAGTCATTCTAGTCCATAGGGAATGGTTCCATGACCTTT CTCTCCCTGGACGTCCCCCTCGAGCACGGCATGGAGAAACAGAGAACTCCTCATGGAATTTGAACAGGCACATGCCACAAA ACAATCCGTCGTAGCTCTTGGGTACAGGAGGGAGGCCTCCATCAAGCGTTGGCAGGAGCCATCGTGGTGGAGTACTCGA GCTCAGTGAAGTTGACATCAGGTACCTGAAATGCAGGCTAAAAATGGACAACTGGCTCTGAAGGGCACGACTTATGGCA TGTGTACAGAAAAATTCTCGTTCGCGAAAAATCCAGCGGACACAGGCCATGGAACAGTTGTCATTGAGCTCACATACTCTGG AAGTGATGGTCCCTGTAAAATTCCGATTGTCTCAGTCGCGAGTTTAAACGACATGACCCCTGTGGGGAGGCTGGTAACAGT AAACCCCTTCGTGCGACATCTAGCTCAAACCTCAAAGGTGCTGGTTGAGATGGAACCTCCCTTCGGAGACTCTTATATCGTG GTTGAAGAGGGGACAAGCAGATTAACCATCACTGGCACAAAGCTGGAAGCACGCTGGGTAAAGCCTTCTCAACAACTTTG AAAGGGGCTCAGAGACTAGCAGCGCTAGGTGACACAGCTTGGGACTTCGGCTCCATTGGAGGGGTATTCAACTCCATAGG GAAAGCTGTTACCAAGTATTTGGCGGTGCATTGAAACGCTCTTCGGGGGAATGTCTTGATCACACAAGGACTAATGGG GGCCTTACTTCTTTGGATGGGTGTCAACGCACGAAACCGGTCAATCGCCCTGGCTTTCTGGCCACGGGAGGTGTGCTCGTG TTTTAGCGACCAATGTGCATGCCCTGATAATAGGCTGGAGCCTCGGTGGCCATGCTTCTTGGCCCTTGGGCCTCCCCCAGCC CCTCCTCCCCTTCTGCACCCGTACCCCGTGGTCTTTGAATAAAGTCTGAGGATCCCTGCAG

Restriction enzymes used for inserting into the plasmid: *EcoRI* and *PstI*

Restriction enzymes used for linearization: *BamHI*

The sequence has been sent to TWIST Bioscience for synthesis (excluding the restriction sites used for ligating the insert into the plasmid; *EcoRI* and *PstI*)





# mRNA synthesis through In-vitro transcription kit

**CELLSCRIPT™**  
RNA for Translation in Cells

## T7-FlashScribe™ Transcription Kit

Cat. Nos. C-ASF3507

### INTRODUCTION

The T7-FlashScribe™ Transcription Kit is specially formulated to enable users to obtain the maximum possible yields of RNA from an *in vitro* transcription (IVT) reaction in just 30 minutes. The standard 30 minute, 20 µl reaction will yield up to 180 µg of RNA from 1 µg of the control template. These yields are made possible by the high-performance properties of the T7-FlashScribe enzyme.

The T7-FlashScribe Transcription Kit produces exceptionally high yields of either long or short transcripts. The standard reaction can be scaled up to produce milligram amounts of RNA.

T7-FlashScribe IVT RNA can be processed into mRNA (5'-end capped and 3'-end poly(A) tailed) through the use of CELLSCRIPT's ScriptCap™ m<sup>7</sup>G Capping System, ScriptCap 2'-O-Methyltransferase Kit and A-Plus™ Poly(A) Polymerase Tailing Kit (available separately).

CELLSCRIPT also offers the INCOGNITO™ line of transcription kits for the production of pseudouridine- & 5-methyl-cytosine-containing (GAΨC and GAΨ5mC) IVT RNA. It has been shown that Ψ-mRNAs and Ψ5mC-mRNAs are translated into protein at higher levels and induce lower innate immune responses in human and other mammalian cells that express various RNA sensors compared to corresponding unmodified mRNAs.<sup>1-4</sup>

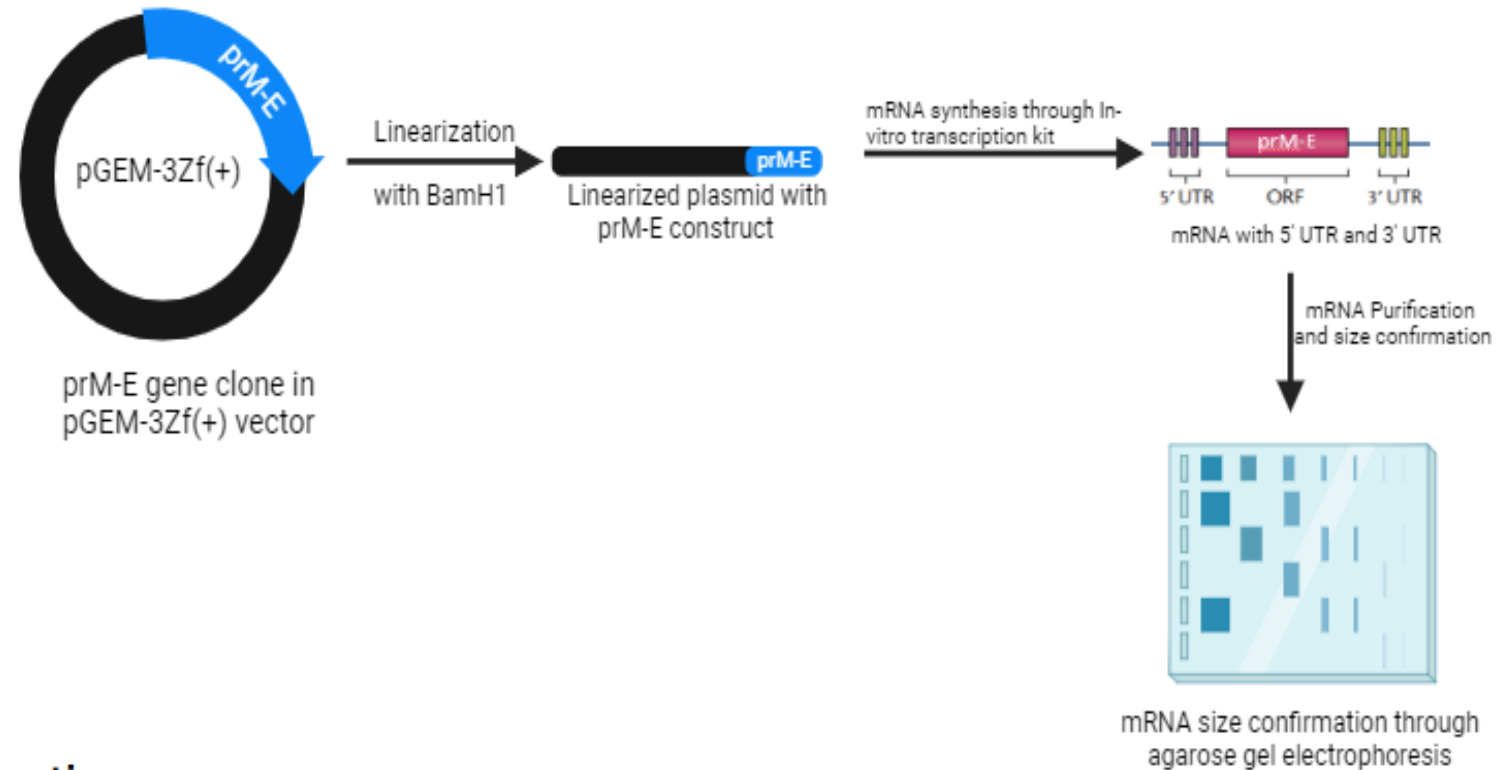
## MEGAclear™ Kit Purification for Large Scale Transcription Reactions

Catalog Number AM1908

Publication Number 1908M Revision C

### Product description

The MEGAclear™ Kit is designed for rapid high-throughput purification of RNA from enzymatic reactions such as *in vitro* transcription. The process is simple and fast, and it recovers from 1 ng to 500 µg of RNA efficiently. The MEGAclear™ Kit is appropriate for purification of ssRNA larger than 100 nt and dsRNA larger than 200 bp.



At this level we will have purified mRNA with 5' and 3' UTR  
Next step is capping of mRNA at 5' position

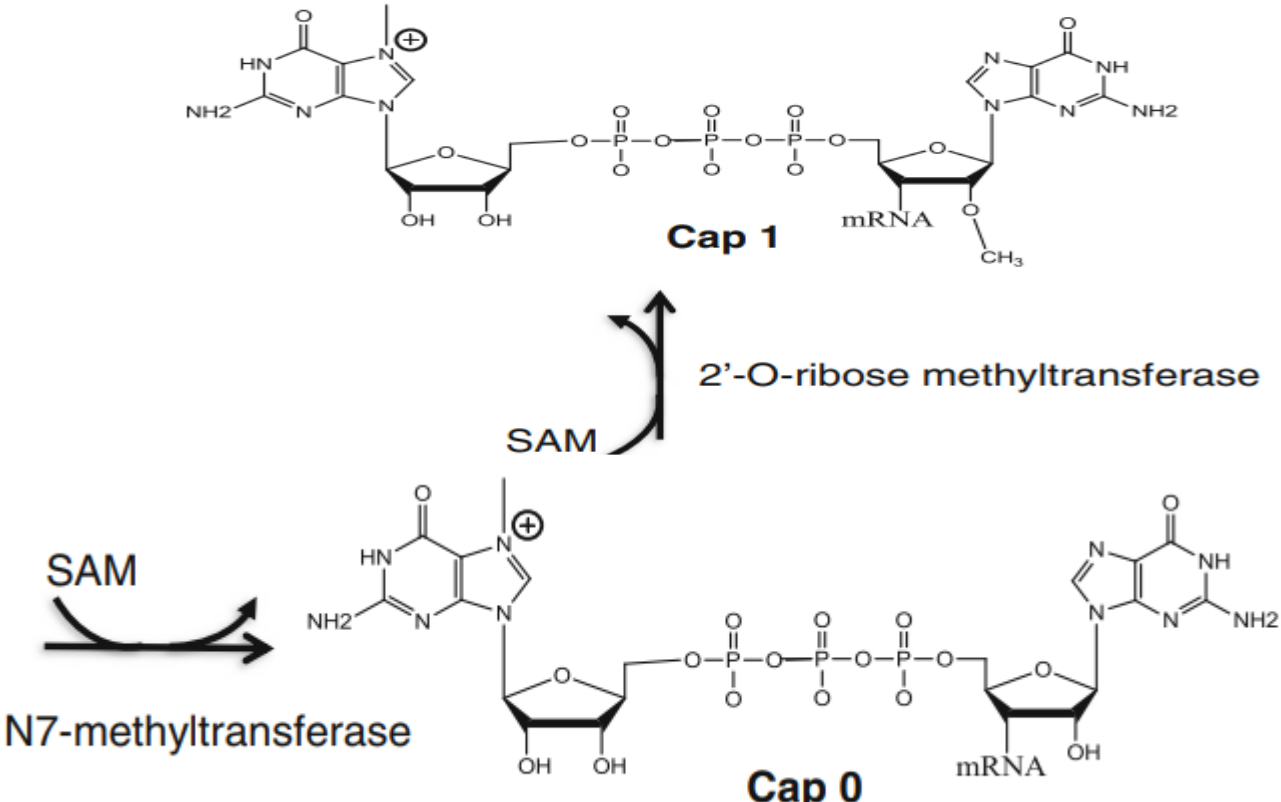
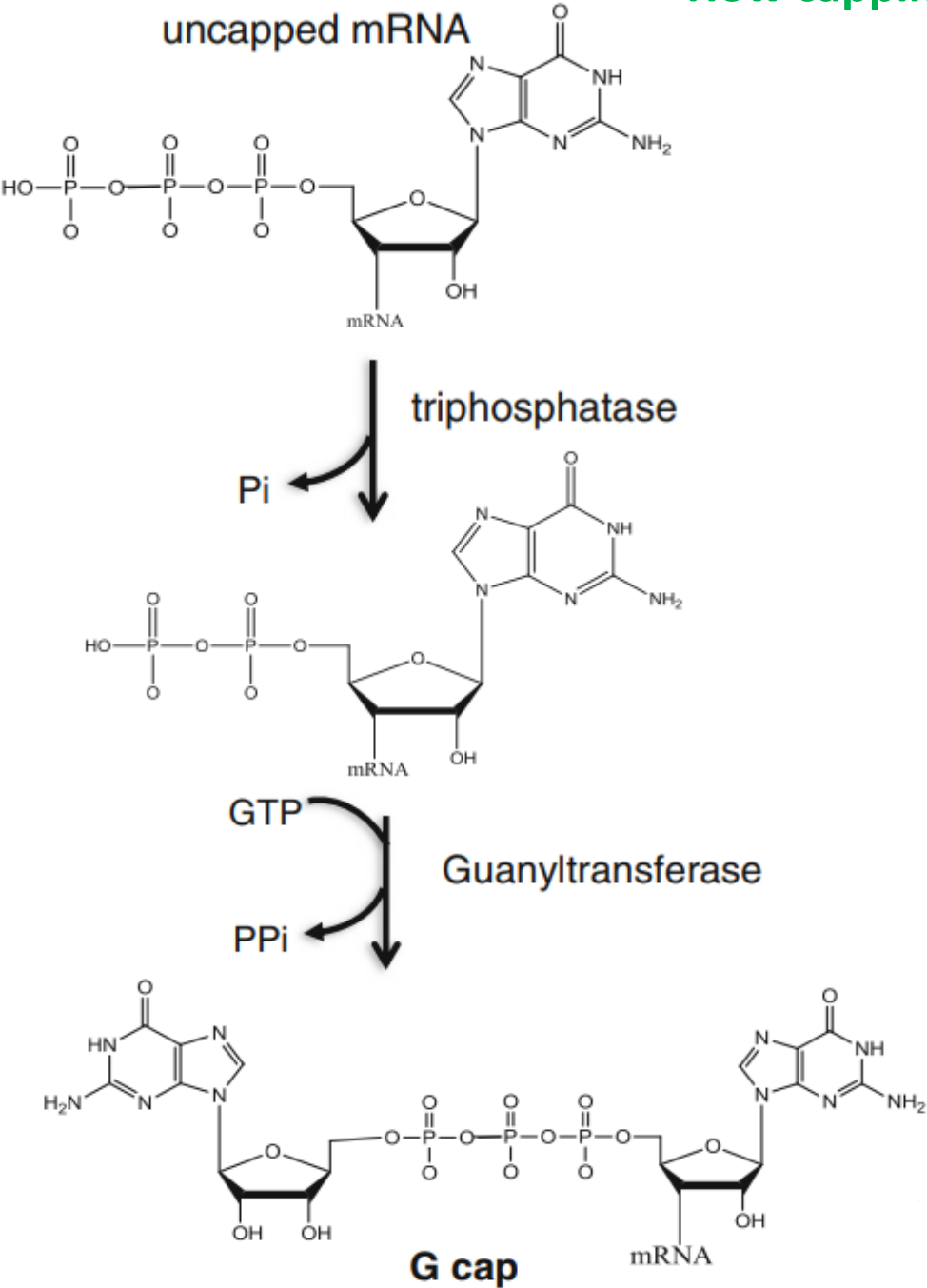
# How capping will be done

MATERIALS

Materials Supplied

**Important** Store at -20°C in a freezer without a defrost cycle. Do not store at -70°C.

ScriptCap™ Cap 1 Capping System Contents		
Kit Component	Reagent Volume	
	C-SCCS1710 10 Reactions	C-SCCS2250 50 Reactions
ScriptCap™ Capping Enzyme, 10 U/μl	40 μl	200 μl
ScriptCap™ 2'-O-Methyltransferase, 100 U/μl	40 μl	200 μl
10X ScriptCap™ Capping Buffer 0.5 M Tris-HCl, pH 8.0, 60 mM KCl and 12.5 mM MgCl <sub>2</sub>	100 μl	500 μl
10 mM GTP Solution	100 μl	500 μl
20 mM S-adenosyl-methionine (SAM)	50 μl	250 μl
ScriptGuard™ RNase Inhibitor, 40 U/μl	25 μl	125 μl
RNase-Free Water	0.67 ml	3.35 ml



# Label-free analysis of mRNA capping efficiency using RNase H probes and LC-MS

Michael Beverly<sup>1</sup> · Amy Dell<sup>1</sup> · Parul Parmar<sup>1</sup> · Leslie Houghton<sup>1</sup>

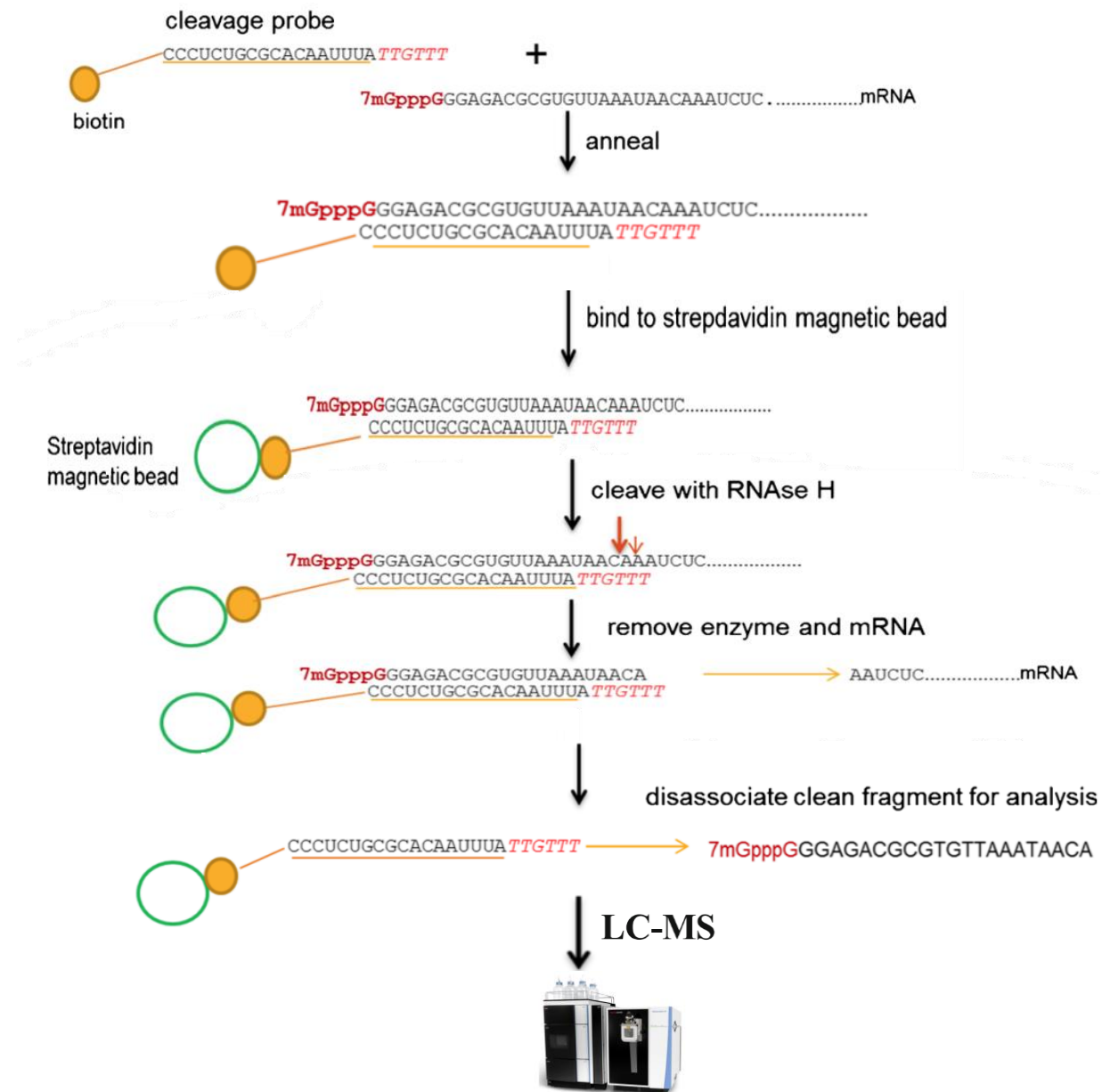
## Materials required for assay

- RNase H cleavage probe conjugated with biotin
- RNase H
- RNase H reaction buffer
- RNA 5' pyrophosphohydrolase (RppH) with NEbuffer2
- Streptavidine-coated magnetic beads
- NaOH
- NaCl
- Tris-HCl
- EDTA
- Methanol
- Hexafluoroisopropanol
- Triethylamine

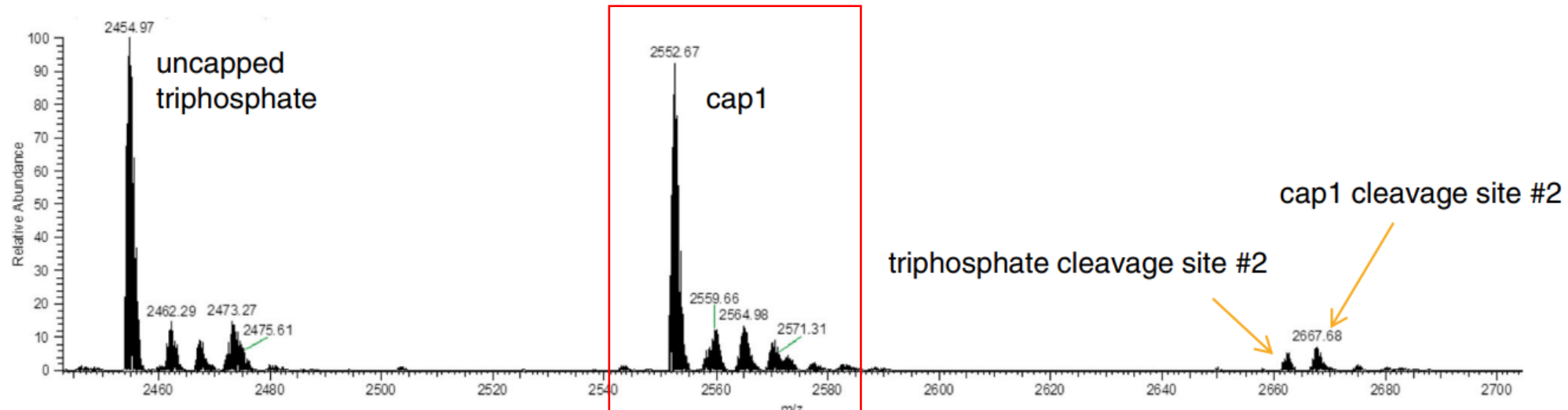
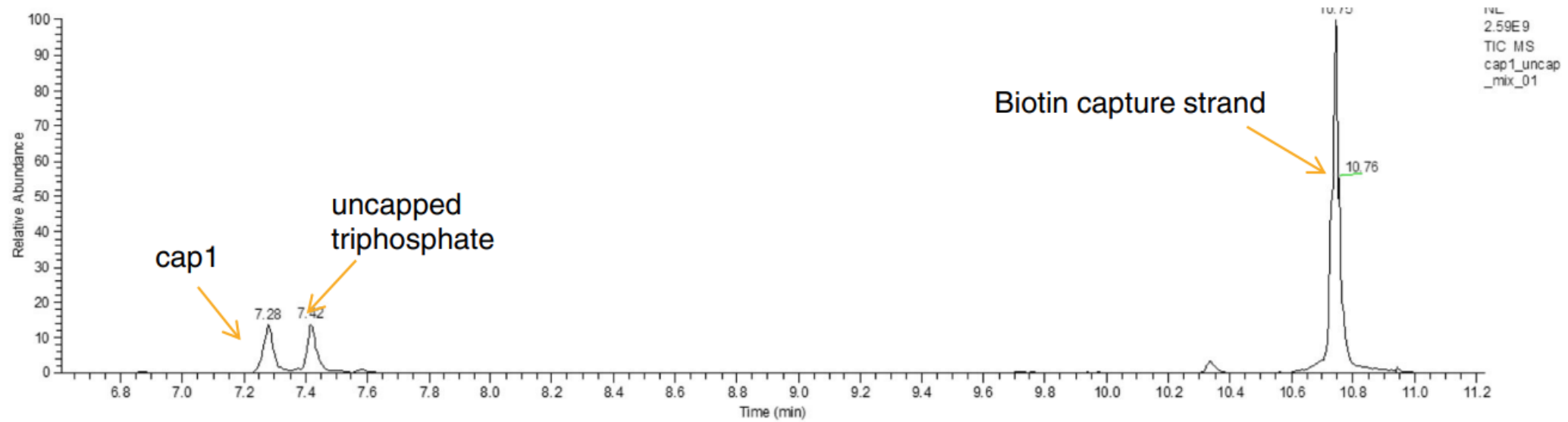
## Instruments

- Vacuum dryer
- LC-MS

## Steps involved in analysis of mRNA capping



# LC-MS Analysis






# Polyadenylation of mRNA

**INTRODUCTION**

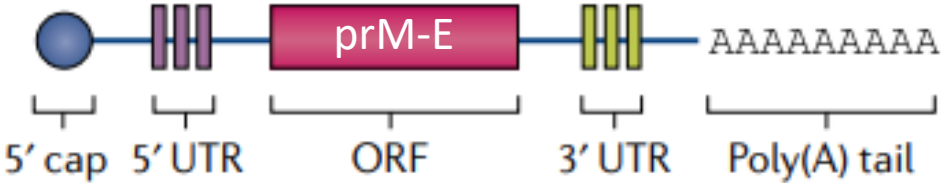
The A-Plus™ Poly(A) Polymerase Tailing Kit uses ATP as a substrate for template-independent addition of adenosine monophosphates to the 3'-hydroxyl termini of RNA. The standard protocol produces a poly(A)-tail length of ~150 b on 40-60 µg of RNA. Polyadenylation increases the stability of RNA in eukaryotic cells and enhances its ability to be translated after transfection or microinjection.<sup>1-3</sup> A Poly(A) tail is useful to provide a priming site for first-strand cDNA synthesis in certain applications, and can be used to end-label<sup>4</sup> or quantify<sup>5</sup> mRNA.

**MATERIALS**

**Materials Supplied**

 Store at -20°C in a freezer without a defrost cycle. Do not store at -70°C.

A-Plus™ Poly(A) Polymerase Tailing Kit Contents (50 reactions)	
Kit Component	Volume
A-Plus™ Poly(A) Polymerase, 4 U/µl in 50% glycerol, 50 mM Tris-HCl, pH 7.5, 0.5 M NaCl, 1 mM dithiothreitol (DTT), 0.1 mM EDTA and 0.1% Triton® X-100.	100 µl
10X A-Plus™ Poly(A) Tailing Buffer 0.5 M Tris-HCl, pH 8.0, 2.5 M NaCl and 100 mM MgCl <sub>2</sub> .	500 µl
10 mM ATP	500 µl
RNase-Free Water	2 x 1.4 ml



mRNA

This is full mRNA which will act as vaccine candidate

## Assays for determining poly(A) tail length and the polarity of mRNA decay in mammalian cells

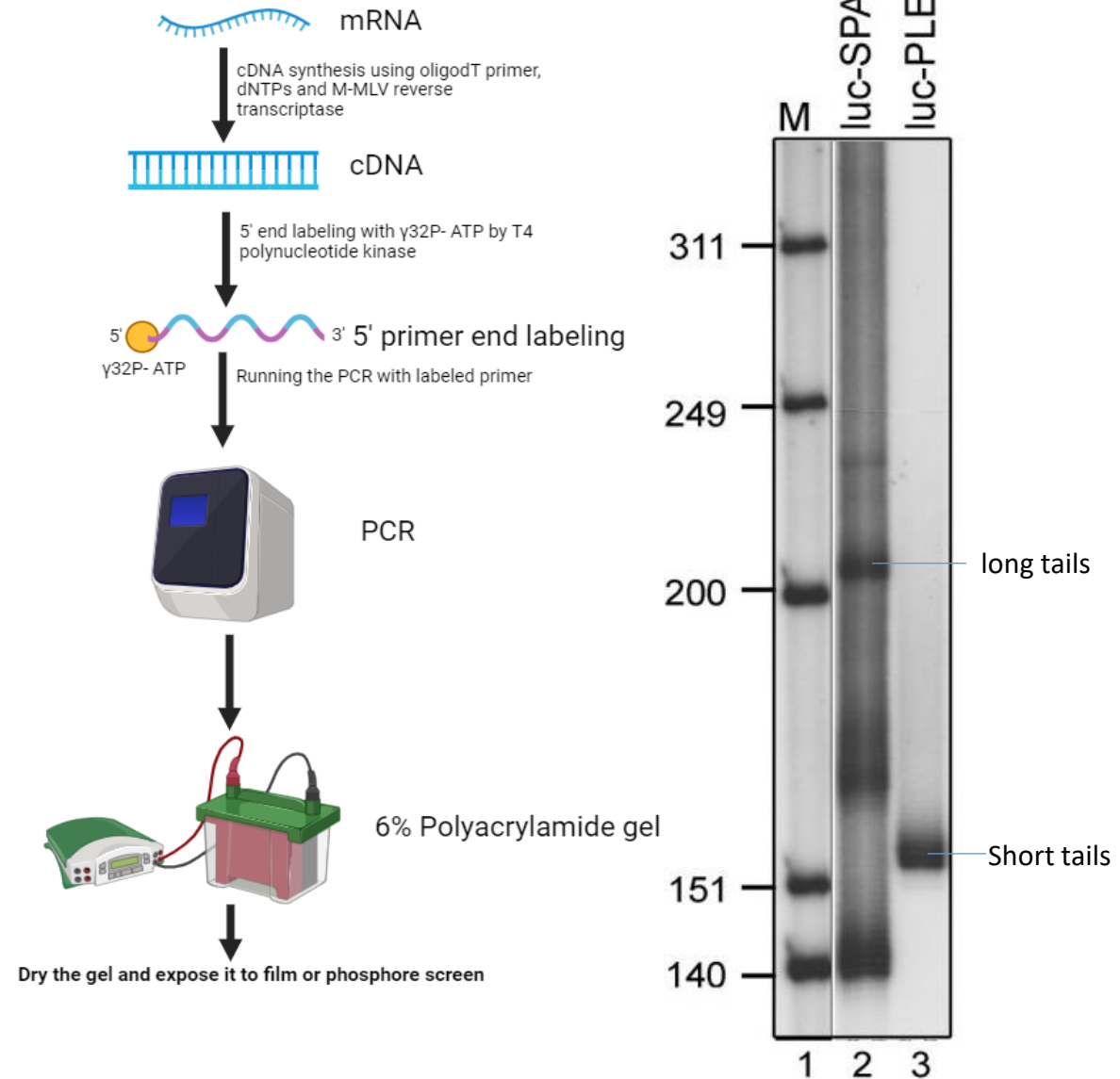
Elizabeth L. Murray and Daniel R. Schoenberg\*

Department of Molecular and Cellular Biochemistry and the RNA Group, The Ohio State University, Columbus, OH 43210-1218

### Materials required for assay

- OligodT primer/adaptor
- mRNA specific primer (Primer should be located 100bp upstream of poly A tail that will give size between 100-300 bases.
- Strand buffer
- DTT
- dNTPs
- RNase Inhibitor
- M-MLV reverse transcriptase
- Polynucleotide kinase buffer
- $\gamma$ 32P-ATP
- T4PNK
- Reagents for PCR
- Reagents to run polyacrylamide gel
- Film or phosphor screen

### Steps involved in assay



mRNAs with short tails yield a compact band while mRNAs with long tails yield PCR products of a variety of lengths which appear as a smear on the gel

## ARTICLE OPEN

## Rational development of a combined mRNA vaccine against COVID-19 and influenza

Qing Ye<sup>1,6</sup>, Mei Wu<sup>1,6</sup>, Chao Zhou<sup>1,6</sup>, Xishan Lu<sup>2,6</sup>, Baoying Huang<sup>3,6</sup>, Ning Zhang<sup>1</sup>, Hui Zhao<sup>1</sup>, Hang Chi<sup>1</sup>, Xiaojing Zhang<sup>2</sup>, Dandan Ling<sup>2</sup>, Rong-Rong Zhang<sup>1</sup>, Zhuofan Li<sup>2</sup>, Dan Luo<sup>1</sup>, Yi-Jiao Huang<sup>1</sup>, Hong-Ying Qiu<sup>1</sup>, Haifeng Song<sup>2</sup>, Wenjie Tan<sup>3,5</sup>, Ke Xu<sup>4</sup>, Bo Ying<sup>2</sup> and Cheng-Feng Qin<sup>1,5</sup>

Cell

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## Article

## A Thermostable mRNA Vaccine against COVID-19

Na-Na Zhang,<sup>1,2,6</sup> Xiao-Feng Li,<sup>1,6</sup> Yong-Qiang Deng,<sup>1,6</sup> Hui Zhao,<sup>1,6</sup> Yi-Jiao Huang,<sup>1,6</sup> Guan Yang,<sup>3,6</sup> Wei-Jin Huang,<sup>4,6</sup> Peng Gao,<sup>5</sup> Chao Zhou,<sup>1</sup> Rong-Rong Zhang,<sup>1</sup> Yan Guo,<sup>1</sup> Shi-Hui Sun,<sup>1</sup> Hang Fan,<sup>1</sup> Shu-Long Zu,<sup>1</sup> Qi Chen,<sup>1</sup> Qi He,<sup>3</sup> Tian-Shu Cao,<sup>1</sup> Xing-Yao Huang,<sup>1</sup> Hong-Ying Qiu,<sup>1</sup> Jian-Hui Nie,<sup>4</sup> Yuhang Jiang,<sup>5</sup> Hua-Yuan Yan,<sup>5</sup> Qing Ye,<sup>1</sup> Xia Zhong,<sup>5</sup> Xia-Lin Xue,<sup>5</sup> Zhen-Yu Zha,<sup>5</sup> Dongsheng Zhou,<sup>1</sup> Xiao Yang,<sup>3</sup> You-Chun Wang,<sup>4,\*</sup> Bo Ying,<sup>5,\*</sup> and Cheng-Feng Qin<sup>1,7,\*</sup>

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Emerging Microbes & Infections  
<https://doi.org/10.1080/22221751.2023.2192815>



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## RESEARCH ARTICLE

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## Rational development of multicomponent mRNA vaccine candidates against mpox

Rong-Rong Zhang<sup>a†</sup>, Zheng-Jian Wang<sup>a†</sup>, Yi-Long Zhu<sup>b,c†</sup>, Wei Tang<sup>a†</sup>, Chao Zhou<sup>a†</sup>, Suo-Qun Zhao<sup>a</sup>, Mei Wu<sup>a</sup>, Tao Ming<sup>a</sup>, Yong-Qiang Deng<sup>a</sup>, Qi Chen<sup>a</sup>, Ning-Yi Jin<sup>b</sup>, Qing Ye<sup>a</sup>, Xiao Li<sup>b</sup> and Cheng-Feng Qin<sup>a,d</sup>

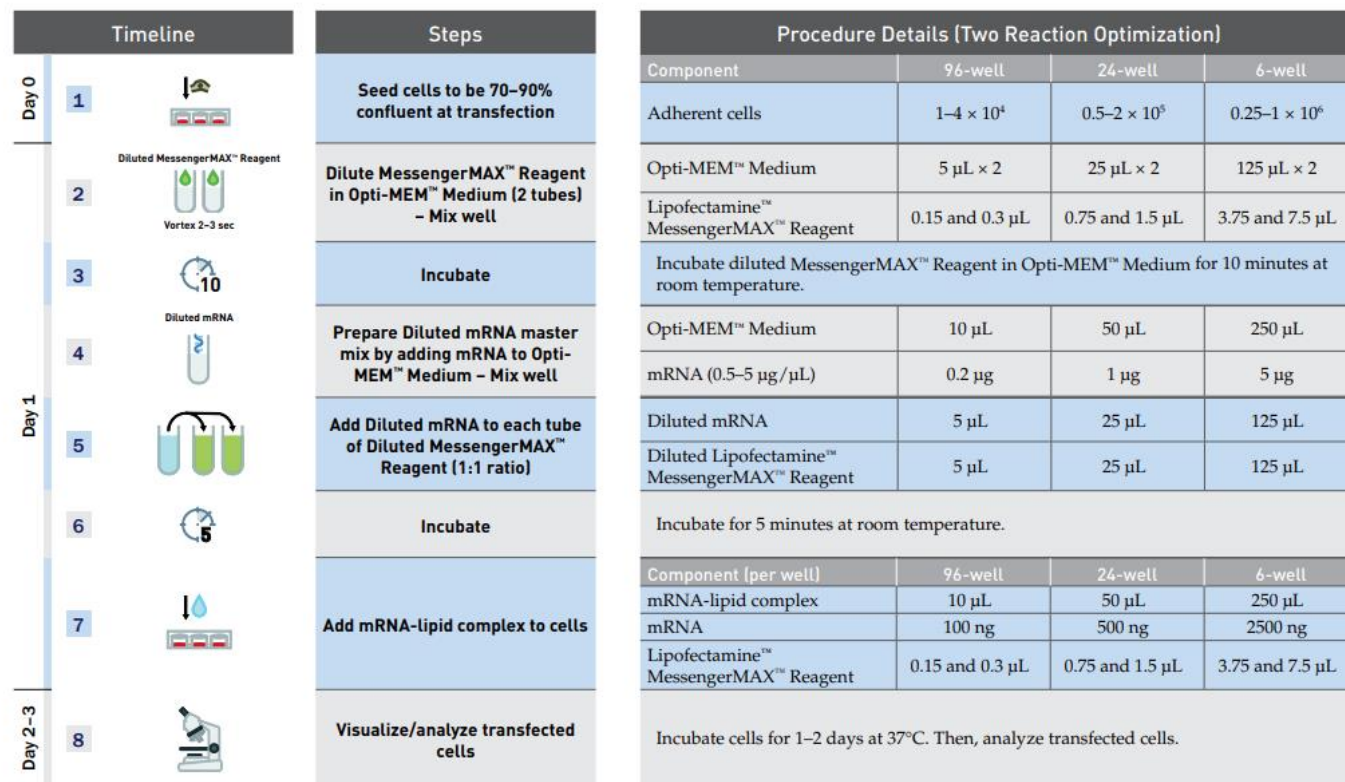
<sup>a</sup>State Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology, Beijing, People's Republic of China; <sup>b</sup>Changchun Veterinary Research Institute, Chinese Academy of Agricultural Sciences, Changchun, People's Republic of China;

<sup>c</sup>Academics Workstation of Jilin Province, Changchun University of Chinese Medicine, Changchun, People's Republic of China;

<sup>d</sup>Research Unit of Discovery and Tracing of Natural Focus Diseases, Chinese Academy of Medical Sciences, Beijing, People's Republic of China

## Transfection of mRNA into cell line

## Lipofectamine MessengerMax reagent transfection protocol (Thermo Fischer)



## mRNA transfection and protein expression

For immunofluorescence staining, HEK293T cells were seeded in 24-well plates at  $4 \times 10^5$  cells per well and cultured at 37 °C in 5% CO<sub>2</sub> for 12 h. mRNA encoding HA was transfected into HEK293T cells using **Lipofectamine MessengerMAX Reagent** (Thermo Fisher Scientific). Transfected cells were fixed with cold methanol/acetone (7:3) 48 h post-transfection and incubated with primary antibody (rabbit polyclonal antibody for H1N1-HA, GeneTex, GTX127357, 1:400) at 37 °C for 1 h. Cells were then washed with PBS three times and then incubated with secondary antibody conjugated to Alexa Fluor 488 (Proteintech, SA00013-2, 1:400). HA-positive cells were examined using a PerkinElmer High Content Analysis System Operetta CLS and processed using Harmony 4.9 software.

48hr post transfection cells will be fixed with cold methanol/acetone (7:3)

Incubate with primary antibody

Wash the cells with PBS three time

Incubate with secondary antibody conjugated with alexa fluor 488

Imaging with microscope



ARTICLE OPEN

## Rational development of a combined mRNA vaccine against COVID-19 and influenza

Qing Ye<sup>1,6</sup>, Mei Wu<sup>1,6</sup>, Chao Zhou<sup>1,6</sup>, Xishan Lu<sup>2,6</sup>, Baoying Huang<sup>3,6</sup>, Ning Zhang<sup>1</sup>, Hui Zhao<sup>1</sup>, Hang Chi<sup>1</sup>, Xiaojing Zhang<sup>2</sup>, Dandan Ling<sup>2</sup>, Rong-Rong Zhang<sup>1</sup>, Zhuofan Li<sup>2</sup>, Dan Luo<sup>1</sup>, Yi-Jiao Huang<sup>1</sup>, Hong-Ying Qiu<sup>1</sup>, Haifeng Song<sup>2</sup>, Wenjie Tan<sup>3,8</sup>, Ke Xu<sup>4</sup>, Bo Ying<sup>2</sup> and Cheng-Feng Qin<sup>1,5,9</sup>

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<sup>1</sup>State Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology, Academy of Military Medical Sciences, Beijing 100071, China

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<https://doi.org/10.1016/j.cell.2020.07.024>

Emerging Microbes & Infections  
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RESEARCH ARTICLE

OPEN ACCESS

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### Rational development of multicomponent mRNA vaccine candidates against mpox

Rong-Rong Zhang<sup>a,†</sup>, Zheng-Jian Wang<sup>a,†</sup>, Yi-Long Zhu<sup>b,c,†</sup>, Wei Tang<sup>a,†</sup>, Chao Zhou<sup>a,†</sup>, Suo-Qun Zhao<sup>a</sup>, Mei Wu<sup>a</sup>, Tao Ming<sup>a</sup>, Yong-Qiang Deng<sup>a</sup>, Qi Chen<sup>a</sup>, Ning-Yi Jin<sup>b</sup>, Qing Ye<sup>a</sup>, Xiao Li<sup>b</sup> and Cheng-Feng Qin<sup>a,d</sup>

<sup>a</sup>State Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology, Beijing, People's Republic of China; <sup>b</sup>Changchun Veterinary Research Institute, Chinese Academy of Agricultural Sciences, Changchun, People's Republic of China;

<sup>c</sup>Academicians Workstation of Jilin Province, Changchun University of Chinese Medicine, Changchun, People's Republic of China;

<sup>d</sup>Research Unit of Discovery and Tracing of Natural Focus Diseases, Chinese Academy of Medical Sciences, Beijing, People's Republic of China

### LNP Formulation of the mRNA

The mRNA vaccine encoding HA protein of H1N1 was prepared in LNP formulations. Briefly, a lipid mixture including ionizable lipids, 1,2-distearoyl-sn-glycero-3-phosphocholine (DSPC), cholesterol and PEG-lipid (molar ratios of 50:10:38.5:1.5) was combined with 20 mM citrate buffer (pH 4.0) containing mRNA at a ratio of 1:2 through a T-mixer. The formulation were then diafiltrated in 10 × volume of PBS (pH 7.4), reduced to the desired concentrations through a tangential flow filtration membrane with 100 kD, passed through a 0.22 μm filter, and stored at 2–8 °C until use. All formulations were tested for particle size, distribution, RNA concentration and encapsulation. The combined mRNA vaccine candidate (AR-CoV/IAV) was developed by mixing ARCoV and ARIAV under the same LNP-mRNA vaccine platform. Empty LNPs were utilized as placebo.



T-mixer (medimix)

### Steps involve in LNP formulation

ionizable lipids, DSPC, cholesterol and polyethylene Glycol (PEG)-lipid (molar ratio of 50:10:38.5:1.5)



Combined with 20mM citrate buffer (pH 4.0) containing mRNA at a ratio of 1:2 and mixed with T-mixer



The formulation will be diafiltrated in 10x volume of PBS



Reduce to desired concentration through tangential flow filtration membrane with 100kD



Pass through 0.22μm filter



stored at 2-8

## Article

# Protective Immune Responses Induced by an mRNA-LNP Vaccine Encoding prM-E Proteins against Japanese Encephalitis Virus Infection

Tao Chen <sup>1,2,3</sup>, Shuo Zhu <sup>1,2,3</sup>, Ning Wei <sup>1,2,3</sup>, Zikai Zhao <sup>1,2,3</sup>, Junjun Niu <sup>1,2,3</sup>, Youhui Si <sup>1,2,3</sup>, Shengbo Cao <sup>1,2,3,\*</sup> and Jing Ye <sup>1,2,3,\*</sup>

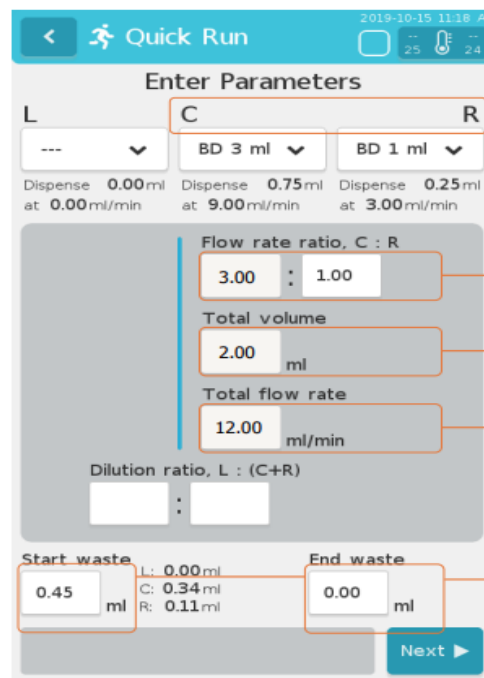
- <sup>1</sup> State Key Laboratory of Agricultural Microbiology, Huazhong Agricultural University, Wuhan 430070, China; ct@webmail.hzau.edu.cn (T.C.); zhushuo@webmail.hzau.edu.cn (S.Z.); weining@webmail.hzau.edu.cn (N.W.); zikaizhao@hotmail.com (Z.Z.); njj@webmail.hzau.edu.cn (J.N.); youhui@mail.hzau.edu.cn (Y.S.)
- <sup>2</sup> Laboratory of Animal Virology, College of Veterinary Medicine, Huazhong Agricultural University, Wuhan 430070, China
- <sup>3</sup> The Cooperative Innovation Center for Sustainable Pig Production, Huazhong Agricultural University, Wuhan 430070, China
- \* Correspondence: sbcao@mail.hzau.edu.cn (S.C.); yej@mail.hzau.edu.cn (J.Y.)

## 2.2. Generation of mRNA and mRNA-LNP

The mRNA contained 5' and 3' UTR and a poly-A tail was produced from a linearized DNA template with a T7 in vitro transcription kit (Cellscript Madison, WI, USA), and pseudouridine was used in place of uridine. Then, the mRNA was enzymatically capped. According to the protocol, the mRNA was dissolved in an aqueous buffer and combined with GenVoy ILM (Precision Nanosystems, Vancouver, BC, Canada) at a flow ratio of 3:1 through a microfluidic mixer (Precision Nanosystems, Vancouver, BC, Canada). The solvent was removed by centrifugation at  $2000 \times g$  using a 100 kDa ultrafiltration tube (Milipore, Billerica, MA, USA) and the size and PDI of mRNA-LNP was measured by dynamic light scattering (DLS) on a Malvern Zetasizer Nano-ZS (Malvern, Westborough, MA, UK). The concentration of mRNA-LNPs was measured by an Invitrogen's Quant-iT Ribogreen RNA assay kit (Invitrogen, Eugene, OR, USA).



NanoAssemblr Ignite



**Quick Run** 2019-10-15 11:18 A

**Enter Parameters**

L	C	R
---	BD 3 ml	BD 1 ml
Dispense 0.00 ml at 0.00 ml/min	Dispense 0.75 ml at 9.00 ml/min	Dispense 0.25 ml at 3.00 ml/min

Flow rate ratio, C : R  
3.00 : 1.00

Total volume  
2.00 ml

Total flow rate  
12.00 ml/min

Dilution ratio, L : (C+R)  
:

Start waste  
L: 0.00 ml  
C: 0.34 ml  
R: 0.11 ml  
0.45 ml

End waste  
0.00 ml

Next

Settings on instrument

## Genovy-ILM reagents (cytiva)

Component	Mol%
PNI Ionizable Lipid	50
DSPC	10
Cholesterol	37.5
PNI stabilizer	2.5
PNI formulation buffer	

- Working Genvoy-ILM- 12.5mM in anhydrous ethanol
- Working RNA concentration- 0.17mg/ml in PNI formulation buffer
- It will be mixed in the ratio of 3:1 (working RNA solution:working lipid mix)
- At least 1.5ml of prepared RNA working solution and at least 0.5ml of prepared Genvoy-ILM solution so total 2ml of RNA-LNP formulation will be made.



## Characterization of mRNA-LNP formulation

- Particle size
  - Zeta potential
- } DLS analyzer (e.g. Malvern Zetasizer)



Malvern Zetasizer

### ➤ RNA concentration in encapsulation

Encapsulation efficiency can be calculated with

Quant-it™ RiboGreen Assay (Thermo Fisher Scientific)

Standard curve will be generated with serial dilution of working RNA



The concentration of unencapsulated RNA ( $U_{[RNA]}$ ) is measured by adding RiboGreen reagent to wells containing intact LNPs



The total amount of RNA ( $T_{[RNA]}$ ) in each sample is measured by adding RiboGreen reagent to wells containing LNPs that have been disrupted using 0.5% Triton X-100 in 1 × TE buffer



Encapsulated RNA ( $E_{[RNA]}$ ) is calculated by subtracting the unencapsulated concentration of RNA from the total concentration of RNA



$$E_{RNA} = T_{RNA} - U_{RNA}$$

$$EE\% = \frac{E_{RNA}}{T_{RNA}} \times 100$$

Excitation wavelength -485nm  
Emission wavelength- 528nm

### scientific reports

OPEN

**A careful look at lipid nanoparticle characterization: analysis of benchmark formulations for encapsulation of RNA cargo size gradient**

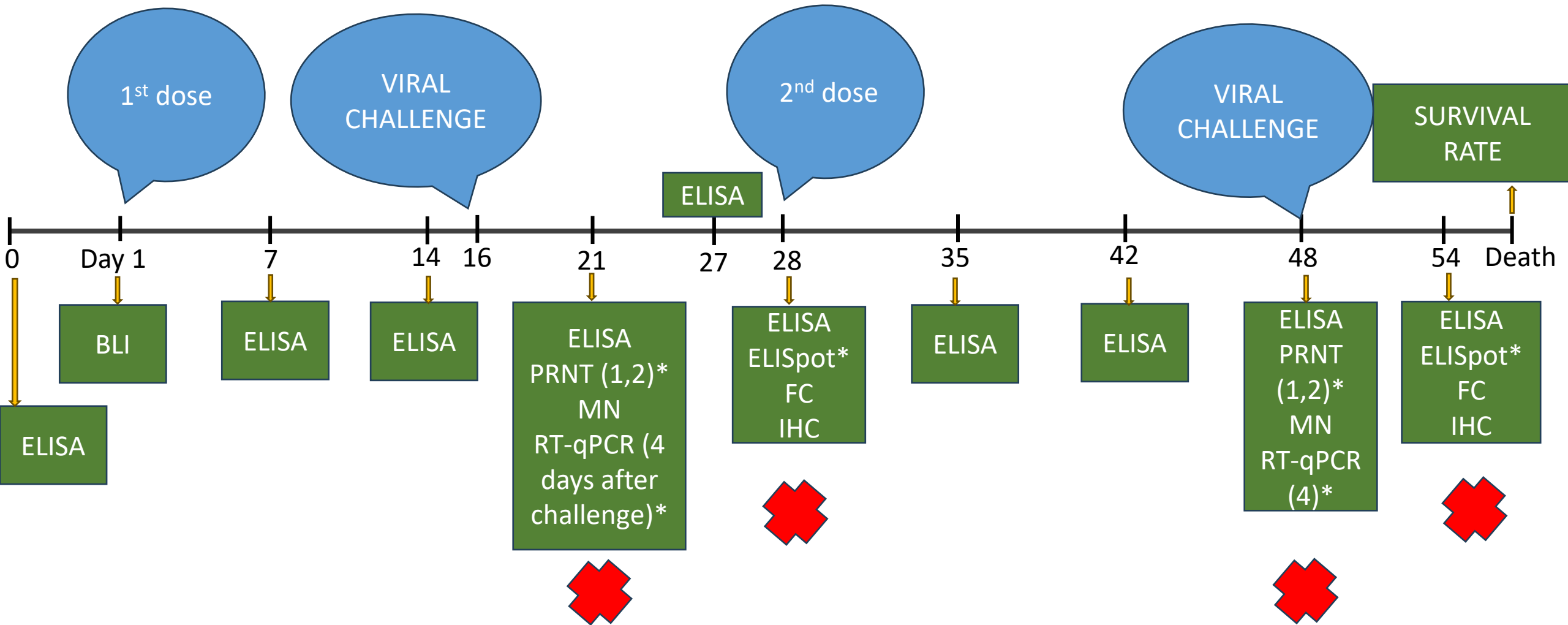
Gretchen B. Schober<sup>1</sup>, Sandra Story<sup>1</sup> & Dev P. Arya<sup>1,2</sup>

Check for updates

Sr. NO.	ASSAY	SAMPLE TYPE	REASON
1.	PLAQUE REDUCTION NEUTRALIZATION TEST (PRNT)	MICE SERUM SAMPLE  Organs (Brain)	To study Neutralizing Ab Titer  Virus Titer
2.	ELISA	MICE SERUM SAMPLE	To study Ab Titer (IgG)
3.	ELISpot assay	Splenocytes	To quantify various cytokines level (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, IL4, IL-6, CCL2, CCL5)*
4.	Flow cytometry (FC)	Splenocytes	To quantify CD4+ & CD8+ Level in spleen, Dendritic cell maturation
5.	MN Assay	MICE SERUM SAMPLE	To study microneutralization
6.	RT-qPCR	Tissue sample (Brain)	To quantify Viral Load
7.	BIOLUMINESCENCE (BLI)	NA	To study distribution and thermostability of LNP-mRNA
8.	H&E/IHC staining	Tissue sample	To study the neural cell distortion level (gliosis and cell morphogenesis)
9.	MICE SURVIVAL RATE	Observation till death	Overall effectiveness of vaccine

4 groups of mice – Negative control, Positive control, LNP, LNP-mRNA

- ✖ Day of animal sacrifice
- Assay to be done
- Day of Dosage



# Plaque reduction neutralization assay

Article | [Open access](#) | Published: 05 August 2020

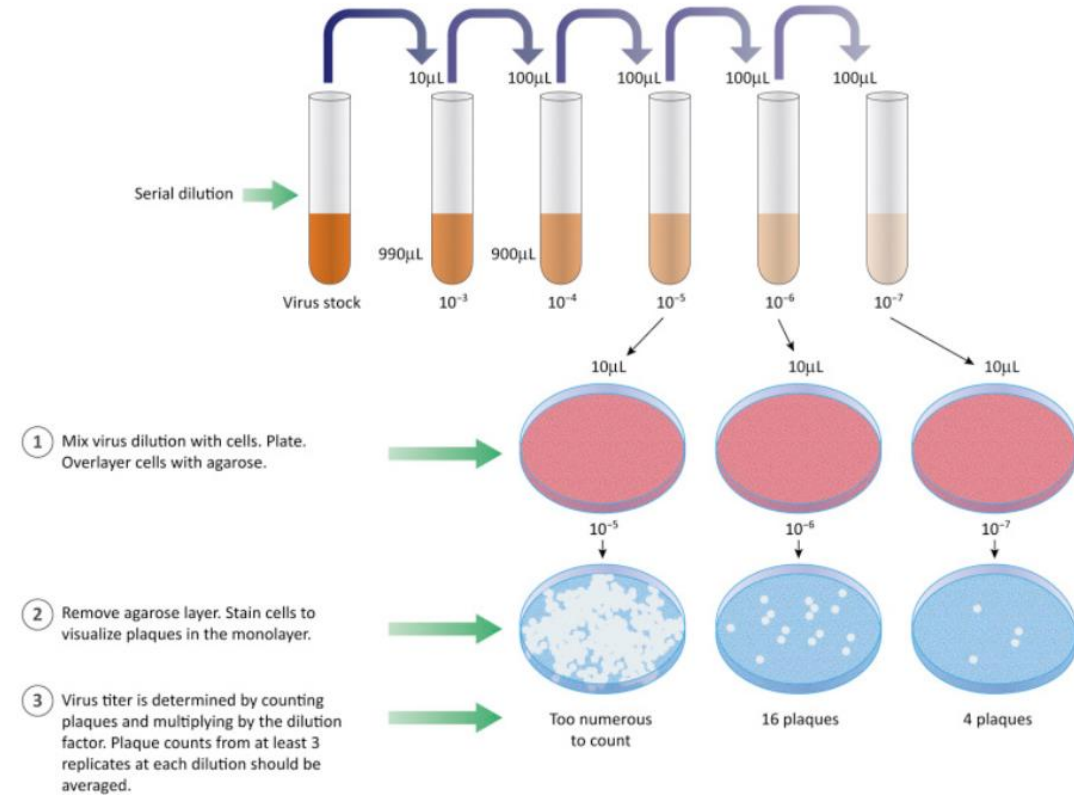
## A replication-defective Japanese encephalitis virus (JEV) vaccine candidate with NS1 deletion confers dual protection against JEV and West Nile virus in mice

[Na Li](#), [Zhe-Rui Zhang](#), [Ya-Nan Zhang](#), [Jing Liu](#), [Cheng-Lin Deng](#), [Pei-Yong Shi](#), [Zhi-Ming Yuan](#), [Han-Qing Ye](#)

& [Bo Zhang](#)

### Requirements

1. BHK21 cells/ Vero cells
2. Serum sample from mice
3. Methyl cellulose
4. Cell specific media
5. FBS
6. Formaldehyde
7. Cell culture plates



# FLOW CYTOMETR

## Requirements

1. FACSCalibur flow cytometer
2. Splenocytes from mice
3. 96 well microtiter plate(round bottom)
4. Peptide mixture (antigen specific)
5. Brefeldin A/GolgiPlug™
6. LIVE/DEAD™ Fixable Aqua Dead Cell Stain Kit (L34966, Invitrogen, MA, USA)
7. PBS
8. Fluorescently Conjugated Ab to CD4, CD8
9. Zombie UV3 fixable viability Kit ( STAIN DAED CELLS)

➤ [Cell](#). 2020 Sep 3;182(5):1271-1283.e16. doi: 10.1016/j.cell.2020.07.024. Epub 2020 Jul 23.

## A Thermostable mRNA Vaccine against COVID-19

Na-Na Zhang <sup>1</sup>, Xiao-Feng Li <sup>2</sup>, Yong-Qiang Deng <sup>2</sup>, Hui Zhao <sup>2</sup>, Yi-Jiao Huang <sup>2</sup>, Guan Yang <sup>3</sup>, Wei-Jin Huang <sup>4</sup>, Peng Gao <sup>5</sup>, Chao Zhou <sup>2</sup>, Rong-Rong Zhang <sup>2</sup>, Yan Guo <sup>2</sup>, Shi-Hui Sun <sup>2</sup>, Hang Fan <sup>2</sup>, Shu-Long Zu <sup>2</sup>, Qi Chen <sup>2</sup>, Qi He <sup>3</sup>, Tian-Shu Cao <sup>2</sup>, Xing-Yao Huang <sup>2</sup>, Hong-Ying Qiu <sup>2</sup>, Jian-Hui Nie <sup>4</sup>, Yuhang Jiang <sup>5</sup>, Hua-Yuan Yan <sup>5</sup>, Qing Ye <sup>2</sup>, Xia Zhong <sup>5</sup>, Xia-Lin Xue <sup>5</sup>, Zhen-Yu Zha <sup>5</sup>, Dongsheng Zhou <sup>2</sup>, Xiao Yang <sup>3</sup>, You-Chun Wang <sup>6</sup>, Bo Ying <sup>7</sup>, Cheng-Feng Qin <sup>8</sup>

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[nature](#) > [npj vaccines](#) > [articles](#) > article

Article | [Open access](#) | Published: 05 June 2023

## Analyzing immune responses to varied mRNA and protein vaccine sequences

[Hyeong-Jun Park](#), [Yoo-Jin Bang](#), [Sung Pil Kwon](#), [Woori Kwak](#), [Sang-In Park](#), [Gahyun Roh](#), [Seo-Hyeon Bae](#), [Jae-Yong Kim](#), [Hye Won Kwak](#), [Yongkwan Kim](#), [Soyeon Yoo](#), [Daeeun Kim](#), [Gyochang Keum](#), [Eun-Kyoung Bang](#) , [So-Hee Hong](#)  & [Jae-Hwan Nam](#) 

[npj Vaccines](#) **8**, Article number: 84 (2023) | [Cite this article](#)



# ELISpot Assay

[Bio Protoc.](#) 2017 Jun 5; 7(11): e2302.

Published online 2017 Jun 5. doi: [10.21769/BioProtoc.2302](https://doi.org/10.21769/BioProtoc.2302)

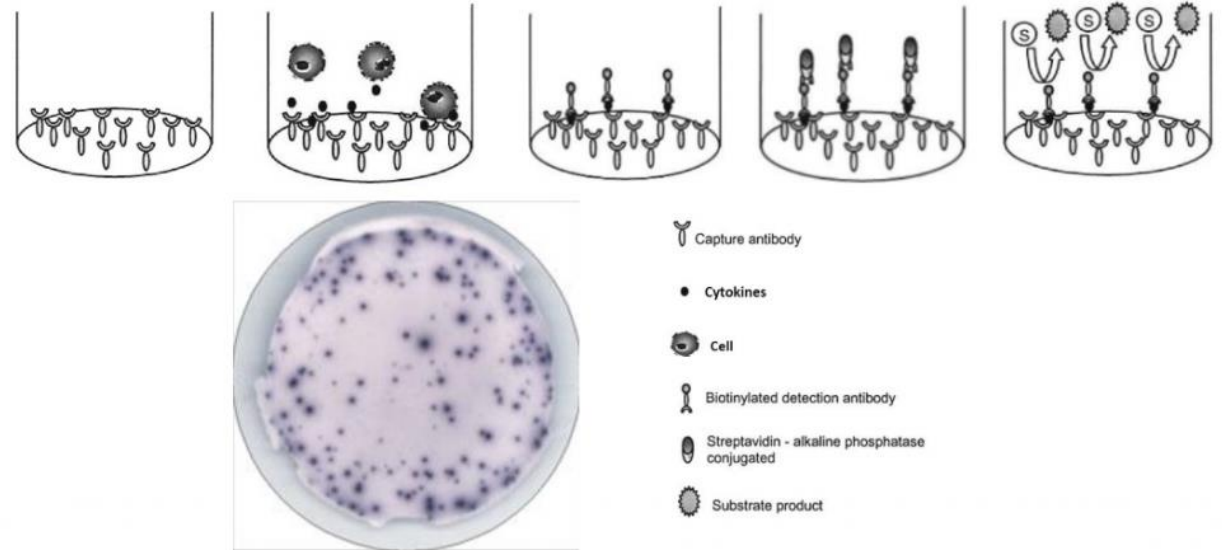
## ELISPOT Assay to Measure Peptide-specific IFN- $\gamma$ Production

[Michelle N. Wykes](#)<sup>1,\*</sup> and [Laurent Renia](#)<sup>2,3,\*</sup>

### Requirements

1. Multiscreen <sub>HTS</sub>-IP plates (PVDF membrane) (Merck, catalog number: MSIPS4510)
2. BSA, FCS
3. Rat anti-mouse IFN $\gamma$  mAb
4. AEC substrate/chromogen
5. Streptavidin-horseradish peroxidase
6. Anti-IFN Ab
7. PBS
8. Peptides specific to antigen
9. Microscope or Automated ELISpot reader

### ELISpot Procedure



# ELISA

[J Immunol.](#) Author manuscript; available in PMC 2022 Jun 1.

PMCID: PMC8165000

*Published in final edited form as:*

NIHMSID: NIHMS1688200

[J Immunol. 2021 Jun 1; 206\(11\): 2596–2604.](#)

PMID: [33972374](#)

Published online 2021 May 10. doi: [10.4049/jimmunol.2100054](#)

## Cellular and humoral immune responses in mice immunized with Vaccinia virus expressing the SARS-CoV-2 spike protein

[Jake C. Harbour,<sup>\\*</sup>](#) [Zoe L. Lyski,<sup>#\\*</sup>](#) [John B. Schell,<sup>#†</sup>](#) [Archana Thomas,<sup>#‡</sup>](#) [William B. Messer,<sup>\\*§¶</sup>](#) [Mark K. Slifka,<sup>‡</sup>](#) and [Jeffrey C. Nolz<sup>\\*||#</sup>](#)

### Requirements




1. ELISA plates
2. Protein specific to antigen
3. PBS
4. Tween
5. Serum sample from mice
6. Citrate buffer
7. *o*-phenylenediamine
8. anti-mouse IgG-HRP
8. Hydrogen peroxide
9. HCL
10. Plate Reader

# MN Assay

[nature](#) > [npj vaccines](#) > [articles](#) > [article](#)

Article | [Open access](#) | Published: 05 June 2023

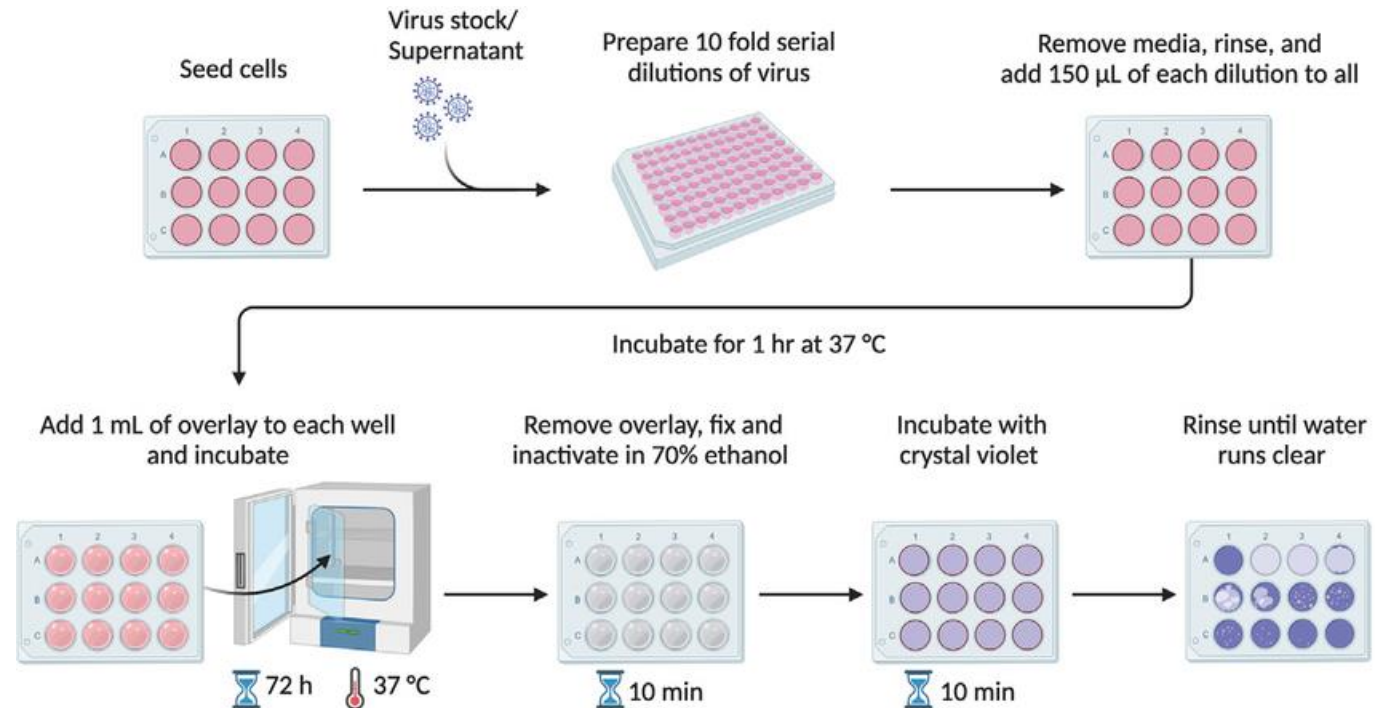
## Analyzing immune responses to varied mRNA and protein vaccine sequences

[Hyeong-Jun Park](#), [Yoo-Jin Bang](#), [Sung Pil Kwon](#), [Woori Kwak](#), [Sang-In Park](#), [Gahyun Roh](#), [Seo-Hyeon Bae](#),  
[Jae-Yong Kim](#), [Hye Won Kwak](#), [Yongkwan Kim](#), [Soyeon Yoo](#), [Daegeun Kim](#), [Gyochang Keum](#), [Eun-Kyoung Bang](#) , [So-Hee Hong](#)  & [Jae-Hwan Nam](#) 

[npj Vaccines](#) **8**, Article number: 84 (2023) | [Cite this article](#)

### Requirements

1. Crystal violet stain
2. Formaldehyde
3. Serum sample from mouse
4. 96-well plate
5. Vero cells
6. Cell specific media and FBS
7. Viral suspension



# **Bioluminescence**

(to study in-vivo distribution of LNP-mRNA and thermostability)

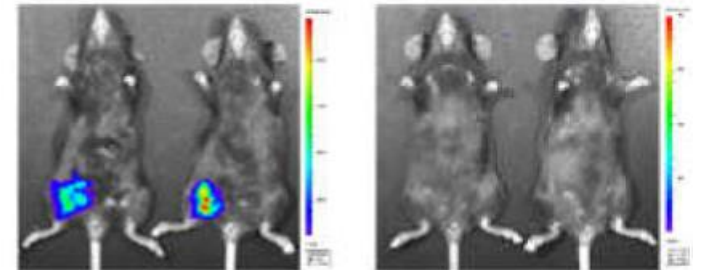
➤ [Cell. 2020 Sep 3;182\(5\):1271-1283.e16. doi: 10.1016/j.cell.2020.07.024. Epub 2020 Jul 23.](#)

## **A Thermostable mRNA Vaccine against COVID-19**

Na-Na Zhang<sup>1</sup>, Xiao-Feng Li<sup>2</sup>, Yong-Qiang Deng<sup>2</sup>, Hui Zhao<sup>2</sup>, Yi-Jiao Huang<sup>2</sup>, Guan Yang<sup>3</sup>, Wei-Jin Huang<sup>4</sup>, Peng Gao<sup>5</sup>, Chao Zhou<sup>2</sup>, Rong-Rong Zhang<sup>2</sup>, Yan Guo<sup>2</sup>, Shi-Hui Sun<sup>2</sup>, Hang Fan<sup>2</sup>, Shu-Long Zu<sup>2</sup>, Qi Chen<sup>2</sup>, Qi He<sup>3</sup>, Tian-Shu Cao<sup>2</sup>, Xing-Yao Huang<sup>2</sup>, Hong-Ying Qiu<sup>2</sup>, Jian-Hui Nie<sup>4</sup>, Yuhang Jiang<sup>5</sup>, Hua-Yuan Yan<sup>5</sup>, Qing Ye<sup>2</sup>, Xia Zhong<sup>5</sup>, Xia-Lin Xue<sup>5</sup>, Zhen-Yu Zha<sup>5</sup>, Dongsheng Zhou<sup>2</sup>, Xiao Yang<sup>3</sup>, You-Chun Wang<sup>6</sup>, Bo Ying<sup>7</sup>, Cheng-Feng Qin<sup>8</sup>

### **Requirements**

1. Luciferase cassette in mRNA
2. Luciferase substrate
3. IVIS Spectrum instrument (for in-vivo)
4. living image 3.0 software



# RT-qPCR

## Requirements

1. Tissue lyser
2. Reverse transcription kit/ABscript II cDNA First Strand Synthesis kit
3. RNA extraction kit /QIAamp Viral RNA Mini Kit
4. QrtPCR kit / One Step PrimeScript RTPCR Kit
5. JEV specific primers and probes



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## mRNA Vaccines Encoding the HA Protein of Influenza A H1N1 Virus Delivered by Cationic Lipid Nanoparticles Induce Protective Immune Responses in Mice

by Xinyu Zhuang<sup>1,†</sup> ✉, Yanxin Qi<sup>2,†</sup> ✉ , Maopeng Wang<sup>3</sup> ✉, Ning Yu<sup>4</sup> ✉, Fulong Nan<sup>5</sup> ✉, He Zhang<sup>1</sup> ✉, Mingyao Tian<sup>1</sup> ✉, Chang Li<sup>1</sup> ✉, Huijun Lu<sup>1</sup> ✉ and Ningyi Jin<sup>1,\*</sup> ✉



# H&E /IHC staining



## Requirements

1. Ethanol
2. Paraffin
3. H&E stain
4. Xylene, H<sub>2</sub>O<sub>2</sub>, Citrate buffer
5. VectaStain ABC kit (Vector Laboratories, Burlingame, CA).
6. Primary rabbit anti-occludin polyclonal antibody (pAb; Santa Cruz, Santa Cruz, CA), rabbit anti-claudin-5 pAb (Invitrogen), and rabbit anti-ZO-1 pAb (Sigma)
7. Diaminobenzidine
8. Fluorescence Microscope

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## Analyzing immune responses to varied mRNA and protein vaccine sequences

[Hyeong-Jun Park](#), [Yoo-Jin Bang](#), [Sung Pil Kwon](#), [Woori Kwak](#), [Sang-In Park](#), [Gahyun Roh](#), [Seo-Hyeon Bae](#), [Jae-Yong Kim](#), [Hye Won Kwak](#), [Yongkwan Kim](#), [Soyeon Yoo](#), [Daeyeun Kim](#), [Gyochang Keum](#), [Eun-Kyoung Bang](#) , [So-Hee Hong](#)  & [Jae-Hwan Nam](#) 

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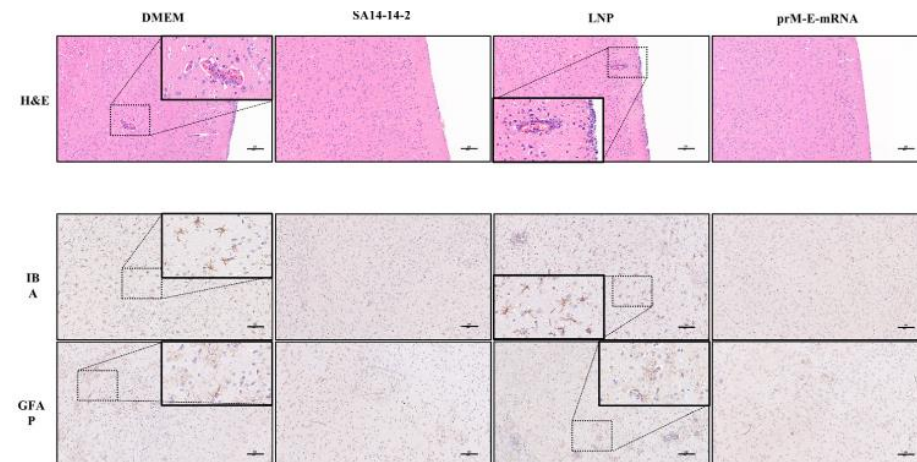
PMCID: PMC4442524

Published online 2015 Mar 11. doi: [10.1128/JVI.00143-15](#)

PMID: [25762733](#)

Viral Infection of the Central Nervous System and Neuroinflammation Precede Blood-Brain Barrier Disruption during Japanese Encephalitis Virus Infection

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Sr. NO.	ASSAY	SAMPLE TYPE	REASON
1.	PLAQUE REDUCTION NEUTRALIZATION TEST (PRNT)	MICE SERUM SAMPLE  Organs (Brain)	To study Neutralizing Ab Titer  Virus Titer
2.	ELISA	MICE SERUM SAMPLE	To study Ab Titer (IgG)
3.	ELISpot assay	Splenocytes	To quantify various cytokines level (IFN- $\gamma$ , TNF- $\alpha$ , IL-2, IL4, IL-6, CCL2, CCL5)*
4.	Flow cytometry (FC)	Splenocytes	To quantify CD4+ & CD8+ Level in spleen, Dendritic cell maturation, TH1 and TH2 response
6.	RT-qPCR	Tissue sample (Brain)	To quantify Viral Load
7.	BIOLUMINESCENCE (BLI)	NA	To study distribution and thermostability of LNP-mRNA
9.	MICE SURVIVAL RATE	Observation till death	Overall effectiveness of vaccine