of the process validation master plan, manufacturing processes conducted PPQ activities with CX-024414 mRNA at a 10 L IVT scale, and mRNA-1273 LNP at a mRNA input scale. The goal is to establish a continuous process validation strategy for the mRNA-1273 manufacturing processes. Phase 3 clinical supplies have been and are being manufactured with representative processes. PPQ activities will be carried out such that the understanding of critical process parameters for mRNA-1273 manufacturing processes, including CX-024414, mRNA-1273 LNP and mRNA-1273 Drug Product will be gained.

# 3.2.P.2.3.7.1 Analytical Assessment for Clinical Trial Material (Phase I through Vial Process Performance Qualification)

Given the manufacturing process changes that have occurred for mRNA-1273 Drug Product through the course of development (see manufacturing history discussion in Section 3.2.P.2.3.4), analytical data have been collected and assessed together in order to gain assurance that the process continues to be robust and consistently produce high-quality material. Release, stability, extended characterization, and impurity characterization data sets have been evaluated and were found to demonstrate that quality attributes of the material are highly similar. Lot release and stability data generated and compared across processes are discussed in Section 3.2.P.2.3.7.6, whereas this subsection focuses on the extended characterization, and impurity characterization. Forced degradation data will be included when available. These data were generated with representative lots from different manufacturing processes:

- Phase 1/2 clinical lots 8520100101, 8520100102, 8520100103, and 8520100104
- Phase 3 clinical lots 6007520001, 6007520002, and 6007520003 (0.20 mg/mL, 5.0 mL fill volume, Ompi 10R vial,
- PPQ lots 6007520004, 6007520005, and 6007520006 (0.20 mg/mL, 5.0 mL fill volume, Ompi 10R vial,

Samples were analyzed in a side-by-side format whenever possible and were evaluated for mRNA-1273 Drug Product physico-chemical properties, particle size, and impurities. A study of the Phase 1/2 and Phase 3 lots was initially executed to evaluate product similarity irrespective of process and scale changes. PPQ lots were later executed to evaluate process consistency at the vial (0.20 mg/mL mRNA-1273) scale. In addition to assessing the various clinical lots against product release criteria, the lots were also examined by a set of extended characterization assays listed in Table 30. The table also provides a summary of the attributes assessed by the characterization assays. There are no pre-defined acceptance criteria for these assays, but the results show that the lots analyzed are similar irrespective of the manufacturing process and scale. The data generated will be used to inform and establish appropriate acceptance criteria for future process consistency and product comparability studies.

In its entirety, the evaluation of analytical data across release, stability, and characterization of mRNA-1273 Drug Product GMP and PPO lots demonstrate a high degree of product quality Attribute Assessment

Testing panel for exploratory characterization of mRNA-1273 Drug Product from PPQ Lots 6007520004, 6007520005, and 6007520006

uct Attribute Method Document #

istribution Nanoparticle tracking analysis DPTM-0039
istribution Asymmetric flow field-flow fractionation

particle count similarity for the manufacturing process and scale changes that occurred throughout process development. The 3 PPQ lots are also highly similar and demonstrate process consistency at the 3,000 vial (0.20 mg/mL mRNA-1273) scale, as summarized in Section 3.2.P.2.3.7.3.

**Table 30:** 

# A.

Product Attribute	Method	Document #	<b>Description</b>
LNP size distribution	Nanoparticle tracking analysis	DPTM-0039	High-resolution LNP size distribution
LNP size distribution	Asymmetric flow field-flow fractionation	DPTM-0103	Fractionation coupled with in-line MALS detection for size determination
Sub-visible particle counts	Coulter counter	DPTM-0035	Measurement of sub-visible particle counts in the grange
Sub-visible particle counts and morphology	Flow microscopy	DPTM-0115	Measurement of sub-visible particle counts and morphology in the range
LNP surface characterization	Isothermal titration calorimetry	DPTM-0119	to LNP surface
LNP charge	Zeta potential	DPTM-0118	Average LNP charge
LNP charge distribution	Capillary isoelectric focusing	DPTM-0068	LNP pI distribution and polydispersity
LNP structure	Dye permeation kinetics	DPTM-0127	Qualitative characterization of LNP surface and encapsulation state based on thionine permeation kinetics
LNP density	Density gradient ultracentrifugation	DPTM-0104	Qualitative assessment of LNP density heterogeneity
LNP surface characterization	Hydrophobic interaction chromatography	DPTM-0096	Qualitative characterization of LNP surface hydrophobicity
LNP surface characterization	Esterase kinetics	DPTM-0130	Qualitative method for distribution
mRNA encapsulation	RiboGreen fluorescence	SOP-0298	Fluorescence based method for mRNA encapsulation

### B. Testing panel for characterization of mRNA-1273 DP from Phase 1/2 and Phase 3 Lots 8520100101, 8520100102, 8520100103, 8520100104, 6007520001, 6007520002, and 6007520003

<b>Product Attribute</b>	Method	Document #	Description
mRNA encapsulation		DPTM-0073	
LNP size distribution	Nanoparticle tracking analysis	DPTM-0039	High-resolution LNP size distribution
LNP size distribution	Asymmetric flow field flow fractionation	DPTM-0103	Fractionation coupled with in-line MALS detection for size determination
Sub-visible particles	Coulter counter	DPTM-0035	Sub-visible particle counts
SVP counts and morphology	Flow microscopy	DPTM-0115	Measurement of SVP counts and morphology in the grange range
LNP surface characterization	Isothermal titration calorimetry	DPTM-0119	to LNP surface
LNP charge	Zeta potential	DPTM-0118	Average LNP charge
LNP charge distribution	Capillary isoelectric focusing	DPTM-0068	LNP pI distribution

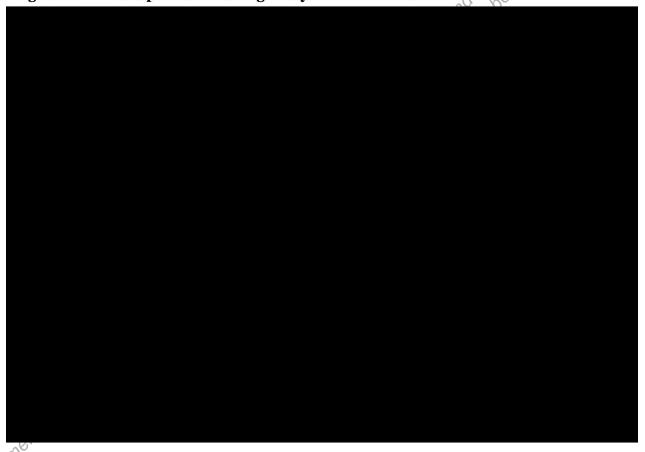
Abbreviations: LNP = lipid nanoparticle; MALS = multi-angle light scattering; pI = isoelectric point(s); PPQ = process performance qualification

The RiboGreen method for RNA encapsulation measurement has been used as a release method to support early development and initial PPQ lots of the mRNA-1273 vaccine.

### 3.2.P.2.3.7.1.1 Nanoparticle Tracking Analysis

Nanoparticle tracking analysis is a technique designed to measure sub-micron particle distributions in liquid solutions. This analysis characterizes nanoparticles from 10-1000 nm in solution. Each particle is individually and simultaneously analyzed by direct observation and measurement of diffusion events. This particle-by-particle methodology produces high-resolution results for nanoparticle size distribution and concentration. Size distribution plots for each mRNA-1273 Drug Product lot are shown in Figure 5. The mode of the size distributions ranged from (Table 31). The breadth of the size distributions, as estimated by full width at half maximum (span), ranged from with an LNP size range broadly covering. These results show similar size distributions across lots and processes.

Figure 5: Nanoparticle Tracking Analysis Size Distribution



**Table 31:** Nanoparticle Tracking Analysis Results

Lot	
85201100101	
85201100102	
85201100103	
85201100104	
6007520001	
6007520002	
6007520003	
6007520004	
6007520005	
6007520006	

# 3.2.P.2.3.7.1.2 Asymmetric Flow Field-Flow Fractionation Analysis

Asymmetric flow field-flow fractionation (aF4) with multi-angle light scattering was employed as an orthogonal measurement of the mRNA-1273 size distributions. aF4 is a one-phase separation that uses a perpendicular flow against a membrane (cross-flow) in conjunction with a channel flow parallel to the membrane to fractionate samples based on their diffusion behavior. The channel flow gives a parabolic profile and the perpendicular flow drives macromolecules toward the boundary layer of the membrane. Diffusion related to Brownian motion moves smaller particles with higher diffusion rates in the channel where longitudinal flow is faster, eluting smaller particles more quickly. Multi-angle light scattering detection enables the particle radius of gyration, which is related to particle mass, to be determined for peaks in the aF4 separation. Results for each mRNA-1273 Drug Product lot are presented in Table 32. The radius of gyration ranged from with polydispersity ranging from All lots demonstrated similar size distribution and polydispersity across processes and scales.

Table 32: Asymmetric Flow Field-Flow Fractionation Results

Lot	
85201100101	
85201100102	
85201100103	
85201100104	
6007520001	
6007520002	
6007520003	
6007520004	
6007520005	
6007520006	

Abbreviations: Mn = number average molecular weight; Mw = weight average molecular weight; Rg = radius of gyration

### **3.2.P.2.3.7.1.3** Coulter Counter

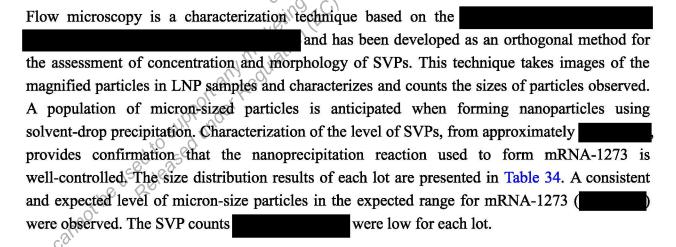
Coulter counter analysis was used to assess sub-visible particle (SVP) content in the range of This technique utilizes electro-zone sensing to determine the size and concentration of particles in the stated size range. Sample in an electrolyte is drawn through an aperture connecting 2 fluid chambers. Particles passing through the aperture are detected by a change in impedance. Since impedance is proportional to the volume of the particle, the size of each particle can be calculated. SVP counts for each mRNA-1273 Drug Product lot are shown in Table 33. SVP counts ranged from particles/mg for all lots, demonstrating consistency across processes.

**Table 33:** Coulter Counter Results

Lot	
85201100101	
85201100102	
85201100103	
85201100104	
6007520001	
6007520002	
6007520003	
6007520004	
6007520005	
6007520006	

Abbreviations: RSD = relative standard deviation; SVP = sub-visible particle

# **3.2.P.2.3.7.1.4** Flow Microscopy



**Table 34:** Flow Microscopy Particle Size Distribution Results

Lot	
	(Particles/mg)
85201100101	
85201100102	
85201100103	
85201100104	
6007520001	
6007520002	
6007520003	
6007520004	
6007520005	
6007520006	

# 3.2.P.2.3.7.1.5 Isothermal Titration Calorimetry

Isothermal titration calorimetry is a calorimetric technique for thermodynamic analysis of binding reactions. When binding occurs, heat is either absorbed or released and this is measured by a sensitive calorimeter during gradual titration of the ligand into the sample cell containing the species of interest. This methodology is used to measure the interaction between The binding affinities, described by the

dissociation constant (Kd), and the binding stoichiometries are reported for each lot in Table 35. The observed binding affinities and binding stoichiometries were similar across lots.

Table 35: Isothermal Calorimetry Results

Lot
85201100101
85201100102
85201100103
85201100104
6007520001
6007520002
6007520003
6007520004
6007520005
6007520006

Abbreviations: Kd = dissociation constant; M = binding stoichiometry (number of

### 3.2.P.2.3.7.1.6 Zeta Potential

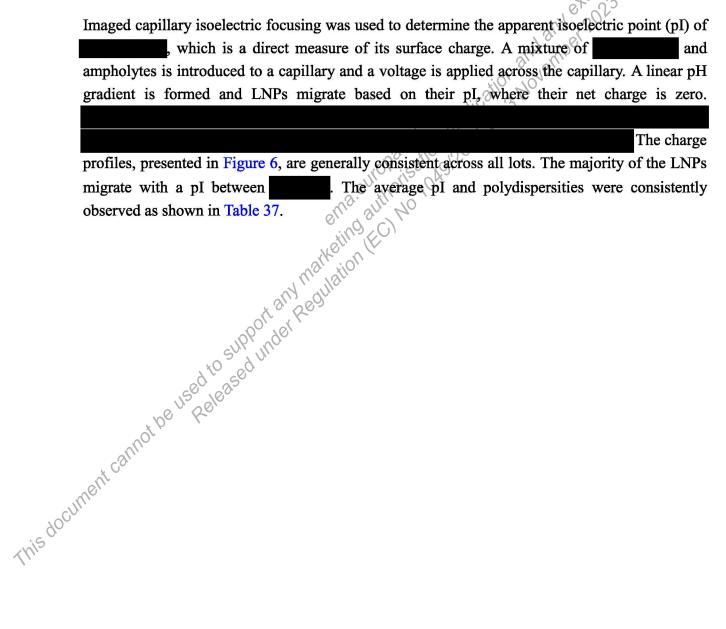
Zeta potential is an indirect measurement of LNP surface charge calculated from the electrophoretic mobility of the particle. The zeta potential of mRNA-1273 was measured using a instrument equipped with

. Zeta potential results shown in Table 36 demonstrate similar surface charge and all mRNA-1273 Drug Product lots are considered approximately neutral.

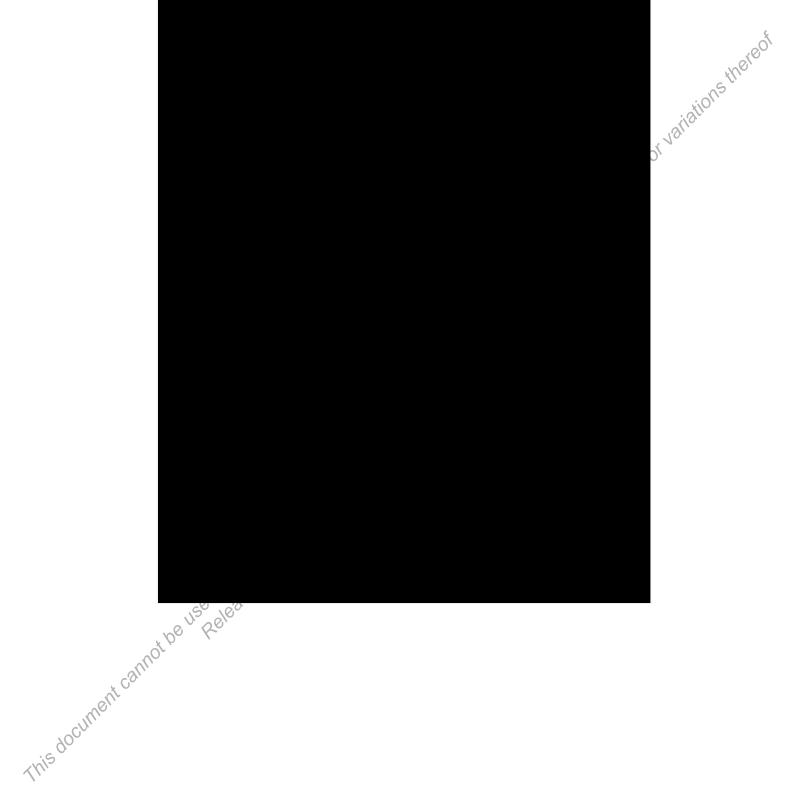
Table 36: Zeta Potential Results

Lot		
85201100101		
85201100102		
85201100103		
85201100104		
6007520001		
6007520002		
6007520003		
6007520004		
6007520005		
6007520006		

## 3.2.P.2.3.7.1.7 Capillary Isoelectric Focusing







**Table 37:** Capillary Isoelectric Focusing Results

Lot	
85201100101	
85201100102	
85201100103	
85201100104	
6007520001	
6007520002	
6007520003	
6007520004	
6007520005	
6007520006	

Abbreviations: pI = isoelectric point

### 3.2.P.2.3.7.1.8 Dye Permeation Kinetics

Dye permeation assay was used to measure the rate at which the cationic phenothiazinium dye, thionine, permeates the LNP from PPQ batches. This is an optical spectroscopic method that monitors with time the change in visible absorption that occurs due to thionine binding to encapsulated mRNA. The kinetic profile of this assay represents a biophysical signature that reports on LNP surface physical properties and mRNA encapsulation state, reported as a first order rate constant. Results shown for each PPQ lot in Table 38 demonstrate similar permeation kinetics.

**Table 38:** Dye Permeation Kinetic Results

Lot				
6007520004				
6007520005	1			
6007520006	7/7			

Abbreviation: CV = coefficient of variation

### 3.2.P.2.3.7.1.9 Density Gradient Ultracentrifugation

Separation using works on the principle of isopycnic separation: an LNP particle of a density will sink during centrifugation until a position is reached where the density of the surrounding solution is exactly the same as the density of the particle. was used as density gradient medium, consisting of Density gradient ultracentrifugation was used to characterize the density distribution of mRNA-1273 Drug Product PPQ batches. A sample tube was placed in an box, to which a was attached. The resulting images (Figure 7) were converted into chromatograms based on pixel intensity and the slope change was monitored. The slope change results shown in Table 39 demonstrate consistent density profiles for these three PPQ batches.

Figure 7: Density Gradient Chromatograms

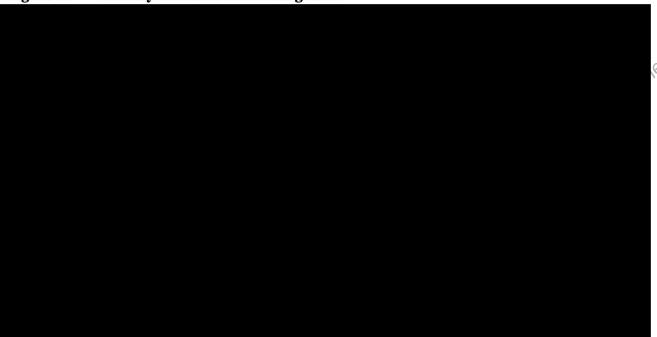


Table 39: Density Gradient Ultracentrifugation Results

Lot	
6007520004	
6007520005	
6007520006	

# 3.2.P.2.3.7.1.10 Hydrophobic Interaction Chromatography

Hydrophobic interaction chromatography (HIC) was used to characterize mRNA-1273 Drug Product PPQ batches. It uses the retention mechanism of "salting out" biologics onto a hydrophobic stationary phase. Salt is then removed, allowing for the elution of the biologic based on its hydrophobicity. LNP samples were loaded onto a column at a flow rate of with UV absorbance monitored at 260 nm. Elution was obtained by

Three chromatographic regions are reported as relative percent areas:

HIC results generated for these three mRNA-1273 Drug Product PPQ batches are reported in Table 40. The relative percent peak areas for each chromatographic region were consistent among these PPQ lots which demonstrate similar hydrophobic profiles.

mRNA-1273

**Hydrophobic Interaction Chromatography Results** 

Lot	
6007520004	
6007520004 6007520005	3,01
6007520006	The.

### **3.2.P.2.3.7.1.11** Esterase Kinetics



across the time course for each PPQ lot is presented in Figure 8 and Table 41. Results demonstrate consistent degradation behavior across lots.

Figure 8:

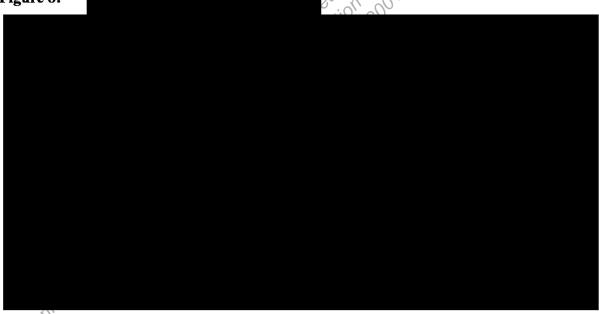
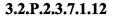
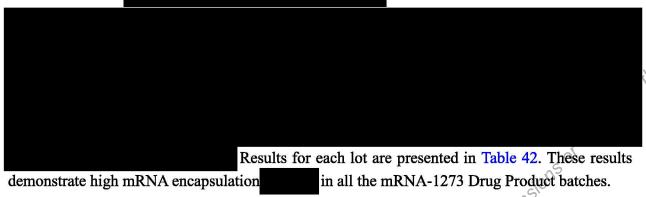


Table 41: **Esterase Kinetics Results** 

4.3		
, ocy	Lot	
:500	6007520004	
(VI)	6007520005	
	6007520006	





**Table 42: Encapsulation Results** 

Lot	Encapsulation (%)	CV (%)
85201100101		
85201100102		
85201100103		
85201100104		
6007520001		
6007520002		
6007520003		

Abbreviations: CV = coefficient of variation

## 3.2.P.2.3.7.1.13

RiboGreen Encapsulation RiboGreen State RiboGreen A fluorescence-based RiboGreen assay was used as a release assay to determine the encapsulation efficiency of mRNA in mRNA-1273 LNP. RiboGreen is a sensitive RNA quantitation reagent with a wide linear dynamic range. RiboGreen can bind to free RNA in solution but is not able to bind to mRNA encapsulated in an LNP. Upon binding to RNA, the Ribogreen dye undergoes fluorescent enhancement and an increase in quantum yield, which results in an emitted quantifiable fluorescent signal that is linearly dependent on available RNA. Concentration determinations of the free mRNA and total mRNA content are measured for each sample to calculate encapsulation efficiency. RNA encapsulation results for the PPQ lots are shown in Table 43. These results agree with the and demonstrate similarly high and consistent mRNA encapsulation for the PPQ lots. The Sponsor is intended to validate as a release assay due to the assay robustness, simplicity, and the usage of a USP-certified assay reagent. The Sponsor will continue to determine encapsulation efficiency of mRNA in mRNA-1273 LNP by RiboGreen assay as an orthogonal characterization assay.

**RiboGreen RNA Encapsulation Results Table 43:** 

Lot	Encapsulation (%)
6007520004	
6007520005	
6007520006	

# 3.2.P.2.3.7.2 Analytical Assessment Across Process Performance Qualifications Vial and Vial)

Given the manufacturing site and scale changes that have occurred for mRNA-1273 Drug Product (DP) through the course of development (see manufacturing history discussion in Section 3.2.P.2.3.4), analytical data have been collected and assessed together in order to gain assurance that the process continues to be robust and consistently produce high-quality material. Release, stability, extended characterization, and impurity characterization data sets have been evaluated and were found to demonstrate that quality attributes of the material are highly similar. Lot release and stability data generated and compared across processes are discussed in Section 3.2.P.2.3.7.6, whereas this subsection focuses on the extended characterization, and impurity characterization. Forced degradation data will be included when available. These data were generated with representative lots from different manufacturing processes:

- ModernaTX, Inc. (Norwood, MA) PPQ lots 6007520004, 6007520005, and 6007520006
   (0.20 mg/mL, 5.0 mL fill volume, Ompi 10R vial, vials)
- Catalent Biologics, LLC (Bloomington, IN) PPQ lots 057G20 (Moderna Lot 6007320001), 062G20 (Moderna Lot 6007320002), 001H20 (Moderna Lot 6007320003) (0.20 mg/mL, 6.3 mL fill volume, Ompi 10R vial, vials)

Samples were analyzed and evaluated for mRNA-1273 Drug Product physico-chemical properties, particle size, and impurities. Initial PPQ lots were executed to evaluate process consistency at the vial (0.20 mg/mL mRNA-1273) scale. Subsequent PPQ lots were executed at the commercial manufacturing facility (Catalent Biologics, LLC, Bloomington, IN) at a vial (0.20 mg/mL mRNA-1273) scale. In addition to assessing all PPQ lots against product release criteria, the lots were also examined by a set of extended characterization assays listed in Table 44. The table also provides a summary of the attributes assessed by the characterization assays. There were no pre-defined acceptance criteria for the characterization assays, but the results show that the lots analyzed are similar irrespective of the manufacturing site and scale.

In its entirety, the evaluation of analytical data across release, stability, and characterization of mRNA-1273 Injection DP PPQ lots demonstrate a high degree of product quality similarity for the manufacturing site and scale changes that occurred. The PPQ lots are also highly similar and demonstrate process consistency at the vial (0.20 mg/mL mRNA-1273) and vial (0.20 mg/mL mRNA-1273) scales, as summarized in Section 3.2.P.2.3.7.3.

Table 44: Attribute Assessment for PPQ Lots

Product Attribute	Method	Document #	Description
LNP size distribution	Nanoparticle tracking analysis	DPTM-0039	High-resolution LNP size distribution
LNP size distribution	Asymmetric flow field-flow fractionation	DPTM-0103	Fractionation coupled with in-line MALS detection for size determination
Sub-visible particle counts	Coulter counter	DPTM-0035	Measurement of sub-visible particle counts in the grant range
Sub-visible particle counts and morphology	Flow microscopy	DPTM-0115	Measurement of sub-visible particle counts and morphology in the
LNP surface characterization	Isothermal titration calorimetry	DPTM-0119	to LNP surface
LNP charge	Zeta potential	DPTM-0118	Average LNP charge
LNP charge distribution	Capillary isoelectric focusing	DPTM-0068	LNP pI distribution and polydispersity
LNP structure	Dye permeation kinetics	DPTM-0127	Qualitative characterization of LNP surface and encapsulation state based on thionine permeation kinetics
LNP density	Density gradient ultracentrifugation	DPTM-0104	Qualitative assessment of LNP density heterogeneity
LNP surface characterization	Hydrophobic interaction chromatography	DPTM-0096	Qualitative characterization of LNP surface hydrophobicity
LNP surface characterization	Esterase kinetics	DPTM-0130	Qualitative method for distribution
mRNA encapsulation	RiboGreen fluorescence (a)	SOP-0298	Fluorescence based method for mRNA encapsulation

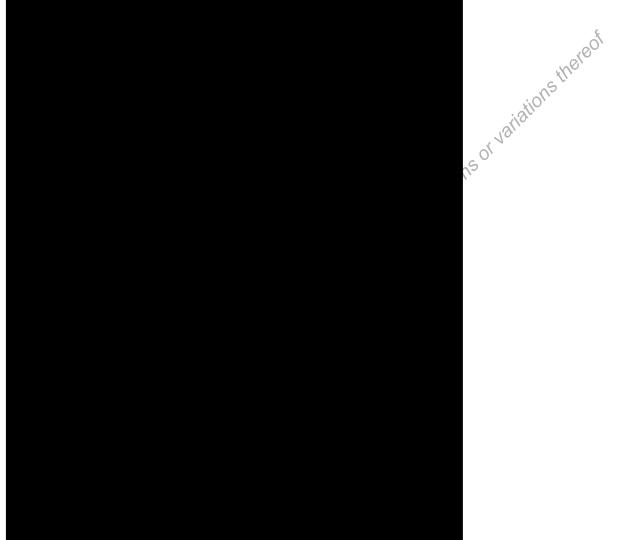
Abbreviations: LNP = lipid nanoparticle; MALS = multi-angle light scattering; pI = isoelectric point(s); pPQ = process performance qualification

# 3.2.P.2.3.7.2.1 Nanoparticle Tracking Analysis

Nanoparticle tracking analysis is a technique designed to measure sub-micron particle distributions in liquid solutions. This analysis characterizes nanoparticles from 10 - 1000 nm in solution. Size distribution plots for each mRNA-1273 Drug Product PPQ lot are shown in Figure 9. The mode of the size distributions ranged from (Table 45). The breadth of the size distributions, as estimated by full width at half maximum, ranged from m with an LNP size range broadly covering the size distributions across lots and processes.

a) The RiboGreen method for RNA encapsulation measurement had been used as a release method to support early development and initial PPQ lots of the mRNA-1273 vaccine.

Figure 9: Nanoparticle Tracking Analysis Size Distribution



Abbreviation: LNP = lipid nanoparticle

Table 45: Nanoparticle Tracking Analysis Results

Lot C	
6007520004	
6007520005	
6007520006	
057G20	
062G20	
001H20	

### 3.2.P.2.3.7.2.2 Asymmetric Flow Field-Flow Fractionation Analysis

Asymmetric flow field-flow fractionation (aF4) with multi-angle light scattering was employed as an orthogonal measurement of the mRNA-1273 size distributions. Results for each mRNA-1273 Drug Product PPQ lot are presented in Table 46. The radius of gyration ranged from with polydispersity ranging from All lots demonstrated similar size distribution and polydispersity across processes and scales.

**Table 46:** Asymmetric Flow Field-Flow Fractionation Results

Lot	
6007520004	
6007520005	
6007520006	
057G20	
062G20	
001H20	

Abbreviations: Mn = number average molecular weight; Mw = weight average molecular weight; Rg or Rz = radius of gyration,

### **3.2.P.2.3.7.2.3** Coulter Counter

Coulter counter analysis was used to assess sub-visible particle (SVP) content in the range of Sub-visible particle (SVP) counts for each mRNA-1273 Drug Product lot are shown in Table 47. SVP counts ranged from for all lots, demonstrating consistency across processes.

Table 47: Coulter Counter Results

Lot	
6007520004	~
6007520005	10
6007520006	ally a
057G20	it Pe
062G20	200.76/
001H20	16,100

Abbreviations: RSD = relative standard deviation; SVP = sub-visible particle

### 3.2.P.2.3.7.2.4 Flow Microscopy

Flow microscopy is technique that takes images of the magnified particles in LNP samples and characterizes and counts the sizes of particles observed. The size distribution results of each PPQ lot are presented in Table 48. A consistent and expected level of micron-size particles in the expected range for mRNA-1273 ( ) were observed. The SVP counts were low for each lot.

Flow Microscopy Particle Size Distribution Results **Table 48:** 

Lot	
	(Particles/mg)
6007520004	
6007520005	
6007520006	
057G20	
062G20	
001H20	

### 3.2.P.2.3.7.2.5 **Isothermal Titration Calorimetry**

Isothermal titration calorimetry is a calorimetric technique used to measure the interaction between with the mRNA-1273 LNP. The binding affinities, described by the dissociation constant (Kd), and the binding stoichiometries (number of are reported for each lot in Table 49. The observed binding affinities and binding stoichiometries were similar across lots.

Table 49: **Isothermal Calorimetry Results** 

Lot			
6007520004			
6007520005			
6007520006			
057G20			
062G20			
001H20			

Abbreviations: Kd = dissociation constant; M = binding stoichiometry (number of

### 3.2.P.2.3.7.2.6 **Zeta Potential**

Zeta potential is an indirect measurement of LNP surface charge calculated from the electrophoretic mobility of the particle. Zeta potential results shown in Table 50 demonstrate similar surface charge and all PPQ lots are considered approximately neutral.

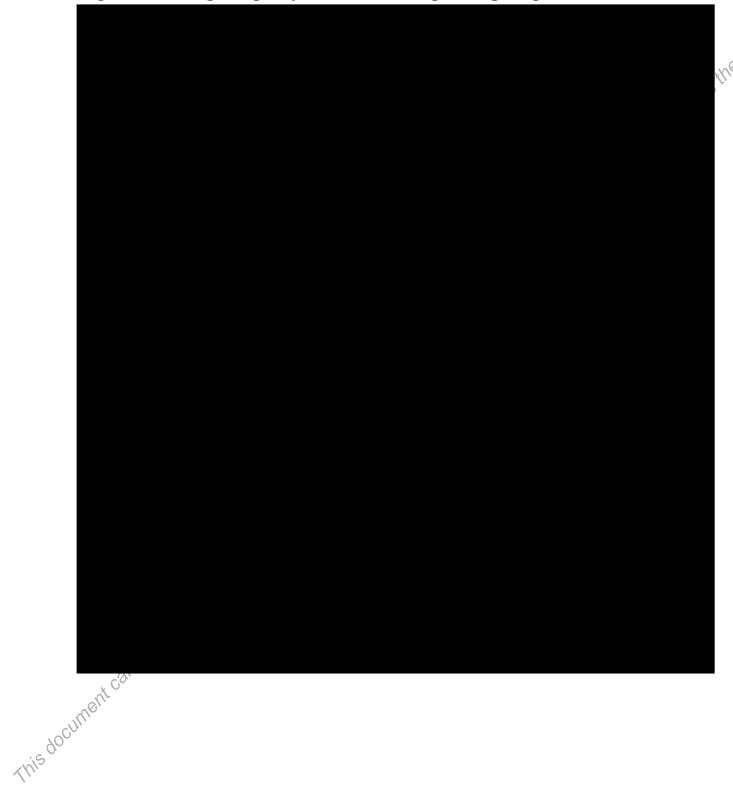
**Zeta Potential Results** Table 50:

Lot	
6007520004	
6007520005	
6007520006	
057G20	
062G20	
001H20	

### **Capillary Isoelectric Focusing**

3.2.P.2.3.7.2.7 Imaged capillary isoelectric focusing was used to determine the apparent isoelectric point (pI) of The charge profiles, presented in Figure 10, are generally consistent across all lots. The majority of the LNPs migrate with a pI between The average pI and polydispersities were consistently observed as shown in Table 51.

Figure 10: Imaged Capillary Isoelectric Focusing Electropherograms



ModernaTX, Inc. 3.2.P.2.3 Manufacturing Process Development

mRNA-1273

**Capillary Isoelectric Focusing Results Table 51:** 

Lot	
6007520004	
6007520005	
6007520006	
057G20	c
062G20	
001H20	
Abbreviations: pI = isoelectric point	, Value
3.2.P.2.3.7.2.8 Dye Permeation Kinetics	,50\
D	iol'

### **Dve Permeation Kinetics** 3.2.P.2.3.7.2.8

Dye permeation assay was used to measure the rate at which the cationic phenothiazinium dye, thionine, permeates the LNP from PPQ batches. The kinetic profile of this assay represents a biophysical signature that reports on LNP surface physical properties and mRNA encapsulation state, reported as a first order rate constant. Results shown for each PPQ lot in Table 52 demonstrate similar permeation kinetics.

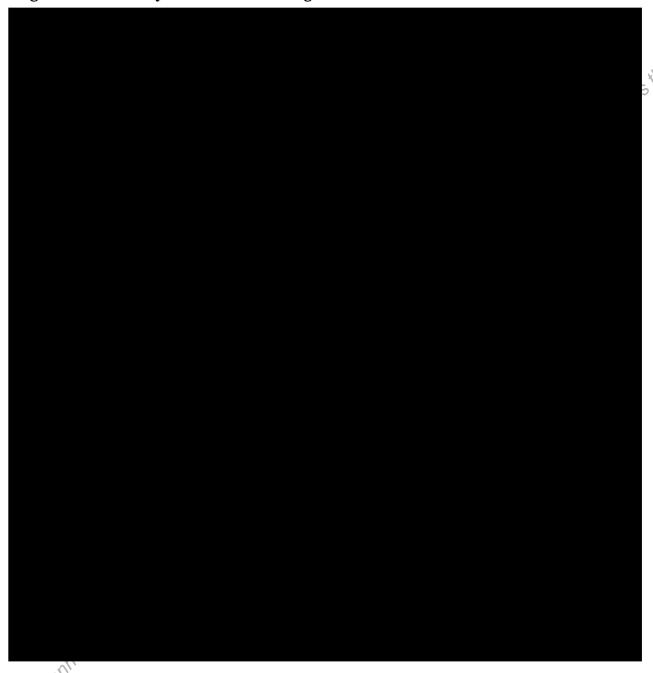
**Dye Permeation Kinetic Results Table 52:** 

Lot		
6007520004		
6007520005		
6007520006		
057G20		
062G20		
001H20		

### **Density Gradient Ultracentrifugation** 3.2.P.2.3.7.2.9

Density gradient ultracentrifugation was used to characterize the density distribution of mRNA-1273 Drug Product PPQ batches. The density gradient ultracentrifugation images (Figure 11) were converted into chromatograms based on pixel intensity and the slope change was monitored. The slope change results shown in Table 53 demonstrate consistent density profiles six PP for each of the six PPQ batches.

Figure 11: **Density Gradient Chromatograms** 



**Density Gradient Ultracentrifugation Results** 

	anli		
	Table 53:	Density Gradient Ul	tracentrifugation Results
c.V		Lot	
400		6007520004	
Mis		6007520005	
Z///		6007520006	
		057G20	
		062G20	
		001H20	

### 3.2.P.2.3.7.2.10 Hydrophobic Interaction Chromatography

Hydrophobic interaction chromatography (HIC) was used to characterize mRNA-1273 Drug Product PPQ batches. Three chromatographic regions are reported as relative percent areas:

HIC results generated for the mRNA-1273

Drug Product PPQ batches are reported in Table 54. The relative percent peak areas for each chromatographic region were consistent among these PPQ lots which demonstrate similar hydrophobic profiles.

Table 54: Hydrophobic Interaction Chromatography Results

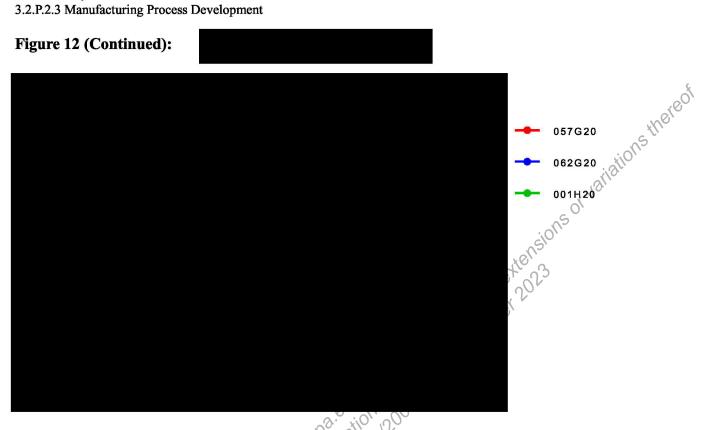
Lot	
6007520004	
6007520005	
6007520006	
057G20	
062G20	
001H20	

# **3.2.P.2.3.7.2.11** Esterase Kinetics

Esterase kinetics was used to determine the in mRNA-1273 Drug Product PPQ batches. The presented in Figure 12 and Table 55. Results demonstrate consistent behavior across lots.



Figure 12 (Continued):



**Table 55:** 

		00, 10, 10, 10, 10, 10, 10, 10, 10, 10,
		1/01/182.40)
		8:11, 0/13 1 0 kg
Table 55:	Esterase Kinetic	s Results
		We do yo
	Lot	6, 20, (1)
	6007520004	till I E
	6007520005	1/4 °U
	6007520006	Way office.
	057G20	11,1/10
	062G2Q	200
	001H20	
A.	.00	

### RiboGreen Encapsulation 3.2.P.2.3.7.2.12

A fluorescence-based RiboGreen assay was used as a release assay to determine the encapsulation efficiency of mRNA in mRNA-1273 LNP. RNA encapsulation results for the PPQ lots are shown in Table 56. These results demonstrate similarly high and consistent mRNA as a release as a release as a release of a USP-certified assay reagent. The Sponsor will continue to determine encapsulation efficiency of mRNA in mRNA-1273 LNP by RiboGreen assay as an orthogonal characterization assay.

**Table 56:** RiboGreen RNA Encapsulation Results

Lot	Encapsulation (%)
6007520004	
6007520005	
6007520006	
057G20	6
062G20	
001H20	

### 3.2.P.2.3.7.3 Analytical Assessment Summary

The evaluation of mRNA-1273 Drug Product GMP and PPQ lots demonstrates that product quality is highly similar for the manufacturing process and scale changes that occurred throughout process development, as shown in Table 57. The 6 PPO lots are also highly similar in vial .ended chara ... understanding con understa terms of product quality and demonstrate process consistency at the vial (Norwood PPO, 0.20 mg/mL mRNA-1273) and vial (Bloomington PPO, 0.20 mg/mL mRNA-1273) scale, as shown in Table 58. Extended characterization results were similar between lots and continue to provide additional understanding of process control together with

Table 57: **Attribute Assessment Results Summary for Clinical Trial Material** 

	Results												
Product Attribute	Method	Document No.	Description	85201100101	85201100102	85201100103	85201100104	6007520001	6007520002	520003	6007520004	6007520005	6007520006
LNP size distribution	Nanoparticle tracking analysis	DPTM-0039	High-resolution LNP size distribution						1/0				
LNP size distribution	Asymmetric flow field-flow fractionation	DPTM-0103	Fractionation coupled with in-line MALS detection for size determination										
Sub-visible particle counts	Coulter counter	DPTM-0035	Measurement of sub-visible particle counts in the range										
Sub-visible particle counts and morphology	Flow microscopy	DPTM-0115	Measurement of sub-visible particle counts and morphology in the range										
LNP surface characterization	Isothermal titration calorimetry	DPTM-0119	to LNP surface										
LNP charge	Zeta potential	DPTM-0118	Average LNP charge										
LNP charge distribution	Capillary isoelectric focusing	DPTM-0068	LNP pI distribution and polydispersity										
LNP structure	Dye permeation kinetics	Research-grade assay	Qualitative characterization of LNP surface and encapsulation state based on thionine permeation kinetics	NT	NT	SWATEL AC	NT	NT	NT	NT			
LNP density	Density gradient ultracentrifugation	DPTM-0104	Qualitative assessment of LNP density heterogeneity	NT	NT	NT	NT	NT	NT	NT			
LNP surface characterization	Hydrophobic interaction chromatography	SOP-0096	Qualitative characterization of LNP surface hydrophobicity	NT	NT NT	Mation	NT	NT	NT	NT			
LNP surface characterization	Esterase kinetics	Research-grade assay	Qualitative method for distribution	NT	Ol Will	NT	NT	NT	NT	NT			
mRNA encapsulation	Ribogreen	SOP-0298	Fluorescence based method for mRNA encapsulation	NT N	NT	NT	NT	NT	NT	NT			
mRNA encapsulation		DPTM-0073									NT	NT	NT
Abbreviations: LNP = 1	ipid nanoparticle; MALS	= multi-angle light so	cattering; NT = not tested; pI =	isoelectric point(s); PE	G = polyethylene glyc	col; PPQ = process performa	ance qualification;	SVP = sub-visible part	icle				
		· s doc'	eattering; NT = not tested; pI =	N. Role									
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Table 58: Attribute Assessment Results Summary for Process Performance Qualifications ( vial and vial scales

				Results					
Product Attribute	Method	Document No.	Description	6007520004	6007520005	6007520006	6007320001 (057G20)	6007320002 (062G20)	6007320003 (001H20)
LNP size distribution	Nanoparticle tracking analysis	DPTM-0039	High-resolution LNP size distribution						
LNP size distribution	Asymmetric flow field-flow fractionation	DPTM-0103	Fractionation coupled with in-line MALS detection for size determination						
Sub-visible particle counts	Coulter counter	DPTM-0035	Measurement of sub-visible particle counts in the range						
Sub-visible particle counts and morphology	Flow microscopy	DPTM-0115	Measurement of sub-visible particle counts and morphology in the range						
LNP surface characterization	Isothermal titration calorimetry	DPTM-0119	to LNP surface						
LNP charge	Zeta potential	DPTM-0118	Average LNP charge						
LNP charge distribution	Capillary isoelectric focusing	DPTM-0068	LNP pI distribution and polydispersity						
LNP structure	Dye permeation kinetics	Research-grade assay	Qualitative characterization of LNP surface and encapsulation state based on thionine permeation kinetics						
LNP density	Density gradient ultracentrifugation	DPTM-0104	Qualitative assessment of LNP density heterogeneity						
LNP surface characterization	Hydrophobic interaction chromatography	SOP-0096	Qualitative characterization of LNP surface hydrophobicity						
LNP surface characterization	Esterase kinetics	Research-grade assay	Qualitative method for distribution						
mRNA encapsulation	Ribogreen	SOP-0298	Fluorescence based method for mRNA encapsulation						

Abbreviations: LNP = lipid nanoparticle; MALS = multi-angle light scattering; NT = not tested; pI = isoelectric point(s);

; PPQ = process performance qualification; SVP = sub-visible particle

# This decement council to be get to get a great and the reading to the property of the property **Summary of Specification Changes – mRNA-1273 Drug Product** 3.2.P.2.3.7.4

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Summary of Specification Revisions – mRNA-1273 Drug Product – PVU Process to Norwood Scale A Table 59:

Test Parameter		Analytical		Rationale for Change				
		Procedure	PVU Process (PN 85201) (SPC-0948) / (Version 2.0/3.0)	Initial Scale A (PN 60075) (SPC-1063) / (Version 1.0)	Scale A PPQ (PN 60075) (SPC-1063) / (Version 2.0)	PVU to Initial Scale A	Initial Scale A to Scale A PPQ	
Appearance		Visual Inspection	White to off-white dispersion. May contain visible, white or translucent product-related particulates	White to off-white dispersion.  May contain visible, white or translucent product-related particulates	White to off-white dispersion.  May contain visible, white or translucent product-related particulates	No Change	No Change	
RNA Content		Anion Exchange- HPLC				Revised based on revised dilution step. Target added for reference.	No Change	
Identity		Reverse Transcription Sanger Sequencing	Sequence matches 100% of the coding region	Sequence matches description	Sequence matches description	No Change	No Change	
Purity		RP-HPLC				No Change	Tightening acceptance criteria to ensure product remains within specification for the duration of shelf-life	
Product-related Impurities			Report % area for each impurity group:  Impurity Group 1 (pre-main peak area) Impurity Group 2 (post-main peak area) Impurity Group 3 (mRNA-adduct species)	Report % area for each impurity group: Impurity Group 1 Impurity Group 2 Impurity Group 3	Report % area for each impurity group: Impurity Group 1 Impurity Group 2 Impurity Group 3	No Change	No Change	
% RNA Encaps	ulation	Fluorescence				No Change	No Change	
Potency		In Vitro Translation Methionine Labelling				Addition of test parameter to ensure SISPQ	No Change	
pН		USP <791>				No Change	No Change	
Osmolality		USP <785>				Revised based on available data	No Change	
Particle Size						No Change	No Change	
Polydispersity		Dynamic Light Scattering	Matches RT of reference	Report result		No Change	Initial establishment of polydispersity acceptance criteria based on analytical capability and process and stability experience.	
	SM-102	201	Matches RT of reference	Matches RT of reference	Matches RT of reference			
Lipid	Cholesterol	UPLC-CAD	Matches RT of reference	Matches RT of reference	Matches RT of reference	No Change	No Change	
Identification	DSPC	ont	Matches RT of reference	Matches RT of reference	Matches RT of reference			
	PEG2000-DMG	Julye	Matches RT of reference	Matches RT of reference	Matches RT of reference			
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Test Parameter		Analytical			Rationale for Change					
		Procedure	PVU Process (PN 85201) (SPC-0948) / (Version 2.0/3.0)		Initial Scale A (PN 60075) (SPC-1063) / (Version 1.0)		Scale A PPQ (PN 60075) (SPC-1063) / (Version 2.0)		PVU to Initial Scale A	Initial Scale A to Scale A PPQ
	SM-102 Cholesterol								0/	Changes in the LSS concentration (refer
Lipid	DSPC PEG2000-DMG								Revised based on revised dilution step.	impact lipid proportions within
Content			Individual Impurities	Report % area and RRT	Individual Impurities	Report % area and RRT	Individual Impurities	Report % area and RRT	No Change	the No Change
	Impurities		Total Impurities	Report % area	Total Impurities	Report % area	Total Impurities	Report % area		
Particulate Matter		USP <788> Method 2							No Change	No Change
Container Content		USP <697>	N/A (	a)					Addition of test parameter to ensure SISPQ	No Change
Bacterial Endo	toxins	USP <85> Ph. Eur. 2.6.14							No Change	No Change
Sterility		USP <71> Ph. Eur. 2.6.1	No Gro	wth	No Gro	wth	No Grow	th	No Change	No Change

Abbreviations: CAD = charged aerosol detector; DSPC = 1,2-Distearoyl-sn-glycero-3-phosphatdylcholfrie; EU = endotoxin unit(s); HPLC = high-performance liquid chromatography; LNP = lipid nanoparticle; LSS = lipid stock solution; N/A = not applicable; RT = retention time; RRT = relative retention time; SISPQ = safety, identity, strength, purity, and quality; UPLC = ultra-high-performance liquid chromatography

a) Test parameters not present on specification at time of testing. Abbreviations: CAD = charged aerosol detector; DSPC = 1,2-Distearoyl-sn-glycero-3-phosphatidylcholine; EU = endotoxin unit(s); HPLC = high-performance liquid chromatography;

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Summary of Specification Revisions – mRNA 1273 Drug Product –Norwood Scale A to Catalent Scale A/Scale B Table 60:

			Acceptano	e Criteria			
Test Parameter	Analytical Procedure	Norwood Scale A PPQ (P (SPC-1063) / (Version	N 60075)	Catalent Scale A PPQ/Scal (SPC-1128) / (Versio		Rationale for Change	
Appearance	Visual Inspection	White to off-white dispersion.  May contain visible, white or translucent product-related particulates		White to off-white dispersion.  May contain visible, white or translucent product-related particulates		No Change	
RNA Content	Anion Exchange-HPLC					Tightened acceptance criteria based on manufacturing and analytical performance results from Norwood Scale A batches	
Identity	Reverse Transcription Sanger Sequencing	Sequence matches descri	ription	Sequence matches de	escription	No Change	
Purity	RP-HPLC					Establishment of purity release acceptance criteria of for increased scale with a shelf life acceptance criteria of	
Product-related Impurities	- M M 20	Report % area for each impurity group: Impurity Group 1 (pre-main peak area) Impurity Group 2 (post-main peak area) Impurity Group 3 (mRNA-adduct species)		Report % area for each impurity group: Impurity Group 1 (pre-main peak area) Impurity Group 2 (post-main peak area) Impurity Group 3 (mRNA-adduct species)		No Change	
% RNA Encapsulation	Fluorescence					No Change	
In Vitro Translation	In Vitro Translation Methionine Labelling					No Change	
pH	USP <791>					No Change	
Osmolality	USP <785>					No Change	
Particle Size	Dynamic Light					No Change	
Polydispersity	Scattering					No Change	
SM-102		Matches RT of refere	ence	Matches RT of ref	erence		
Lipid Cholesterol		Matches RT of refere		Matches RT of ref	erence	No Change	
Identification DSPC		Matches RT of refere		Matches RT of reference		No Change	
PEG2000-DMG		Matches RT of refere	ence	Matches RT of ref	erence		
Lipid Cholesterol Content DSPC PEG2000-DMG	UPLC-CAD	:				No Change	
Lipid Impurities	569,0	Individual Impurities	Report % area and RRT	Individual Impurities		Establishing acceptance criteria based on ICH M7 guidance and	
	(*) (V)	Total Impurities	Report % area	Total Impurities		manufacturing experience.	
Particulate Matter	USP <788> Method 2					No Change	
Container Content	USP <697>					To enable a 10-dose multiple-dose vial	
Bacterial Endotoxins	USP <85> Ph. Eur. 2.6.14					No Change	
Sterility	USP <71> Ph. Eur. 2.6.1	No Growth		No Growth	No Change		