



INX100594280: PF-07302048 (Comirnaty) Residual DNA Characterization Report

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The following responsible areas have reviewed and approved this memorandum:

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Level	Change.	Rationale for Change.
1.0	Original Version	N/A



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1. EXECUTIVE SUMMARY

This report presents data characterizing the residual DNA template present in the Comirnaty vaccine. The findings presented herein are consistent with the Residual DNA Risk Assessment, which established that the presence of residual DNA in Comirnaty, which is below the internationally recommended threshold levels [4], and the non utilized sequence elements (like SV40 promoter) found at trace levels in the residual DNA, pose no safety risk to vaccinees. The summary findings are as follows:

- Comirnaty drug substance (DS) residual DNA template results are similar across all
 manufacturing sites and the manufacturing process consistently yields DS that complies
 with the established specification, which is based on WHO recommendations of not more
 than 10 ng per dose [4].
- Residual DNA testing on drug product (DP) yields similar ordower residual DNA compared to direct assessment in DS. Thus, testing for residual DNA template in DS provides an appropriate assessment of DNA levels present in DP.
- All Comirnaty DP dosage forms for adult, adolescent and pediatric populations comply with WHO recommendations of not more than 10 ng residual DNA per dose.
- Residual DNA size distribution data demonstrate the Comirnaty DS manufacturing process, which includes an enzymatic digestion of DNA template followed by a 2-step purification, yields short residual DNA template fragments. Most digested fragments are smaller than CCI

 Fragments larger than CCI
- All Comirnaty DS batches were confirmed positive for SV40 sequence elements, as expected based on the understanding of the residual DNA composition.
- CCI



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BACKGROUND INFORMATION

Comirnaty is an mRNA-based vaccine indicated for active immunization to prevent coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). COVID-19 vaccines have prevented appreximation deaths, of which Comirnaty contributed a large event of more than 1 billion doses worldwide. Corrected the component is a component in the contributed and a corrected appreximation of the contributed and contributed a component is made using a linearized plasmid DNA template enzymatically cut from a circular DNA plasmid. The plasmid DNA contains sequences that support its generation as well as regions important for production of mRNA. CC

In subsequent steps of the vaccine manufacturing process, the bulk of the DNA template is further removed by enzymatic digestion followed by a 2-step purification. The resulting purified mRNA, known as the Drug Substance (D\$), is later formulated into lipid nanoparticles (LNP) for the final vaccine Drug Product (DP). No DNA material is used or introduced in the manufacturing process other than the initial use of the DNA plasmid.

The DNA template (circular plasmid and linear DNA) is a well characterized starting material, manufactured under Good Manufacturing Practices (GMP) conditions which ensures a high level of control and quality in the production process. DNA template is released against the globally registered specifications. The mRNA DS is tested for several quality attributes, including residual DNA, which is considered an intrinsic impurity, routinely found in all cellderived biological products, such as vaccines based on inactivated or attenuated microorganisms, recombinant protein vaccines, DNA plasmid and viral vector vaccines [2, 3]. All Comirnaty doses released for use globally, meet the residual DNA requirements as defined by approved specifications and the World Health Organization (WHO) standards [4].



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Purpose of this document is to present data characterizing the residual DNA template and its non utilized SV40 sequence elements that may be present in the vaccine. The data presented in this document address the following:

• Evaluation of residual of the sequence of the sequence elements that may be present in the vaccine. The data presented in this document address the following:

- Evaluation of residual DNA template batch analysis data from the registered DS manufacturing sites
- Assessment of residual DNA template quantitation in DP samples
- Estimated residual DNA template content in DP
- Characterization of the size distribution of residual DNA template fragments
- Assessment of the presence of SV40 sequence elements in residual DNA template

4. EVALUATION OF RESIDUAL DNA BATCH ANALYSIS DATA

The World Health Organization (WHO) has established specific acceptance criterion recommendations for residual DNA levels, considered safe for human use in biological products, including mRNA vaccines (using DNA template as starting material). The acceptance criterion for residual DNA in Comirnaty's DS (\leq 330 ng DNA/mg RNA) aligns with WHO recommendations of not more than 10 ng per dose, translating to a total of 9.9 ng or less per 30 µg dose.

Residual DNA is routinely controlled in the DS material using an appropriate validated, quantitative Polymerase Chain Reaction (PCR) assay. The PCR assay is a widely recommended standard for residual DNA testing [5, 6] and approved by regulatory authorities worldwide. The assay used in Comirnaty testing is highly sensitive, enabling detection and quantification of trace amounts of DNA. To date, all commercialized DS batches of Comirnaty meet the regulatory approved acceptance criterion for residual DNA.

Residual DNA release testing results for 236 commercial Comirnaty DS batches were evaluated. The summary of this evaluation is presented in Table 1.



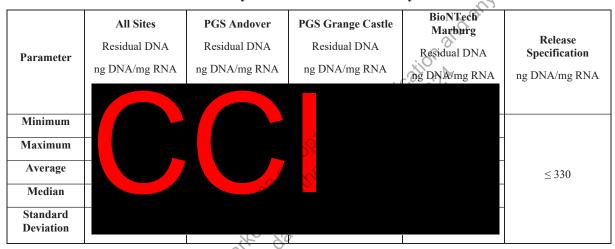
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The commercial DS batches were comprised of:

- Batches manufactured at three commercial manufacturing sites: Pfizer Global Supply (PGS) Andover, PGS Grange Castle, and BioNTech Marburg

 Four unique variants (Wildtype/Original and Original And Orig

Table 1 Residual DNA Batch Analysis Evaluation Summary



Conclusion:

This evaluation demonstrates residual DNA template results are similar across all manufacturing sites and the Comirnaty DS manufacturing process consistently yields DS which complies with the established specification.

5. ASSESSMENT OF RESIDUAL DNA IN DRUG PRODUCT SAMPLES

Purified DS is forward processed into DP through the combination mRNA DS with lipids to form lipid nanoparticles (LNP), which encapsulate the mRNA. Residual DNA template



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present in the DS is expected to be present in the DP as there are no purification steps for further removal of residual DNA within the DP manufacturing process. Therefore, routine quantitation of residual DNA template in DS provides an accurate assessment of residual DNA present in DP. An assessment of residual DNA in DP samples was performed for characterization and to demonstrate the suitability of testing at DS.

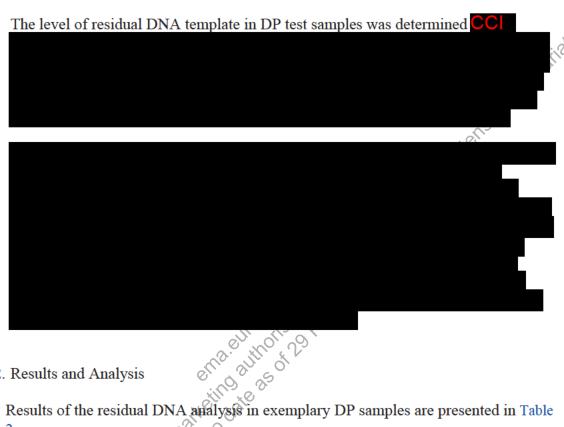
Each DP sample was analyzed with and without sample extraction step. Two separate nucleic acid extraction techniques were employed for DNA (and RNA) extraction from the DPs in three DP samples.





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5.1. Method



5.2. Results and Analysis

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The results demonstrate the qPCR assay is capable of residual DNA template quantification in DP samples, with or without sample extraction treatments. This observation is consistent with other PCR-based methods used in the analysis of DP samples, which do not require RNA extraction prior to analysis

. Additionally, both manual and automated nucleic acid extraction methods yielded residual DNA template results comparable to analysis without nucleic acid extraction.



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The DP results are compared to the residual DNA template results for the corresponding DS batches used in the manufacture of each DP lot. For the two bivalent DP lots (GN0005 and 22-DP-01012), the Wildtype (WT/Original) and Omicron BA.4/5 DS mRNA are mixed at a CC to formulate the final bivalent DP. Therefore, the average residual DNA result for the two DS batches is used for comparison. The residual DNA template measured in DP samples (see Table 2, column 'Residual DNA Measured in DP') are comparable to, or, lower than residual DNA template results measured during DS release testing (see Table 2, column 'Residual DNA for Input DS Batches').

Table 2 Drug Product Residual DNA Analysis

		. 60	
Drug Product Sample	Residual DNA Measured in DP ng DNA/mg RNA	Residual DNA for Input DS Batch(es) ng DNA/mg RNA	% Recovery of Average DS Result
GN0005 (WT / BA.4/5 bivalent) CC GN0005 (WT / BA.4/5 bivalent) CC 22-DP-01012 (WT / BA.4/5 bivalent) CC 22-DP-01012 (WT / BA.4/5 bivalent) CC HD9364Z (XBB.1.5)			
HD9364Z (XBB.1.5)	No.	1'-1-(DD1-(116-	0/ P

¹Average of the two input DS batch residual DNA results for bivalent DP batches used for % Recovery calculation. ²Monovalent DP batch, one input DS batch only.

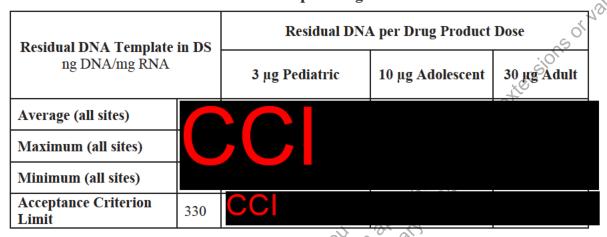
Using the residual DNA template batch analysis data presented in Table 1, the estimated residual DNA template content within each DP dose is calculated and summarized in Table 3. All DP dosage forms comply with WHO recommendations of not more than 10 ng per dose.



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Table 3 Estimated Residual DNA Content per Drug Product Dose



Conclusion:

Testing at DP yields a similar but more variable assessment of residual DNA in the vaccine, which is attributed to the requirement for low levels of residual DNA to be liberated from the LNP for detection and quantification. Thus, testing for residual DNA template in DS provides an appropriate assessment of DNA levels present in DP.

All DP dosage forms adult, adolescent and pediatric populations comply with WHO recommendations of not more than 10 ng per dose. As described in the Residual DNA Risk Assessment, the residual plasmid DNA template in Comirnaty is not considered to pose a safety risk.

6. DRUG SUBSTANCE SELECTED FOR RESIDUAL DNA CHARACTERIZATION

Subsequent characterization of residual DNA in Comirnaty is performed using DS samples, which is consistent with the routine batch testing performed for the vaccine. Twelve DS batches were selected to include the three commercial manufacturing sites (PGS Andover, PGS Grange Castle, and BioNTech Marburg), four unique variants (Wildtype/Original, Omicron BA.4/BA.5, and XBB.1.5) and residual DNA template CCI



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. Table 4 describes the DS batches and

characterization analyses.

Table 4 Drug Substance Samples for Residual DNA Characterization

Drug Substance Sample Description				Residual DNA Characterization Analyses		
Batch	Manufacturing Site	Strain	Residual DNA ¹ ng DNA/mg R	Size Dist. by TapeStation	SV40 presence by PCR	
GA5174	Andover	BA.1				
GF1001	Andover	WT				
GH5745	Andover	BA.4/5		-3		
GJ3638	Andover	BA.4/5				
GJ6907	Andover	BA.4/5				
HD1999	Andover	XBB.1.5	oP.	· Oli		
AB00050	Marburg	XBB.1.5				
AB00051	Marburg	XBB.1.5				
AB00060	Marburg	XBB.1.5				
HG3789	Grange Castle	XBB.1.5	(O)			
HG4918	Grange Castle	XBB 1.5	D			
HG4921	Grange Castle	XBB.1.5				

Residual DNA template measured at DS batch release.

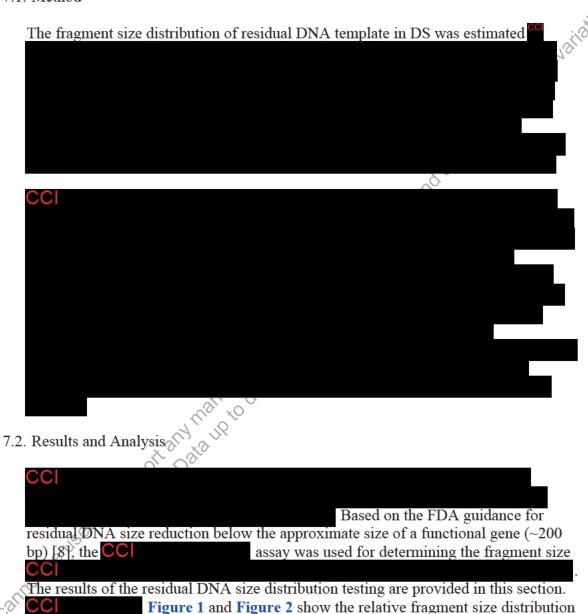
Results from DS batches determined to be sufficient and representative CC



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7. CHARACTERIZATION OF RESIDUAL DNA SIZE DISTRIBUTION

7.1. Method



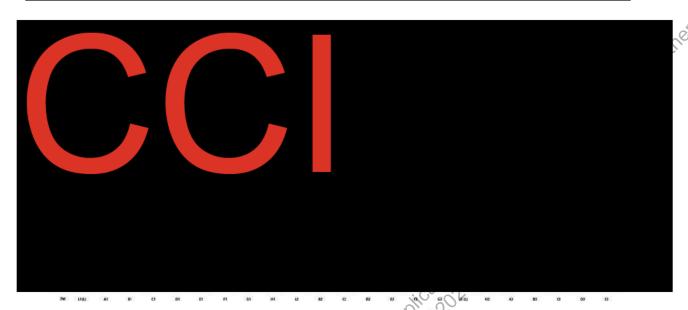


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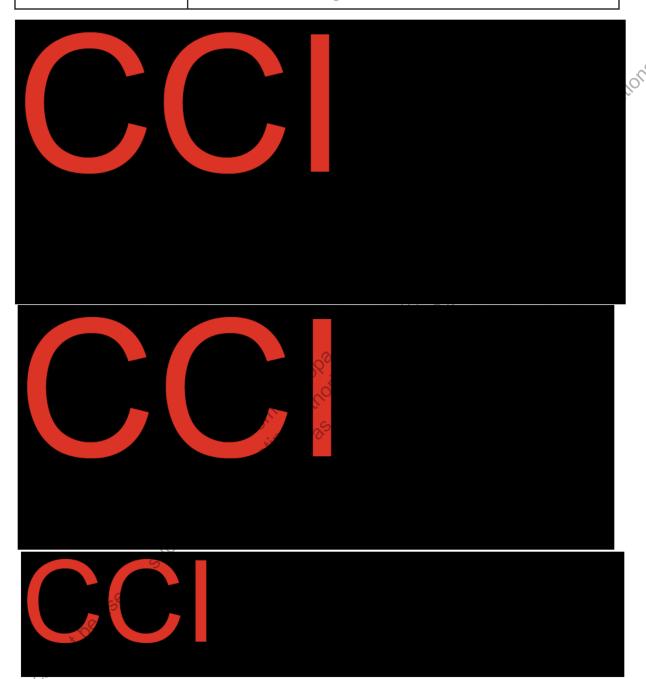








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Table 5. Percent Region Molarity (pmol/l) of Fragments CC DS Batch % Region % Region % Region Standard % RSD Average Molarity (pmol/l) Molarity (pmol/l) Molarity (pmol/l) Deviation (%) C GA5174 GF1001 GH5745 GJ3638 GJ6907 HD1999 AB00050 AB00051 AB00060 **HG3789 HG4918**

Table 6. Percent Region Molarity (pmol/l) of Fragments CC DS Batch % Region % Region % Region Average Standard % RSD Molarity (pmol/l) Molarity (pmol/l) Molarity (pmol/l) (%) Deviation CCI **GA5174** GF1001 GH5745 **GJ3638** GJ6907 HD1999 AB00050 AB00051 AB00060 HG3789 HG4918 HG4921

Conclusion:

The size distribution data demonstrate the Comirnaty DS manufacturing process, which includes an enzymatic digestion of DNA template followed by a 2-step purification, yields short residual DNA template fragments.



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This is confirmed using a CCI assay CCI

8. CHARACTERIZATION OF SV40 SEQUENCE ELEMENTS IN RESIDUAL DNA, All The plasmid-derived linear DNA template, serving as a starting material for synthesis.

Comirnaty vaccine's DS mRNA, incorporates additional DNA multipurpose plasmid DNA a construction of the plasmid DNA a construction. elements which are not utilized for DS mRNA production. These typical DNA sequence elements include restriction and cloning sites, as well as sequences aiding transcription and polyadenylation in eukaryotic cells such as the

properties of each sequence element are well known, and the functional roles have been documented (Section 3.2.S.2.3).

Based on the size distribution data presented in Section 7, the composition of residual DNA template fragments **CC**

designed to specifically detect

8.1. Method

within residual DNA template in DS The presence of **CC** samples was determined CCI



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Table 7. PCR Primer Design for Detection of SV40 Sequence Elements

Sequence Element	Origin	Starting Position	Ending Position
		(bp)) \	(bp)
CCI	Backbone cloning vector sequence	CCI	

8.2. Results and Analysis

The results of the PCR testing for presence of elements are shown in. All DS batches were confirmed positive for these SV40 sequence elements, as expected based on understanding of the residual DNA composition.



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Table 8. CC Results for Detection of CC

DS Batch/Sample	Result	Mean Ct Value
G3789		
G4918		
G4921		10
AB00050		
AB00051		
AB00060		
GA5174		7
GF1001		70,
GH5745		
GJ3638		
GJ6907		
HD1999		
High Assay		
Control		
Low Assay		
Control	is used as a reference to indicate positive of heatiful countries	

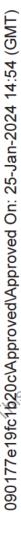
¹The Low Assay Control is used as a reference to indicate positive or negative result for samples.

Estimation of SV40 Sequence Element Content in Residual DNA:



Conclusion:

All DS batches were confirmed positive for SV40 sequence elements, as expected based on the understanding of the residual DNA composition.





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Or variations thereof Enzymatic digestion and filtration methods used in the DS process are highly effective at linearizing and removing the majority of DNA. As the approved Linearization Efficiency (Plasmid Topology) acceptance criterion CC (Section 3.2.S.2.3), CC

As the plasmid DNA contains sequence elements required for amplification in bacteria, an CC assay is a highly sensitive means of detecting CC

9.1. Method





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Table 9. DS Batches Selected for CC

	Characterization Repo	ort	
Table 9. DS Batches Sele	ected for CCI	DS Concentration mg/ml	
DS Batch	Strain	DS Concentration mg/ml	
GA5174	BA.1	CCI Jaffe	
GF1001	WT		
GH5745	BA.4/5	75	
GJ3638	BA.4/5	ė' į	
GJ6907	BA.4/5	100	
HD1999	XBB.1.5	, ⊗t	

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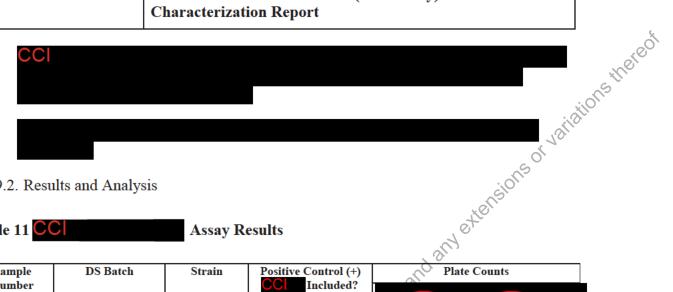
Table 10. CC Experimental Design

Sample Number	DS Batch	DS V	olume	Positive (+) Control	TE or Positive (+) Control
1	GA5174	13	μL		
2	GF1001	13	μL		o'
3	GH5745	13	μL		
4	GJ3638	13	μL		
5	GJ6907	13	μL		<u> </u>
6	HD1999	13	μL		
7	Null (TE)	13	μL		1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
8	GA5174	13	μL		
9	GF1001	13	μL		
10	GH5745	13	μL		
11	GJ3638	13	μL		
12	GJ6907	13	μL		
13	HD1999	13	μL		
14	Null (TE)	13	μL		





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9.2. Results and Analysis

Table 11 CC **Assay Results**

Sample Number	DS Batch	Strain	Positive Control (+ CC Included:	
1	GA5174	BA.1		
2	GF1001	WT		<u> </u>
3	GH5745	BA.4/5		
4	GJ3638	BA.4/5		
5	GJ6907	BA.4/5	JIO.	
6	HD1999	XBB.1.5		
7	Null (TE)	None	2	
8	GA5174	BA.1	~ C	
9	GF1001	MI 8	0	
10	GH5745	BA.4/5		
11	GJ3638	BA.475		
12	GJ6907	BA.4/5		
13	HD1999	XBB.1.5		
14	Null (TE)	None		



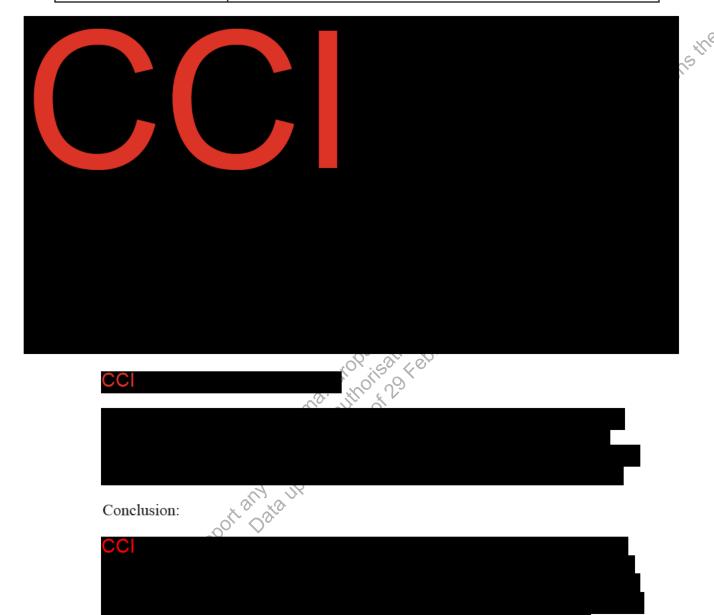


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3K istions thereof . The plasmid DNA used in Comirnaty production does not contain genes associated with causing cancer (oncogenes) and is not replication competent in mammalian cells [7], i.e., it cannot amplify in the human body. Therefore, it is not considered to pose a safety risk (See Residual DNA – Risk Assessment).

10. CONCLUSION

The safety profile of Comirnaty is well characterized after administration of more than 1 billion doses to individuals worldwide over the last 3 years. The safety profile of Comirnaty is described in the product labeling. The presence of residual DNA in Comirnaty, CC infectious, non-oncogenic, and is below the recommended limits set by the WHO guidelines. Additionally, the plasmid DNA used in Comirnaty production is not replication competent in mammalian cells [7], i.e., it cannot amplify in the human body. In summary, residual DNA in Comirnaty does not pose a safety fisk to vaccinees.

11. REFERENCES

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- 7. Emergent Human Pathogen Simian Virus 40 and Its Role in Cancer. Vilchez RA and Butel JSClin
- is docurring, It Guidance for Industry - Characterization and Qualification of Cell Substrates and Other Biological Materials Used in the Production of Viral Vaccines for Infectious Disease Indications, U.S. DHHS, FDA, CBER,

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