

Table IR15.C - 14.2.1.1.2.1.4.1  
Summary of Variant Groups in SARS-CoV-2 RT-PCR Positive  
Per-Protocol Set

	Placebo (N=14164) n (%)	mRNA-1273 (N=14287) n (%)
Number of Subjects with SARS-CoV-2 RT-PCR Positive, n (%)	885 (6.2)	80 (0.6)
Number of Events by Lineage, n (%)		
B.1	5 (<0.0)	-
B.1.1	1 (<0.0)	-
B.1.1.128	1 (<0.0)	-
B.1.1.186	2 (<0.0)	-
B.1.1.207	1 (<0.0)	-
B.1.1.222	8 (0.1)	-
B.1.1.316	1 (<0.0)	-
B.1.1.337	1 (<0.0)	-
B.1.1.432	1 (<0.0)	-
B.1.1.434	1 (<0.0)	-
B.1.1.519	2 (<0.0)	-
B.1.177.50	1 (<0.0)	-
B.1.2	427 (3.0)	15 (0.1)
B.1.232	1 (<0.0)	-
B.1.234	6 (<0.0)	-
B.1.240	1 (<0.0)	-
B.1.243	24 (0.2)	1 (<0.0)
B.1.311	6 (<0.0)	-
B.1.349	1 (<0.0)	-
B.1.369	2 (<0.0)	-

\*a subject is included if sequencing data from all available SARS-CoV-2 RT-PCR positive NP samples collected from July 2020 through May 2021 from participants in the blinded portion of the study

\*\*Haplotype is available without Lineage

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Summary of Variant Groups in SARS-CoV-2 RT-PCR Positive  
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	Placebo (N=14164) n (%)	mRNA-1273 (N=14287) n (%)
Number of Events by Lineage, n (%) (Cont.)		
B.1.375	1 (<0.0)	-
B.1.382	1 (<0.0)	-
B.1.396	1 (<0.0)	-
B.1.404	2 (<0.0)	-
B.1.427	6 (<0.0)	-
B.1.429	9 (0.1)	4 (<0.0)
B.1.517	2 (<0.0)	-
B.1.526.3	1 (<0.0)	-
B.1.544	2 (<0.0)	-
B.1.551	1 (<0.0)	-
B.1.561	5 (<0.0)	-
B.1.564	3 (<0.0)	-
B.1.587	8 (0.1)	-
B.1.595	2 (<0.0)	-
B.1.596	13 (0.1)	-
B.1.599	2 (<0.0)	-
B.1.605	1 (<0.0)	-
B.1.609	1 (<0.0)	-
P.1	1 (<0.0)	-
P.2	2 (<0.0)	-
R.1	3 (<0.0)	-
WILD TYPE	1 (<0.0)	-

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Summary of Variant Groups in SARS-CoV-2 RT-PCR Positive  
Per-Protocol Set

	Placebo (N=14164) n (%)	mRNA-1273 (N=14287) n (%)
Number of Events by Lineage, n (%) (Cont.)		
None**	20 (0.1)	-
No Sequencing Data Available	304 (2.1)	60 (0.4)
By First Detected, n (%)		
Brazil	1 (<0.0)	-
P.1	1 (<0.0)	-
California	15 (0.1)	4 (<0.0)
B.1.427	6 (<0.0)	-
B.1.429	9 (0.1)	4 (<0.0)
Variant of Concern	16 (0.1)	4 (<0.0)
B.1.427	6 (<0.0)	-
B.1.429	9 (0.1)	4 (<0.0)
P.1	1 (<0.0)	-
Variant of Interest	2 (<0.0)	-
P.2	2 (<0.0)	-

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