

Appendix Table 2. SARS-CoV-2 epitopes predicted by the ElliPro Antibody Epitope Prediction server in the NTD domain of the Spike rprotein.

	Start	End	Peptide	Number of Residues	Score	IR	RDR
6VXX	64	83	WFHAIHDNPV	10	0.957	IR-2	RDR1 - Δ 60-75
	94	103	STEKSNIRG	10	0.956	-	-
	109	114	TLDSKT	6	0.908	-	-
	118	130	LIVNNATNVVIKV	13	0.924	-	-
	132	165	EFQFCNDPFLGVN	13	0.964	IR-3	RDR2 - Δ 139-146
	168	188	FEYVSFKN	8	0.919	-	-
	209	216	PINLVRDL	8	0.933	IR-4	RDR3 - Δ 210-212
	239	265	QTLLALHAAY	10	0.958	IR-5	RDR4 - Δ 242-248
6VSB	63	83	TWFHDNPV	8	0.932	IR-2	RDR1 - Δ 60-75
	94	103	STNIIRG	7	0.949	-	-
	110	113	LDSK	4	0.903	-	-
	120	127	VNNATNVV	8	0.938	-	-
	132	165	EFQFCNDPFFRVYSSANN	18	0.959	IR-3	RDR2 - Δ 139-146
	173	188	QPFLKN	6	0.921	-	-
	209	216	PINLVRDL	8	0.928	IR-4	RDR3 - Δ 210-212
	240	265	TLLALHGAAAY	11	0.969	IR-5	RDR4 - Δ 242-248