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Erratum/corregendum

Corrigendum to "Design of a peptide-based subunit vaccine against novel coronavirus SARS-CoV-2" [Microbial Pathog. 145 (2020) 104236]

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The authors regret the incorrect publication of the amino acid sequence of the epitopes of the surface spike glycoprotein in Table 3. However, it should be noted that they were correctly used for the generation of the vaccine construct (as mentioned in the supplementary Table and supplementary Fig. 2), and so there is no change in the data/results. The authors would like to apologize for any inconvenience caused.

Incorrect Table 3. Predicted B-cell binding epitopes with their probable score and start position.

Sl. No.	Protein Name	Epitope	Position	Score
1	Nucleocapsid protein	TRRIRGGDGKMKDLSP	91	0.94
		KSAAEASKKPRQKRTA	249	0.93
		EGALNTPKDHIGTRNP	136	0.93
2	Membrane glycoprotein	RSMWSFNPETNILLNV	107	0.89
		SFRLFARTRSMWSFNP	99	0.88
3	**Surface spike glycoprotein**	**YACWHHSIGFDYVYNP**	**6149**	**0.96**
		VVKIYCPACHNSEVGP	**365**	**0.96**
		TLKGGAPTKVTFGDDT	**814**	**0.95**
		TSRYWEPEFYEAMYTP	**5304**	**0.94**

Correct Table 3. Predicted B-cell binding epitopes with their probable score and start position.

Sl. No.	Protein Name	Epitope	Position	Score
1	Nucleocapsid protein	TRRIRGGDGKMKDLSP	91	0.94
		KSAAEASKKPRQKRTA	249	0.93
		EGALNTPKDHIGTRNP	136	0.93
2	Membrane glycoprotein	RSMWSFNPETNILLNV	107	0.89
		SFRLFARTRSMWSFNP	99	0.88
3	**Surface spike glycoprotein**	**AGTITSGWTFGAGAAL**	**888**	**0.97**
		GVSVITPGTNTSNQVA	**603**	**0.95**
		GWTAGAAAYYVGYLQP	**266**	**0.95**
		PQIITTDNTFVSGNCD	**1121**	**0.95**

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