Protein Similarity of the SARS-CoV and Sars-CoV2 genes.

Protein SARS1	AS	Protein SARS2	AS	Similarity [%]
ND 020000 2 mars4	100	VD 000735307.4	100	1
NP_828860.2 nsp1	180	YP_009725297.1	180	84,40
ND 020064.2 2	620	leader_protein	620	60.24
NP_828861.2 nsp2	638	YP_009725298.1 nsp2	638	68,34
NP_828862.2 <u>nsp3</u>	1922	YP_009725299.1 <u>nsp3</u>	1945	75,97
NP_904322.1 nsp4	500	YP_009725300.1 nsp4	500	80,00
NP_828864.1 nsp6	290	YP_009725302.1 nsp6	290	88,15
NP_828865.1 <u>nsp7</u>	83	YP_009725303.1 <u>nsp7</u>	83	98,80
NP_828866.1 <u>nsp8</u>	198	YP_009725304.1 <u>nsp8</u>	198	97,47
NP_828867.1 <u>nsp9</u>	113	YP_009725305.1 nsp9	113	97,35
NP_828868.1 <u>nsp10</u>	139	YP_009725306.1 nsp10	139	97,12
NP_904321.1 nsp11	13	YP_009725312.1 nsp11	13	84,62
NP_828870.1 nsp13	601	YP_009725308.1 <u>helicase</u>	601	99,83
NP_828873.2 nsp16	298	YP_009725311.1	298	93,29
		<u>methyltransferase</u>		
NP_828863.1	306	YP_009725301.1 <u>3c-</u>	306	96,08
3c_like_proteinase		<u>like_proteinase</u>		
NP_828872.1 <u>endornase</u>	346	YP_009725310.1 endornase	346	88,73
NP_828871.1 exonuclease	527	YP_009725309.1exonuclease	527	95,07
NP_828869.1	932	YP_009725307.1	932	96,35
rna_polymerase		rna_polymerase		
NP_828855.1 matrix protein	221	YP_009724393.1	222	90,54
		membrane_glycoprotein		
NP 828858.1	422	YP 009724397.2	419	90,52
nucleocapsid_protein		nucleocapsid_protein		
NP_828851.1	1255	YP_009724390.1	1273	75,96
surface_glycoprotein		surface_glycoprotein		
NP 828854.1 protein e	76	YP_009724392.1 protein e	75	94,74
NP 828857.1				
hypothetical_protein_sars7a	122			
NP 849177.1 hypothetical	84			
protein sars8b				
NP 849176.1 hypothetical	39			
protein sars8a				
NP 849175.1 hypothetical	44			
protein sars7b				
NP 828852.2 hypothetical	221			
protein sars3a	-			
		YP_009724391.1 ORF3a	275	
		protein		
	1	 	1	1

[&]quot;protein>" are proteins which have PDB Structures in the repository, underscored ist the name we used in the repo.

Gene reference for CoV: NC_004718.3 Gene reference for CoV2: NC_045512.2