

POS	REF	ALT	COV026530	COV026531	COV026532	Tipo de cambio	Gen/región	Cambio aa
241	C	T	T	T	T	intergenic_region	CHR_START-ORF1ab	
2706	C	T,Y	T	C/T	T	missense_variant	ORF1ab	Thr814Ile
3037	C	T	T	T	T	synonymous_variant	ORF1ab	Phe924Phe
3267	C	Y	C	C/T	C	missense_variant	ORF1ab	Thr1001Ile
6954	T	Y	T	T/C	T	missense_variant	ORF1ab	Ile2230Thr
9165	C	Y	C	C	C/T	missense_variant	ORF1ab	Thr2967Ile
10533	G	K	G	G	G/T	missense_variant	ORF1ab	Cys3423Phe
11417	G	T	T	T	T	missense_variant	ORF1ab	Val3718Phe
13012	A	M	A	A	A/C	synonymous_variant	ORF1ab	Thr4249Thr
14103	T	Y	T	T	T/G	synonymous_variant	ORF1ab	Asp4613Asp
14247	T	C,Y	C	T/C	C	synonymous_variant	ORF1ab	Asp4661Asp
14408	C	T	T	T	T	missense_variant	ORF1ab	Pro4715Leu
15279	C	Y	C	C/T	C	synonymous_variant	ORF1ab	His5005His
16176	T	Y	T	T/C	T	synonymous_variant	ORF1ab	Thr5304Thr
20268	A	G,R	G	G/A	G	synonymous_variant	ORF1ab	Leu6668Leu
22673	T	Y,C	T/C	T	C	missense_variant	S	Ser371Pro
23271	C	M	C	C/A	C	missense_variant	S	Ala570Asp
23403	A	G	G	G	G	missense_variant	S	Asp614Gly
24506	T	K	T	T/G	T	missense_variant	S	Ser982Ala
25038	A	T	T	T	T	missense_variant	S	His1159Leu
25152	T	A	T	T	A	missense_variant	S	Leu1197His
25731	C	T,Y	T	C/T	T	synonymous_variant	ORF3a	Tyr113Tyr
26257	G	T	G	G	T	missense_variant	E	Val5Phe
28253	C	T,Y	T	C/T	T	synonymous_variant	ORF8	Phe120Phe
28254	A	M	A/C	A	A	missense_variant	ORF8	Ile121Leu
28833	C	T	T	T	T	missense_variant	N	Ser187Leu
29870	C	N,A	N	A	N	intergenic_region	ORF10-CHR_END	