

Table S3. Mutations present in the consensus sequences of the virus clones indicated

Virus clones ¹	Mutations ²				
	Non-coding regions	A2	CP	A1	Replicase
Qβ _{Anc} (1)					
Qβ _{Anc} (2)					
Qβ _{Anc} (3)					U2972C
L1C1			U1665C (F107L)	C2001U (L219F)	U4001C
L1C2		G1312A (V417I)	U1665C (F107L), C1689U	A2187C (S281G)	U2847C (Y165H), U4001C
L1C3		C420U	U1665C (F107L)		G3341A, U4001C
L1C4			U1665C (F107L)		U4001C
L2C1		U969C, G1312A (V417I)		A1930G (Q195R)	
L2C2		U192C	G1494A (V50I)	A1930G (Q195R), G2223A (V293I)	G2798A
L2C3		G1312A (V417I)		A1930G (Q195R)	U2740C (M129T)
L2C4		U393C, C1101U		C1953U (L203F), C2201U	
L3C1		A1065G		A1930G (Q195R)	G2468A , U2951C
L3C2		U354C, A1065G		A1930G (Q195R)	G2468A , U2847C (Y165H)
L3C3	A54G	A1065G		A1930G (Q195R)	G2468A , U3314C
L3C4		A258G, A1065G		A1930G (Q195R)	G2468A

¹The virus clones indicated are described in the section “Isolation of biological clones” of Materials and Methods.

²For each mutation, the protein where it is located is indicated. Non synonymous mutations are shown in bold letters and the corresponding amino acid changes are indicated in brackets. Common mutations between clones and the populations from which they were isolated (see Table S2) are shown in blue.