



(c)

	<u>Region of homology</u>
SOCS1	<u>VRPLQELC</u>
SOCS3	<u>VATLQHLC</u>
VHL	<u>VYTLKERC</u>
NS1	<u>VALLKITC</u>
Consensus	<u>VXXLXXXC</u>
Mutation in NS1	<u>AALAKITA</u>

(d)

	V	A	L	L	K	I	T	C
	160		170		180			
rRSVNS1F	AGTAGCATTG	TTAAAAATAA	CATGCTATAC					
rRSVNS1F-ELCmut	AGCAGCATTG	GCAAAAATAA	CAGCCTATAC					
	A	A	L	A	K	I	T	A