Table S6: Number of sequences of each sub-type making up the negative training set used for determining whether a given sub-type (X) is present in the *gag-pol* region of a CRF.

X=A		**X=B**		**X=C**		**X=D**		**X=F**		**X=G**	
subtype	No.	subtype	No.	Subtype	No.	subtype	No.	subtype	No.	subtype	No.
A	0	A	2	A	10	A	2	A	2	A	6
B	2	B	0	B	2	B	31	B	2	B	2
C	2	C	2	C	0	C	2	C	2	C	2
D	2	D	22	D	2	D	0	D	2	D	2
F	2	F	2	F	2	F	2	F	0	F	2
G	6	G	2	G	2	G	2	G	2	G	0
H	2	H	2	H	2	H	2	H	2	H	2
J	2	J	2	J	2	J	2	J	2	J	2