

Supplementary Table S6. Range of conservation of individual Rev domain across different HIV-1 group M clades.

Clade ^a	Conservation (%)										
	64.5 – 72.1	72.2 – 74.9	75.0-77.9	78.0-80.9	81.0-83.9	84.0-86.9	87.0-89.9	90.0-92.9	93.0-95.9	96.0-98.9	99.0-100
Group M	C-term			Turn	OD2, ONL	NES	N-term, OD1			ARM	
A1				C-term	Turn	OD1	N-term, OD2, NES			ARM	
A6					OD2, ONL	N-term, OD1, Turn, ARM, C-term			NES		
B	OD2		ONL	OD1, C-term	Turn		N-term		ARM, NES		
C	C-term			ONL	OD2	Turn, NES		N-term, OD1, ARM			
D					OD1, C-term	N-term, OD2, NES, Turn			ONL	ARM	
F1					N-term, OD1, OD2, ONL, NES		Turn		ARM		
F2	ONL, C- term				OD2		N-term, Turn, NES, OD1		ARM		
G	ONL, C- term			OD1, OD2, NES	N-term, Turn		ARM				
H	N-term, OD1, OD2, ONL			Turn, C-term						ARM, NES	
01_AE				Turn, ONL	OD2, C-term	OD1, ARM, NES		N-term			
02_AG	C-term			OD2, ONL	OD1, NES	N-term, Turn		ARM			
06_cpx				Turn	ONL	N-term, OD1, OD2, NES, C-term			ARM		
07_BC	N-term			C-term	OD1, OD2, ONL, NES		Turn		ARM		
08_BC					ARM, C-term		Turn, OD2, ONL		N-term, OD1, NES		
11_cpx	Turn		OD1, ONL, C- term			OD2, NES		N-term			
12_BF					N-term, OD1, OD2, ONL		Turn, C-term		ARM, NES		
13_cpx	OD1, OD2, C- term				Turn		ONL, NES		N-term		
14_BG					OD1, ONL		OD2, C-term		Turn, ARM, NES		
22_01A1	Turn, ONL			OD1, NES	N-term, C-term		ARM, OD2				
35_A1D	OD1			Turn	ONL, NES, C-term		N-term, ARM		OD2		

42_BF1				OD2	N-term, Turn, ONL, C-term	OD1, ARM, NES
63_02A6		OD2, ONL	N-term, OD1, NES	Turn, C-term	ARM	
71_BF1	OD2	OD1, ONL	Turn, C-term	N-term	ARM, NES	
85_BC				OD1,OD2	N-term, Turn, ARM	ONL, NES, C-term
91_cpx				OD2	OD1, ONL	N-term, Turn, ARM,NES, C-term
103_01B	Turn	OD2, ONL		OD1	C-term	ARM, NES N-term

Annotated protein domains: N-term, N-terminal region; OD, Oligomerization domain; ARM, Arginine-Rich Motif; NES, Nuclear Export Signal; C-term, C-terminal region; ONL, OD-NES Linker.