Gene	HXB2 coordinates	Sequence Logo and degeneracy	Function
Regions with predicted functions			
gag Figure 3a 'INS1'	898 - 903	*GCAAGC	Within (INS1) first Inhibitory sequence region
Figure3a 'INS1'	985 - 996	CCA-CTCTTSAG	Within INS1
Figure 3a 'INS2'	1309 - 1314	FÇÇATŢA	Within INS2
pol Figure 3b 'p1'	4764 - 4776	GCAGTATTCAT	Within nuclease- hypersensitive region
Figure 3b: 'p1'	4777 - 4790	0 2 0 0 2 0 0 2 0 0 2 0 0 2 2 0 2 0 2 0	potential binding motif for Oct-1
Figure 3b 'p2'	4864 - 4866	AAT	In nuclease- hypersensitive region
Novel regions			
vpu Figure 4b 'n2'	6101 - 6106	GTAGC	novel conserved region
Figure 34b 'n3'	6143 - 6151	TAGAATAT.	novel conserved region
Figure 3b 'n4'	6167 - 6178	0 0 2 2 0 4 0 0 2 0 0 3	novel conserved region
env Figure 4c 'n1'	7656 - 7667	AATTGGAGAAGT	novel region, analyzed in fitness assays