

Supplementary information for:
Coevolution Analysis of HIV-1 Envelope Glycoprotein Complex

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**Table S3: Coevolving pairs between residues in V3 and other structural regions
in 361 the HIV-1 Env.**

	i (HXB2)	i (Domain)	j (HXB2)	j (Domain)	GREMLIN	TP
15	295	C2	297	V3	0.78	1
79	325	V3	419	C4	0.45	1
90	330	V3	334	C3	0.43	0
107	330	V3	332	C3	0.41	1
129	300	V3	442	C4	0.38	1
154	154	V1	300	V3	0.35	1
231	172	V2	305	V3	0.30	1
281	330	V3	415	V4	0.28	1
298	293	C2	297	V3	0.27	0
397	173	V2	305	V3	0.24	1
414	303	V3	440	C4	0.24	1