

Table S1. 2G12 epitope in sequential R5 Env sequences and 2G12 IC50 for corresponding R5 isolates

		2G12 epitope ^b						2G12 IC50 ug/ml
Patient ^a	Isolate	Clinical status	295	332	339	386	392	
G	1228	Chronic	+	+	+	+	-	>25
	4481	End-stage	+	+	-	-	+	3,95
H	624	Chronic	+	+	+	+	+	>25
	3899	End-stage	+	+	+	+	+	20,16
I	5013	Chronic	-	+	+	-	+	>25
	8616	End-stage	-	+	+	- (+ 1/4 ^c)	+	>25
J	1372	Chronic	+	+	- (+ 1/4 ^c)	+ (- 1/4 ^d)	+	4,87
	5714	End-stage	+	+	-	+	+	14,24
M	668	Chronic	+	+	+	+	+	>25
	7363	End-stage	+	+	+	+	+	13,47
R	6322	Chronic	+	+	+	+	+	1,16
	8004	End-stage	+	+	+	+	+	0,36

a) Patient code according to [10]

b) Data based on sequences of four clones; + presence of PNGS in 4/4 clones, - absence of PNGS in 4/4 clones.

c) The sequence of one clone out of four had the PNGS

d) The sequence of one clone out of four lacked the PNGS