Supplementary Table S4A. Patient AM: Intrapatient Mutations in HIV gp120 Sequences of Various Tissues and Organs Compared to the Liver

			*a	*a		*a	*a												
Position in HXB2	130	132	136	137	139	141	142	-	144	145	-	-	-	146	147	148	151	152	175
Residue in HXB2	K	T	N	D	N	N	S	-	S	G	-	-	-	R	M	I	K	G	F
Liver	D	T	Α	I	K	С	S	T	G		T	N	Н	Т	C		M	E	
Diaphragm	N	S	V	T	A	S	T	-	V		S	G	Q	L	N		I	G	
Threshold	D	T	Α	I	K	C	S	T	G		T	N	H	T	C		M	E	
Liver	D										T		Η	T		N			V
Kidney	E										N		Q	N		S			L
Threshold	D										T		H	T		N			V
Liver	D										T			T		N			V
Left lymph node	E										N			N		S			L
Threshold	D										T			T		N			V
Liver		T																	
Right lymph node		I																	
Threshold		T																	
Liver	D										T			T		N			V
Liver tumor (1)	E										N			N		S			L
Threshold	D										T			T		N			V
Liver	D								G		T			T		N			V
Liver tumor (2)	E								R		N			N		S			L
Threshold	D								G		T			T		N			V
Liver		T		•									Η			N			
Prostate		I											Q			S			
Threshold		T											Η			N			
Liver	D			•							T			T		N			V
Spleen	E			•							N			N		S			L
Threshold	D			•							T			T		N			V
Liver	D										T			T		N			V
Stomach lymphoma	E										N			N		S			L
Threshold	D										T			T		N			V

^{*}a=glycosite in HXB2.

Amino acid differences between gp120 in the liver and other tissues were included only if the amino acid was conserved at a frequency of 0.8 in the liver. Dots represent no difference between residues of the liver and the other tissue. Residues not shown are conserved in all tissues. Dashes represent insertions and/or deletions relative to HXB2. Sequence data were originally reported in Lamers *et al.*¹⁸

Supplementary Table S4B. Patient AM: Intrapatient Mutations in HIV gp120 Sequences OF VARIOUS TISSUES AND ORGANS COMPARED TO THE LIVER

D W 1 1 1111D	4	470	*b	*b	405	*a				*a	100	400							404
Position in HXB2 Residue in HXB2	177 Y	178 K	179 L	181 I	185 D	187 D	-	-	-	188 T	189 T	190 S	-	-	-	-	-	-	191 Y
RESIDUE IN TIADZ	1	N	L	1	D	D				1	1	3	_						
Liver		K	P	L	N	N		Y	V		Q							R	
Diaphragm		E	L	Ι	E	D		K	-		N							-	
Threshold		K	P	L	N	N		Y	V		Q							R	
Liver	Н		P	L	N	N	K		V		Q		-	-	-	-	-		
Kidney	Y		I	I	D	D	E		-		S		I	N	N	K	T		
Threshold	Η		P	L	N	N	K		V		Q		-	-	-	-	-		
Liver	Н		P	L	N	N	K		V		Q		-	-	-	-	-		
Left lymph node	Y		I	I	D	D	E		-		S		I	N	N	K	T		
Threshold	Н		P	L	N	N	K		V		Q		-	-	-	-	-		
Liver																			
Right lymph node																			
Threshold																			
Liver	Н		P	L	N	N	K		V		Q		-	-	-	-	-		
Liver tumor (1)	Y		I	I	D	D	E		-		$\tilde{\mathbf{s}}$		I	N	N	K	T		
Threshold	Н		P	L	N	N	K		V		Q		-	-	-	-	-		
Liver	Н		P	L	N	N	K		V		õ		_	_	-	_	_		
Liver tumor (2)	Y		I	I	D	D	E		-		Q S		I	N	N	K	T		
Threshold	Н		P	L	N	N	K		V		Q		-	_	-	_	_		
Liver											Q								
Prostate											s								
Threshold											Q								
Liver	Н		P	L	N	N			V		~								
Spleen	Y		I	I	D	D			_										
Threshold	Н		Р	L	N	N			V										
Liver	Н		P	L	N	N	K		V		Q		_	_	_	_	_		
Stomach lymphoma	Y		I	Ī	D	D	E		_		ŝ		Ι	N	N	K	T		
Threshold	H	•	P	Ĺ	N	N	K		V	•	Q	•	-	-	-	-	-		

^{*}a=glycosite in HXB2, *b=LDI/LDV tripeptide in V2 binds integrin $\alpha_4\beta_7$. Amino acid differences between gp120 in the liver and other tissues were included only if the amino acid was conserved at a frequency of 0.8 in the liver. Dots represent no difference between residues of the liver and the other tissue. Residues not shown are conserved in all tissues. Dashes represent insertions and/or deletions relative to HXB2. Sequence data were originally reported in Lamers *et al.* ¹⁸

Supplementary Table S4C. Patient AM: Intrapatient Mutations in HIV gp120 Sequences of Various Tissues and Organs Compared to the Liver

		*c	*a	*a	*a		*d		*d		*e	*d	*d		*f	*f	*f	
Position in HXB2	194	198	289	290	291	305	306	307	308	311	312	318	322	328	332	333	334	336
Residue in HXB2	T	T	N	T	S	K	R	I	R	R	G	V	K	Q	N	Ι	S	A
Liver	I	S				K	G		R		G		K	Q	N	L	S	Α
Diaphragm	R	T				R	S		Н		A		E	ĸ	K	I	N	T
Threshold	I	S				K	G		R		G		K	Q	N	L	S	Α
Liver		S	N		S		G		R				K			L		
Kidney		T	D		F		S		Н				E			I		
Threshold		S	N		S		G		R				K			L		
Liver		S	N		S		G		R				K			L		
Left lymph node		T	D		F		S		Н				E			I		
Threshold		S	N		S		G		R				K			L		
Liver																		
Right lymph node																		
Threshold																		
Liver		S	N		S		G		R				K			L		
Liver tumor (1)		T	D		F		S		Н				E			I		
Threshold		S	N		S		G		R				K			L		
Liver		S	N		S		G		R				K			L		
Liver tumor (2)		T	D		F		S		Н				E			I		
Threshold		S	N		S		G		R				K			L		
Liver										I		Y						
Prostate										M		I						
Threshold										I		Y						
Liver		S	N		S		G		R				K			L		
Spleen		T	D		F		S		Н				E			I		
Threshold		S	N		S		G		R				K			L		
Liver		S	N		S		G		R				K			L		
Stomach lymphoma		T	D		F		S		Η				E			I		
Threshold		S	N		S		G		R				K			L		

Sequence data were originally reported in Lamers et al. 18

^{*}a=glycosite in HXB2, *c=CD4 contact residue, *d=coreceptor binding site, *e=GPGR tip of V3 loop, neutralizing antibody (447-52D) binding, coreceptor binding site, *f=monoclonal Ab 2G12 glycosylation binding site.

Amino acid differences between gp120 in the liver and other tissues were included only if the amino acid was conserved at a frequency of 0.8 in the liver. Dots represent no difference between residues of the liver and the other tissue. Residues not shown are conserved in all tissues. Dashes represent insertions and/or deletions relative to HXB2.

Supplementary Table S4D. Patient AM: Intrapatient Mutations in HIV gp120 Sequences OF VARIOUS TISSUES AND ORGANS COMPARED TO THE LIVER

								*8	*h			*a		*d	*c		
Position in HXB2	340	343	346	347	359	360	362	373	386	388	396	397	415	440	461	467	496
Residue in HXB2	N	K	A	S	Ι	Ι	K	T	N	T	F	N	T	S	S	I	V
Liver	Т	Т	V	K		I	K		N	S	K	Т		S	S		
Diaphragm	Ñ	K	À	A	•	V	N		D	A	S	Î	•	K	I	•	•
Threshold	T	T	V	K		İ	K	-	N	S	K	Ť	·	S	Š		
Liver	T	T		K	I		K	R			K	T	T			T	I
Kidney	N	K		Q	_		T	M			N	S	I			I	M
Threshold	T	T		\tilde{K}	I		K	R			K	T	T			T	I
Liver	T	T		K	I		K	R			K	T	T			T	
Left lymph node	N	K		Q	-		T	M			N	N	I			I	
Threshold	T	T		K	I		K	R			K	T	T			T	
Liver												T					
Right lymph node												I					
Threshold												T					
Liver	T	T		K	I		K	R			K	T	T			T	
Liver tumor (1)	N	K		Q	-		T	M			N	N	I			I	
Threshold	T	T		K	I		K	R			K	T	T			T	
Liver	T	T		K	I		K	R			K	T	T			T	
Liver tumor (2)	N	K	•	Q	-		T	M			N	N	I			I	
Threshold	T	T	•	K	Ι		K	R			K	T	T			T	
Liver			•					•				T					
Prostate			•					•				I					
Threshold			•					•				T					
Liver	T	T		K			K	R		•	K	T	T		•	T	
Spleen	N	K	•	Q			T	M			N	N	I			I	
Threshold	T	T		K			K	R		•	K	T	T		•	T	
Liver	T	T		K	I		K	R		•	K	T	T			T	
Stomach lymphoma	N	K		Q	-		T	M			N	N	I			I	
Threshold	T	T		K	I		K	R		•	K	T	T			T	

^{*}a=glycosite in HXB2, *c=CD4 contact residue, *d=coreceptor (CCR5/CXCR4) binding site, *g=CD4 binding loop: SSGGDPEIVTH, *h=lectin DC-SIGN binding site (N386-T388), glycosite, monoclonal Ab 2G12 glycosylation binding site (N386-T388). Amino acid differences between gp120 in the liver and other tissues were included only if the amino acid was conserved at a frequency of 0.8 in the liver. Dots represent no difference between residues of the liver and the other tissue. Residues not shown are conserved in all tissues. Dashes represent insertions and/or deletions relative to HXB2. Sequence data were originally reported in Lamers *et al.* ¹⁸