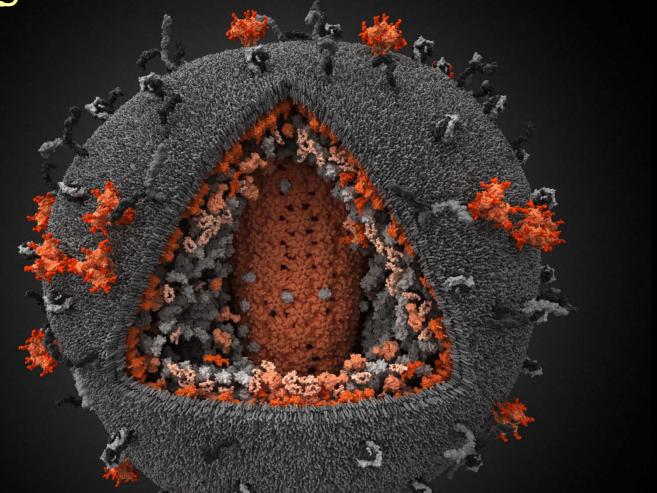
Finding Antibodies that Neutralize HIV



Beatrix Ueberheide, Sunnie Myung, David Fenyo, Brian Chait

Ivan Konstantinov, Yury Stefanov, Aleksander

Johannes Scheid, Michel Nussenzweig Science Company)

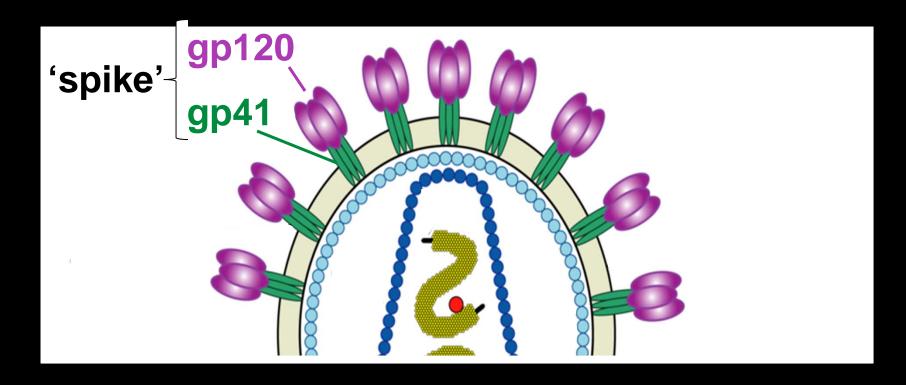
HIV: Human Immunodeficiency Virus



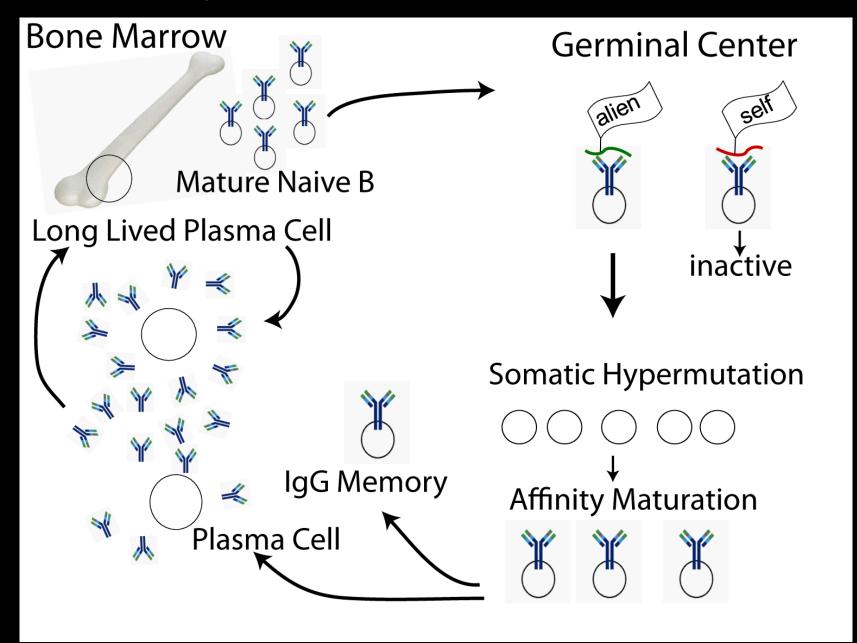
Killed over 25 million people since 1981

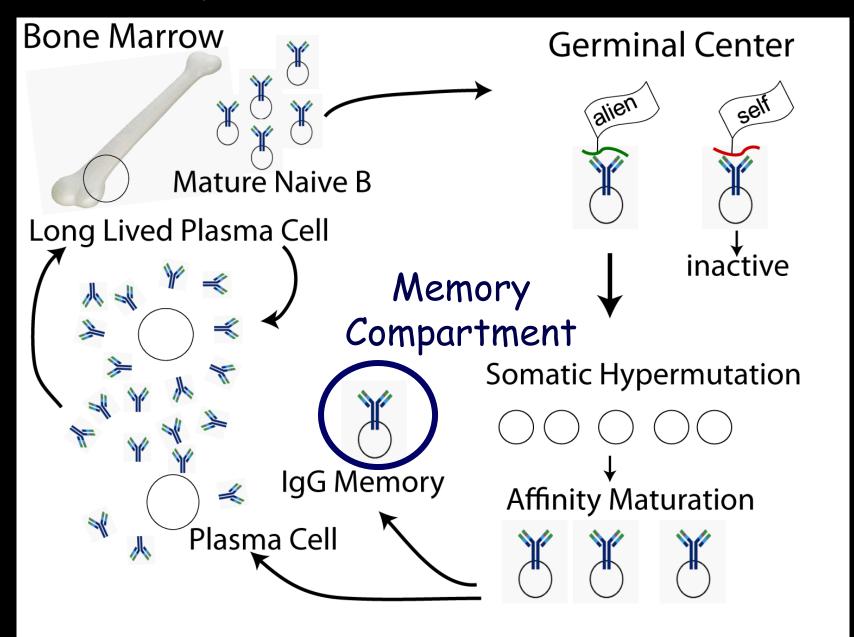
Nan Konstantinov, Yury Stefanov, Aleksander Currently 34 million people are infected Science Company)

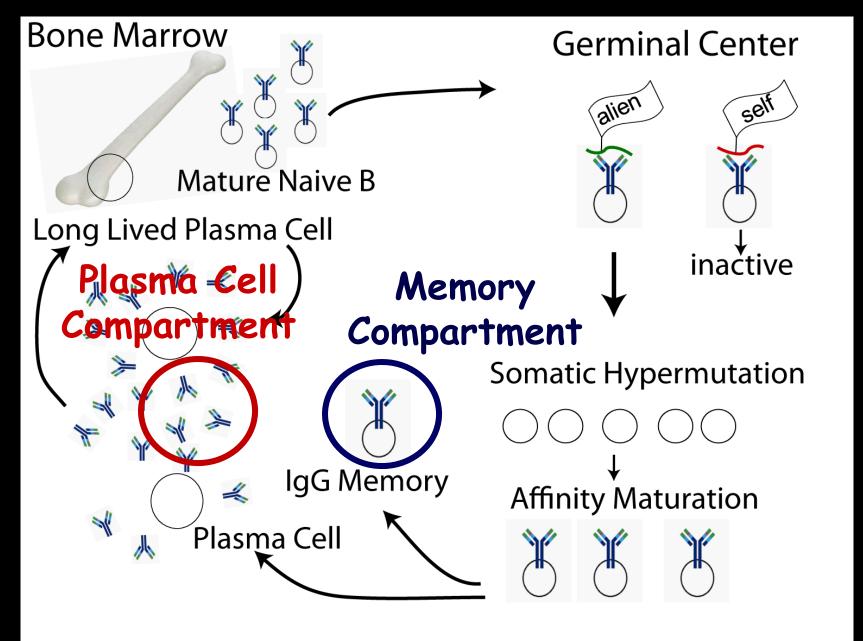
Long-Term Non-Progressors & Elite Controllers of HIV



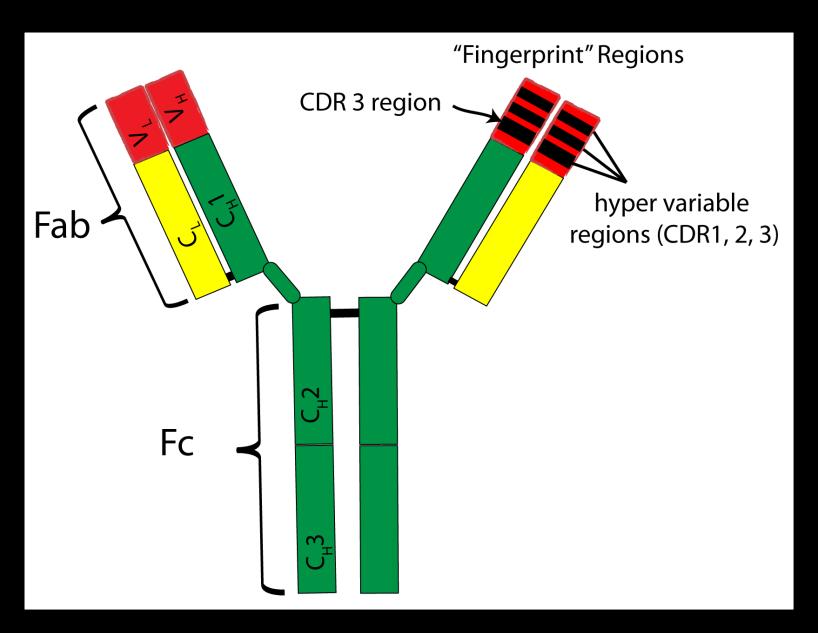
Individuals who control viral replication to low or undetectable levels without anti-retroviral therapy



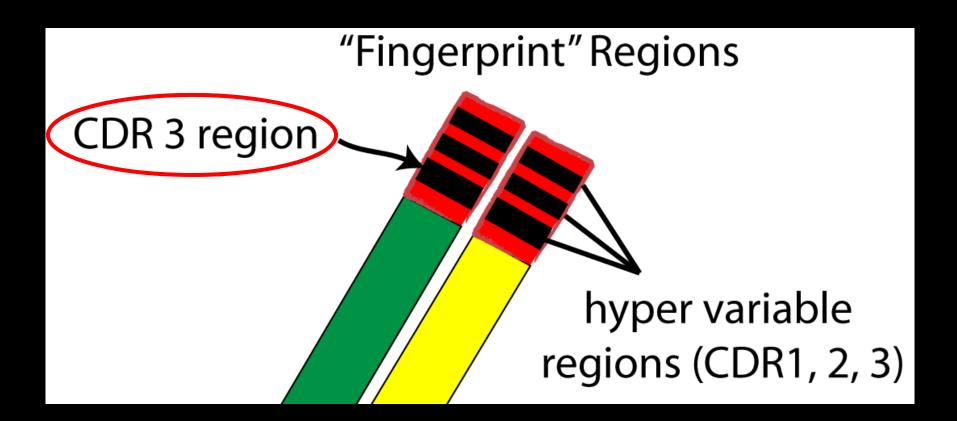




The IgG Molecule



The IgG Molecule



Representative Examples of CDR 3 regions

	FWR3	CDR3	J	CH1
Patient '	1			
Clone 1	CARC	GFRGSPFSSGSMYFDSV	VGQGSQVIVSA	ASTKGPS
Clone 2	CAT	GFRGSPFSTGSFYFDS	WGQGTLVTVSS	ASTKGPS
Clone 3	CARAVITDLHTF	GDYELEDPSYYYMDV	WGEGTTVTVSS	ASTKGPS
Clone 4	CARDTTTF	GAFGGSPNMGGLDP	WGHGTLVTVSS	ASTKGPS
Patient :	2			
Clone 1	CAR	GRLLMQWPPRGGFDP	WGQGTPVIVSS	ASTKGPS
Clone 2	CAR	GRLFVQWPPQGGFDP	WGQGTLVIVSS	ASTKGPS
Clone 3	CA	RTRNTGNSLPYWFDL	WGQGTLVTVSS	ASTKGPS
Clone 4		CVRHIAVGGREEE	WGQGILVAVSS	ASTKGPS

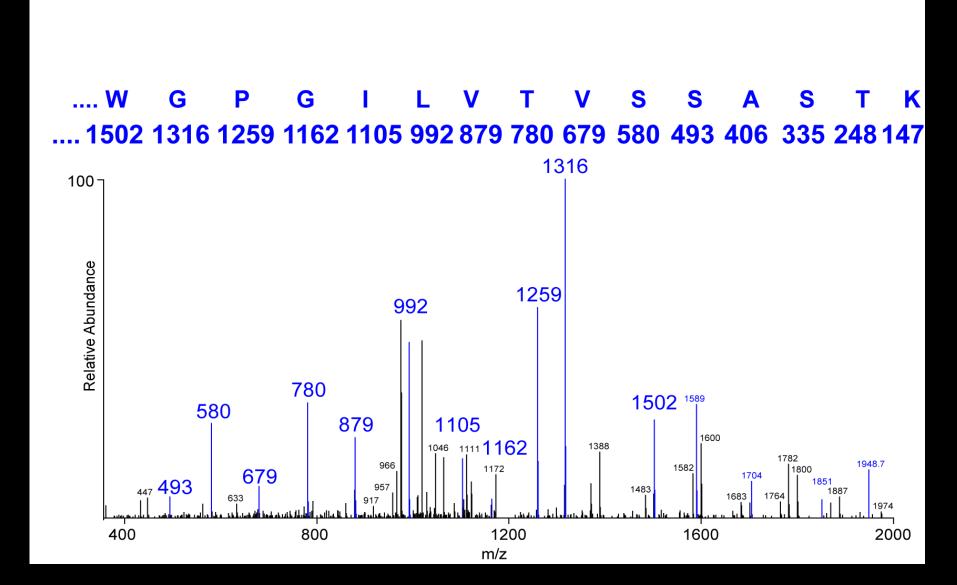
Representative Examples of CDR 3 regions Trypsin Cleavage

	FWR3	CDR3	J	C	H1
Patient	1				
Clone 1	CA <mark>R</mark> G	F <mark>R</mark> GSPFSSGSMYFDS	WGQGSQVIVSA	AST K	GPS
Clone 2	CATO	GF <mark>R</mark> GSPFSTGSFYFDS	WGQGTLVTVSS	AST K	GPS
Clone 3	CARAVITDLHTF	GDYELEDPSYYYMDV	WGEGTTVTVSS	AST K	GPS
Clone 4	CARDTTTF	GAFGGSPNMGGLDP	WGHGTLVTVSS	AST K	GPS
Patient 2	2				
Clone 1	CARC	RLLMQWPP <mark>R</mark> GGFDF	PWGQGTPVIVS	AST K	GPS
Clone 2	CAR	G <mark>R</mark> LFVQWPPQGGFDF	PWGQGTLVIVS	AST K	GPS
Clone 3	CA	RTR <u>N</u> TGNSLPYWFDL	WGQGTLVTVSS	AST K	GPS
Clone 4		CVRHIAVGGREE	EWGQGILVAVS	AST K	GPS

Representative Examples of CDR 3 regions Trypsin Cleavage

	FWR3	CDR3	J	CH1
Patient	1			
Clone 1	CARC	GF <mark>R</mark> GSPFSSGSMYFDS	WGQGSQVIVSAA:	ST <mark>K</mark> GPS
Clone 2	CAT	GF <mark>R</mark> GSPFSTGSFYFDS	WGQGTLVTVSSA:	st <mark>k</mark> gps
Clone 3	CARAVITDLHTF	GDYELEDPSYYYMDV	WGEGTTVTVSSA:	st <mark>k</mark> gps
Clone 4	CARDTTTF	GAFGGSPNMGGLDP	WGHGTLVTVSSA:	st <mark>k</mark> gps
Patient :	2			
Clone 1	CARC	G <mark>R</mark> LLMQWPP <mark>R</mark> GGFDF	WGQGTPVIVSSA	ST <mark>K</mark> GPS
Clone 2	CAR	G <mark>R</mark> LFVQWPPQGGFDI	PWGQGTLVIVSSA	ST <mark>K</mark> GPS
Clone 3	CA	RTRNTGNSLPYWFDL	WGQGTLVTVSSA	ST <mark>K</mark> GPS
Clone 4		CVRHIAVGGREE	WGQGILVAVSSA	ST <mark>K</mark> GPS

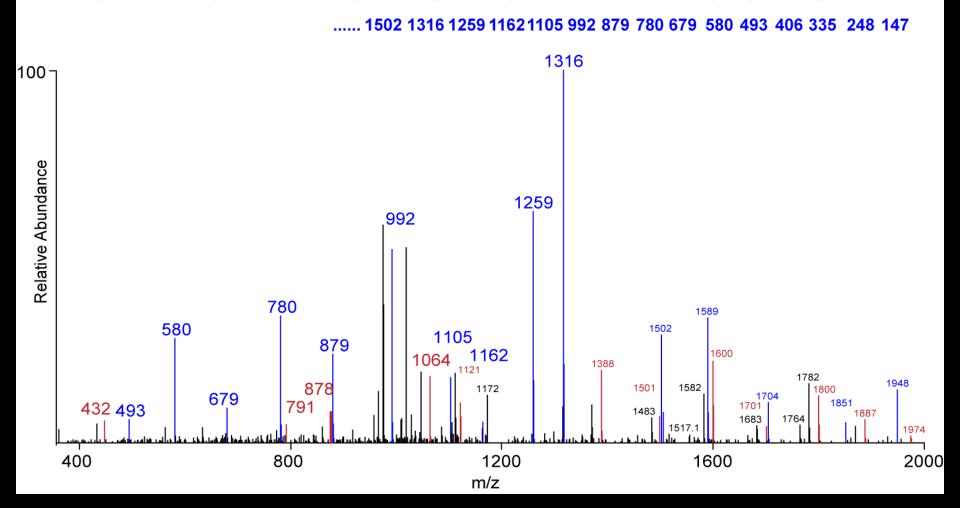
C-terminal Fragment Ions Serve as Markers for most CDR3 Peptides



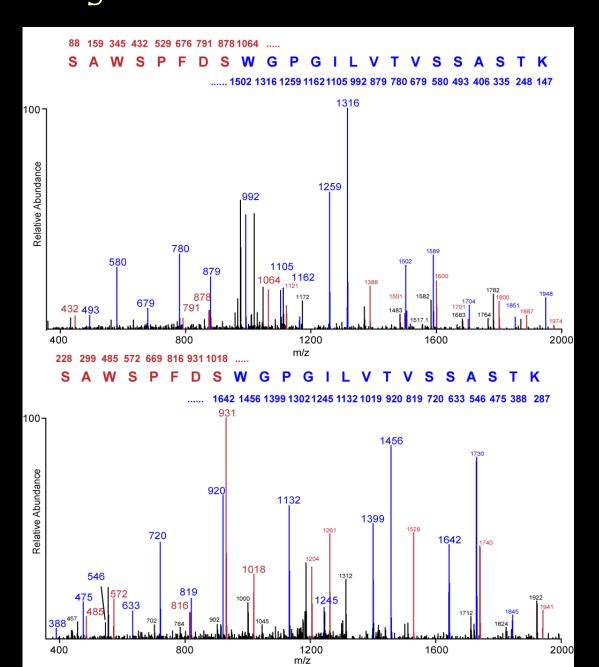
N-terminal Fragment Ions Identify the CDR3 Region

88 159 345 432 529 676 791 878 1064

SAWSPFDSWGPGILVTVSSASTK

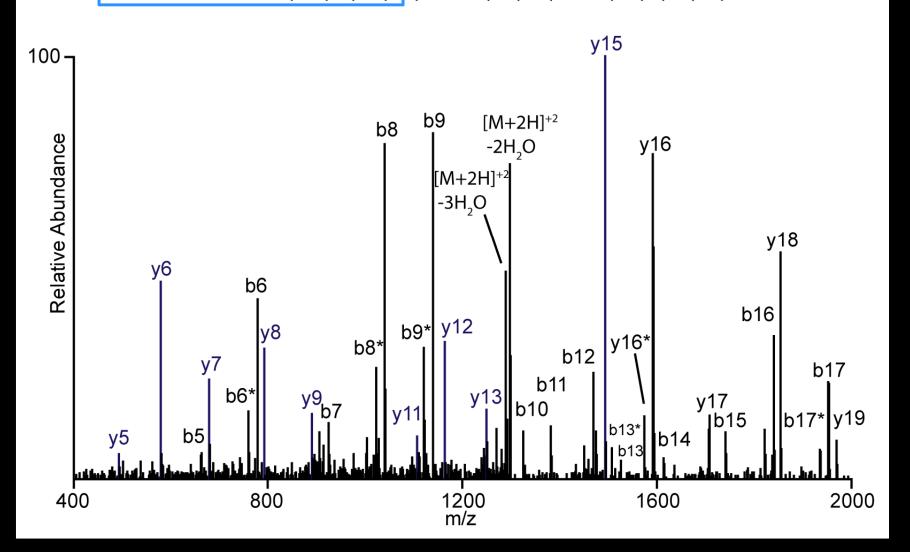


mTRAQ tag enhances N-terminal ion series

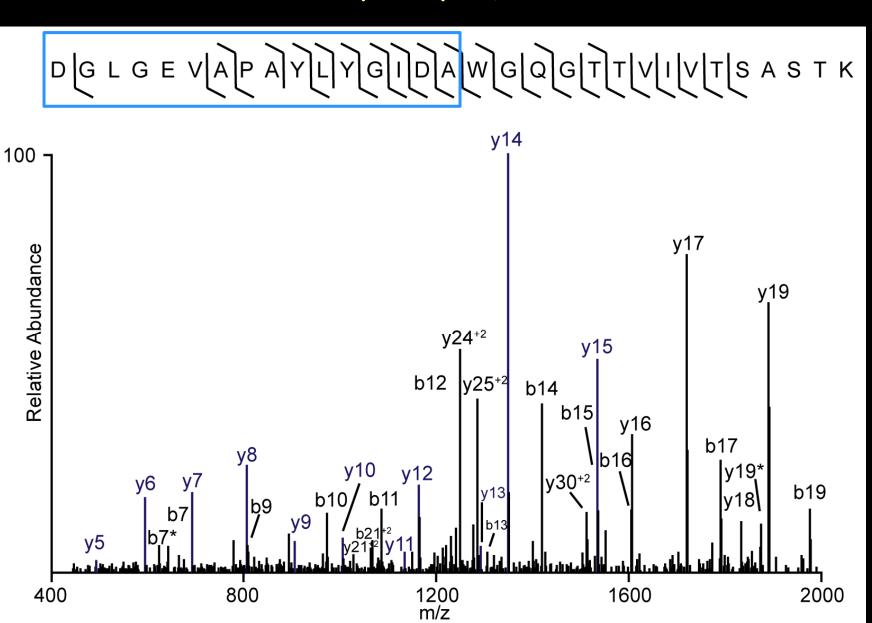


Individual 1

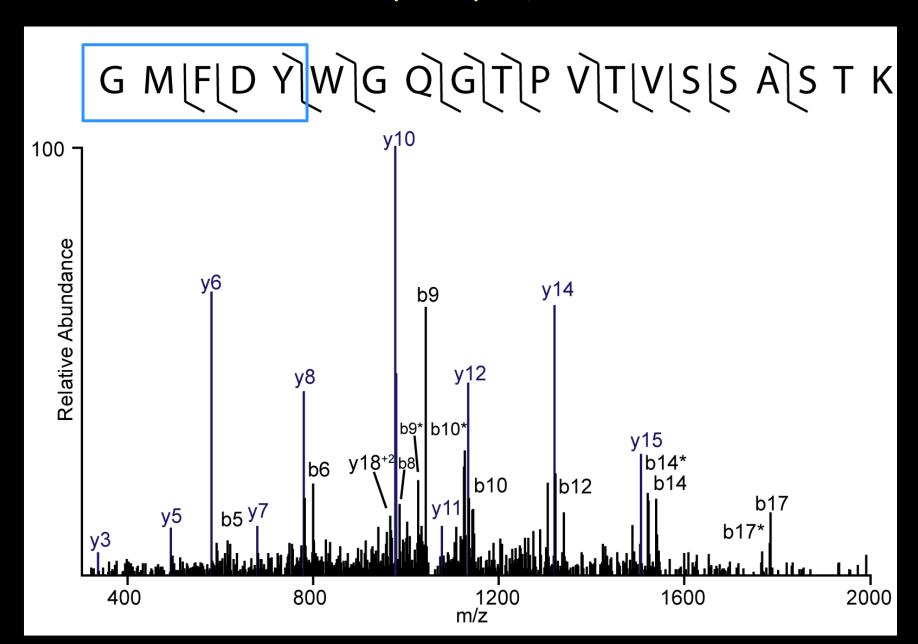
H S D Y CDDFDVWGSGSGSQVIVVSS A S T K



Individual 2

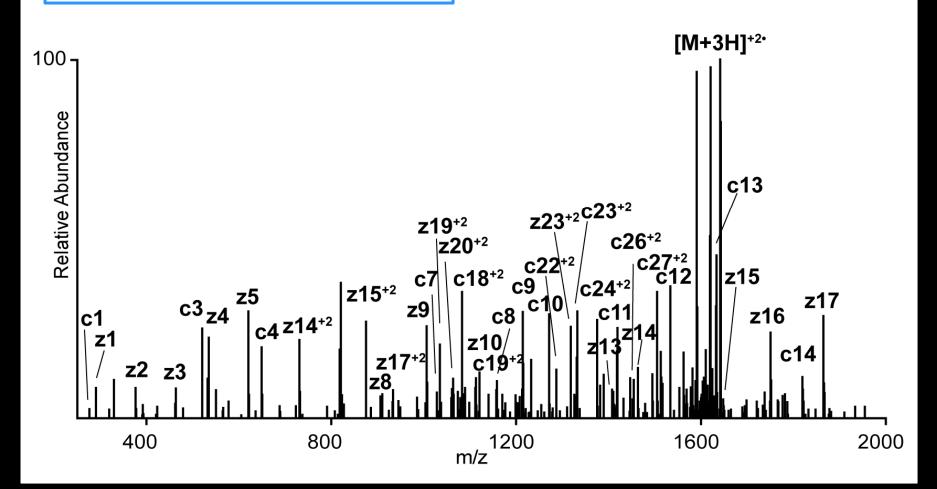


Individual 3



ETD after mTRAQ labeling

LIFIVIQUE PPOLGIGIFIDIT WIGIOIG TILIVIT VISISIAISITIK



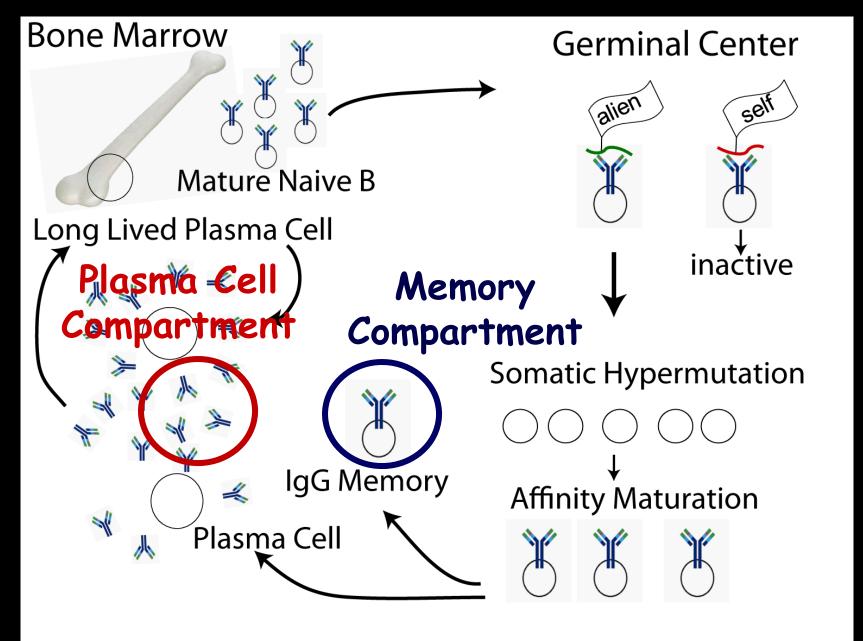
Protocol: Elucidating Memory B Cells

Enrich Patients B Cells

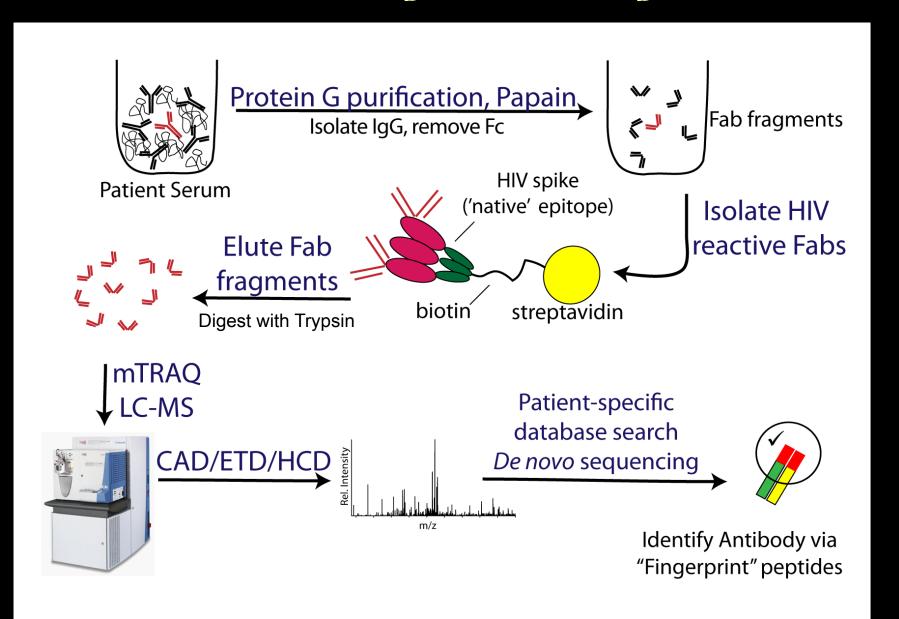
Isolate HIV Spike Binding IgG+ B Cells

Amplify & Sequence Heavy & Light Chains of IgG from these HIV Spike Binding IgG⁺ B Cells

Patient Specific HIV Spike Binding Memory B Cell Antibody Database



Protocol: Elucidating Circulating Antibodies



First Findings

Unamigiously Identified Clones

'ASTK' peptides

AD27HC LSS 167 LC LSS_319_HC LSS 479LC LSS 491 HC LSS 521 HC LSS 527 HC LSS 540 HC LSS 684 KC LSS 68HC LSS 693 HC LSS 705 HC LSS 711_HC LSS 774 KC LSS_779_KC LSS 79 HC LSS 809 HC LSS 81 LC LSS_82_HC LSS 880 HC LSS 89 KC

GGPNMGGFDPWGQGTLVTVSSASTK QIGDYWGQGTLVTVSSASTK FTSPFDSWGQGTLVTVSSASTK QIGDYLGQGTLVTVSSASTK MGGLDPWGQGTLVTVSSASTK FTSPFDSWGQGTLVTVSSASTK GHSFTSPFDSWGQGTLVTVSSASTK DAFDIWGPGTMVTVSSASTK SPFDSWGPGILVTVSSASTK **NPLDIWGQGTMVTVSSASTK** LDVWGQGTTVTVSSASTK STPLVWPPANGLDVWGQGTTVTVSSASTK **EGWGQGTLVTVSSASTK** LDHWGOGTLVTVSSASTK DTTTFTTFGGGPNMGGFDPWGQGTLVTVSSASTK YYHYGLDVWGQGTMVTVSSASTK



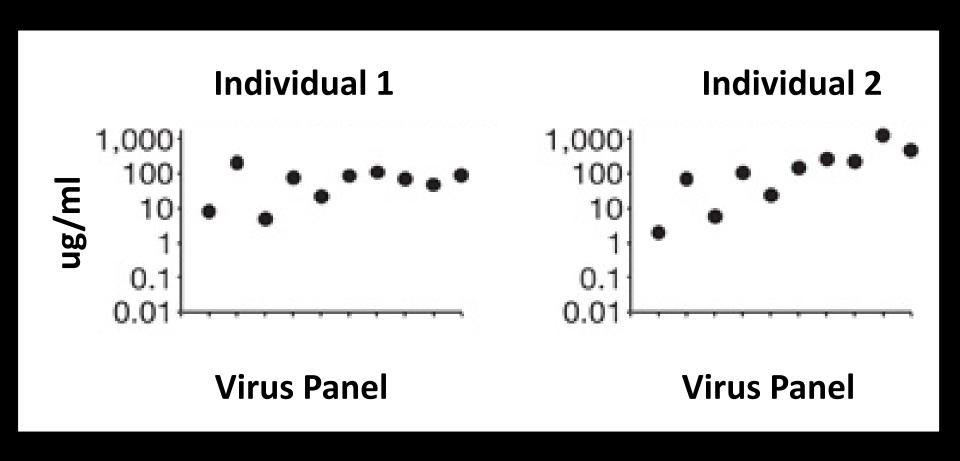
additional variants

Additional Clonal Variants (not in the database)



about 25% of identified peptides carry further mutations

Neutralization (IC50) of IgG Clones

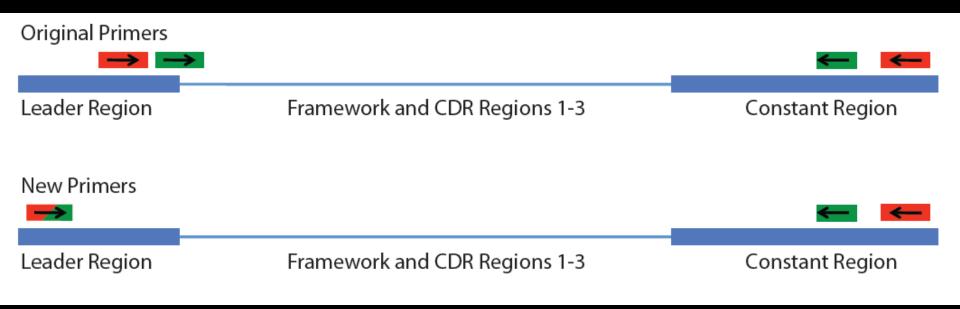


Is there bias in our data?

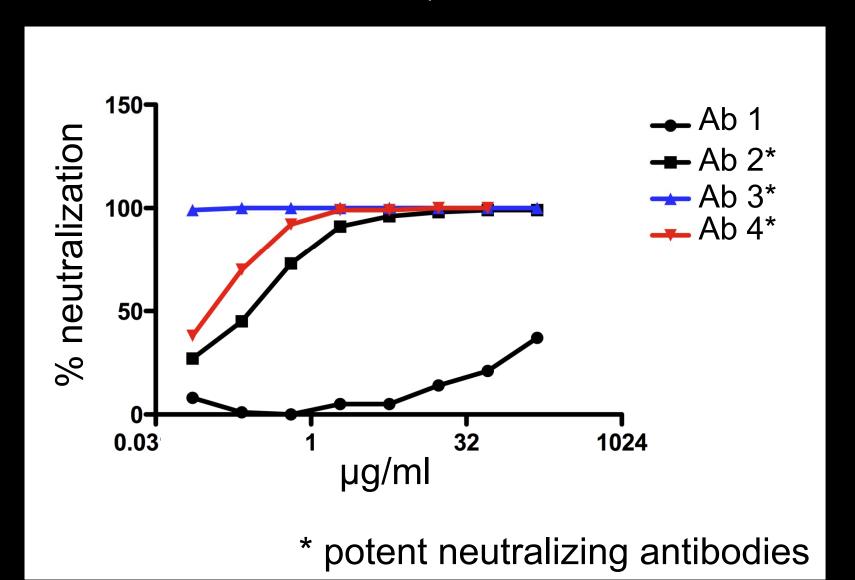
Is there bias in our data?

⇒ primers used for cloning the memory B cells

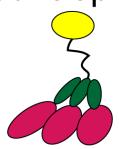
Highly mutated antibodies captured with new primers & amplification strategy



One Potent Broadly Neutralizing Clone per Elite Controller



'Native Spike'

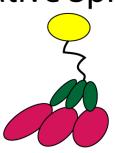


Identified Clones

3white 3A816HC 3white 3B8HC 3white 3A816KC 3white 3B93HC 3white_3A93HC 3white_3BNC149HC 3white_3ANC3HC 3white 3BNC48HC 3white_3ANC66HC 3white_3BNC58HC 3white_3b129kc 3white_3BNC78HC 3white_3B186HC 3white_3A125HC 3white_3A144HC 3white_3b27kc 3white_3B2HC 3white 3A160HC 3white 3B44HC 3white 3A18KC 3white 3B45HC 3white 3A204KC 3white 3b45kc 3white 3A228HC 3white 3b46HC 3white 3A244HC 3white 3b46kc 3white 3A255HC 3white 3B51HC 3white 3A296HC

3white 3A296KC 3white 3A334LC 3white_3A650HC 3white_3A67HC 3white 3A67KC 3white_3A381HC 3white_3A384HC 3white_3a426kc 3white 3A474KC 3white 3A518KC 3white 3A539HC 3white_3A539LC 3white 3A576HC 3white 3A64HC 3red 3bnc105kc 3red_3B106HC 3red 3BNC81HC

'Native Spike'



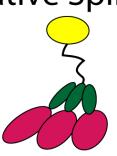
Identified Clones

3white 3A816HC 3white 3B8HC 3white 3A816KC 3white 3B93HC 3white_3A93HC 3white_3BNC149HC 3white_3ANC3HC 3white 3BNC48HC 3white_3ANC66HC 3white_3BNC58HC 3white_3b129kc 3white_3BNC78HC 3white_3B186HC 3white_3A125HC 3white_3A144HC 3white_3b27kc 3white_3B2HC 3white 3A160HC 3white 3B44HC 3white 3A18KC 3white 3B45HC 3white 3A204KC 3white 3b45kc 3white 3A228HC 3white 3b46HC 3white 3A244HC 3white 3b46kc 3white 3A255HC 3white 3B51HC 3white 3A296HC

3white 3A296KC 3white 3A334LC 3white_3A650HC 3white_3A67HC 3white 3A67KC 3white_3A381HC 3white_3A384HC 3white_3a426kc 3white 3A474KC 3white 3A518KC 3white 3A539HC 3white_3A539LC 3white 3A576HC Swnite 3A64HC 3red 3bnc105kc 3red 3B106HC

3red_3BNC81HC

'Native Spike'



3white_3A296KC 3white 3A334LC

3white_3A650HC

3white_3A67HC

3white 3A67KC

3white_3A381HC

3white_3A384HC

3white_3a426kc

3white 3A474KC

3white 3A518KC

3white 3A539HC

3white 3A539LC

3white 3A576HC

Swnite 3A64HC

3red_3bnc105kc 3red_3B106HC 3red_3BNC81HC

Identified Clones

3white 3B8HC 3white 3A816HC 3white 3A816KC 3white 3B93HC 3white_3A93HC 3white 3BNC149HC 3white_3ANC3HC 3white 3BNC48HC 3white 3ANC66HC 3white 3BNC58HC 3white_3b129kc 3white_3BNC78HC 3white_3B186HC 3white_3A125HC 3white_3b27kc 3white 3A144HC 3white_3B2HC 3white 3A160HC 3white 3B44HC 3white 3A18KC 3white 3B45HC 3white 3A204KC 3white 3b45kc 3white 3A228HC 3white 3b46HC 3white 3A244HC 3white 3b46kc 3white 3A255HC 3white 3B51HC 3white 3A296HC

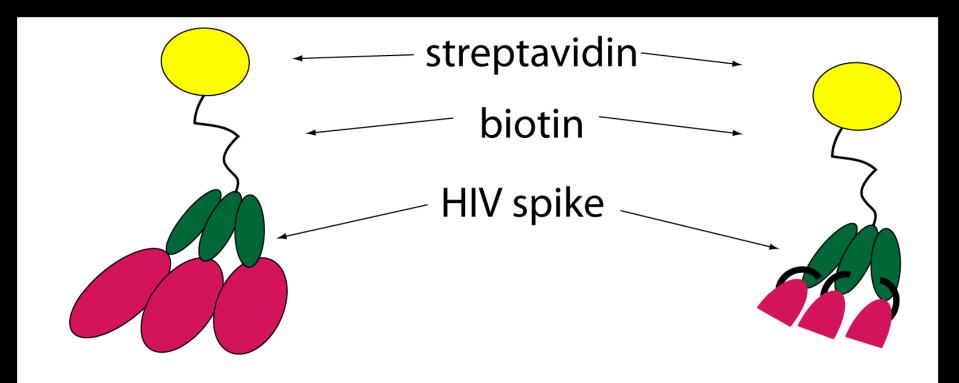
Identified characteristic peptides from potent neutralizers

LLIYDGSR TGQPNNPR DTFQEILFMDLR

Is there any further bias in our data?

- ⇒ primers used for cloning the memory B cells
- ⇒ Bait used for enriching HIV reactive IgGs

Bait Construct



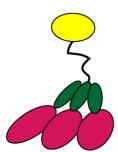
artificially stabilized

HIV spike

trimmed down gp120 artificially exposing **CD4 binding site**

Chen et al Nature 433, 834 (2005)
Dey et al PLoS Pathog. 5, e1000445 (2009)

'Native Spike'



3white 3A125HC 3white_3A144HC 3white 3A160HC 3white_3A18KC 3white 3A204KC 3white 3A228HC 3white 3A244HC 3white_3A255HC 3white 3A296HC 3white_3A296KC 3white 3A334LC 3white_3A381HC 3white_3A384HC 3white_3a426kc 3white_3A474KC 3white 3A518KC 3white_3A539HC 3white_3A539LC 3white 3A576HC 3white_3A64HC 3white 3A650HC 3white_3A67HC

3white 3A67KC

3white_3A816HC 3white_3A816KC 3white 3A93HC 3white_3ANC3HC 3white 3ANC66HC 3white 3b129kc 3white 3B186HC 3white_3b27kc 3white 3B2HC 3white_3B44HC 3white 3B45HC 3white_3b45kc 3white_3b46HC 3white 3b46kc 3white_3B51HC 3white 3B8HC 3white_3B93HC 3white_3BNC149HC 3white 3BNC48HC 3white_3BNC58HC 3white 3BNC78HC 3red_3bnc105kc 3red 3B106HC 3red 3BNC81HC

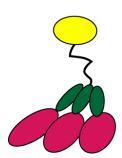
'exposed CD4 site'



LSSwhite LSSNEC106HC LSSwhite_LSSNEC77HC LSSwhite LSSNEC91HC LSSwhite 154 LSSwhite LSS 167 LC LSSwhite LSS 187 KC LSSwhite LSS 191 KC LSSwhite_LSS_309_HC LSSwhite LSS 380 LC LSSwhite_LSS_408_LC LSSwhite LSS 41 LC LSSwhite_LSS_479LC LSSwhite LSS 494 LC LSSwhite_LSS_505_KC LSSwhite_LSS_531_KC LSSwhite LSS 612 LC LSSwhite_LSS_687_LC LSSwhite LSS 693 HC LSSwhite_LSS_695_LC LSSwhite_LSS_706_LC LSSwhite LSS 71 HC LSSwhite_LSS_73_LC LSSwhite LSS 74HC LSSwhite LSS 750 KC LSSwhite LSS 752 LC LSSwhite LSS 763 HC LSSwhite LSS 774 KC LSSwhite_LSS_81_LC

LSSwhite_LSS_863_HC LSSwhite_LSS_880_HC LSSwhite LSS 91 LC LSSadd LSSB2066HC LSSgreen LSSB2364LC LSSgreen LSSB2609HC LSSgreen LSSB2612HC LSSgreen_LSSNEC108HC LSSgreen LSSNEC108LC LSSgreen_LSSNEC109HC LSSgreen LSSNEC110HC LSSgreen_LSSNEC110HC LSSgreen_LSSNEC117HC LSSgreen_LSSNEC122HC LSSgreen_LSSNEC123HC LSSgreen LSSNEC18HC LSSgreen_LSSNEC24HC LSSgreen_LSSNEC24LC LSSgreen_LSSNEC3HC LSSgreen_LSSNEC46HC LSSgreen LSSNEC46HC LSSgreen_LSSNEC46HC LSSgreen LSSNEC60HC LSSgreen LSSNEC70LC LSSgreen LSSNEC7HC LSSgreen LSSNEC89LC

'Native Spike'



3white 3A125HC 3white_3A144HC 3white 3A160HC 3white_3A18KC 3white 3A204KC 3white 3A228HC 3white 3A244HC 3white_3A255HC 3white 3A296HC 3white 3A296KC 3white 3A334LC 3white_3A381HC 3white_3A384HC 3white 3a426kc 3white_3A474KC 3white 3A518KC 3white_3A539HC 3white_3A539LC 3white 3A576HC 3white_3A64HC 3white 3A650HC 3white_3A67HC 3white 3A67KC

3white_3A816HC 3white_3A816KC 3white 3A93HC 3white_3ANC3HC 3white 3ANC66HC 3white 3b129kc 3white 3B186HC 3white_3b27kc 3white 3B2HC 3white_3B44HC 3white 3B45HC 3white_3b45kc 3white_3b46HC 3white 3b46kc 3white_3B51HC 3white 3B8HC 3white_3B93HC 3white_3BNC149HC 3white_3BNC48HC 3white_3BNC58HC 3White_SDNC78HC 3red_3bnc105kc 3red 3B106HC 3red 3BNC81HC

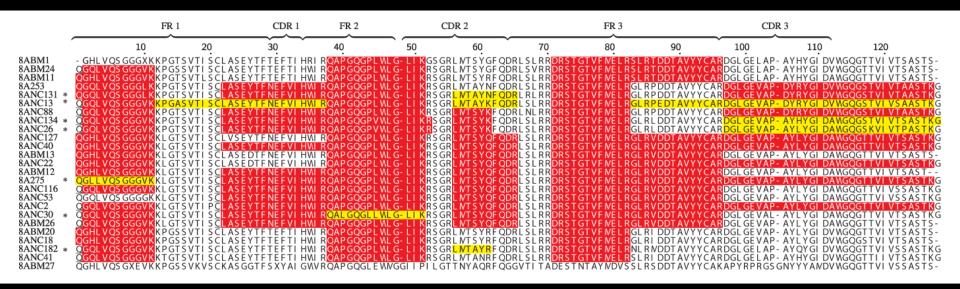
'exposed CD4 site'



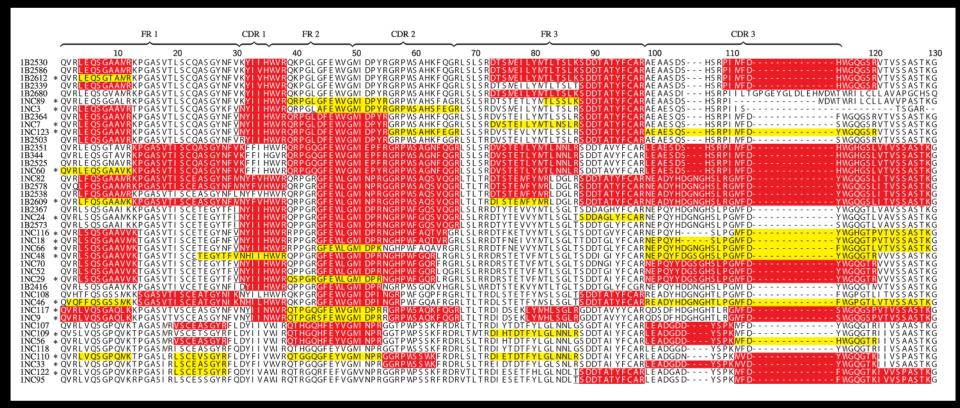
LSSwhite LSSNEC106HC LSSwhite_LSSNEC77HC LSSwhite LSSNEC91HC LSSwhite 154 LSSwhite LSS 167 LC LSSwhite LSS 187 KC LSSwhite LSS 191 KC LSSwhite LSS 309 HC LSSwhite LSS 380 LC LSSwhite_LSS_408_LC LSSwhite LSS 41 LC LSSwhite_LSS_479LC LSSwhite LSS 494 LC LSSwhite_LSS_505_KC LSSwhite_LSS_531_KC LSSwhite LSS 612 LC LSSwhite_LSS_687_LC LSSwhite LSS 693 HC LSSwhite_LSS_695_LC LSSwhite_LSS_706_LC LSSwhite LSS 71 HC LSSwhite_LSS_73_LC LSSwhite LSS 74HC LSSwhite LSS 750 KC LSSwhite LSS 752 LC LSSwhite LSS 763 HC LSSwhite LSS 774 KC LSSwhite LSS 81 LC

LSSwhite_LSS_863_HC LSSwhite_LSS_880_HC LSSwhite LSS 91 LC LSSadd LSSB2066HC LSSgreen LSSb2364LC LSScreen LSSB2609HC LSSgreen LSSB2612H SSgreen LSSNEC108H LSSgreen LSSNEC108LC LSSgreen_LSSNEC109HC LSSgreen LSSNEC110HC LSSgreen_LSSNEC110HC LSSgreen_LSSNEC117HC LSSgreen_LSSNEC122HC LSSgreen_LSSNEC123HC LSSgreen LSSNEC18HC LSSgreen_LSSNEC24HC LSSgreen_LSSNEC24LC LSSgreen_LSSNEC3HC LSSgreen_LSSNEC46HC LSSgreen LSSNEC46HC LSSgreen_LSSNEC46HC LSSgreen LSSNEC60HC LSSgreen_LSSNEC70LC SSgreen LSSNEC7HC LS. green_LSSNEC89L

Elite Controller 1



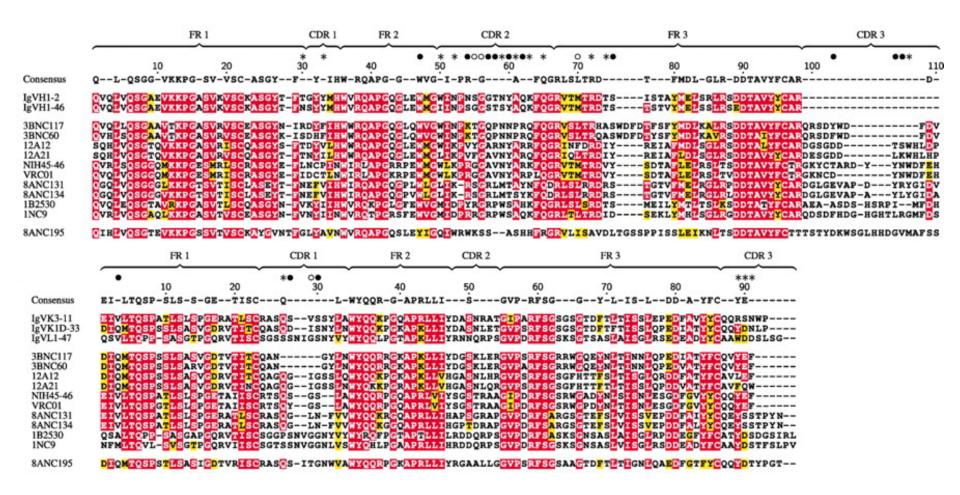
Elite Controller 2



Long-Term Nonprogressor 1



Sequence conservation of highly active agonistic CD4-binding site antibodies



Conclusions

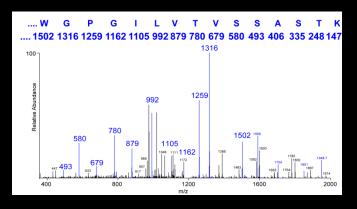
Cloned 576 new HIV Antibodies from 4 unrelated Individuals

All 4 individuals produced expanded clones of potent broadly neutralizing antibodies that mimic binding to CD4

Despite extensive hypermutation, the new antibodies share a consensus of 68 IgH chain amino acids

Conclusions

Our Marker Ion Strategy enables rapid estimation of the different clones present in patient sera with/without a database



For each individual under study we have provided the first direct proof that potent broadly neutralizing antibodies are circulating in the blood (<1% of all HIV reactive IgG)

These technologies are applicable to any disease in which antibodies are induced against a known antigen

Acknowledgements



Beatrix Ueberheide



David Fenyő



Sunnie Myung



Johannes Scheid



Michel Nussenzweig

NIH RR00862 NIH RR022220

