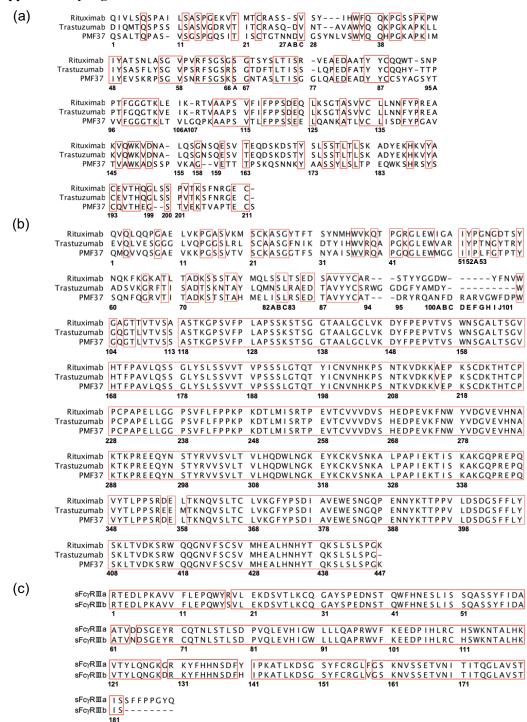
Supplementary information

The Fab portion of immunoglobulin G contributes to its binding to Fcy receptor III

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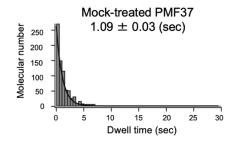
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Supplementary Figures



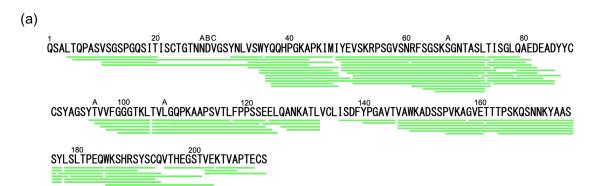
Supplementary Figure 1

Sequence alignments of (a) the light chains and (b) the heavy chains of rituximab, trastuzumab, and PMF37 and (c) sFc γ RIIIa and sFc γ RIIIb. The numbering system used in the present study for the constant region is based on human myeloma protein Eu 1 . The convention of Kabat et al. (1987) has been followed for the numbering of the variable regions 2 . Conserved residues are boxed in red.

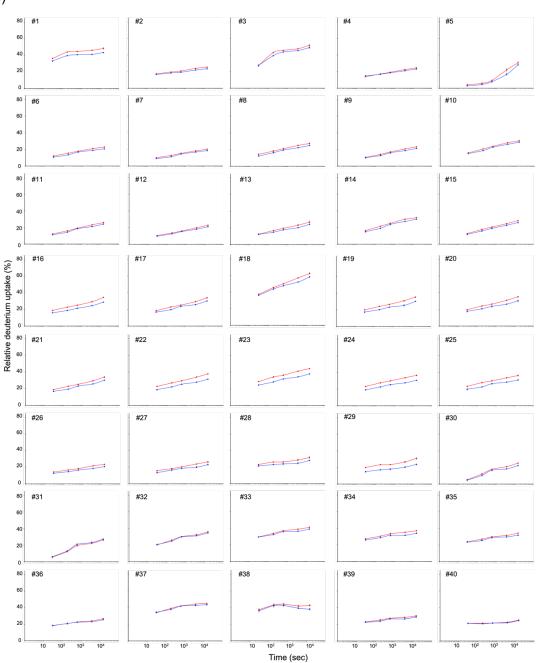


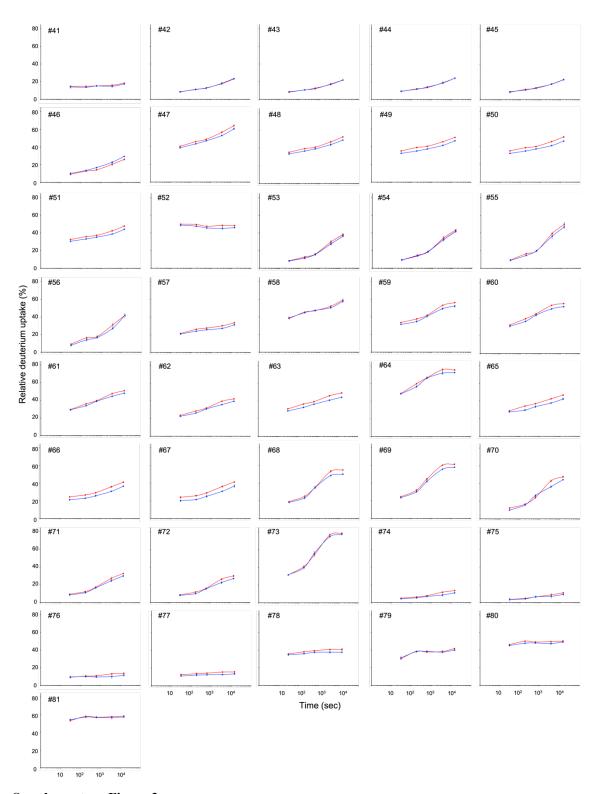
Supplementary Figure 2

Dwell time of mock-treated PMF37 on sFc γ RIIIa.









Supplementary Figure 3

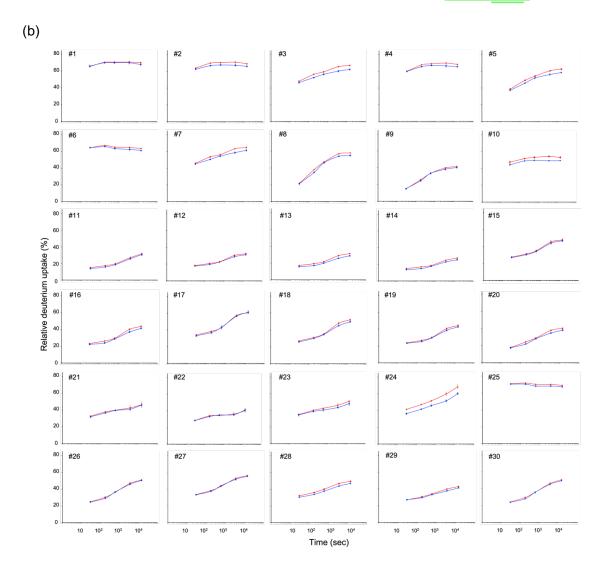
HDX-MS data of the light chain. (a) Coverage map of identified peptides and (b) time-dependent deuterium uptake graphs of all peptides used for deuterium uptake comparison between free (red) and complexed states (blue).

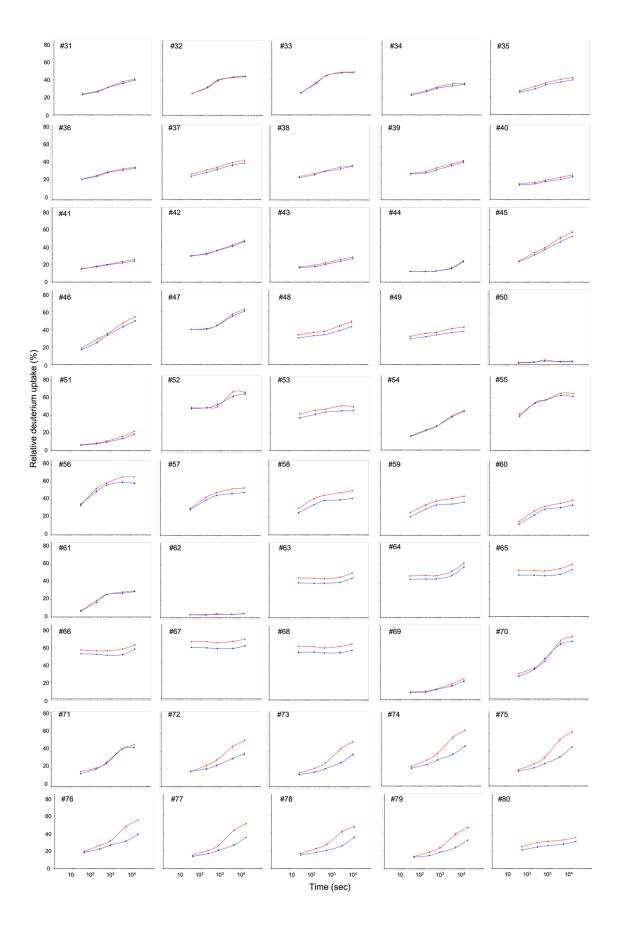
YYCATDRYRQANFDRARVGWFDPWGQGTLVTVSSASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQS

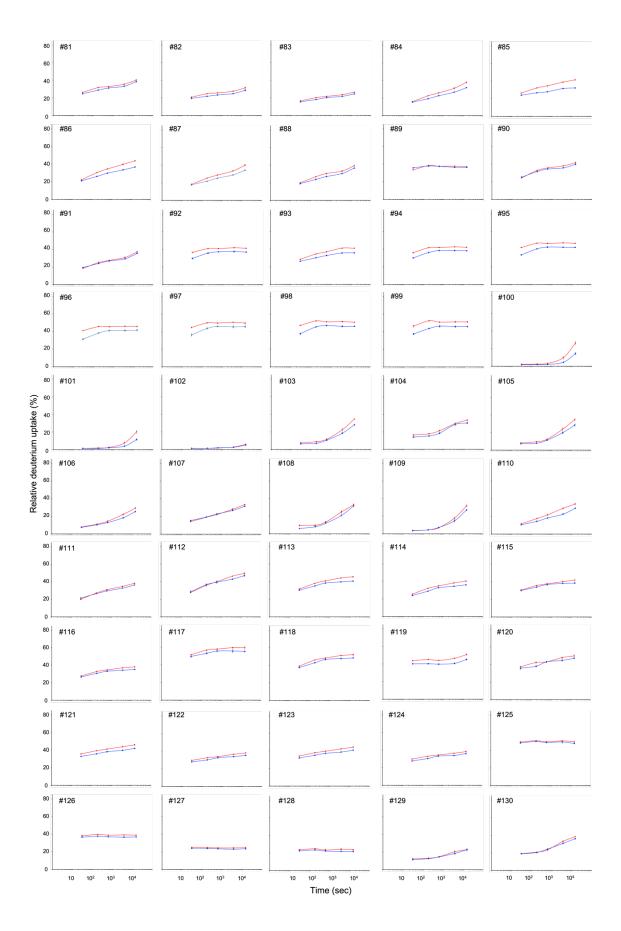
180 SGLYSLSSVVTVPSSSLGTQTY I CNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTLM I SRTPEVTCVVVDVSHE

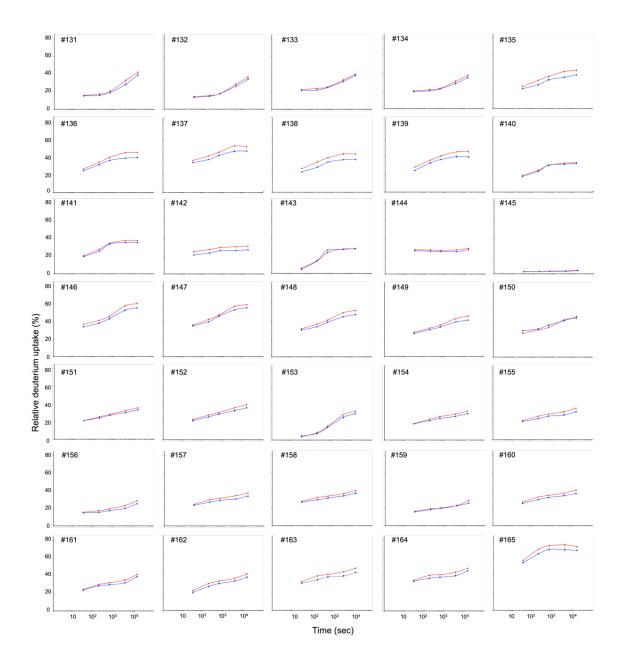
DPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQ

 $\underline{\textbf{VSL}} \underline{\textbf{TCLVKGFYPSD}} \underline{\textbf{IAVEWESNGQPENNYKTTPPVL}} \underline{\textbf{DSDGSFFLYSKL}} \underline{\textbf{TVDKSRWQQG}} \underline{\textbf{NVFSCSVMHEALHNHYTQKSLSLSPGK}} \underline{\textbf{VSL}} \underline{\textbf{TCLVKGFYPSD}} \underline{\textbf{IAVEWESNGQPENNYKTTPPVL}} \underline{\textbf{DSDGSFFLYSKL}} \underline{\textbf{TVDKSRWQQG}} \underline{\textbf{NVFSCSVMHEALHNHYTQKSLSLSPGK}} \underline{\textbf{VSL}} \underline{\textbf{TVDKSRWQQG}} \underline{\textbf{NVFSCSVMHEALHNHYTQKSLSLSPGK}} \underline{\textbf{VSL}} \underline{\textbf{VSL}$









Supplementary Figure 4

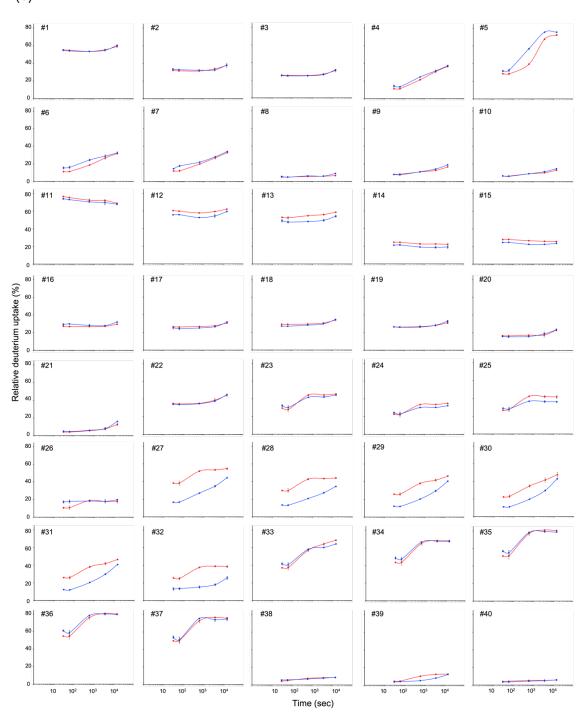
HDX-MS data of the heavy chain. (a) Coverage map of identified peptides and (b) time-dependent deuterium uptake graphs of all peptides used for deuterium uptake comparison between free (red) and complexed states (blue).

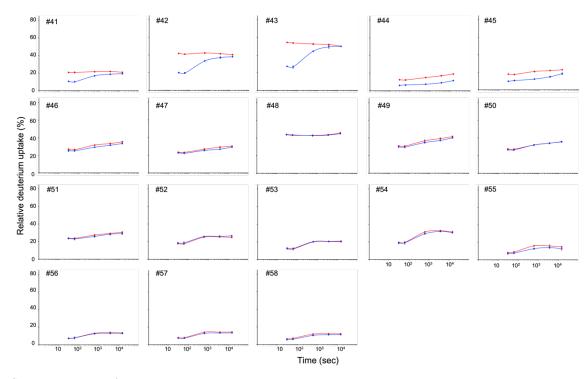
(a)

3
EDLPKAVVFLEPQWYRVLEKDSVTLKCQGAYSPEDNSTQWFHNESL ISSQASSYFIDAAT VDDSGEYRCQTQLSTLSDPVQLEVHIGWLLLQAPRWVF

 $\underbrace{^{120}}_{\text{KEEDPIHL}R\text{CHSWKNTAL}} \underbrace{^{120}}_{\text{IRO}} \underbrace{^{140}}_{\text{IRO}} \underbrace{^{140}}_{\text{IRO}} \underbrace{^{160}}_{\text{IRO}} \underbrace{^{180}}_{\text{IRO}} \underbrace{\text{IRO}} \underbrace{^{180}}_{\text{IRO}} \underbrace{^{180}}_{\text{IRO}} \underbrace$

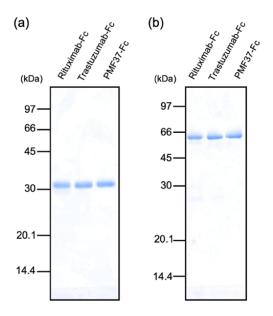
(b)





Supplementary Figure 5

HDX-MS data of $sFc\gamma RIIIa$. (a) Coverage map of identified peptides and (b) time-dependent deuterium uptake graphs of all peptides used for deuterium uptake comparison between free (red) and complexed states (blue).



Supplementary Figure 6

SDS-PAGE of the Fc preparations under (a) reducing and (b) non-reducing conditions.

Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence
#1	L4-T20	#23	V51-S71	#45	V97-L104	#67	T163-Y177
#2	T5-A43	#24	V51-L72	#46	G99-L104	#68	Y177-Q18
#3	V11-T20	#25	V51-T73	#47	T105-L117	#69	L178-Q18
#4	V11-A43	#26	R61-L72	#48	T105-S122	#70	L178-S19
#5	Y30-W35	#27	R61-T73	#49	T105-E123	#71	L178-C19
#6	V33-M47	#28	S63-L72	#50	T105-E124	#72	L178-Q19
#7	V33-I48	#29	S65-L72	#51	T105-L125	#73	S179-Q18
#8	W35-M47	#30	L72-L78	#52	F118-E123	#74	W185-C19
#9	W35-I48	#31	T73-Q79	#53	E123-L132	#75	W185-Q1
#10	Y36-M47	#32	T73-A80	#54	E124-L132	#76	W185-T2
#11	Y36-I48	#33	T73-E81	#55	Q126-L132	#77	W185-E2
#12	Y36-E50	#34	T73-D82	#56	A127-L132	#78	V195-E20
#13	Y36-L72	#35	T73-E83	#57	I136-T145	#79	V202-A20
#14	Q37-M47	#36	T73-D85	#58	Y140-T145	#80	V202-S21
#15	Q37-I48	#37	174-E81	#59	V146-A157	#81	K204-E21
#16	148-L72	#38	174-D82	#60	V146-E160		
#17	148-T73	#39	174-D85	#61	V146-S176		
#18	Y49-N60	#40	Q79-D85	#62	V146-L178		
#19	Y49-L72	#41	Q79-Y86	#63	W148-Y177		
#20	Y49-T73	#42	Y95-L104	#64	P154-E160		
#21	Y49-I74	#43	T95 _A -L104	#65	G158-Y177		
#22	E50-L72	#44	V96-L104	#66	T161-Y177		

)	Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence
	#1	M2-A9	#35	R66-M80	#69	Y198-T209	#103	L306-E318	#137	W381-N390
	#2	M2-E10	#36	R66-L82	#70	L235-V240	#104	T307-W313	#138	W381-L398
	#3	M2-S17	#37	T70-M80	#71	L235-F241	#105	T307-E318	#139	E382-L398
	#4	Q3-E10	#38	T70-L82	#72	F241-L251	#106	T307-E333	#140	N389-L398
	#5	Q3-T19	#39	A71-M80	#73	F241-M252	#107	T307-V348	#141	Y391-L398
	#6	V4-E10	#40	A71-L82	#74	L242-L251	#108	L309-E318	#142	Y391-F404
	#7	V4-S17	#41	A71-A100	#75	L242-M252	#109	H310-E318	#143	K392-L398
	#8	V11-T19	#42	D72-M80	#76	F243-L251	#110	Y319-E333	#144	D399-F404
	#9	V11-C22	#43	D72-L82	#77	F243-M252	#111	Y319-V348	#145	F405-L410
	#10	V20-Y32	#44	R83-V89	#78	P244-L251	#112	K326-V348	#146	T411-G420
	#11	Y32-L45	#45	D100c-L108	#79	P244-M252	#113	K334-V348	#147	T411-V422
	#12	A33-G44	#46	R100o-L108	#80	M252-C261	#114	K334-L358	#148	T411-F423
	#13	A33-L45	#47	V109-A118	#81	I253-T260	#115	K334-S364	#149	T411-C425
	#14	A33-E46	#48	V109-F126	#82	1253-C261	#116	K334-L365	#150	K414-F423
	#15	W36-G44	#49	V109-L142	#83	1253-V262	#117	T335-V348	#151	S424-G446
	#16	W36-L45	#50	V109-T155	#84	V262-W277	#118	1336-V348	#152	S426-G446
	#17	V37-G44	#51	S119-N159	#85	V263-V273	#119	Y349-D356	#153	V427-L432
	#18	V37-L45	#52	P127-S132	#86	V263-F275	#120	Y349-N361	#154	V427-G446
	#19	V37-E46	#53	P127-L142	#87	V263-W277	#121	Y349-S364	#155	M428-G446
	#20	V37-M48	#54	V146-W158	#88	D265-W277	#122	Y349-T366	#156	H429-L441
	#21	E46-N62	#55	V156-A162	#89	V266-V273	#123	P352-S364	#157	H429-G446
	#22	W47-L53	#56	V156-L163	#90	V266-F275	#124	P352-L365	#158	H429-K447
	#23	W47-N62	#57	V156-L174	#91	V266-W277	#125	E357-S364	#159	A431-L443
	#24	F54-N62	#58	N159-L174	#92	N276-T299	#126	E357-L365	#160	A431-G446
	#25	S60-G65	#59	A162-L174	#93	Y278-E283	#127	T359-L365	#161	H433-S444
	#26	F63-T70	#60	T164-L174	#94	Y278-Y296	#128	T359-T366	#162	H433-L443
	#27	F63-A71	#61	H168-L174	#95	Y278-T299	#129	T366-D376	#163	H433-G446
	#28	F63-M80	#62	Y180-V185	#96	Y278-Y300	#130	V369-D376	#164	H433-K447
	#29	F63-L82	#63	S183-T197	#97	D280-T299	#131	V369-I377	#165	L441-G446
	#30	Q64-A71	#64	V185-L193	#98	V282-T299	#132	V369-A378		
	#31	Q64-L82	#65	V185-T197	#99	V282-Y300	#133	V369-V379		
	#32	R66-A71	#66	V186-L193	#100	Y300-V305	#134	V369-E380		
	#33	R66-D72	#67	V186-T197	#101	Y300-L306	#135	1377-L398		
	#34	R66-A78	#68	P189-T197	#102	R301-L306	#136	E382-L398		

Peptide number	Sequence	Peptide number	Sequence	Peptide number	Sequence
#1	E3-A8	#23	S76-Q83	#45	L124-F139
#2	E3-V10	#24	S76-L84	#46	F139-S15
#3	E3-F11	#25	L78-Q83	#47	F139-Y15
#4	V9-T26	#26	S79-L84	#48	Y140-L146
#5	F11-W16	#27	L84-W90	#49	Y140-S15
#6	L12-D23	#28	L84-L91	#50	Y140-Y15
#7	L12-T26	#29	L84-L92	#51	Y140-F15
#8	Y17-D23	#30	L84-L93	#52	T167-H18
#9	Y17-T26	#31	E85-L93	#53	V168-H18
#10	Y17-L27	#32	H87-L92	#54	I170-H18
#11	I49-S55	#33	L91-F100	#55	T171-H18
#12	149-Y56	#34	L92-F100	#56	I172-H18
#13	149-F57	#35	L93-F100	#57	T173-H18
#14	Y56-T62	#36	Q94-F100	#58	Q174-H18
#15	F57-T62	#37	A95-F100		
#16	158-E68	#38	K101-L108		
#17	A60-E68	#39	K101-C110		
#18	A61-E68	#40	K101-S112		
#19	A61-L75	#41	K101-L118		
#20	T62-E68	#42	R109-L118		
#21	V63-E68	#43	H111-L118		
#22	Y69-L75	#44	H119-F139		

Supplementary Table

Peptide identifications in (a) the light chain and (b) the heavy chain of PMF37 and (c) $sFc\gamma RIIIa$.

Supplementary References

- Edelman GM, Cunningham BA, Gall WE, Gottlieb PD, Rutishauser U, Waxdal MJ. The covalent structure of an entire gammaG immunoglobulin molecule. *Proc Natl Acad Sci U S A* 63, 78-85 (1969).
- 2. Rabat, E. A., Wu, T. T., Reid-Miller, M., Perry, M., & Gottesman, K. S. Sequences of Proteins of Immu-nological Interest, 4th ed., U.S. Department of Health and Human Services, National Institutes of Health, Washington, DC. (1987)