

Supplementary information for

Quadrivalent influenza nanoparticle vaccines induce broad protection

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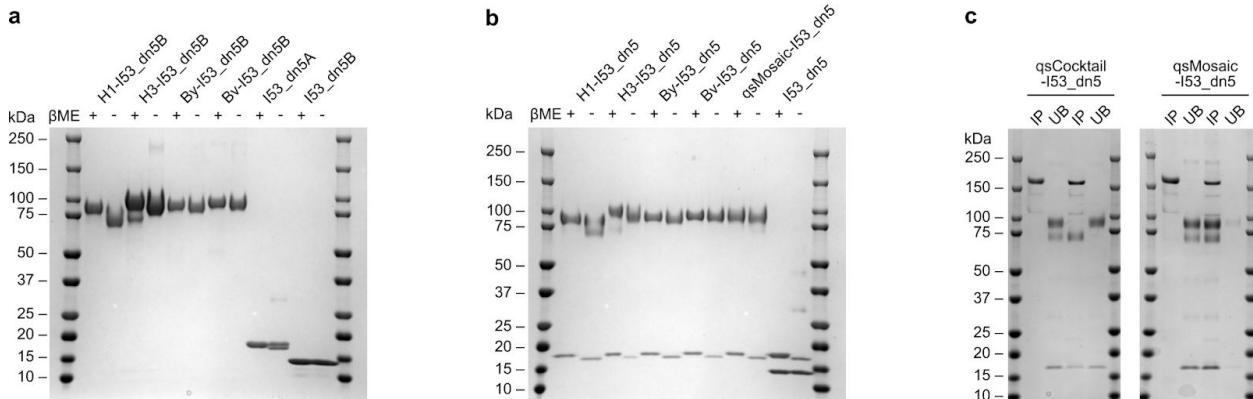
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¹⁰These authors contributed equally: Neil P. King and Masaru Kanekiyo.

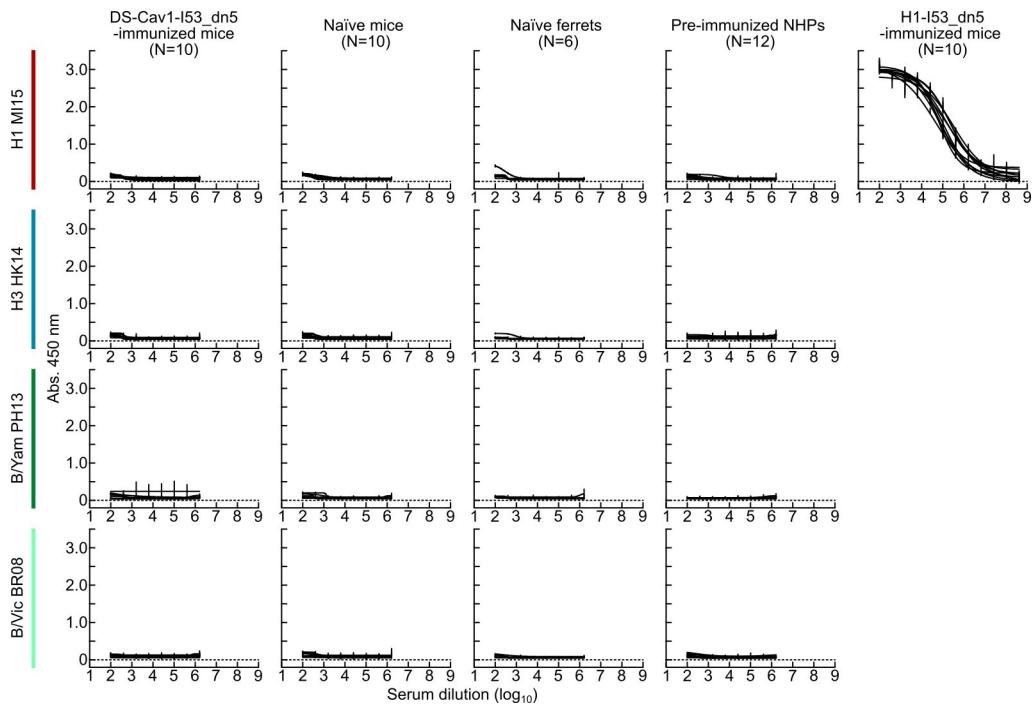
*email: bgraham@nih.gov (B.S.G.); neil@ipd.uw.edu (N.P.K.); kanekiyom@nih.gov (M.K.)

Table of contents

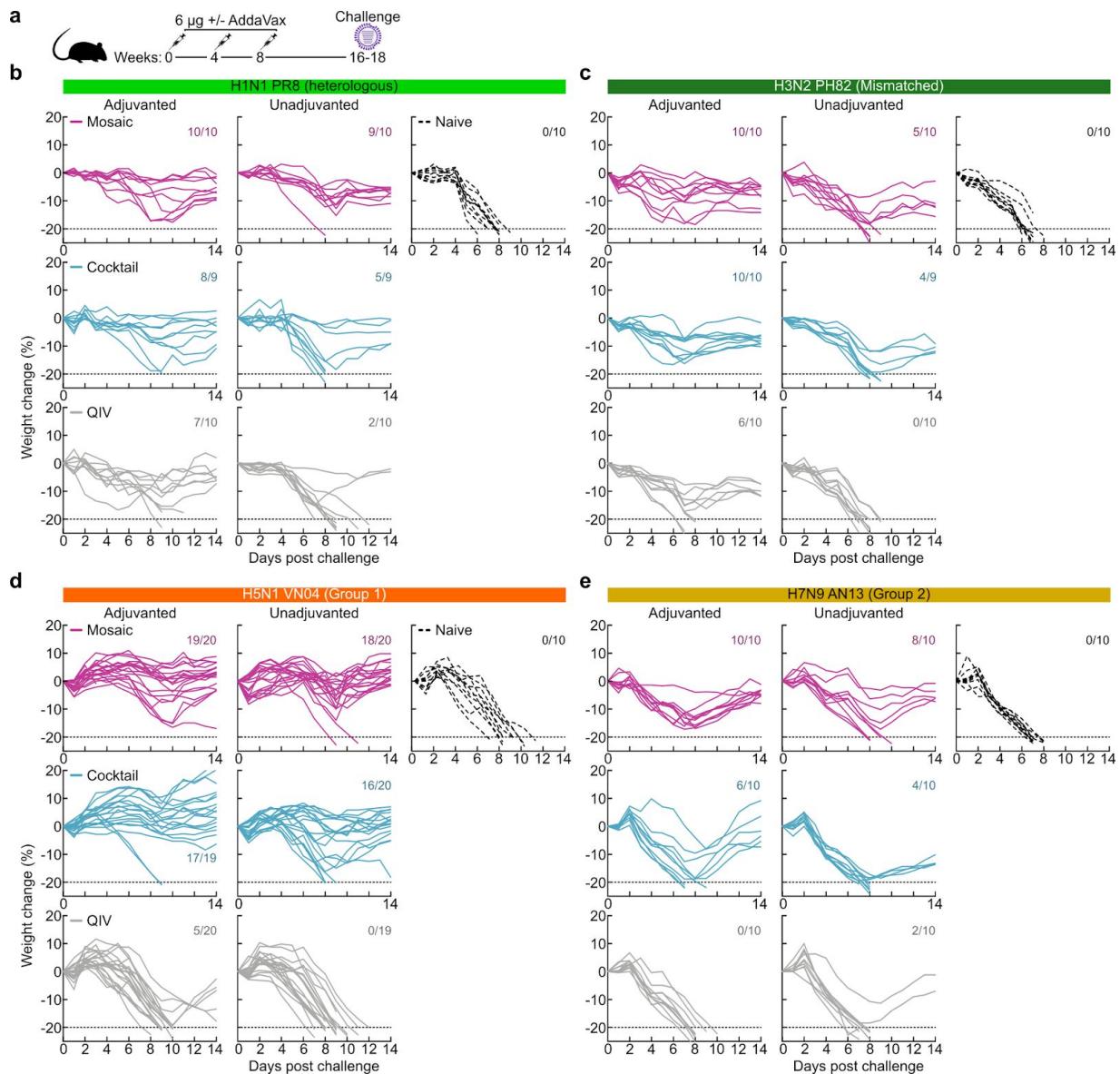
Supplementary fig. 1	2
Supplementary fig. 2	3
Supplementary fig. 3	4
Supplementary fig. 4	5
Supplementary fig. 5	6
Supplementary table 1	7
Supplementary table 2	12
Supplementary table 3	13
Supplementary table 4	14



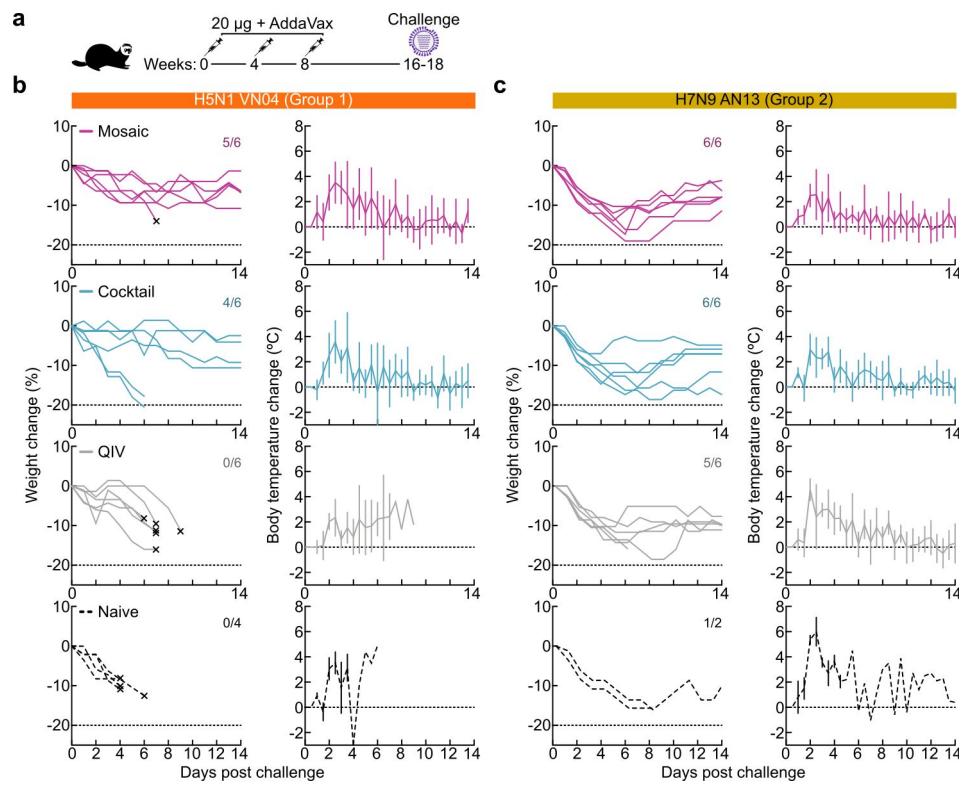
Supplementary Fig. 1 | Uncropped SDS-PAGE of H1-I53_dn5B fusions, purified nanoparticles and immunoprecipitations. **a**, Uncropped SDS-PAGE gel from [Extended Data Fig. 1b](#). **b**, Uncropped SDS-PAGE gel from [Extended Data Fig. 1d](#). **c**, Uncropped SDS-PAGE gel of qsCocktail-I53_dn5 and qsMosaic-I53_dn5 immunoprecipitations from [Fig. 1d](#). IP, immunoprecipitated; UB, unbound.



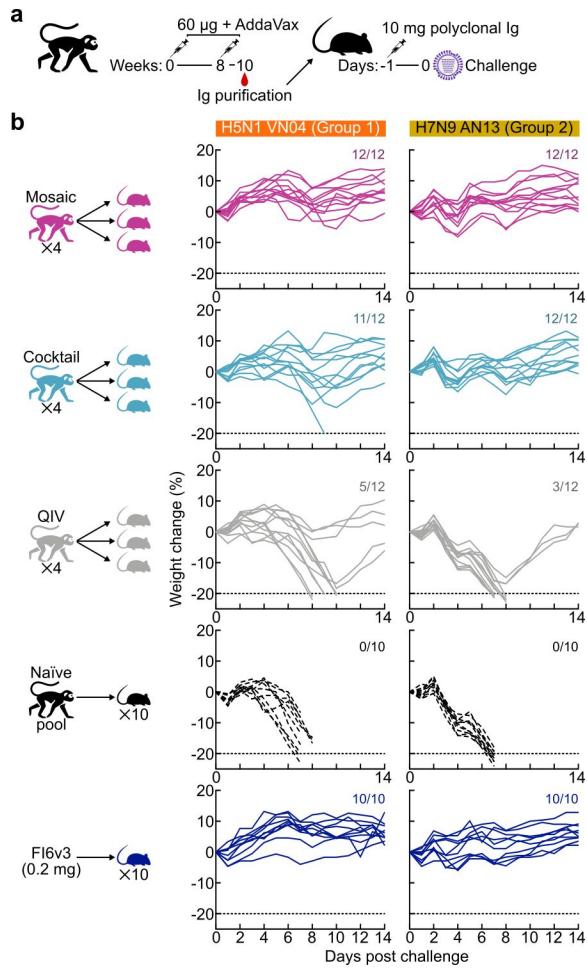
Supplementary Fig. 2 | ELISA binding curves of serum samples obtained from unimmunized, PBS- or irrelevant antigen-immunized animals. Mice and ferrets were immunized intramuscularly with PBS, DS-Cav1-I53_dn5, or H1-I53_dn5 plus AddaVax. NHP samples taken prior to immunization with our vaccines. Groups of BALB/cJ mice ($N = 10$), Finch ferrets ($N = 6$), and rhesus macaques ($N = 12$) were used. ELISA binding curves for HAs of H1 MI15, H3 HK14, B/Yam PH13 and B/Vic BR08 are shown. All animal experiments except for NHP were performed at least twice and representative data are shown.



Supplementary Fig. 3 | Body weight changes for mouse challenge studies. **a**, Experimental scheme of mouse challenge studies. Changes in body weight after heterologous H1N1 (**b**), mismatched H3N2 (**c**), heterosubtypic H5N1 (**d**) or H7N9 (**e**) virus infection in immunized BALB/cJ mice ($N=10$; $N=9$ for qsCocktail-I53_dn5 groups in **b** and unadjuvanted qsCocktail-I53_dn5 group in **c**; $N=19$ for adjuvanted qsCocktail-I53_dn5 and unadjuvanted QIV group in **d**; and $N=20$ for all other groups in panel **d**). Viruses were given intranasally on day 0 and mice were monitored for 14 days post infection. Body weight curves are plotted for each individual animal. Animals suffered $>20\%$ of body weight loss from day 0 were humanely euthanized.



Supplementary Fig. 4 | Body weight and temperature changes for ferret challenge studies. **a**, Experimental scheme of ferret challenge studies. Body weight (left) and temperature (right) changes after heterosubtypic H5N1 (**b**) or H7N9 (**c**) virus infection in immunized ferrets. Viruses were given intranasally on day 0 and animals were monitored for 14 days post infection. Changes in body weight and temperature are plotted for each individual animal. Predetermined humane endpoints include >20% of body weight loss from day 0 or other symptoms such as neurological abnormalities and moribound state. X symbols indicate animals euthanized based on the symptom-based humane endpoints. These animals exhibited one or more of the following symptoms: neurological abnormalities, bilateral hind leg paresis or paralysis, one-sided head tilt and side rolling, and moribound state. One animal was found dead.



Supplementary Fig. 5 | Body weight changes for mouse challenge studies after passive transfer of hyper immunoglobulins. **a**, Experimental scheme of the study. Immunoglobulins were purified from immunized NHPs and given to recipient BALB/cAnNHsd mice ($N = 3$ per NHP; $N = 12$ per immunization group). Control groups of mice ($N = 10$) were given either immunoglobulins obtained prior to immunization (Naïve pool) or anti-HA stem monoclonal antibody Fl6v3. **b**, Changes in body weight after heterosubtypic H5N1 (left) or H7N9 (right) virus infection in recipient mice. Immunoglobulins were given intraperitoneally at day -1. Viruses were given intranasally on day 0 and mice were monitored for 14 days post infection. Body weight curves are plotted for each individual animal. Animals suffered $>20\%$ of body weight loss from day 0 were humanely euthanized.

Supplementary Table 1 | Amino acid sequences for novel proteins used in this study.

Non-antigen-bearing nanoparticle components

>I53_dn5A

MGKYDGSKLRIGILHARWNAEILALVLGALKRLQEFGVKRENIIETVPGSFELPYGSKLFVEKQKRLGKPLDAIIPIG
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HHHHHH

>I53_dn5B

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HA-I53_dn5B trimers

>H1-I53_dn5B: A/Michigan/45/2015 HA 1-676 Y98F no lkr dn5B.SA.WELQ-H

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HA-foldon trimers (ELISA antigens and EMPEM)

>H1 MI15: A/Michigan/45/2015 HA 1-676 Y98F FAH

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>H3 HK14: A/Hong Kong/4801/2014 HA 1-676 Y98F FAH

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>B/Yam PH13: B/Phuket/3073/2013 HA 1-676 FAH

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>B/Vic BR08: B/Brisbane/60/2008 HA 1-676 FAH

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>H5 IN05: A/Indonesia/05/2005 HA 1-676 Y98F delF FAH

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GNPMCDEFINVPEWSYIVEKANPTNDLCFPGSFNDYEELKHLLSRINHFEKIQIIPKSSWSDHEASSGVSSACPYLG
SPSFRNVVWLIKKNSTYPTIKSYNNNTQEDLLVLWGIHPNDAAEQTRLYQNPTTYISIGTSLNQRLVPKIATRS
KVNGQSGRMEFFWTILKPNDAINFESNGNFIAPERAYKIVKKGDSAIMKSELEYGNCNTKCQTPMGAINSSMPFHNI
HPLTIGECPKYVKSNRVLATGLRNSPQRERGLFGAIAGFIEGGWQGMVDGWYGYHHSNEQGSGYAADKESTQ
KAIDGVTNKVNSIIDKMNTQFEAVGREFNNLERRIENLNKKMEDGFLDVWTYNAELLVLMENERTLDFHDSNVKNL
YDKVRLQLRDNAKELGNGCFYHKCDNECMESIRNGTYNYPQYSEEARLKREEISVGGSGYIPEAPRDGQAYVR
KDGEWVLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H6 TW13: A/Taiwan/2/2013 HA 1-676 Y98F FAH

MIAIVVIALASAGKSDKICIGYHANNSTTVQDVTLEKNVTVHSVELLENQKEKRFCKIMNKAPLDLKDCIEGWILGN
PKCDLLLGDQSWSYIVERPNAQNGICFPGVNEEELKAFIGSGERVERFEMFPKSTWAGVDTSRGVTNACPSYI
DSSFYRNLWIVKTDSATYPVIKGTYNNTGTQPILEFWGVHHPLDTTVQDNLYGSGDKYVRMGTESMNFAKSPEIA
ARPAVNGQRSRIDYYWSVLRPGETLNVESNGNLIAPWYAYKFVSTNKKAVFKSDLPIENCATCQITGVLRTNK
TFQNVSPWLIGECPKYVKSesrlLATGLRNVPQIATRGIFGAIAGFIEGGWTGMIDGWYGYHHENSQGSGYAADR
ESTQKAIDGITNKVNSIINKMNTQFEAVDHEFSNLERRIGNLNKRMEDGFLDVWTYNAELLVLENERTLDLHDANV
KNLYEKVKSQRLDNANDLNGNCFEFWHKCDNECMESVKNGTYDYPKYQKESKLNQRQIESVGGSGYIPEAPRDGQ
AYVRKDGEWVLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H7 AN13: A/Anhui/1/2013 HA 1-676 Y98F FAH

MNTQILVFALIAIPTNADKICLGHHAVSNGTKVNTLTERGVEVVNATEVERTNIPRICKSGKRTVDLGQCGLLTIT
GPPQCDQFLEFSADLIERRREGSDVCPGKFVNEEALRQILRESGGIDKEAMGFTYSGIRTNGATSACRRSGSSFY
AEMKWLLSNTDNAAFPQMTKSYKNTRKSPALIVWGIHHSVSTAEQTKLYGSGNKLVTVGSSNYQQSFVPSPGARP
QVNGLSGRIDFHMLNPNNDTFTSFNGAFIAPDRASFLRGKSMGIQSGVQVDANCEGDCYHSGGTIISNLPFQNI
DSRAVGKCPRYVKQRSLLLATGMKVNPEIPKGRGLFGAIAGFIENGWEGLIDGWYGRHQNAQGEGTAADYKSTQ
SAIDQITGKLNRRIEKTNQQFELIDNEFNEVEKQIGNVINWTRDSITEWWSYNAELLVAMENQHTIDLADSEMDKLYE
RVKRQLRENAEEDGTGCFEIFHKCDDDCMASIRNNTYDHSKYREEAMQNRIQIDPVGGSGYIPEAPRDGQAYVR
KDGEWVLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H10 JD13: A/Jiangxi Donghu/346/2013 HA 1-676 Y98F FAH

MYKIVVIIALLGAVKGLDKICLGHHAVANGTIVKTLTNEQEEVTNATEVESTGINRLCMKGRKHKDLGNCHPIGMLIG
TPACDLHLTGMWDTLIERENAIAYCPGATVNVEALRQKIMESGGINKISTGFTYGSINSAGTTRACMRNGGNSFY
AELKWLVSKSQGQNFQPTTNTYRNTDAEHLIMWGIHPSSTQEKNDLYGTQSLISVGSSTYRNNFVPVVGARP
QVNGLSGRIDFHWTLVQPGDNITFSHNGGLIAPSRSVSKLIGRGLGIQSDAPINNCESKCFWRGGSINTRLPFQNL
SPRTVGQCPKYVNRRSLMLATGMRNPVPELIQGRGLFGAIAGFLENGWEGMVDGWYGRHQNAQGTTGQAADYKS
TQAAIDQITGKLNRRIEKTNTEFESIESEFSEIEHQIGNVINWTKDSITDIWTYQAELLVAMENQHTIDLADSEMLNY
ERVRKQLRQNAEEDGKGCFEIYHACDDSCMESIRNNTYDHSQLYREEALLNRLNINPVGGSGYIPEAPRDGQAYVR
KDGEWVLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H1 stem: A/California/04/2009 stabilized stem FAH

MKAILVVLLYTFATANADTLCIGYHANNSTDVTDTVLEKNVTVHSVNLGSGRLATGLRNIPQRETRGLFGAIAGFI
EGGWTGMVDGWYGYHHQNEQGSGYAADLKSTQNAIDEITNMVNSVIEKMGSGGGTDLAELLVLLNQWTLLYH

DSNVKNLYEKVRSQLKNNAKEIGNGCFEFYHKCDNTCMESVKNGTYDYPKYSEEAKLNREEIDGSGYIPEAPRDG
QAYVRKDGEWLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H3 stem: A/Finland/486/2004 stabilized stem FAH

MKTIIALSYILCLVFAQKLPGNNDNSTATCLGHHAVPNGTIVKTITNDQIEVTNATELVFPGCGVLKLATGMRNVPEK
QTRGIFGAIAGFIENGWEGMVDGWYGFRHQNSEGIGQAADLKSTQAQAINQINGMVNRVIALMAQGGPDCYLAELL
VALLNQHVIDLTDSEMRKLFERTKKQLRENAEDMGNGCFKIYHKCDNACIGSIRNGTYDHDVYRDEALNNRFQIKG
GPGSGYIPEAPRDGQAYVRKDGEWLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>H3 SG16: A/Singapore/INFIMH-16-0019/2016 HA 1-676 Y98F FAH

MKTIIALSYILCLVFAQKIPGNDNSTATCLGHHAVPNGTIVKTITNDRIEVTNATELVQNSSIGEICDSPHQILDGENC
TLIDALLGDPQCDGFQNKKWDLFVERSKAYSNCFPYDVPDYASRLSVASSGTLEFKNESFNWTGVTQNGTSSAC
IRGSSSSFFSRLNLTHLNYYTYPALNVTPNKEQFDKLYIWGVHHPGTDKDQIFLYAQSSGRITVSTKRSQQAVIPN
IGSRPRIRDIPSRSISIYWTIVKPGDILLINSTGNLIAPRGYFKIRSGKSSIMRSDAPIGKCKSECITPNGSIPNDKPFQNV
NRITYGACPRYVKHSTLKLATGMRNVPEKQTRGIFGAIAGFIENGWEGMVDGWYGFRHQNSEGRGQAADLKSTQ
AAIDQINGKLNRLIGKTNEKFHQIEKEFSEVEGRVQDLEKYVEDTKIDLWSYNAELLVALENQHTIDLTDSEMNLFE
KTKKQLRENAEDMGNGCFKIYHKCDNACIESIRNETYDHNVYRDEALNNRFQIKGVGSGYIPEAPRDGQAYVRKD
GEWLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

>B/Vic CO17: B/Colorado/06/2017 HA 1-676 FAH

MKAIIVLLMVVTSSADRICTGITSSNSPHVVKTATQGEVNVTGVIPLTTPTKSHFANLKGTTETRGKLCPKCLNCTDL
DVALGRPCKTGKIPSARVSILHEVRPVTSFCGCFIMHDRTKIRQLPNLLRGYEHVRLSTHNVINAEGAPGGPYKIGTS
GSCPNITNGNGFFATMAWAVPDKNKTATNPLTIEVPYVCTEGEDQITVWGFHSDNETQMAKLYGDSKPQKFTSSA
NGVTTHYVSQIGGFPNQTEDGGLPQSGRIVDYMVQKSGKTGTTYQRGILLPKVWCASGRSKVIK GSLPLIGEA
DCLHEKYGGLNKS PKPYYTGEHAKAIGNCPIWKPLKLANGTKYRPPAKLLKERGFFGAIAGFLEGGWEGMIAGW
HGYTSHGAHVAVAADLKSTQEAINKITKNLNSLSELEVKNLQRLSGAMDELHNEILELDEKVDDLRA DTISSQIELA
VLLSNEGIINSEDEHLLALERKLKKMLGPSAVEIGNGCFETKHKCNQTCLDKIAAGTFDAGEFSLPTFDNLITAASL
GSGYIPEAPRDGQAYVRKDGEWLLSTFLGSGLNDIFEAQKIEWHEGHHHHHH

Appended sequences including the foldon trimerization domain, WELQuT protease recognition sites, Avi tags, and hexa-histidine tags are underlined.

Supplementary Table 2 | Cryo-EM data collection, refinement and validation statistics.

	H1-I53_dn5 nanoparticle (EMD-22935)	H1 HA/MI15 Localized reconstruction (EMD-22937; PDB 7KNA)	H5 bound to 3 polyclonal Fabs (EMD-22940)	H5 bound to 2 polyclonal Fabs (EMD-22939)	H5 bound to 1 Fab (EMD-22938)
Data collection and processing					
Magnification					
Magnification	130,000	130,000	130,000	130,000	130,000
Voltage (kV)	300	300	300	300	300
Electron exposure (e ⁻ /Å ²)	70	70	70	70	70
Defocus range (μm)	0.5-3.0	0.5-3.0	0.5-3.0	0.5-3.0	0.5-3.0
Pixel size (Å)	0.525	0.525	0.525	0.525	0.525
Symmetry imposed	I	C3	C1	C1	C1
Initial particle images (no.)	22,563	397,700 (sym-exp)	115,722	115,722	115,722
Final particle images (no.)	19,885	148,916	36,783	19,849	22,211
Map resolution (Å)	6.6	3.3	3.6	4.1	4.0
FSC threshold	0.143	0.143	0.143	0.143	0.143
Refinement					
Model resolution (Å)		3.4			
FSC threshold		0.5			
Map sharpening <i>B</i> factor (Å ²)		-174			
Model composition					
Nonhydrogen atoms		11,593			
Protein residues		1,449			
Ligands		24 (NAG)			
<i>B</i> factors (Å ²)					
Protein		37.5			
Ligand		44.2			
R.m.s. deviations					
Bond lengths (Å)		0.015			
Bond angles (°)		1.4			
Validation					
MolProbity score		0.66			
Clashscore		0.4			
Poor rotamers (%)		0.5			
Ramachandran plot					
Favored (%)		97.91			
Allowed (%)		1.88			
Disallowed (%)		0.21			

Supplementary Table 3 | Peptides used in label-free quantitation of HA content in mosaic nanoparticles and data for post-SEC quantitation of HA content in qsMosaic-I53_dn5 (related to [Extended Data Fig. 3](#)).

HA con- struct	Unique peptide sequence ^a	Standard equimolar mix (area)	qsMosaic-I53_dn5 post-SEC (area)	Relative ratio (qsMosaic-I53_dn5 /standard)	Average relative ratio and standard deviation	qsMosaic-I53_dn5 post-SEC composition ^b
H1	NLIWLVK	1.07×10^7	5.35×10^6	0.502		
	GVTAACPHAGAK	1.44×10^6	9.59×10^5	0.668	0.558 +/- 0.028	21.2 +/- 2.8 %
	KFKPEIATRPK	1.18×10^6	6.22×10^5	0.529		
	FKPEIATRPK	2.46×10^6	1.32×10^6	0.536		
H3	KWDLFVERSK +WDLFVERSK ^c	5.88×10^6	4.87×10^6	0.827		
	SSIMRSDAPIGK ^d	1.05×10^7	7.45×10^6	0.707	0.720 +/- 0.030	27.3 +/- 3.0 %
	LATGMNRNVPEK ^d	6.24×10^6	3.99×10^6	0.638		
	EFSEVEGRIQDLEK ^e	8.62×10^6	6.11×10^6	0.709		
B/Yam	IRLSTQNVIDAEK	5.44×10^6	3.77×10^6	0.693		
	SLYGDSNPQK	5.01×10^6	3.11×10^6	0.621	0.632 +/- 0.012	24.0 +/- 1.6 %
	SYFANLK	6.73×10^6	4.00×10^6	0.595		
	IRQLPNLLRGYEK	1.30×10^7	8.07×10^6	0.620		
B/Vic	GSLPLIGEADCLHEK	7.38×10^6	5.19×10^6	0.703		
	LYGDSKPQK	4.92×10^5	3.60×10^5	0.731	0.727 +/- 0.013	27.6 +/- 1.3 %
	TGTITYQRGILLPQK	6.72×10^6	4.70×10^6	0.700		
	VWCASGRSK	1.99×10^6	1.54×10^6	0.773		

^aOnly abundant peptides that were unique to each specific HA were included.

^bNormalized to the total sum of the average relative ratios.

^cA high degree of missed cleavage was observed for this peptide and the sum of the two peptides was used for the calculations.

^dThe oxidized methionine forms were also included in these calculations.

^eThe pyro-glutamic acid form of the peptide was also included in these calculations.

Supplementary Table 4 | Viruses used in HAI, microneutralization, and challenge experiments.

Subtype	Virus	Short name	Type of experiment	Figure
H1N1	A/Michigan/45/2015	MI15; 2015	HAI	2a; ED 4e; ED 5c
			Microneutralization	2b; 3a; ED 4e; ED 5d; 5b; ED 7f; ED 10
	A/Boston/YGA-01050/2012	2012	Microneutralization	3a; ED 6
	A/California/07/2009	2009	Microneutralization	3a; ED 6
	A/Weiss/1943	1943	Microneutralization	3a; ED 6
	A/Malaysia/1954	1954	Microneutralization	3a; ED 6
	A/USSR/90/1977	1977	Microneutralization	3a; ED 6
	A/Memphis/4/1987	1987	Microneutralization	3a; ED 6
	A/New York/146/2000	2000	Microneutralization	3a; ED 6
	A/New Caledonia/20/1999	1999	Microneutralization	3a; ED 6
H3N2	A/Solomon Islands/03/2006	2006	Microneutralization	3a; ED 6
	A/Puerto Rico/8/1934	PR8	Challenge (mice)	3b; S3b
			Pseudovirus neutralization	3b
	A/Hong Kong/4801/2014	HK14; 2014	HAI	2a
			Microneutralization	2b; 3a; ED 4e; ED 7f
	A/Switzerland/9715293/2013	2013	Microneutralization	3a; ED 6
	A/Texas/50/2012	2012	Microneutralization	3a; ED 6
	A/Victoria/361/2011	2011	Microneutralization	3a; ED 6
	A/Brisbane/10/2007	2007	Microneutralization	3a; ED 6
	A/Wisconsin/67/2005	2005	Microneutralization	3a; ED 6
H5N1	A/Fujian/411/2002	2002	Microneutralization	3a; ED 6
	A/Moscow/10/1999	1999	Microneutralization	3a; ED 6
	A/Shangdong/9/1993	1993	Microneutralization	3a; ED 6
	A/Philippines/2/1982	PH82	Challenge (mice)	3c; S3c
			Pseudovirus neutralization	3c
	A/Singapore/INFIMH-16-0019/2016	SG16	Microneutralization	ED 5d; ED 10
	A/Vietnam/1203/2004	VN04	Challenge (mice, ferrets)	4d–f; S3d; S4b; S5b
			Microneutralization	5b
H7N9	A/Anhui/1/2013	AN13	Challenge (mice, ferrets)	4d–f; S3e; S4c
	B/Phuket/3073/2013	PH13	HAI	2a; ED 4e; ED 5c
B/Vic			Microneutralization	2b; ED 4e; ED 5d; ED 7f; ED 10
	B/Colorado/06/2017	CO17	HAI	2a; ED 4e; ED 5c
			Microneutralization	2b; ED 5d; ED 7f; ED 10