

1                                   **An egg-derived sulfated N-Acetylactosamine glycan is an**  
2                                   **antigenic decoy of influenza virus vaccines**

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18 **Abstract**

19 Influenza viruses grown in eggs for the purposes of vaccine generation often acquire mutations during  
20 egg adaptation or possess differential glycosylation patterns than viruses circulating amongst humans.  
21 Here, we report that seasonal influenza virus vaccines possess an egg-derived sulfated N-  
22 acetylglucosamine (LacNAc) that is an antigenic decoy. Half of subjects that received an egg-grown  
23 vaccine mounted an antibody response against this egg-derived antigen. Egg-binding monoclonal  
24 antibodies specifically bind viruses grown in eggs, but not viruses grown in other chicken derived cells,  
25 suggesting only egg-grown vaccines can induce anti-LacNAc antibodies. Notably, antibodies against the  
26 sulfated LacNAc utilized a restricted antibody repertoire and possessed features of natural antibodies,  
27 as most antibodies were IgM and have simple heavy chain complementarity determining region 3. By  
28 analyzing a public dataset of influenza virus vaccine induced plasmablasts, we discovered egg-binding  
29 public clonotypes that were shared across studies. Together, this study shows that egg-grown vaccines  
30 can induce antibodies against an egg-associated glycan, which may divert the host immune response  
31 away from protective epitopes.

## 32 **Introduction**

33           Influenza viruses have been historically grown in embryonated chicken eggs as a way to culture  
34 large quantities of virus, and as a result, most influenza virus vaccines are still generated using viruses  
35 grown in eggs. However, this process has the potential to change the immunogenicity of the virus, as the  
36 viruses may mutate their major surface glycoproteins hemagglutinin (HA) and neuraminidase (NA) to  
37 increase infectivity in eggs (1-5). Moreover, influenza viruses grown in eggs are often less immunogenic  
38 than viruses grown in mammalian cells (6-8) and have been shown to be less effective than mammalian  
39 cell-based influenza vaccines and recombinantly expressed HA vaccines (9, 10). Due to the inherent  
40 difference in avian versus mammalian glycosylation patterns, egg-grown vaccines may lack certain  
41 glycans that would be expressed on influenza viruses transmitted among humans. Notably, vaccine  
42 effectiveness against recent H3N2 viruses may be reduced due to the lack of a glycan on HA of H3N2  
43 viruses grown in eggs (1).

44           However, slight differences in viral sequences and glycosylation patterns of HA do not fully explain  
45 why vaccine effectiveness is low, as serum from vaccinated subjects can have similar antibody titers  
46 against egg adapted strains and viruses circulating in the population (11). Poor immunogenicity against  
47 HA may explain reductions in vaccine effectiveness rather than egg-adapted mutations (11). It is possible  
48 that egg-grown vaccines are preferentially inducing antibodies against non-protective viral antigens,  
49 therefore reducing vaccine effectiveness and seroconversion against protective epitopes on the HA head  
50 domain. Similar to subjects receiving egg-grown vaccines (12, 13), HA-reactive antibodies induced by  
51 vaccines grown in mammalian cells and insect cells largely induced antibodies mostly targeting the head  
52 domain of HA (14). However, whether different vaccine platforms induced antibodies against distinct  
53 influenza virus antigens, other than HA, is not known. As internal antigens, such as the nucleoprotein  
54 (NP), were shown to provide limited protection against infection (12), it remains to be determined how  
55 different vaccine formulations drive antibodies against distinct protective and non-protective antigens.

56           To address whether these egg-grown influenza virus vaccines induced antibodies against  
57 potentially non-protective antigens, we cloned monoclonal antibodies (mAbs) from plasmablasts (PBs),  
58 a transient antibody secreting cell population, isolated from subjects following vaccination with egg-grown

59 influenza virus vaccines. We show that 50% of subjects generated a PB response against an egg-derived  
60 antigen present in the vaccine. Subjects that mounted a response against the egg-associated antigen  
61 seroconverted against HA to similar levels as subjects that did not mount an anti-egg response, indicating  
62 egg-grown vaccines did not reduce overall secreted antibody responses against HA. We identified that  
63 the egg-derived antigen was a sulfated N-acetylglucosamine (LacNAc) glycan and was only present in  
64 viruses grown in the allantois of eggs, but not in viruses grown in chicken embryo cell line or primary  
65 chicken fibroblasts. Antibodies binding the egg-derived glycan utilized a restricted repertoire and  
66 resembled natural antibodies, as antibodies were largely IgM and had short heavy chain complementarity  
67 determining region 3 (H-CDR3). Moreover, we identified that egg-binding antibodies identified in our  
68 study were public clonotypes, indicating the same antibodies were found across individual subjects that  
69 had been vaccinated with an egg-grown vaccine. Together, our study shows that egg-grown vaccines  
70 can induce antibodies against an egg related glycan and that these glycan-binding mAbs resemble those  
71 produced by innate-like B cells.

## 72 **Results**

### 73 **Influenza virus vaccination induces antibodies against an egg-associated antigen**

74 To address the antigen specificity of memory B cells recalled by egg-grown influenza virus vaccines, we  
75 generated mAbs from sorted PBs 7 days following vaccination. The transient PB population are highly  
76 specific to the components of the vaccine and are recalled from pre-existing memory B cells (15-17). We  
77 focused our studies on mAbs generated from subjects following vaccination with the 2009 monovalent  
78 pandemic H1N1 inactivated influenza virus vaccine (MIV) and the 2010 trivalent inactivated influenza  
79 virus vaccine (TIV). From the vaccine induced PBs, we found that 75% of mAbs bound HA (Figure S1A  
80 and B). Notably, 27 mAbs generated from multiple subjects receiving either of the egg-grown vaccines  
81 bound all influenza virus strains tested (Figure 1A; Table S1). To confirm these mAbs were specific to  
82 influenza viruses and not an artifact of vaccine preparation in eggs, we tested mAb binding to  
83 A/California/7/2009 H1N1 virus grown in eggs or in mammalian Madin-Darby Canine Kidney (MCDK)  
84 cells. Strikingly, these mAbs only bound to A/California/7/2009 grown in eggs, but not virus propagated  
85 in mammalian cells (Figure 1B and C), indicating these broadly reactive mAbs were binding to an egg-  
86 related antigen. Moreover, these broadly reactive mAbs bound to allantoic fluid from both uninfected eggs  
87 and A/California/7/2009 H1N1 infected eggs (Figure 1D-E), indicating these mAbs were specific to an  
88 egg associated antigen.

89 50% of all subjects analyzed generated egg-specific antibody responses (Figure S1C), with 1 out  
90 of 5 of the subjects receiving the MIV and 6 out of 9 of subjects receiving the TIV generating an egg-  
91 specific antibody response (Table S1). Within the subjects that mounted an antibody response against  
92 this egg-derived antigen, 31% of mAbs generated specifically bound the egg-derived antigen (Figure 1F).  
93 The range of antibodies per subject ranged from 8% to 58% of isolated vaccine induced PBs (Figure 1G).  
94 Additionally, we found that subjects that had egg-reactive mAbs had a larger fold increase in serum IgG  
95 responses against uninfected allantoic fluid relative to subjects who did not have detectable mAbs against  
96 the egg-associated antigen (Figure 1H). Despite this, subjects that mounted an antibody response  
97 against the egg-associated antigen had a similar fold increase in serum IgG titers against  
98 A/California/7/2009 recombinant HA and hemagglutination inhibition (HAI) titers against

99 A/California/7/2009 H1N1, relative to subjects that did not generate a PB response against the egg  
100 antigen (Figure S1D-F). Together, these data indicate that some subjects following influenza virus  
101 vaccination generate an antibody response against an egg-derived antigen.

## 102 **Viruses grown in allantoic fluid, but not other parts of the egg, possess the egg antigen**

103 Starting in 2018, the United States Center for Disease Control began recommending that people  
104 with egg allergies could receive egg-grown influenza virus vaccines, suggesting the major egg allergens  
105 were removed from the vaccine (<https://www.cdc.gov/flu/prevent/egg-allergies.htm>). Although subjects  
106 within our cohorts had not experienced an allergic response to influenza virus vaccination or reported a  
107 history of egg allergies, we next tested whether the identified egg-specific mAbs could bind to more recent  
108 inactivated influenza virus vaccines grown in eggs that lack the egg allergens. Notably, the egg-specific  
109 mAbs could bind old TIVs and quadrivalent inactivated influenza virus vaccines (QIVs) to a similar degree  
110 as recent egg-grown QIVs, including the 2020 Fluarix QIV (Figure 2A), indicating the egg-specific antigen  
111 identified still persists in recent egg-grown vaccines. Additionally, some QIVs are also made from viruses  
112 grown in mammalian MDCK cells (Flucelvax) and recombinant HA generated in insect cells (Flublok).  
113 The egg-binding mAbs specifically bound to QIV viruses grown in eggs, but not viruses grown in MDCK  
114 or recombinant HA grown in insect cells (Figure 2B). We also confirmed that these egg-binding mAbs  
115 could bind other viruses grown in the allantoic fluid of eggs, as these mAbs bound as strongly to  
116 Newcastle Disease Virus grown in eggs as it did to A/California/7/2009 H1N1 grown in eggs (Figure 2C).  
117 To understand how ubiquitous the egg antigen was across chicken cell- and egg-grown vaccines, we  
118 next tested for mAb binding to other vaccines grown in chicken cells. Notably, the mumps and measles  
119 viruses of the Measles/Mumps/Rubella vaccine (MMR) are both grown in a chicken embryo cell line and  
120 the rabies virus in Rabavert is grown in primary chicken fibroblasts. As a control, we also tested the egg  
121 binding mAbs for binding to non-egg or chicken grown vaccines including the Japanese Encephalitis  
122 Virus Vaccine (Ixiaro) grown in Vero cells and the Pneumovax 23 vaccine that contains purified capsular  
123 polysaccharides from 23 distinct *Streptococcus pneumoniae* serotypes. The egg-binding mAbs only  
124 bound to vaccines grown in eggs, but not to vaccines grown in a chicken embryo cell line (MMR), primary  
125 chicken fibroblasts (Rabavert), or vaccines not produced in eggs (Ixiaro and Pneumovax 23; Figure 2D).

126 Together, these data reveal that the egg-associated antigen is only present in viruses grown in allantoic  
127 membrane, but not in those grown in cells isolated from chicken embryos or chicken fibroblasts.

### 128 **Sulfated LacNAc is the egg-derived antigen**

129 Egg allergies are typically caused by antibody responses against ovalbumin and ovomucoid (18),  
130 which are present in both the egg white and allantoic fluid (19, 20). However, the egg-specific mAbs did  
131 not bind to ovalbumin or ovomucoid purified from egg whites (data not shown), further indicating these  
132 mAbs were likely not specific to known egg allergens. As the antigen did not seem to be protein in nature,  
133 we next tested whether egg-binding mAbs were binding an egg-associated glycan. To test this, we  
134 deglycosylated the 2020 Fluarix QIV with a deglycosylating enzyme that removes N-linked glycans. MAbs  
135 had reduced binding to deglycosylated vaccine relative to untreated vaccine (Figure 3A). To investigate  
136 the particular glycan these mAbs were binding, we tested two mAbs (029-09 3A04 and 034-10 4G02) for  
137 binding to a glycan microarray that included 585 distinct glycans (Table S2). Both mAbs specifically bound  
138 to two sulfated LacNAc antigens, (6S)(4S)Gla $\beta$ 1-4GlcNAc $\beta$  and (4S)Gla $\beta$ 1-4GlcNAc $\beta$  (Figure 3  
139 B and C). Treatment of purified egg-grown A/Hong Kong/485197/2014 H3N2 with a sulfate ester  
140 sulfatase significantly reduced egg-specific mAb binding (Figure 3D). Moreover, both mAbs only bound  
141 to LacNAc with a sulfate group on the hydroxyl group of 4C' of galactose, and not the hydroxyl group on  
142 6C' of the galactose of LacNAc only, or an unsulfated LacNAc (Figure 3E and F; Table S3). Together,  
143 these data reveal that B cells induced by influenza viruses grown in eggs are binding a sulfated LacNAc.

### 144 **Egg binding antibodies utilize a restricted repertoire and resemble natural antibodies**

145 Of the egg-binding mAbs, we identified strong repertoire biases on the usage of particular heavy  
146 and light chain variable genes, with the vast majority of mAbs using VH3-07 and VL1-44 (Figure 4A and  
147 B). However, there was a lot of diversity in the H-CDR3 sequences, with no consensus on DH gene  
148 usage (Figure S2A). H-CDR3s and light chain CDR3s (L-CDR3) of egg-binding mAbs preferentially used  
149 JH4 and JL3, suggesting there was some selection for certain J genes (Figure S2B and C). Moreover,  
150 H-CDR3s of egg-binding mAbs were significantly shorter than those of vaccine-specific mAbs, whereas  
151 the L-CDR3 were significantly longer than vaccine-specific mAbs (Figure 4C). Concordantly, H-CDR3s

152 of egg-binding mAbs had fewer non-templated DNA insertions, or N-nucleotides, relative to vaccine-  
153 specific mAbs (Figure 4D; Figure S2D).

154 Analysis of clonal expansions revealed that the light chain of egg-binding mAbs across individual  
155 mAbs were highly clonal, with one light chain clone occupying over one-third of all egg-binding mAbs  
156 (Figure S2E; Table S2 and S5). Additionally, we identified 4 distinct clonal expansions, accounting for  
157 one-third of all antibodies identified (Figure S2E). However, these paired heavy and light chain clones  
158 were private, with individual clones only identified in one subject each (Table S2). PBs induced by  
159 vaccination derive from memory B cells and therefore are usually class-switched to IgG and highly  
160 mutated (17). However, egg-binding mAbs were largely IgM and had fewer mutations in the heavy chain  
161 relative to vaccine-specific mAbs (Figure 4E and F). In combination, egg-binding mAbs resemble natural  
162 antibodies produced by innate-B cells, as they express simple and short H-CDR3s, do not commonly  
163 class-switch, and have fewer mutations than vaccine-specific antibodies (21, 22). In addition, natural  
164 antibodies commonly target glycans (23) and are polyreactive (24, 25). However, egg-binding mAbs were  
165 not enriched for polyreactivity relative to vaccine-induced antibodies (Figure S2F). Despite having fewer  
166 mutations than vaccine-induced antibodies, germline (GL) reverted egg-binding mAbs had reduced  
167 binding affinity for influenza viruses grown in eggs relative to the affinity-matured (AM) mAbs generated  
168 from PBs (Figure 4G). Furthermore, we identified a clonal expansion from one subject over two influenza  
169 virus vaccine seasons (2010 TIV and 2011 TIV), with the mAbs from 2011 having higher affinity for  
170 influenza virus strains relative to the mAb from 2010 (Figure S2G-I). With the highly restricted VH/VL  
171 repertoire, short H-CDR3 sequences and reduced N-nucleotide additions, lack of class-switching, and  
172 fewer mutations, mAbs targeting sulfated LacNAc resemble natural antibodies produced by innate-like B  
173 cells (26).

#### 174 **Influenza virus vaccines commonly induce PBs with repertoire features of egg-binding mAbs**

175 We next addressed whether antibodies with repertoire features of egg-binding antibodies are commonly  
176 induced after influenza virus vaccination. To dissect this question, we utilized a dataset of 7,777 B cell  
177 receptor sequences from influenza virus vaccine induced PBs from subjects that received the egg-grown  
178 2016-2017 Fluzone QIV (27). From this dataset, we identified that 2% (175 total) of all IgG<sup>+</sup> PBs

179 expressed VH3-7 with a H-CDR3 length of equal to or less than 12 amino acids that was paired with VL1-  
180 44 or VL1-51 (potential egg-binding mAbs; Figure 5A). Notably, 13 out of 17 total subjects had PBs with  
181 these repertoire features, occupying 0.2-17.4% of the PB response per subject (Figure 5B). Notably, this  
182 dataset was specifically generated from IgG<sup>+</sup> PBs. As we identified most egg-binding mAbs were IgM  
183 (Figure 4E), the true number of potential egg-specific PBs induced within these subjects may be  
184 substantially higher. Of the 175 heavy and light chain pairings identified, we discovered 6 public  
185 clonotypes (Figure 5C; Table S5), which comprised 66.9% of the total response (117/175 paired  
186 sequences). Strikingly, 3 of the public clones were shared between our study and the dataset (Figure 5C;  
187 Table S5), suggesting the PBs induced in subjects in the Forgacs et al. study are specific to the egg  
188 glycan. Moreover, we identified that egg-binding mAbs from our study shared at least a heavy chain or  
189 light chain clone with the potential egg-binding PBs from the IgG<sup>+</sup> PB dataset (Figure 5D; Table S5).  
190 Together, these data suggest that PBs targeting an egg-associated glycan are commonly induced by  
191 influenza virus vaccines grown in eggs.

## 192 **Discussion**

193 In this report, we identified that egg-grown viruses possess an egg-associated glycan that is  
194 immunogenic in humans. Egg-specific mAbs from PBs had evidence of prior affinity-maturation and likely  
195 derived from memory B cells that were primed by earlier vaccination with egg-grown vaccines. Moreover,  
196 most subjects within the 2010 TIV cohort had previously been vaccinated with the 2009 MIV, suggesting  
197 prior exposure to the egg antigen from the 2009 MIV generated memory B cells against this antigen.  
198 Despite some subjects mounting a response against the egg antigen, the same subjects seroconverted  
199 against the H1N1 component of the vaccine to similar levels as subjects that did not mount a response  
200 against the egg-associated antigen. Moreover, 13 out of 17 subjects from Forgacs et al. had detectable  
201 PBs with repertoire features of egg-binding mAbs. Notably, this study specifically recruited subjects that  
202 had not been vaccinated in the prior 3 seasons (27). Prior research has indicated that pre-existing serum  
203 antibodies against the strains included in the vaccine inversely correlate with the induction of PBs by  
204 vaccination (28). Therefore, subjects that mounted an anti-egg response perhaps had more B cell  
205 activation relative to subjects that did not and as a result has similar anti-HA antibody responses.

206 Despite no differences in serum antibodies against the H1N1 component of the vaccine, B cells  
207 mounted against the egg antigen may compete within germinal centers with B cells targeting protective  
208 epitopes of HA and NA, which could perturb the generation of plasma cells and memory B cells against  
209 protective epitopes. In accordance, we identified the same egg-specific clone was recalled over multiple  
210 vaccine years, suggesting subjects can repeatedly recall memory B cells against the egg-derived antigen  
211 upon repeated vaccination and further indicating egg-specific antibodies are fixed in the memory B cell  
212 repertoire. Therefore, our study suggests that repeatedly vaccinated subjects can preferentially recall  
213 memory B cells targeting irrelevant antigens associated with egg-based vaccine production, which could  
214 come at the cost of affinity-maturation and generation of plasma cells and memory B cells specific for  
215 protective epitopes that provide long-lived protection against influenza viruses. However, the precise role  
216 of egg-specific B cells competing with virus-specific B cells remains to be determined.

217 The egg-binding mAbs demonstrated features of natural antibodies produced by innate-like B  
218 cells, including their glycan specificity and repertoire features. Natural antibodies are typically raised

219 against self-antigens and evolutionarily conserved antigens such as lipids and glycans (29). In humans,  
220 natural antibodies are largely elicited against glycans, including the blood group A and B antigens and  
221 xenoglycans from other mammals, including the “ $\alpha$ -gal” epitope of galactose- $\alpha$ -1,3-galactose expressed  
222 by most non-primate mammals (30, 31). Our studies reveals that the egg-binding antibodies were  
223 specifically targeting a secondary sulfate structure of LacNAc, a common glycan found across all life  
224 forms. Although the egg-binding mAbs identified in this study share key characteristics of natural  
225 antibodies, the precise cellular origins of these antibodies require further analysis.

226 LacNAcs are a critical component of glycosaminoglycans (GAG), including keratan sulfate and  
227 the Lewis blood group determinants (32, 33). LacNAcs are also the main ligand for galectins and mediate  
228 a variety of cellular functions including cell adhesion, migration, proliferation, and apoptosis (34-36).  
229 However, most mammals do not commonly sulfonate the galactose 4C' of LacNAc, and instead  
230 commonly sulfate the 6C' of galactose of LacNAc and 6C' of N-acetylglucosamine of LacNAc (37).  
231 Therefore, humans may mount a response specifically against the sulfated 4C' of LacNAc as it is not  
232 normally sulfated in humans. However, inflammation is associated with aberrant glycosylation patterns,  
233 including during cancer and autoimmunity (38). Moreover, antibodies against a sulfated 4C' LacNAc, the  
234 same antigen identified in this study, were elevated in subjects with systemic sclerosis and was  
235 associated with a higher prevalence of pulmonary hypertension (39). Despite these findings, the precise  
236 role and function of anti-sulfated-LacNAc antibodies in the development and severity of systemic  
237 sclerosis remain unknown. Furthermore, it is unknown if the antibodies induced by egg-grown influenza  
238 virus vaccines and those observed during systemic sclerosis share similar repertoire features and could  
239 be derived from the same B cell precursors. Lastly, it remains unknown how the 4C' sulfate LacNAc is  
240 conjugated to influenza viruses grown in eggs, as a study of the glycosylation patterns of HA and NA  
241 isolated from egg grown viruses did not have this glycan (40). In summary, this study identifies that  
242 antibodies with features of natural antibodies can be induced against a sulfated LacNAc glycan present  
243 in egg-grown vaccines.

## 244 **Methods**

### 245 **Monoclonal antibody production and sequence analysis**

246 Monoclonal antibodies were generated as previously described (15, 41, 42). Peripheral blood was  
247 obtained from each subject approximately 7 days after vaccination or infection. Lymphocytes were  
248 isolated and enriched for B cells using RosetteSep. PBs (CD3<sup>-</sup>CD19<sup>+</sup>CD27<sup>hi</sup>CD38<sup>hi</sup>) were single-cell  
249 sorted into 96-well plates. Immunoglobulin heavy and light chain genes were amplified by reverse  
250 transcriptase polymerase chain reaction (RT-PCR), sequenced, cloned into human IgG1, human kappa  
251 chain, or human lambda expression vectors, and co-transfected into HEK293T cells. Secreted mAbs  
252 were purified from the supernatant using protein A agarose beads. B cell clones were determined by  
253 aligning all the V(D)J sequences sharing identical progenitor sequences, as predicted by IgBLAST using  
254 our in-house software, Vgenes. For germline mAbs, germline sequences were synthesized (IdT) and  
255 cloned into antibody expression vectors, as described above.

### 256 **Antibody sequences and clonal analyses**

257 Previously published IgG<sup>+</sup> PB sequences (27) were downloaded from NCBI GenBank (KEOV000000000  
258 and KEOU000000000). V(D)J gene usage from our study and Forgacs et al. were analyzed using IgBlast  
259 and clones were determined using our in-house software, Vgenes, based on germline sequences. For  
260 identification of egg-like mAbs from Forgacs et al. we selected B cell clones that specifically used VH3-7  
261 with a H-CDR3 of 12 or fewer amino acids that was paired with VL1-44 or VL1-51. MAb sequence  
262 alignments were made using ClustalOmega (EMBL-EBI). Non-templated nucleotide insertions at the V-  
263 D and D-J junctions of the heavy-chain gene were identified using partis v0.15.0, a Hidden Markov Model-  
264 based tool for annotating B cell receptor sequences (43). Custom code was used for processing the  
265 output (available at [https://github.com/cobeylab/egg\\_antibodies](https://github.com/cobeylab/egg_antibodies)). Visualization of clones in Figure 5D  
266 was performed in R using circlize v0.4.12 (44).

### 267 **Viruses, proteins, and vaccines.**

268 Influenza viruses used in all assays were grown in-house in specific pathogen free (SPF) eggs,  
269 harvested, purified, and titered. Allantoic fluid was harvested from both infected and uninfected eggs. For

270 MDCK cell grown virus, A/California/7/2009 H1N1 was grown in MDCK-SIAT1 cells, concentrated, and  
271 chemically inactivated with beta-propiolactone. Newcastle disease virus was grown in eggs and allantoic  
272 fluid was harvested and subsequently inactivated with beta-propiolactone, purified, and quantified.  
273 Vaccines used for mAb binding assays are outlined in Table S6. Recombinant HA (rHA) from  
274 A/California/7/2009 was expressed in HEK293T cells.

### 275 **Antigen-Specific ELISA**

276 High protein-binding microtiter plates (Costar) were coated with 8 hemagglutination units (HAU) of virus  
277 or allantoic fluid diluted 1:500 in carbonate buffer overnight at 4°C. Plates were coated with recombinant  
278 HA from A/California/7/2009 at 1 µg/ml in PBS overnight at 4°C. For testing egg-binding mAb binding to  
279 various vaccines, influenza virus vaccines were diluted to 5 µg/ml, rabavert was diluted to 0.05 UI/ml,  
280 MMR was diluted 1:100, Ixiaro (JEV) was diluted to 0.05 antigen units per ml, and Pneumovax 23 was  
281 diluted to 5 µg/ml. All tested vaccines were diluted in PBS and coated overnight at 4°C. NDV was diluted  
282 to 5 µg/ml in carbonate buffer and plates were coated overnight at 4°C.

283 Plates were washed the next morning with PBS 0.05% Tween and blocked with PBS containing  
284 20% fetal bovine serum (FBS) for 1 hour at 37°C. MAbs were then serially diluted 3-fold starting at 10  
285 µg/ml and incubated for 1.5 hour at 37°C. For serum ELISAs, serum was diluted 1:50 and further diluted  
286 2-fold. Horseradish peroxidase (HRP)-conjugated goat anti-human IgG antibody diluted 1:1000 (Jackson  
287 Immuno Research) was used to detect binding of mAbs and serum antibodies, and plates were  
288 subsequently developed with Super Aquablue ELISA substrate (eBiosciences). Absorbance was  
289 measured at 405 nm on a microplate spectrophotometer (BioRad). To standardize the assays, control  
290 antibodies with known binding characteristics were included on each plate, and the plates were  
291 developed when the absorbance of the control reached 3.0 OD units. For the other vaccines used in  
292 Figure 2D, anti-sera against the various vaccines were used to confirm antigenicity of vaccines.  
293 Polyreactivity was determined using a polyreactive ELISA protocol, as previously described (13). Briefly,  
294 mAbs were tested for binding to 6 antigens (cardiolipin, dsDNA, flagellin, insulin, KLH, and LPS) starting  
295 at 1 µg/ml for 1.5 hours at 37°C. HRP-conjugated goat anti-human IgG antibody (Jackson Immuno  
296 Research) diluted 1:2000 in PBS/0.05% Tween/0.1mM EDTA was used to detect binding of mAbs, and

297 plates were subsequently developed with Super Aquablue ELISA substrate (eBiosciences). All mAbs or  
298 serum samples were tested in duplicate and all assays were performed 2-3 times. To determine mAb  
299 affinity, a non-linear regression was performed on background subtracted ODs and area under the curve  
300 (AUC) values were reported. Serum samples used in Figure 1h and Figure S1d-f are listed in Table S7.

### 301 **Hemagglutination inhibition assays**

302 For serum HAI assays, 1 part serum was treated with 3 parts Receptor Destroying Enzyme II (Seiken,  
303 Hardy Diagnostics) for 18 hours at 37°C, followed by 30 minutes at 56°C. Serum was further diluted to  
304 1:10 with PBS and serially diluted 2-fold in PBS in duplicate in a 96-well round-bottom plate. Serially  
305 diluted serum was mixed with an equal volume of A/California/7/2009 virus (4 HAU/25 µl), and  
306 subsequently incubated at room temperature for 1 hour. 50 µl of 0.5% turkey red blood cells (Lampire  
307 Biological) were added to each well and incubated for 45 minutes at room temperature. HAI titers were  
308 determined based on the final dilution of serum for which hemagglutination inhibition was observed. All  
309 experiments were performed in duplicate twice. The fold change in HAI serum titers of post-vaccination  
310 samples relative to pre-vaccination samples are shown in Figure S1F.

### 311 **Virus deglycosylation and sulfatase treatment**

312 To deglycosylate the vaccine, 25 µg of the 2020 QIV was denatured for 10 minutes at 75°C and treated  
313 with the Protein Deglycosylation Mix II (New England Biolabs) for 30 minutes at 25°C and 1 hour at 37°C.  
314 For sulfatase treatment, A/Hong Kong/485197/2014 H3N2 virus was diluted to 160 HAU in sodium  
315 acetate (pH 5.0) and treated with 20 units/ml of sulfatase from abalone entrails (Sigma-Aldrich) for 1 hour  
316 at 37°C. As a control, equal quantities were diluted and incubated but did not receive the Deglycosylation  
317 Mix II or sulfatase enzymes and are referred to as untreated. After treatment, preparations underwent  
318 buffer exchange with PBS to remove freed glycans and sulfate groups. ELISA plates were coated at 1  
319 µg/ml for the 2020 QIV or 8 HAU for the virus.

### 320 **Glycan microarray**

321 MAbs 029-09 3A04 and 034-10 4G02 were sent to the Protein-Glycan Interaction Resource of the Center  
322 for Functional Glycomics at the Beth Israel Deaconess Medical Center, Harvard Medical School. Printed  
323 arrays consist of 585 glycans in replicate of 6. All glycans used in the microarray are listed in Table S3

324 and S4. MAbs were diluted to 50  $\mu\text{g/ml}$  and run on the array. The highest and lowest replicates from each  
325 set of 6 replicates were removed and the average  $\pm$  S.D. was calculated from the middle 4 replicates.  
326 Structure of 6S,4S-LacNAc was made using ChemDraw JS (PerkinElmer).

### 327 **Statistics**

328 All statistical analysis was performed using Prism software (Graphpad Version 9.0). *P* values less than  
329 or equal to 0.05 were considered significant. \*  $P \leq 0.05$ , \*\*  $P \leq 0.01$ , \*\*\*  $P \leq 0.001$ , \*\*\*\*  $P < 0.0001$ .  
330 Specific number of mAbs shown in each Figure are listed in the corresponding figure legends.

### 331 **Subjects, cohorts, and study approval**

332 Written informed consent was received from participants prior to inclusion in the study. All studies were  
333 performed with the approval of the University of Chicago and Emory University institutional review boards.  
334 Subjects were recruited to receive the Sanofi Pasteur 2009 pandemic H1N1 MIV or the 2010 Novartis  
335 Fluvirin TIV. Subjects labeled with SFV were recruited and vaccinated at Emory University, as previously  
336 described (45). All other subjects were recruited and vaccinated at the University of Chicago. Subject  
337 demographics are detailed in Table S1.

338 **Author contributions**

339 J.J.G. designed the study, characterized mAbs, analyzed the data, and wrote the manuscript. H.A.U.  
340 characterized mAbs. C.H. assisted with the vaccine comparison study. L.L. assisted with data analysis.  
341 N.-Y.Z. grew and purified influenza viruses. W.S. provided inactivated Newcastle Disease Virus samples.  
342 M.C.V. performed non-templated DNA sequence analyses. S.Z. and S.H. provided MDCK cell grown  
343 virus. M.H. performed mAb cloning. S.C. and P.P. provided critical feedback and discussion on the  
344 manuscript. P.C.W. supervised the work and edited the manuscript. All authors reviewed and edited the  
345 manuscript.

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359 **Declaration of Interests**

360 The authors have declared no conflicts of interest.

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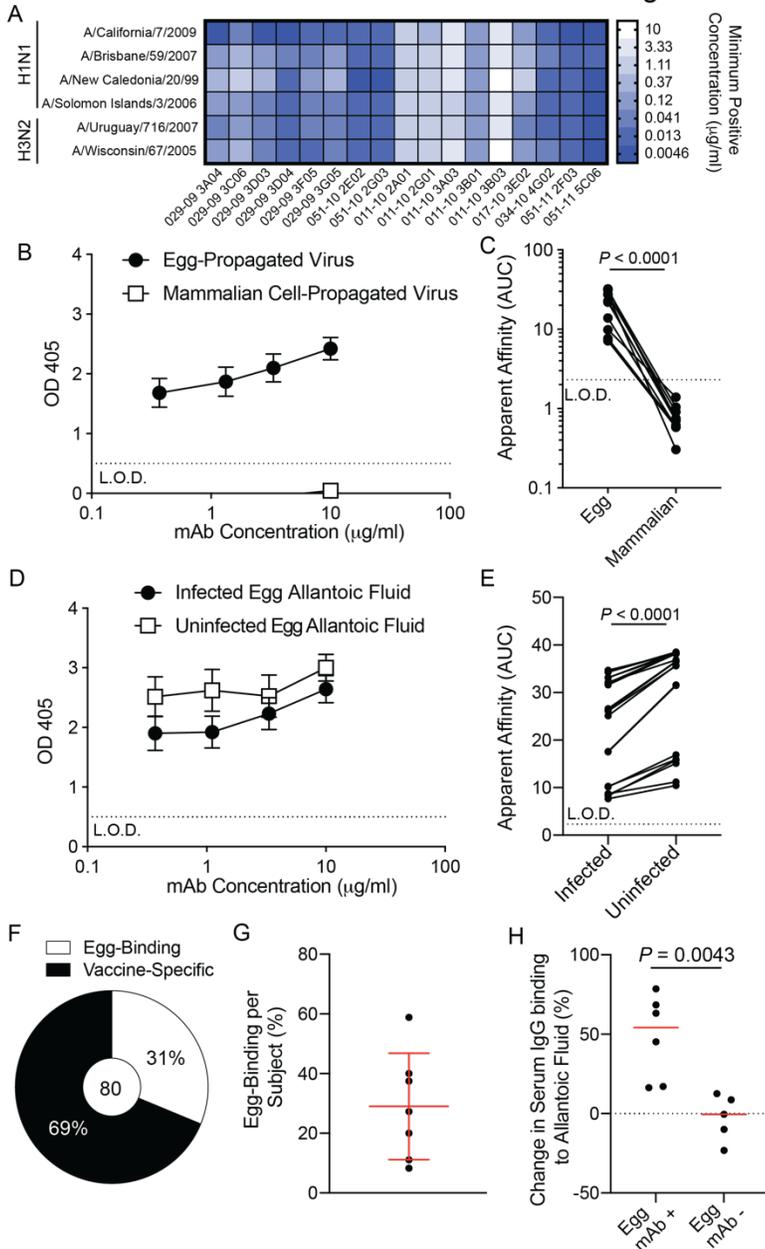
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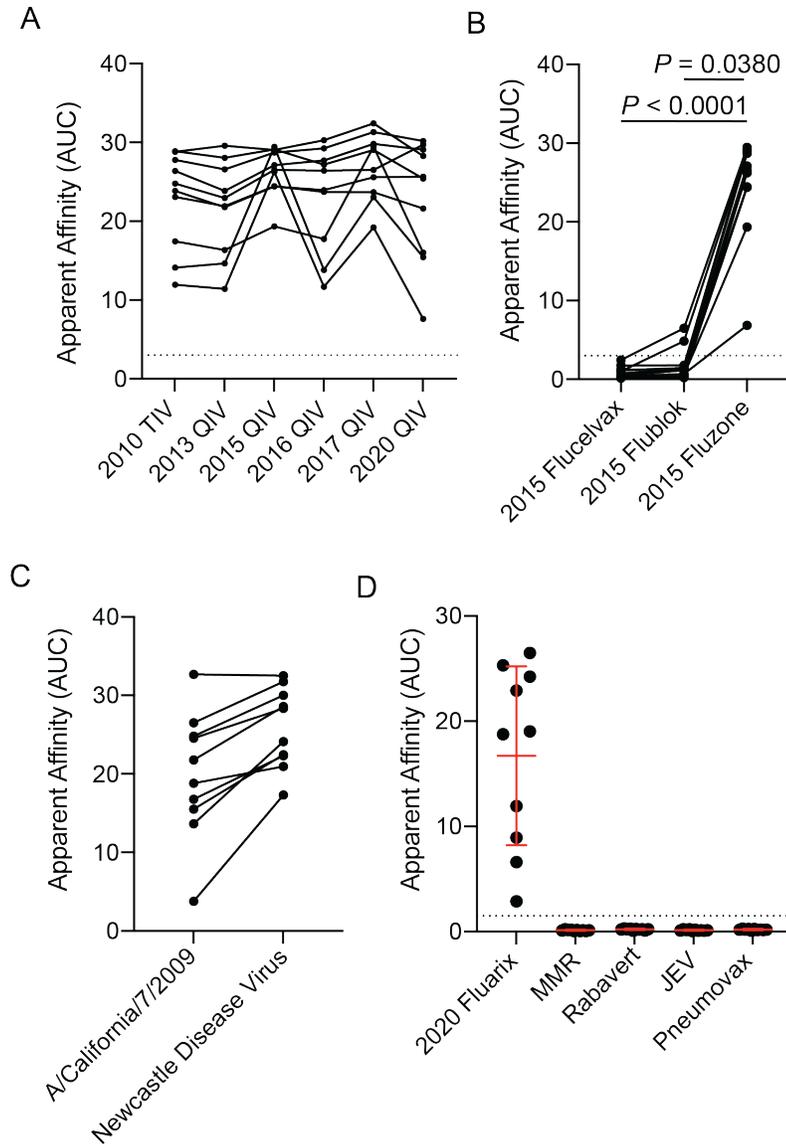
Figure 1



478

479 **Figure 1: Identification of mAbs binding an egg-specific antigen.** **A**, Heatmap of selected mAbs  
 480 binding all influenza virus strains tested. Data are representative of all 27 identified antibodies. **B** and **C**,  
 481 Broadly reactive mAbs (n=22) binding to egg-propagated and MDCK cell (mammalian) propagated  
 482 A/California/7/2009 H1N1 (**B**) and apparent affinity, as calculated as area under the curve (AUC), of mAb  
 483 binding (**C**; n=11 mAbs). **D** and **E**, Broadly reactive mAb binding to A/California/7/2009 H1N1 infected  
 484 allantoic fluid and uninfected allantoic fluid (**D**) and AUC of mAb binding (**E**; n=21). **F**, Proportion of mAbs  
 485 from subjects with egg mAbs that are egg-binding or are specific to the vaccine. **G**, Proportion of total  
 486 mAbs that are egg-binding per subject. **H**, Serum was isolated from subjects with or without isolated egg-  
 487 binding mAbs before and 14-21 days after vaccination. Relative change in serum IgG binding to  
 488 uninfected allantoic fluid represented as a percentage. Red line represented median. Data in **B**, **D**, and  
 489 **G** are mean  $\pm$  S.D. Data in **C** and **E** were analyzed using a two-tailed Wilcoxon matched-pairs signed  
 490 rank test. Data in **h** were analyzed using a two-tailed Mann-Whitney test.

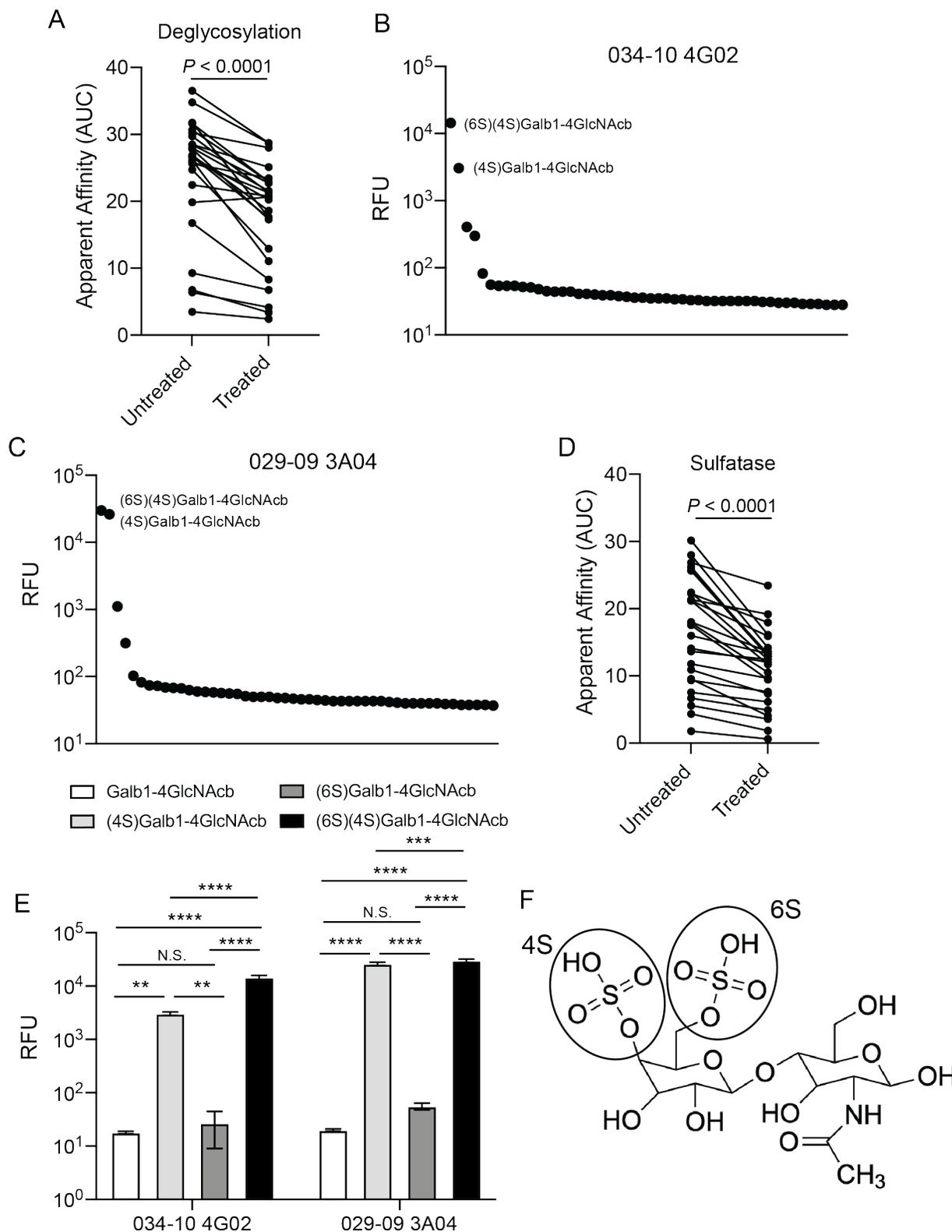
Figure 2



491

492 **Figure 2: Binding is specific to viruses grown in eggs, but not chicken cells.** **A**, Apparent affinity  
493 (AUC) of egg-specific mAbs binding to multiple years of influenza virus vaccines. Each line connects the  
494 same mAb binding a different vaccine (n=10 mAbs). **B**, Apparent affinity of mAbs binding to the  
495 mammalian cell grown vaccine (Flucelvax), insect cell grown vaccine (Flublok), and egg-grown vaccine  
496 (Fluzone). Each line connects the same mAb binding a different vaccine (n=11 mAbs). **C**, Egg-specific  
497 mAb binding to Newcastle Disease Virus grown in eggs and egg-grown A/California/7/2009 H1N1 virus  
498 (n=10). Each line connects the same mAb. **D**, Egg-specific mAbs (n=10) binding to the 2020/2021 Fluarix  
499 vaccine (egg-grown), measles/mumps/rubella vaccine (MMR; chicken cell line grown), rabavert (Rabies  
500 vaccine; chicken cell grown), Japanese Encephalitis Virus Vaccine (JEV, Ixiaro, Vero-cell grown), and  
501 pneumovax-23 vaccine (polysaccharides from bacteria). Data in **D** are mean  $\pm$  S.D. Data in **B** were  
502 analyzed using a non-parametric Friedman Test. Data in **C** were analyzed using a two-tailed Wilcoxon  
503 matched-pairs signed rank test.

Figure 3



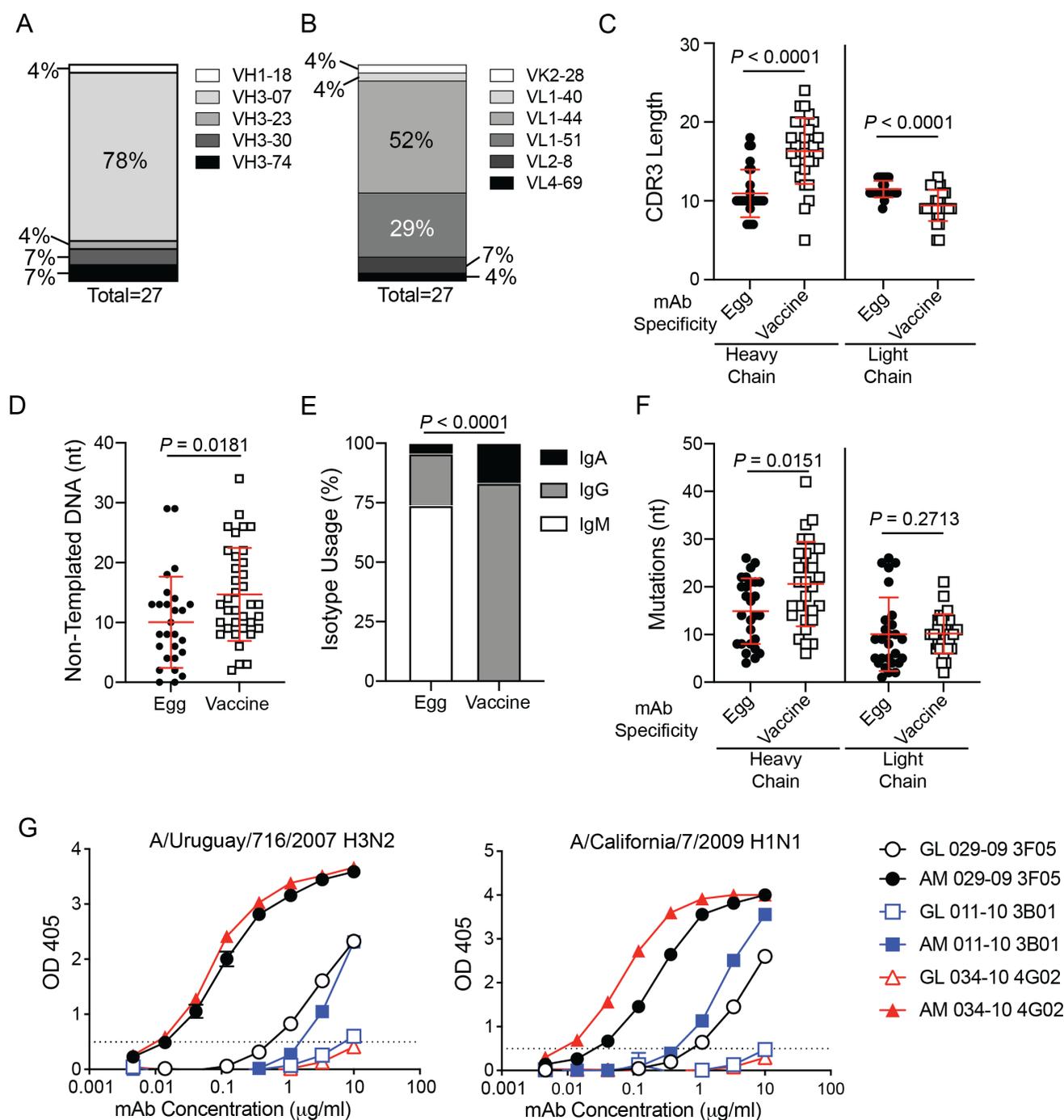
504

505 **Figure 3: Egg-specific mAbs are binding to a sulfated LacNAc.** **A**, Egg-specific mAb (n=24) binding to deglycosylated or untreated 2020 FluAriX QIV. **B** and **C**, 034-10 4G02 (**B**) and 029-09 3A04 (**C**) were tested for binding to glycans on a microarray. Data represent the top 50 glycan hits. **D**, Egg-specific mAb (n=26) binding to sulfatase treated or untreated A/Hong Kong/485197/2014 H3N2. **E**, 034-10 4G02 and

508

509 029-09 3A04 binding to non-sulfated and sulfated LacNAc glycans in glycan microarray. **F**, Structure of  
510 (6S)(4S)Galb1-4GlcNac (LacNAc). Data in **B**, **C**, and **E** are averaged RFU of 4 replicates. Each line in  
511 **A** and **D** connects the same mAb. Data in **E** are mean  $\pm$  S.D. Data in **A** and **D** were analyzed using a  
512 two-tailed Wilcoxon matched-pairs signed rank test and data in **E** were analyzed using an ordinary two-  
513 way ANOVA.

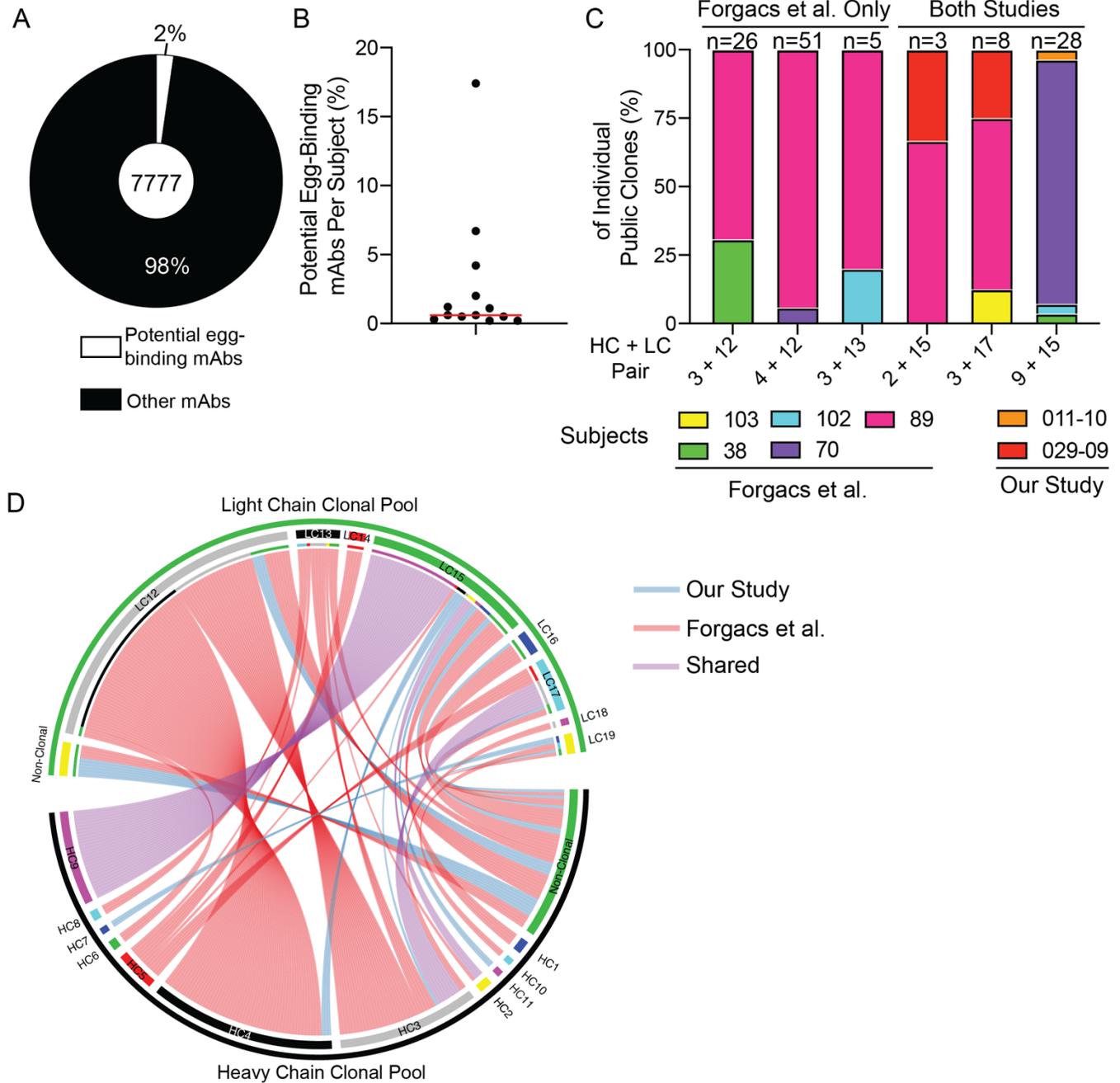
Figure 4



514

515 **Figure 4: Egg-binding mAbs resemble natural antibodies.** **A** and **B**, VH (**A**) and VK/VL (**B**) gene  
 516 usage of egg-binding mAbs. **C** and **D**, heavy chain and light chain CDR3 lengths (**C**) and non-templated  
 517 DNA (N-nucleotides; **D**) of egg-binding and vaccine-specific mAb heavy chains. **E** and **F**, isotype usage  
 518 (**E**) and somatic hypermutations (nucleotide mutations; nt; **F**) of egg-binding and vaccine-specific mAbs.  
 519 **G**, affinity-matured (AM) 029-09 3F05, 011-10 3B01, and 034-10 4G02 were reverted back to germline  
 520 (GL) and were tested for binding to egg-grown A/Uruguay/716/2007 H3N2 and A/California/7/2009 H1N1  
 521 relative to their affinity-matured (AM) counterparts. Data in **C**, **D**, **F**, and **G** are mean  $\pm$  S.D. Data in **C**, **D**,  
 522 and **F** were analyzed using two-tailed Mann-Whitney tests. Data in **E** were analyzed using chi-squared  
 523 test.

Figure 5



524

525 **Figure 5: PBs with repertoire features of egg-binding antibodies are common after vaccination.**  
 526 B cell receptor sequences from 7,777 IgG<sup>+</sup> PBs induced by influenza virus vaccination were analyzed  
 527 for repertoire features of egg-binding mAbs (VH3-7 with H-CDR3 of ≤ 12 amino acids paired with VL1-  
 528 44 or VL1-51). **A** and **B**, proportion of sequences with egg-binding mAb repertoire features out of all  
 529 sequences (**A**) and by subject (**B**). Red line in **B** represents the median. **C**, subject makeup (%) of  
 530 public clones, including public clones specific to Forgacs et al. and shared across studies. The number  
 531 above each column represents the number of clonal members per clone. **D**, circos plot of heavy and  
 532 light chain clones distinct to each study (our study: blue; Forgacs et al. red) or shared across studies  
 533 (purple).