

## **Supplementary information for**

### **Broadly protective murine monoclonal antibodies against influenza B virus target highly conserved neuraminidase epitopes**

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**Supplementary Figure 5. IBV anti-NA mAbs reduce viral lung titers in mice, activate ADCC, inhibit NA activity of drug-resistant IBV, and demonstrate superior effectiveness to oseltamivir.**

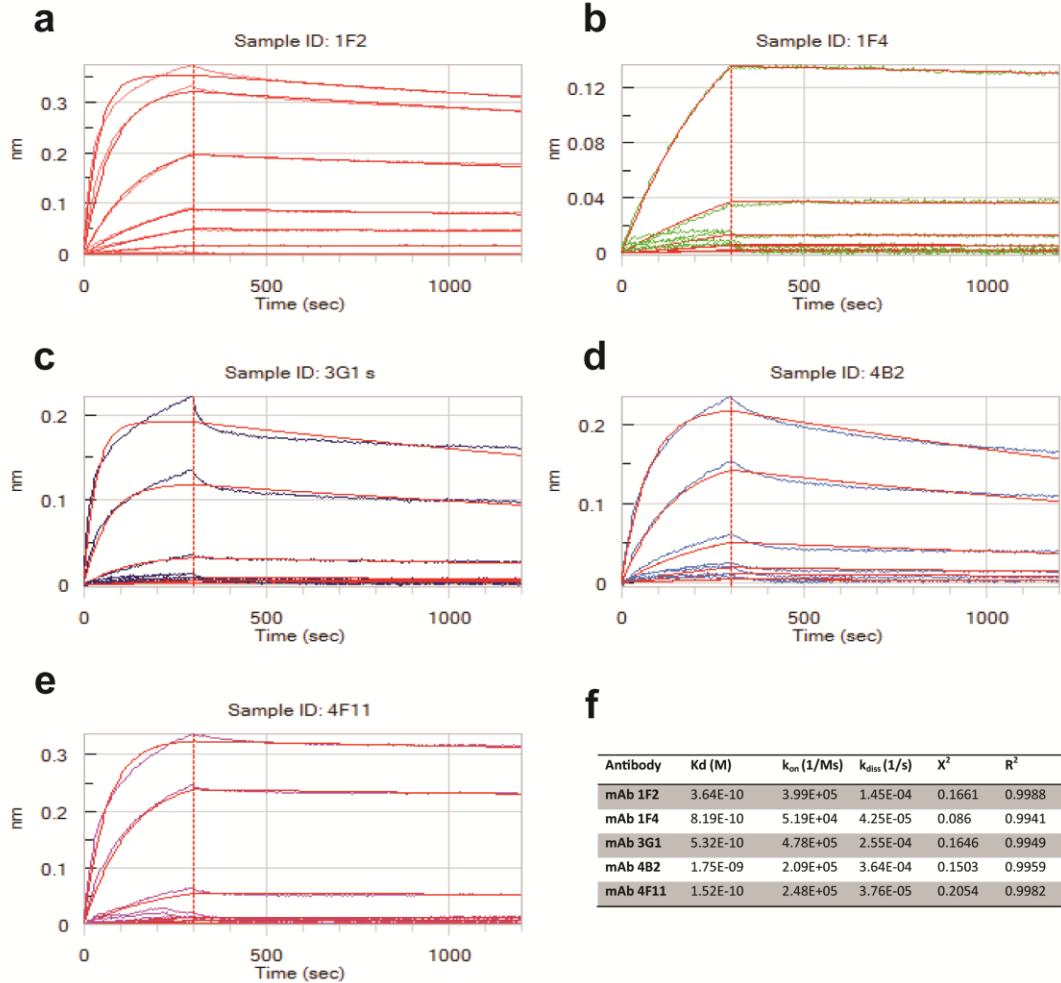
**Supplementary Figure 6. Neuraminidase inhibition assay against the escape mutants raised with the five mAbs.**

**Supplementary Figure 7. Oseltamivir treatment of mice initiated 48 hours post infection with B/Malaysia/2506/04 virus does not protect from weight loss but leads to survival of the infection.**

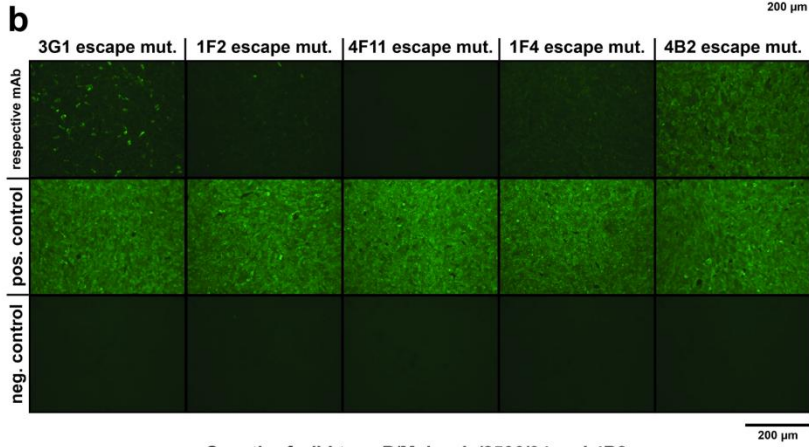
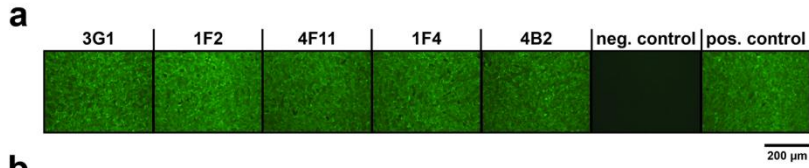
#### **Supplementary notes**

**IgG sequences and comparison to IgG germlines**

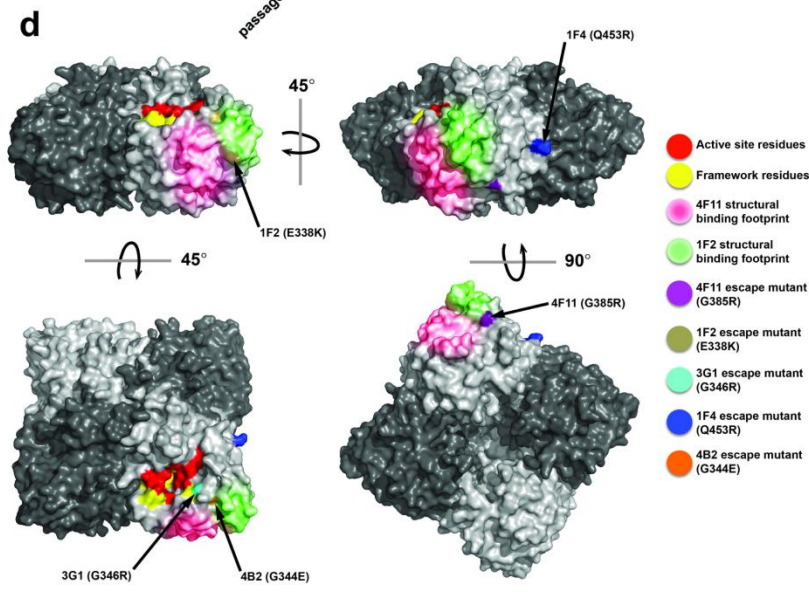
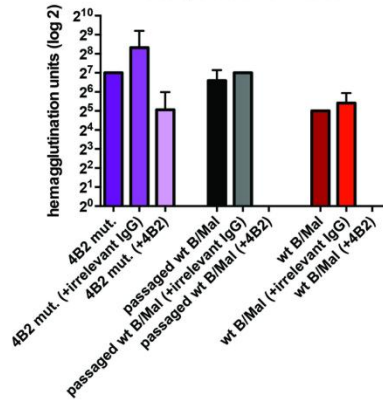
## Supplementary Figures



**Supplementary Figure 1. Antibody affinity as measured by bio-layer interferometry.** Antibody binding kinetics to recombinant NA from B/Malaysia/2406/04 of mAbs 1F2 (**a**), 1F4 (**b**), 3G1 (**c**), 4B2 (**d**) and 4F11 (**e**) as measured by bio-layer interferometry. (**F**) shows a summary of the measured  $K_d$ ,  $k_{on}$ ,  $k_{diss}$  as well as curve fitting parameters.



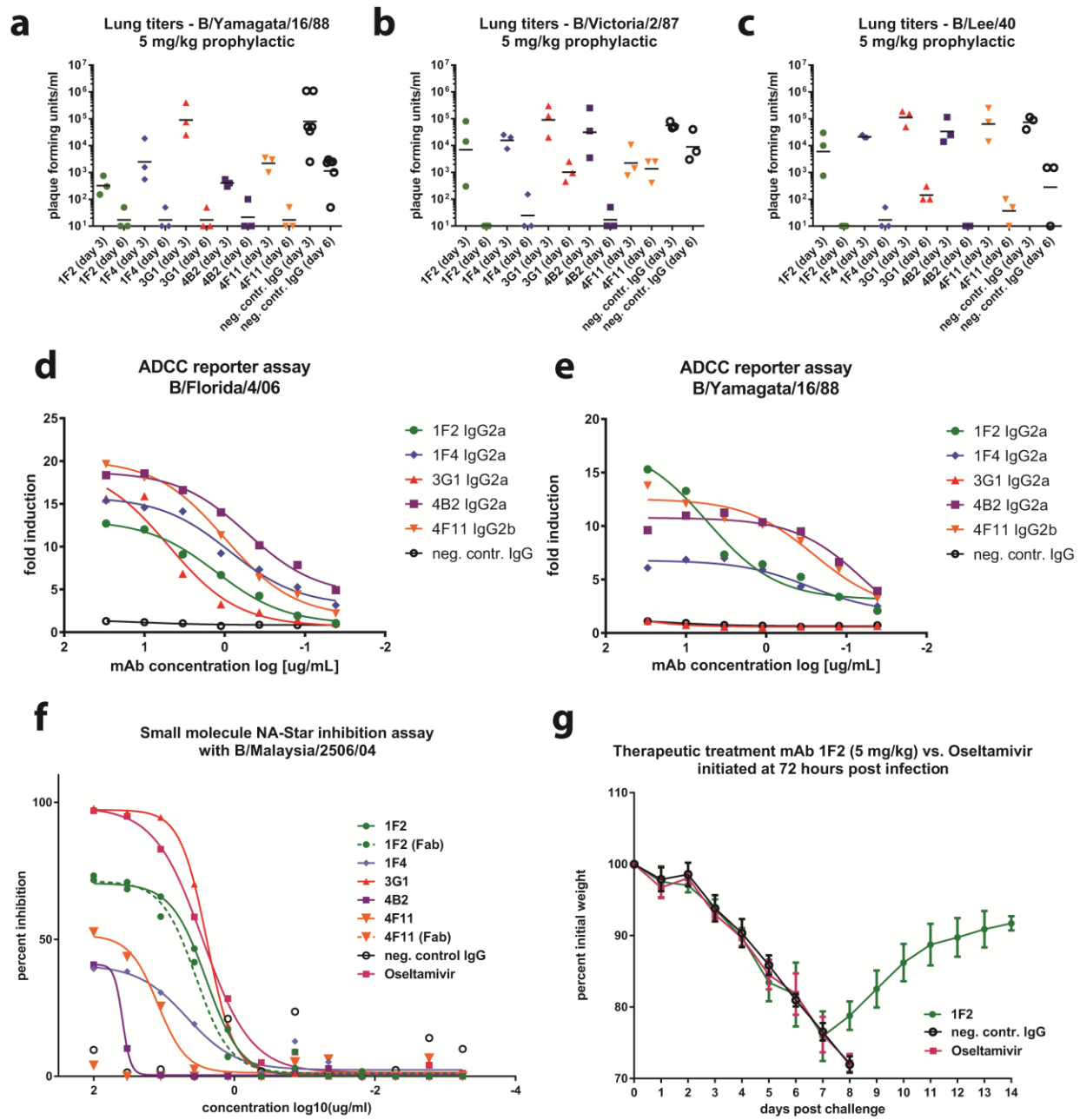
**c** Growth of wild type B/Malaysia/2506/04 and 4B2 escape mutant viruses



**e**

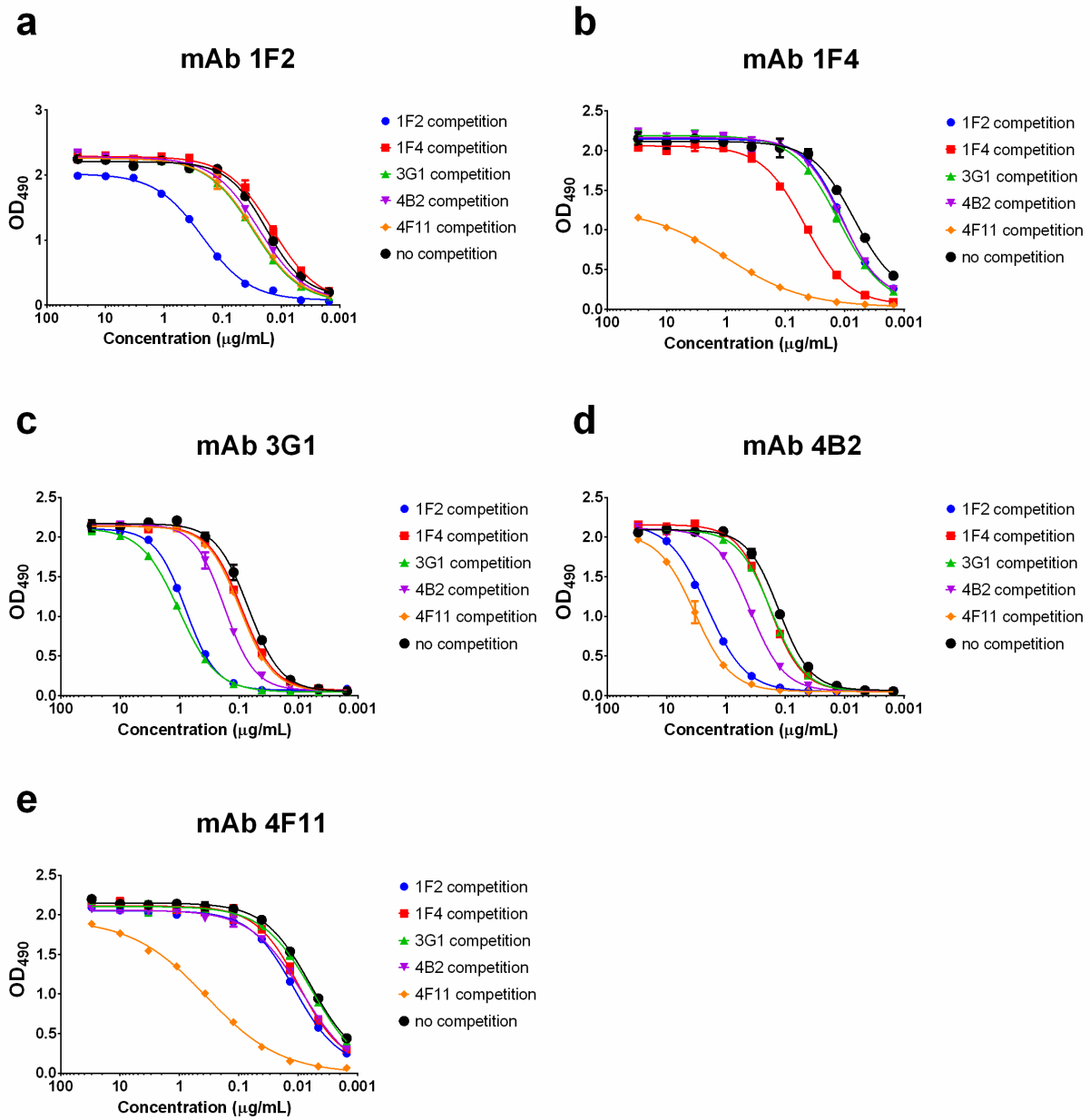
	Position	Amino Acid Residue	% Conservation
4F11 escape mutant	385	G	100
1F2 escape mutant	338	E	99.36
3G1 escape mutant	346	G	99.89
1F4 escape mutant	453	Q	99.89
4B2 escape mutant	344	G	99.89

**Supplementary Figure 2. IBV escape mutants reveal amino acid residues critical for mAb binding.** **(a)** To demonstrate mAb reactivity via immunofluorescence, MDCK cells were infected with wt B/Malaysia/2506/04 virus (MOI = 10), fixed, and stained using anti-NA mAbs (30 ug/ml). All mAbs displayed clear surface staining, allowing this assay to be used as a screen for potential binding mutants. A polyclonal cocktail of purified mouse mAb IgGs against the IBV HA was used as a positive infection control (pos. control). An irrelevant mouse mAb, 8H9 was used as a negative control (neg. control). **(b)** MDCK cells were infected with the generated B/Malaysia/2506/04 escape mutant viruses and stained with the respective mAb to which the escape mutant was generated, in a similar fashion to panel A. All mutant viruses – except that generated to mAb 4B2 – displayed clear loss of binding to the corresponding mAb. Scale bars represent 200 um. **(c)** HA titers of wt B/Malaysia/2506/04 virus (wt B/Mal), 4B2 escape mutant virus (4B2 mut.), and passaged wt B/Malaysia/2506/04 virus (passaged wt B/Mal) in the presence of mAb 4B2 at 10 ug/ml, irrelevant mouse mAb 3C12 (anti-N8) at 10 ug/ml, or no mAb at 72 hpi. Only the generated 4B2 escape mutant virus grew to detectable titers in the presence of 4B2. Passaged wt B/Malaysia/2506/04 virus, as explained in detail in the materials and methods section, is a control virus produced by serially passaging wt B/Malaysia/2506/04 virus in MDCK cells in the presence of irrelevant mouse mAb 3C12 alongside wt B/Malaysia/2506/04 virus in the presence of increasing concentrations of 4B2. The mean and standard error of the mean, obtained from biological triplicates, are displayed graphically. **(d)** Critical binding residues identified in IBV escape mutants – along with the structurally defined binding footprints from Fig. 2 and the NA enzymatic active site/framework residues – were mapped on one of the four monomers of the 3D structure of the NA from B/Brisbane/60/2008 virus (PDB ID: 4CPL). The remaining three monomers of the tetramer are shown in either light or dark grey. Degrees of rotation are approximate. **(e)** List of amino acid residues (position, identity, and percent conservation) identified as critical binding residues by escape mutant generation. Percent conservation was determined using 944 subsampled IBVs. B/Malaysia/2506/04 numbering is used throughout.

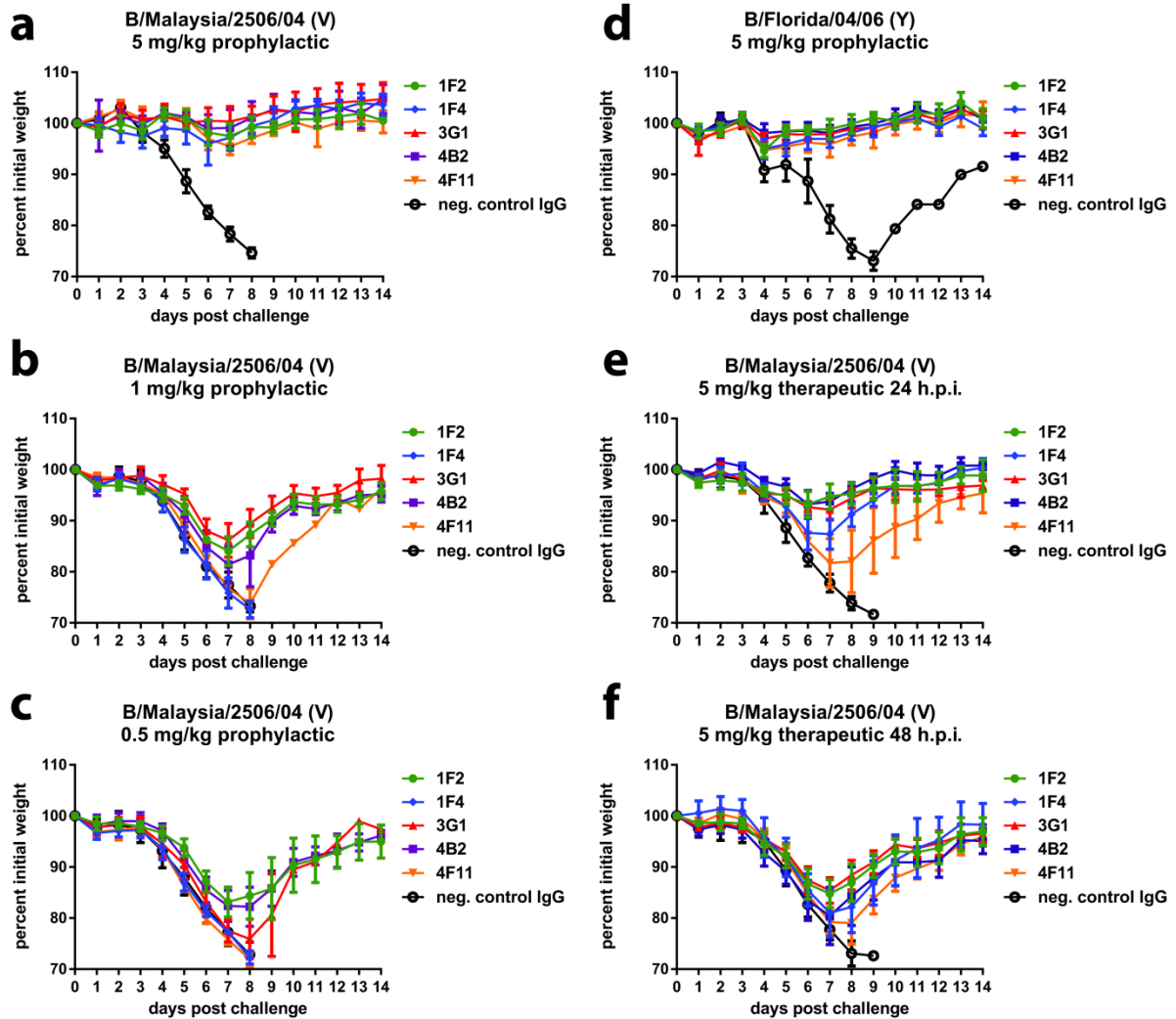


**Supplementary Figure 3. IBV anti-NA mAbs reduce viral lung titers in mice, activate ADCC, inhibit NA activity of drug-resistant IBV, and demonstrate superior effectiveness to oseltamivir.** (a-c) Female BALB/c mice (3 per group) were administered 5 mg/kg antibody prophylactically, challenged with B/Yamagata/16/88, B/Victoria/2/87, or B/Lee/40 viruses respectively and in identical fashion to Figure 4A. Mice were sacrificed on day 3 or 6 post-infection for lung titer analysis. Lung titers in groups treated with anti-NA mAbs are most reduced on day 6 post-infection compared to negative control mAb 8H9. (d, e) Anti-NA mAb incubated with MDCK cells infected with B/Florida/04/06 or B/Yamagata/16/88 viruses, respectively (MOI = 3), are able to engage Fc receptors and activate ADCC *in vitro*. Fold induction is defined as RLU

(induced by antibody)/ RLU (no antibody control background). Murine mAb 2G12 (anti-Ebolavirus Gp) is used as a negative control. Technical triplicates were performed. **(f)** NI activity against B/Malaysia/2506/04 using the NA-*Star* assay. Data points are presented as percent inhibition. Technical duplicates were performed. **(g)** Female BALB/c mice (5 per group) were administered either 5 mg/kg of mAb 1F2 intraperitoneally, 5 mg/kg negative control mAb 8H9 intraperitoneally, or placed on a twice daily, 20 mg/kg, 6 day-long regimen of oseltamivir delivered via oral gavage and initiated at 72 hpi with B/Malaysia/2506/04. Percent weight is shown and is calculated based on the initial body weight on day 0.

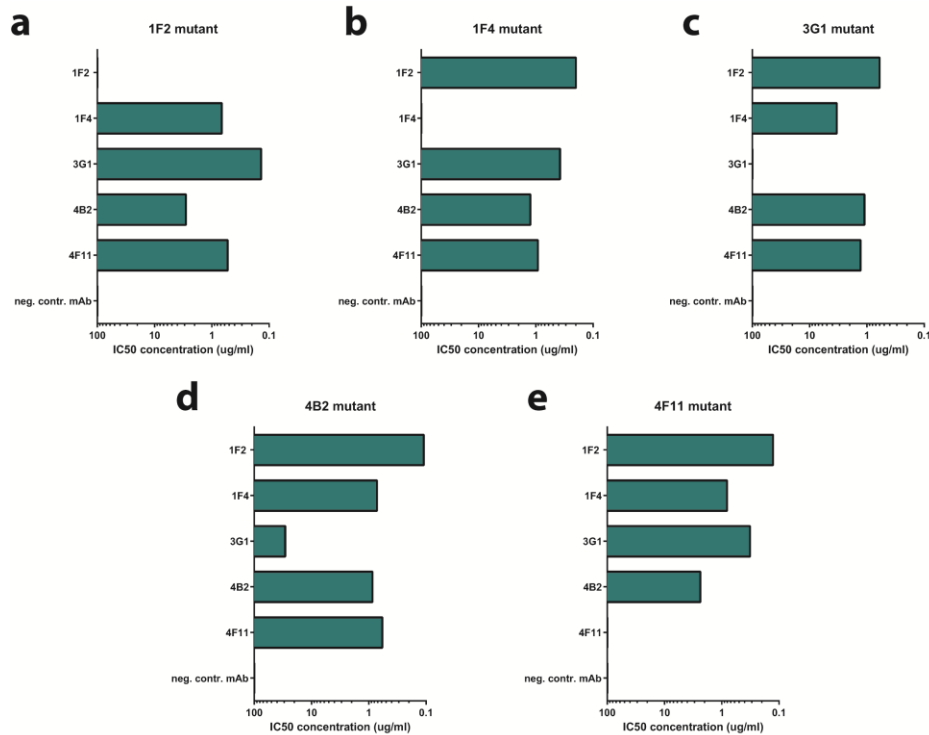


**Supplementary Figure 4. Competition between different mAbs for binding to B NA.** Antibodies 1F2 (a), 1F4 (b), 3G1 (c), 4B2 (d) and 4F11 (e) were competed against themselves and each other using an ELISA-based assay. Technical duplicates were performed.

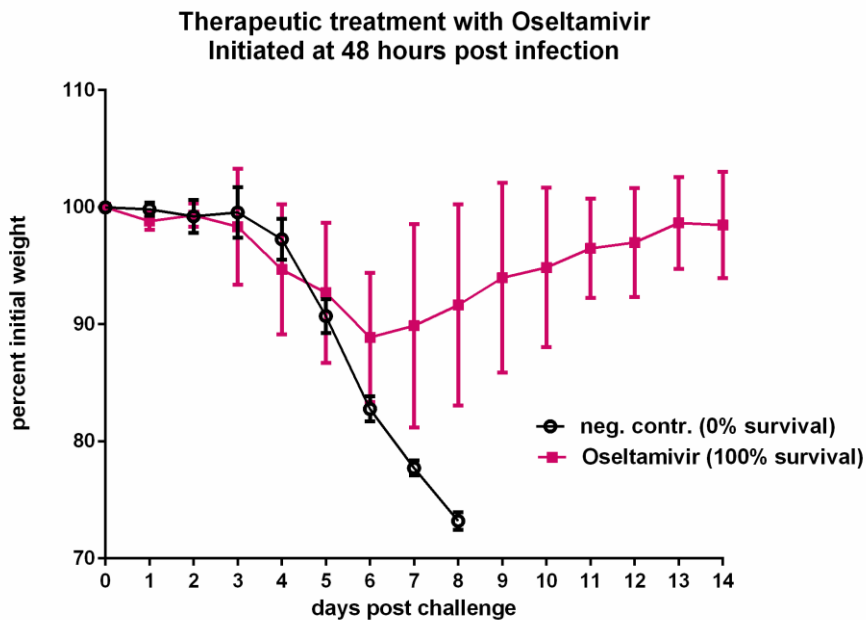


**Supplementary Figure 5. IBV anti-NA mAbs protect mice from morbidity when administered prophylactically or therapeutically.** Displayed are the weight loss curves corresponding to the survival curves in **Fig. 3**. Mice (5 per group) were administered either 5, 1, or 0.5 mg/kg of mAb intraperitoneally 2 h prior to a 5 mLD<sub>50</sub> challenge with B/Malaysia/2506/04 virus (**a-c**) or administered 5 mg/kg of mAb intraperitoneally 2 h prior to a 5 mLD<sub>50</sub> challenge with B/Florida/04/06 virus (**d**). In therapeutic studies, mice were administered 5 mg/kg of each antibody either 24 (**e**) or 48 (**f**) h after challenge with 5 mLD<sub>50</sub> of B/Malaysia/2506/04 virus. Percent weight is calculated based on the initial body weight on day 0. Mice that lost more than 25% of their initial body weights were humanely euthanized, and the remaining curves were generated from the weights of the surviving mice only. Murine mAb 8H9 (anti-H6) was used as a negative control in all experiments.





**Supplementary Figure 6. Neuraminidase inhibition assay against the escape mutants raised with the five mAbs.** Escape mutants generated with 1F2 (a), 1F4 (b), 3G1 (c), 4B2 (d) and 4F11 (e) were each tested for sensitivity to the panel of five mAbs. The means obtained from technical duplicates are displayed graphically.



**Supplementary Figure 7. Oseltamivir treatment of mice initiated 48 hours post infection with B/Malaysia/2506/04 virus does not protect from weight loss but**

**leads to survival of the infection.** Female BALB/c mice (5 per group) were administered 5 mg/kg negative control mAb 8H9 intraperitoneally or were placed on a twice daily, 20 mg/kg, 6 day-long regimen of oseltamivir delivered via oral gavage and initiated at 48 hpi. Percent weight is shown and is calculated based on the initial body weight on day 0. Percentages next to the legend indicate survival.

## Supplementary notes

### **IgG sequences and comparison to IgG germlines**

#### **mAb 1F2**

##### IGHV-1 77\*01

Query	1	QVHLQQSGPEVARPGASVKLSCKASGYTFTDYILNWVKQRPRQGLEWIGQIHPGSTNTYY	60
		QV L+QSG E+ +PGASVK+SCKASGYTFTDYY+NWVKQRP QGLEWIG+I PGS +TYY	
Germ.	1	QVQLKQSGAELVKPGASVKISCKASGYTFTDYYINWVKQRPGQGLEWIGKIGPGSGSTYY	60
Query	61	NEKFKGKATLTADKSSSTAYMQLSSLTFEDSAVYFCA	97
		NEKFKGKATLTADKSSSTAYMQLSSLT EDSA VYFCA	
Germ.	61	NEKFKGKATLTADKSSSTAYMQLSSLTSEDSAVYFCA	97

##### IGHJ4\*01

Query	107	YAMVCWGQGTAVTVSS	122
		YAM WGQGT+VTVSS	
Germ.	2	YAMDYWGQGTSVTVSS	17

##### IGHD3\*01

Query	112	WGQGTAVTVSS	122
		WGQGT VTVS+	
Germ.	5	WGQGTLVTVSA	15

##### IGKV6-15\*01

Query	1	DIVMTQSQKFMSTSVGDRVSVTCKASQNVVTNVVWYQQKPGQSPKPLIYSASYRYSYSGVPD	60
		DIVMTQSQKFMSTSVGDRVSVTCKASQNV TNV WYQQKPGQSPK LIYSASYRYSYSGVPD	
Germ.	1	DIVMTQSQKFMSTSVGDRVSVTCKASQNVGTNVAWYQQKPGQSPKALIIYSASYRYSYSGVPD	60
Query	61	RFTGSGSGTDFTLTISNVQSEDLAEYCCQQYHSYP	95
		RFTGSGSGTDFTLTISNVQSEDLAEY CQQY+SYP	
Germ.	61	RFTGSGSGTDFTLTISNVQSEDLAEYFCQQYNSYP	95

##### IGKJ4\*01

Query	96	FTFGSGTKLEVK	107
		FTFGSGTKLE+K	
Sbjct	1	FTFGSGTKLEIK	12

#### **mAb 1F4**

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Query 1 QVHLQQSGSELRSPPGSSVKLSCKDFDSEVFPIVYMRWIRQKPGHGFIEWIGDILPSFGRTI 60  
QVHLQQSGSELRSPPGSSVKLSCKDFDSEVFPI YM W+RQKPGHGFIEWIGDILPS GRTI  
Germ. 1 QVHLQQSGSELRSPPGSSVKLSCKDFDSEVFPIAYMSWVRQKPGHGFIEWIGDILPSIGRTI 60

Query 61 YGEKFEDKATLDADTVSNTAYLELNSLTSEDSAIYYCAR 99  
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Germ. 61 YGEKFEDKATLDADTVSNTAYLELNSLTSEDSAIYYCAR 99

IGHJ3\*01

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W AYWQGTLVTVSA  
Germ. 1 WFAYWQGTLVTVSA 15

IGHD2-1\*01

YYGNY  
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IGKV6-25\*01

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DIVMTQSHKFMSTSVGDRV+ITCKASQDVST VAWYQQKPGQSPKLLIYWASTRHTGVP+  
Germ. 1 DIVMTQSHKFMSTSVGDRVSITCKASQDVSTAVAWYQQKPGQSPKLLIYWASTRHTGVPD 60  
Query 61 RFTGIIISGTDYTLTISSVQAEDRALYYCQQHYSAP-----WTFGGGTKLEIK 107  
RFTG SGTDYTLTISSVQAED ALYYCQQHYS P WTFGGGTKLEIK  
Germ. 61 RFTGSGSGTDYTLTISSVQAEDLALYYCQQHYSTPIGKJ\*WTFGGGTKLEIK 112

IGKJ1\*01

Query 96 WTFGGGTKLEIK 107  
WTFGGGTKLEIK  
Germ. 1 WTFGGGTKLEIK 12

**mAb 3G1**

IGHV1-9\*01

Query 1 QVQLQQSGAELMKPGASVKISCKATGKYKFTSYWIGWVKQRPGHGLEWCGEIFPGSGSINY 60  
QVQLQQSGAELMKPGASVK+SCKATGY FT YWI WVKQRPGHGLEW GEI PGSGS NY  
Germ. 1 QVQLQQSGAELMKPGASVKLSCKATGYTFTGYWIEWVKQRPGHGLEWIGEILPGSGSTNY 60  
Query 61 NEKFKGKATFTADTSSNTAYLQLTSLTSEDSAVYYCAR 98  
NEKFKGKATFTADTSSNTAY+QL+SLT+EDSA+YYCAR  
Germ. 61 NEKFKGKATFTADTSSNTAYMQLSSLTTEDSAIYYCAR 98

IGHJ4\*01

Query 107 YGAMDYWGQGTSLTVSS 123  
Y AMDYWGQGT+TVSS  
Germ. 1 YYAMDYWGQGT+SVTVSS 17

IGHD1-1\*01

Query 102 YYGSSY 107  
YYGSSY  
Germ. 2 YYGSSY 7

IGKV16-104\*01

Query 1 DVQITQSPSYLAASPGETITINCRASKSISKYVAWYQEKPGRTNKVLIYSGSILSFGNPS 60  
 DVQITQSPSYLAASPGETITINCRASKSISKY+AWYQEKPG+TNK+LIYSGS L G PS  
 Germ. 1 DVQITQSPSYLAASPGETITINCRASKSISKYLAWYQEKPGKTNKLLIYSGSTLQSGIPS 60

Query 61 RFSGSGSGTDFTLTISSELEPEDFAMYYCQQHNEYPW 96  
 RFSGSGSGTDFTLTISSELEPEDFAMYYCQQHNEY+  
 Germ. 61 RFSGSGSGTDFTLTISSELEPEDFAMYYCQQHNEYPY 96

IGKJ1\*01

Query 96 WTFGGGKLEIK 107  
 WTFGGGKLEIK  
 Germ. 1 WTFGGGKLEIK 12

**mAb 4B2**

IGHV9-2-1\*01

Query 1 QIQLVQSGPELKKPGETVKISCKASGFTFTDYPMHVVKQAPGKSLKWMGWINTETEEPTY 60  
 QIQLVQSGPELKKPGETVKISCKASG+TFTDY MHVVKQAPGK LKWMGWINTET EPTY  
 Sbjct 1 QIQLVQSGPELKKPGETVKISCKASGYTFTDYSMHVVKQAPGKGLKWMGWINTETGEPTY 60

Query 61 SDDFKGRSPLSLETSASTTYLQINNLKNEEDTSTYFCVR 98  
 +DDFKGR SLETSAST YLQINNLKNEEDT+TYFC R  
 Sbjct 61 ADDFKGRFAFSLETSASTAYLQINNLKNEEDTATYFCAR 98

IGHJ3\*01

Query 109 WFGYWGQGTTLVTVSA 123  
 WF YWGQGTTLVTVSA  
 Sbjct 1 WFAYWGQGTTLVTVSA 15

IGHJ3\*02

Query 109 WFGYWGQGTTLVTVSA 123  
 WFG WGQGTTLVTVSA  
 Sbjct 1 WFG\*WGQGTTLVTVSA 15

IGHD1-1\*01

Query 101 YYYGSTY 107  
 YYYGS+Y  
 Sbjct 1 YYYGSSY 7

IGKV1-110\*01

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 Sbjct 1 DVVMTQTPLSLPVSLGDQASISCRSSQSLVHNSNGNTYHLHWYLQKPGQSPKLLIYKVSNRF 60

Query 61 SGVPDRFTGGGSGTDFTLKISRVEAEDLGIYFCSQSALFP 100  
 SGVPDRF+G GSGTDFTLKISRVEAEDLG+YFCSQS P  
 Sbjct 61 SGVPDRFSGSGSGTDFTLKISRVEAEDLGVYFCSQSTHVP 100

IGKJ2\*01

Query 101 YTFGGGTNLEIK 112  
 YTFGGGT LEIK  
 Sbjct 1 YTFGGGKLEIK 12

**mAb 4F11**

IGHV5-6-4\*01

Query 1 DVKLVESGGDLVKPGGSLKLSAASGFTFSAYSMSWVRQTPERRLEWVATINTGGSFTYY 60  
DVKLVESGG LVKPGGSLKLSAASGFTFS+Y+MSWVRQTPE+RLEWVATI++GGS+TTY  
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Query 61 PDSVKGRFTISRDNANTLYLQMSLSEDTAMYFCTR 98  
PDSVKGRFTISRDNANTLYLQMSLSEDTAMY+CTR  
Sbjct 61 PDSVKGRFTISRDNANTLYLQMSLSEDTAMYYCTR 98

IGHJ3\*01

Query 107 YFPYWGQGTLVIVFA 121  
+F YWGQGLV V A  
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IGHD1-1\*01

YYYGSSY  
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IGKV4-68\*01

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Sbjct 1 QIVLTQSPALMSASPGEKVTMTCSASSSVSYMYWYQQKPRSSPKPWIYLT SNLASGVPAR 60  
Query 61 FSGSGSGTSYSLTISSMEAEDVATYYCQQWSSDP 94  
FSGSGSGTSYSLTISSMEAED ATYYCQQWSS+P  
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IGKJ2\*01

Query 96 TFGGGTKVEIK 106  
TFGGGTK+EIK  
Sbjct 2 TFGGGTKLEIK 12