

## Supplementary Material

## **Supplementary Figures**

Victoria Lineage (B-V)				
Use	Strain	Sequence		
Vaccine	B/Maryland/15/2016	MKAIIVLLMVVTSNADRICTGITSSNSPHVVKTATQGEVNVTGVIPLTTTPTKSHFANLKGTETRGKLCP KCLNCTDLDVALGRPKCTGKIPSARVSILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEHVRLSTHNVI NAEDAPGGPYKIGTSGSCPNITMGNGFFATMAWAVPDKNKTATRPLTRUPVPVCTEGEDQITVWGFHSDN EIQMAKLYGDSKPQKFTSSANGVTTHYVSQIGGFPNQTEDGGLPQSGRIVVDYMVQKSGKTGTITYQRGI LLPQKVWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGTKY RPPAKLKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGGVAVADLKSTQEAINITKNLINSLSELEV KNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPSA VEIGNGCFETKHKCNQTCLDKIAAGTFDAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLAV TILMIEIVYMVMYSRDNYSCSICL		
	B/lowa/06/2017	MKAIIVLLMVVTSNADRICTGITSSNSPHVVKTATQGEVNVTGVIPLTTTPTKSHFANLKGTETRGKLCP KCLNCTDLDVALGRPKCTGKIPSARVSILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEHVRLSTHNVI NAEGAPGGPYKIGTSGCPNITMGNGFFATMAWAVPDKNKTATNPLTIPLVPYVCTEGEDQITVWGFHSDN ETQMAKLYGDSKPQKFTSSANGVTTHYVSQIGGFPNQTEDGGLPQSGRIVVDYMVQKSGKTGTITYQRGI LLPQKVWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGTKY RPPAKLKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGGVAVADLKSTQEAINITKNLNISLSELEV KNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPSA VEIGNGCFETKHKKNQTCLDKIAAGTFDAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLAV TLMIAIFVVYMVSRDNYSCSICL		
ELISA	B/Brisbane/60/2008	MKAIIVLLMVVTSNADRICTGITSSNSPHVVKTATQGEVNVTGVIPLTTTPTKSHFANLKGTETRGKLCP KCLNCTDLDVALGRPKCTGKIPSARVSILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEHIRLSTHINVI NAENAPGGPYKIGTSGSCPNITNGNGFFATMAWAVPKNDKNKTATNPLTIEVPYICTEGEDQITVWGFHS DNEAQMAKLYGDSKPQKFTSSANGVTTHYVSQIGGFPNQTEDGGLPQSGRIVVDYMVQKSGKTGTITYQR GILLPQKVWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGT KYRPPAKLLKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGVAVAADLKSTQEAINKITKNLNSLSEL EVKNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGP SAVEIGNGCFETKHKCNQTCLDRIAAGTFDAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSL AVTLMIAIFVVYMVSRDNYSCSICL		
Neutralization	B/Colorado/06/2017	MKAIIVLLMVVTSSADRICTGITSSNSPHVKTATQGEVNVTGVIPLTTTPTKSHFANLKGTETRGKLCP KCLOTTDLDVALGRPKCTGKIPSARVSILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEHVRLSTHNVI NAEGAPGGPYKIGTSGSCPNITNGNGFFATMAWAVPDKNKTATNPLTIEVPYVCTEGEDQITVWGFHSDN ETQMAKLYGDSKPQKFTSSANGVTTHYVSQIGGFPNQTEDGGLPQSGRIVVDYMVQKSGKTGTITYQRGI LLPQKVWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGTKY RPPAKLIKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGVAVAADLKSTQEAINKITKNLNSLSELEV KNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPSA VEIGNGCFETKHKCNQTCLDKIAAGTFDAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLAV ITLMIAIFVVYMVSRDNVSCSICL		

Yamagata Lineage (B-Y)				
Use	Strain	Sequence		
Vaccine	B/Singapore/INFTT-16-0610/2016	MVVTSNADRICTGITSSNSPHVVKTATGGEVNVTGVIPLTTTPTKSYFANLKGTRTRGKLCPD CLNCTDLDVALGRPMCVGTTPSAKASILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEKIRLSTQNVI DAEKAPGGPYRLGTSGSCPNATSKIGFFSTMAWAVPKDNYKNATNPQTVEVPYICTEGEDQITVWGFHSDN KTQMKSLYGDSNPQKFTSSANGVTTHVVSQIGDFPDQTEDGGLPQSGRIVVDYMVQKPGKTGTIVYQRG VLLPQKVWCASGRSKVIKGSLPLIGEADCLHEEYGGLNKSKPYYTGKHAKAIGNCPIWVKTPLKLANGTK YRPPAKLLKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGVAVADLKSTQEAINKITKNLNSLSELE VKNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPS AVDIGNGCFETKHKCNQTCLDRIAAGTFNAGEYSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLA WTLMLAISVYMVSRDNYSCSICL		
ELISA/ Neutralization	B/Phuket/3073/2013	MKAIIVLLMVVTSNADRICTGITSSNSPHVVKTATQGEVNVTGVIPLTTTPTKSYFANLKGTRTRGKLCP DCLNCTDLDVALGRPMCVGTTPSAKASILHEVRPVTSGCFPIMHDRTKIRQLPNLLRGYEKIRLSTQNVI DAEKAPGGPYRLGTSGSCPNATSKIGFFATMAWAVPKDNYKNATNPLTVEVPYICTEGEDQITVWGFHSDN KTQMKSLYGDSNPQKFTSSANGVTTHYVSQIGDFPDQTEDGGLPQSGRIVVDYMMQKPGKTGTIVVQRG VLLPQKVWCASGRSKVIKGSLPLIGEADCLHEEVGGLNKSKPYYTGKHAKAIGNCPIWVKTPLKLANGTK YRPPAKLLKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGVAVAADLKSTQEAINKITKNLNSLSELE VKNLCRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPS AVDIGNGCFETKHKCNQTCLDRIAAGTFNAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLA VTLMLAIFIVYMVSRDNVSCSICL		
Challenge	B/Florida/04/2006	MKAIIVILMVVTSNADRICTGITSSNSPHVKTATQGEVNVTGVIPLTTTPTKSYFANLKGTRTRGKLCP DCLNCTDLDVALGRPMCVGTTPSAKASILHEVKPVTSGCEPIMDDTKIRQLPNLLRGYENIRLSTQNVI DAEKAPGGPYBLGTSGSCPNATSKSGFFATTMAWAVPKDNNKNATNPLTVEVPVICTEGEDQITVWGFHSD DKTQMKNLYGDSNPQKFTSSANGVTTHYVSQIGSFPDQTEDGGLPQSGRIVVDYMMQKPGKTGTIVVQRGVLLPQKVWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGTK YNLPQKKWCASGRSKVIKGSLPLIGEADCLHEKYGGLNKSKPYYTGEHAKAIGNCPIWVKTPLKLANGTK YNPPAKLIKERGFFGAIAGFLEGGWEGMIAGWHGYTSHGAHGVAVADLKSTQEAINKITKNLNSLSELE VKNLQRLSGAMDELHNEILELDEKVDDLRADTISSQIELAVLLSNEGIINSEDEHLLALERKLKKMLGPS AVEIGNGCFETKHKCNQTCLDRIAAGTFNAGEFSLPTFDSLNITAASLNDDGLDNHTILLYYSTAASSLA VTILMLAIFIVYMVSRDNVSCSICL		

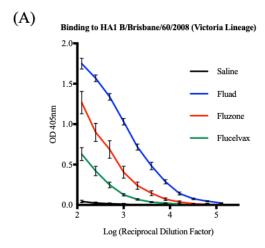
## Supplementary Fig. S1. Sequences of influenza B viral HA proteins used in this study.

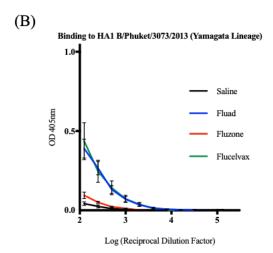
Sequences are for influenza B HA in commercial vaccines, and proteins/viruses used in ELISA and neutralization assays and challenge experiments. Sequences were downloaded from different databases such the NCBI GenBank. Influenza B HA of Victoria lineage is denoted as B-V, while Influenza B HA of Yamagata-lineage is denoted as B-Y.

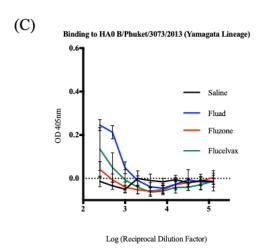
These are all the pairwise sequence identity percentages (from chimera), and also attached is a graphic of the multiple sequence alignment.

AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)] vs. AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)]: 100.00% identity AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)] vs. ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)]: 93.49% identity AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)] vs. ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)]: 92.97% identity AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)] vs. AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)]: 93.14% identity AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)] vs. ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)]: 92.97% identity ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)] vs. AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)]: 93.49% identity ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)] vs. ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)]: 100.00% identity ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)] vs. ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)]: 99.14% identity ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)] vs. AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)]: 99.14% identity ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)] vs. ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)]: 98.97% identity ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)] vs. AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)]: 92.97% identity ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)] vs. ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)]: 99.14% identity ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)] vs. ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)]: 100.00% identity ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)] vs. AQY15026.1 hemagglutinin [Influenza B virus (B/lowa/06/2017)]: 99.66% identity ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)] vs. ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)]: 99.49% identity AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)] vs. AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)]: 93.14% identity AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)] vs. ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)]: 99.14% identity AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)] vs. ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)]: 99.66% identity AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)] vs. AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)]: 100.00% identity AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)] vs. ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)]: 99.83% identity ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)] vs. AGM53847.1 hemagglutinin [Influenza B virus (B/Florida/04/2006)]: 92.97% identity ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)] vs. ANC28539.1 hemagglutinin [Influenza B virus (B/Brisbane/60/2008)]: 98.97% identity ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)] vs. ASW32353.1 hemagglutinin [Influenza B virus (B/Maryland/15/2016)]: 99.49% identity ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)] vs. AQY15026.1 hemagglutinin [Influenza B virus (B/Iowa/06/2017)]: 99.83% identity ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)] vs. ASK81305.1 hemagglutinin [Influenza B virus (B/Colorado/06/2017)]: 100.00% identity

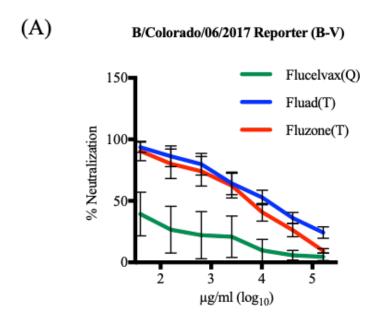
Supplementary Fig. S2. Sequence comparison and sequence accession numbers of influenza B viral HA proteins used in this study. Accession numbers and strain names are given with the percent identity at the end of each line for the sequence comparison.

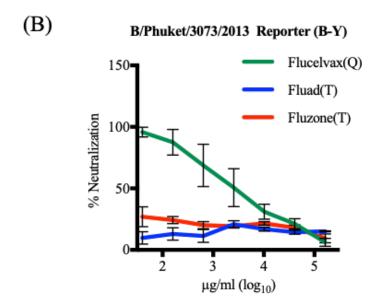






Supplementary Fig. S3. Dilution curves of the antibody binding from vaccinated mouse sera to influenza B HA1 proteins. Serially diluted sera from mice immunized with commercial vaccines Fluad (T), Flucelvax (Q), and Fluzone (T) were analyzed by ELISA for binding to antigenically matched and mismatch influenza HA1 proteins representing (A) B/Brisbane/60/08 (Victoria-lineage), (B) B/Phuket/307/13 (Yamagata lineage) viruses.





Supplementary Fig. S4. Titration curves for neutralization of influenza B reporter viruses by sera from mice immunized with commercial influenza vaccines. Reporter influenza B viruses were (A) B/Colorado/17 (B-V) and (B) B/Phuket/307/13 (B-Y). Influenza B HA of Victoria lineage is denoted as B-V, while Influenza B HA of Yamagata-lineage is denoted as B-Y.

(A) Histologic Score Bronchiole lumina Exudate in bronchiole lumina. None (score 0 points), Exudate present (score 1 point); moderate (score 2 points); Severe-partial to complete occlusion of 1 or more central bronchioles (score 3 points) Terminal bronchiole lumina Exudate in terminal bronchiolar lumina. None (score 0 points), mild (score 1 point), moderate (score 2 points), Severe-partial to complete occlusion of 1 or more central bronchioles (score 3 points) Bronchiolar/TB epithelial None (score 0 points); Mild (score 1 point); Moderate (score 2 points); necrosis/degeneration Severe (score 3 points) Perivascular inflammation/fibrin None (score 0 points); Mild (score 1 point); Moderate (score 2 points); Severe (score 3 points) Alveoli/interstitium edema None (score 0 points); <25% (score 1 point); 25-50% (score 2 points); >50% (score 3 points) Alveoli inflammatory cells None-few (score 0 points); mild increase (score 1 point); moderate (score 2 points); Marked (score 3 points) Interstitial inflammatory cells None-few (score 0 points); mild increase (score 1 point); moderate (score 2 points); Marked (score 3 points) Anywhere: hemorrhage/necrosis/fibrin None (score 0 points); up to 5% of section (score 1 point); 5-50% of

(B)	Influenza B Immunohistochemical Score			
(2)		None (score 0 points), Up to 5% (score 1 point), 5-50% (score 2 points), >50% (score 3 points)		
		None (score 0 points), Up to 5% (score 1 point), 5-50% (score 2 points), >50% (score 3 points)		
	Alveolar septa (with immunoreactive interstitial cells) immunoreactivity	None (score 0 points); up to 5% (score 1 point), 5-50% (score 2 points), >50 % (score 3 points)		
	Bronchiolar exudate	Immunostaining absent (score 0 points); Immunostaining present score 1 point)		
	Alveolar exudate (+/- immunoreactive alveolar macrophages)	Immunostaining absent (score 0 points); Immunostaining present score 1 point)		

section (score 2 points); >50% of section (score 3 points)

Supplementary Fig. S5. Scoring rubrics for pathology of lungs in mice challenged with influenza. (A) Histologic scoring. (B) Influenza B immunohistochemical scoring. Yamagata-lineage challenge virus was mouse-adapted B/Florida/04/2006.