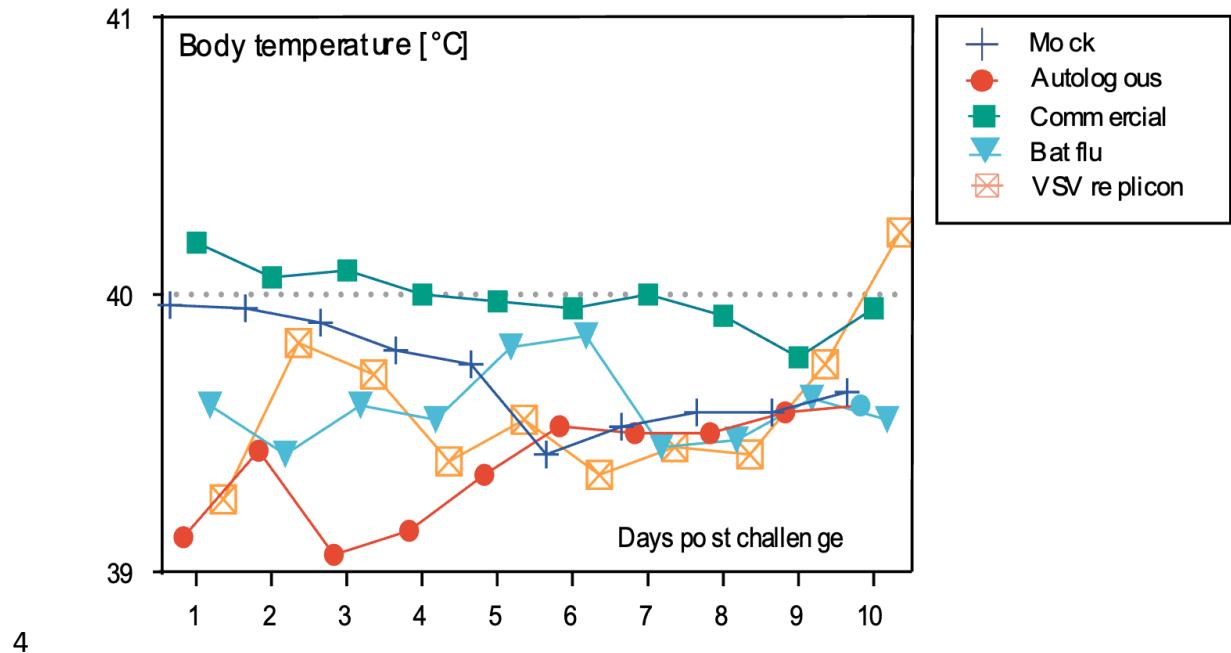
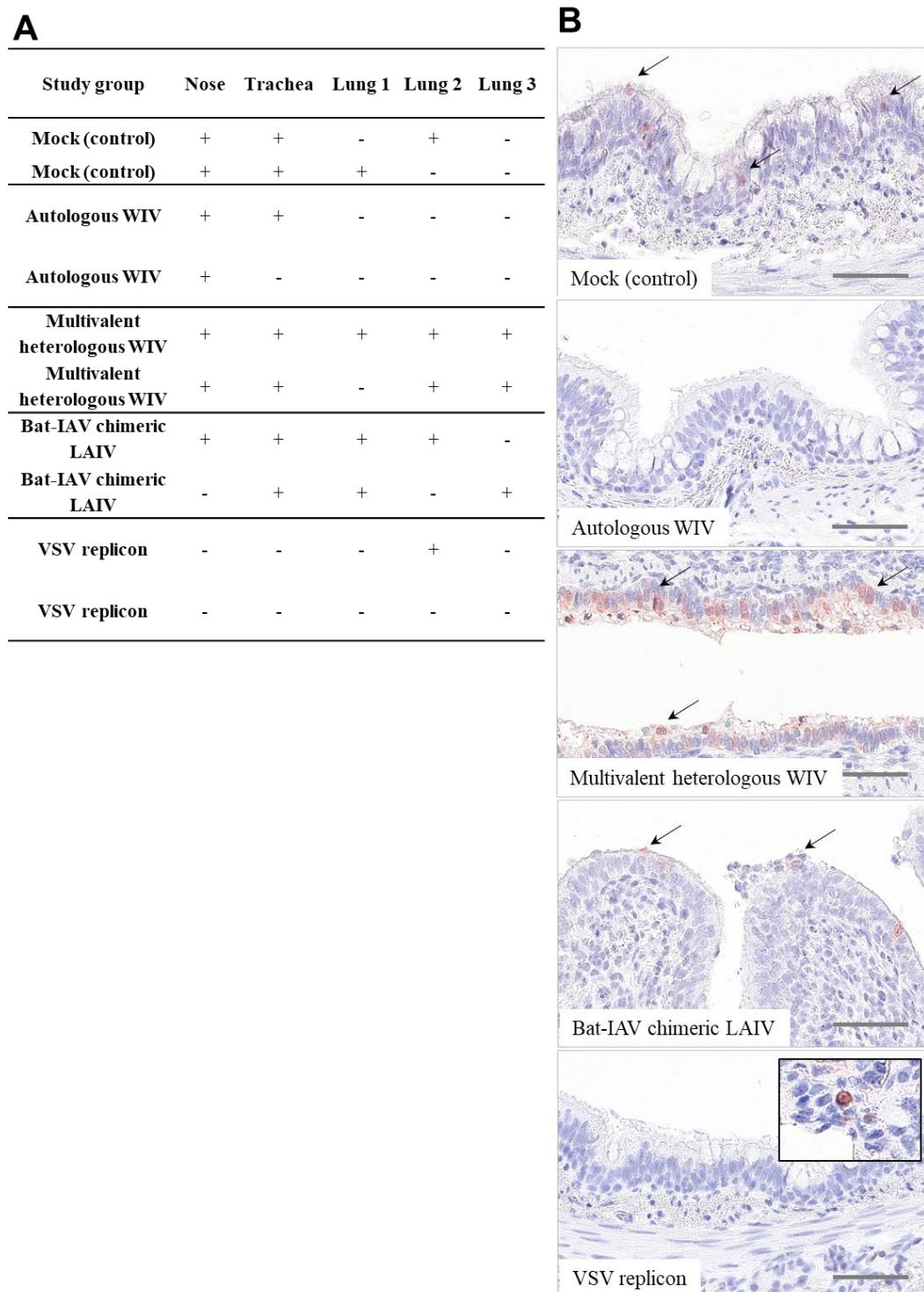


1 **Supplementary Figure 1.** Median rectal temperature of pigs challenged with swIAV isolate
2 H1pdmN2 AI03362 by contact to inoculated seeder animals. Dashed line shows the 40°C fever
3 threshold.



5 **Supplementary Figure 2.** Histopathological results of representative WIV and LAIV
6 vaccinated pigs after challenge infection with H1pdmN2. A) Immunohistochemical results of
7 the nose, trachea and lung of individual pigs selected based on positive viral genome detection.
8 (+) = positive, (-) = negative for viral matrix protein. B) Immunohistochemistry of the lung
9 illustrating IAV matrix-1 protein-positive bronchial respiratory epithelial cells (arrow) and foci
10 of alveolar cells (VSV-LAIV only, inlay) in the different groups. Scale bar = 50 µm.



12 **Supplementary Table 1.** Post mortem findings, 4 dpc, pulmonary atelectasis given in percent
13 affected area (%).

14

Study group	Status	pulmonary atelectasis (%) and affected lung lobe
Mock (control)	Seeder	<5 % cranial part of left cranial lobe, right middle lobe, right caudal lobe
	vaccinated contact	-
Autologous WIV	Seeder	-
	vaccinated contact	-
Multivalent heterologous WIV	Seeder	-
	vaccinated contact	-
Bat-IAV chimeric LAIV	Seeder	<5 % caudal part of left cranial lobe, right middle lobe
	vaccinated contact	-
	vaccinated contact	-
	vaccinated contact	-
	vaccinated contact	<5 % right middle lobe
VSV replicon	Seeder	30% right middle lobe
	vaccinated contact	-

15

16 **Supplementary Table 2.** Significance levels calculated by the Kruskal-Wallis test using the Dunn-Bonferoni correction for pairwise comparisons.
 17 Highly significant differences of vaccine groups versus the Mock group are indicated by asterisks in Figures 2 and 3 in the main text.

18

Vaccine groups	Lower respiratory tract (Fig. 3b)				Upper respiratory tract		Nasal swabs by day post challenge (Fig. 2b)					
	Trachea	Bronchial swab	Lung	Lymphnode	Conchae	Tonsil	2	3	4	5	6	7
Bat flu-Mock	0.0364	0.03138	0.002964	0.3095	0.0486	0.6113	0.1262	0.003633	0.001152	0.001886	0.002807	0.00207
Bat flu-WIV Comm	0.5108	0.1201	0.03092	0.4026	0.0486	0.3095	0.0004001	0.01311	0.02762	0.01018	0.01207	0.3858
Bat flu-VSV replicon	0.135	0.1884	0.01698	0.08297	0.8577	0.7423	0.9318	0.8979	0.5637	0.2562	0.232	0.05193
Bat flu-WIV auto	0.4026	0.5499	0.7723	0.4731	0.05583	0.5108	0.01708	0.0795	0.000728	0.03144	0.1693	0.08285
Mock-WIV Comm	0.1513	0.5499	0.3894	0.8577	1	0.6113	0.04436	0.6689	0.2947	0.5907	0.6326	0.02691
Mock-VSV replicon	0.0003345	0.0005254	1,12E-04	0.005958	0.03647	0.8577	0.1488	0.005435	0.007515	0.0486	0.073	0.2558
Mock-WIV auto	0.2093	0.1201	0.007177	0.08297	0.9049	0.2437	0.3922	0.2482	0.8979	0.339	0.1066	0.1784
WIV Comm-VSV replicon	0.03138	0.00411	0.000005484	0.01015	0.0486	0.4918	0.0005513	0.01866	0.1041	0.1515	0.1886	0.2817
WIV Comm-WIV auto	0.8577	0.3388	0.06165	0.1201	0.9049	0.09414	0.2481	0.4672	0.2395	0.6757	0.2562	0.3858
VSV replicon-WIV auto	0.01972	0.05574	0.007438	0.3095	0.03647	0.3239	0.02148	0.1041	0.005088	0.3097	0.8577	0.8342

Significant
(p=0,049-
0,01)

highly significant
(p<0,01)

19

20 **Supplementary Table 3.** Detection of swIAV in nasal swabs of piglets in the selected pig herd
21 by RT-qPCR at different weeks of age. A herd with at least one positive sample ($cq \leq 30$) is
22 marked in bold.

	<i>age (week(s))</i>	<i>herd tested positive (cq ≤ 30)</i>
<i>suckling piglets</i>	1	-
	2	-
	3	-
<i>weaned piglets</i>	4	yes
	5	yes
	6	yes
	7	-
	8	yes
	9	-
	10	-