S3 Table. Summary of the results from various assays in this study.

al No.	Type of mutation	Gene (No. of mutations)	Mutation	Inter-segment combination with mutation	In vitro polymerase activity in: †					In vitro virus growth in: <sup>‡</sup>			In vivo virus titers in: *		
ai NO.					vRNA at 37 °C	293T cells tra vRNA at 33 °C	ensfected with cRNA at 37 °C	cRNA at 33 °C	QT-6 cells tra	ensfected with cRNA at 37 °C	SAE 37°C	cells at 33°C	CEF cells at 37°C	Mouse 3 d post-infection	lungs at 6 d post-infectio
1			K80R		=	=	=	+	=	=	ND*	ND	ND	ND	ND
2	Single mutations		T129N D195N		_	_ _	_		_ =	_	ND ND	ND ND	ND ND	ND ND	ND ND
4			Q257L		_	<del>-</del>	<del>-</del>	<del>-</del>	_	_	ND ND	ND ND	ND ND	ND ND	ND ND
5			1292M		_	_	_	_	=	=	ND	ND	ND	ND	ND
6			R299K R389K	Inter-5 Inter-8	= _	=	= _	=	= _	=	=	_	_	ND ND	ND ND
8		PB2 (14)	N456D	inter 0		=	+	+	=	=	ND	ND	ND	ND	ND
9			I463M	Inter-3	_	=	_	=	_	=	=	_ ND		ND	ND
10 11			M473V M570I		= =	=	=	=	= =	=	ND ND	ND ND	ND ND	ND ND	ND ND
12			K617E		_	_	_	_	_	-	ND	ND	ND	ND	ND
13			T637I T683I		_	_	_ _	_ _	_	_	ND ND	ND ND	ND ND	ND ND	ND ND
15			K54R		=	=	+	+	=	=	ND	ND	ND ND	ND	ND
16			D175N			=	_	=   =	_ _	-	ND	ND	ND =	ND ND	ND ND
1 <i>7</i> 18			T182I K198R	Inter-3,4,5	=   +	_ =	++	++	=	+	+	++	+	ND +	ND ++
19			K214R		+	=	=	+	=	=	ND	ND	ND	ND	ND
20		PB1 (13)	A374T L384S		_	=	_ +	=	_ _	_	ND ND	ND ND	ND ND	ND ND	ND ND
22		1 51 (10)	K386R		=	=	+	=	_	<u> </u>	ND	ND	ND	ND	ND
23			M616R		_	_	_	_	_	_	ND ND	ND ND	ND	ND ND	ND ND
24 25			E618D P627L		=	+++	=   =	= +++	_	=	ND =	ND ++	ND =	ND ND	ND ND
26			N642K			<del>-</del>	_	<del>-</del>	_	-	ND	ND	ND	ND	ND
27 28			P756S D55N		<u> </u>	<u> </u>	<u> </u>	<u> </u>	_ _	=	ND ND	ND ND	ND ND	ND ND	ND ND
29		PA (17)	N321S		_	=	=	=	=	=	ND	ND	ND	ND	ND
30			N321I	Inter-6	=	=	=	=	=	=	_	=	_ =	ND ND	ND ND
32			E327K V341L	Inter-3 Inter-4	=	=	=	++	=	=	_	_	=	ND ND	ND ND
33			E382D		-	=	_	=	_	- 1	ND	ND	ND	ND	ND
34			S400T T608S		=	=	=	=	_	=	ND ND	ND ND	ND ND	ND ND	ND ND
36			K615R		_	_	_	=	_	=	ND	ND	ND	ND	ND
37			E629H		_	_	<u>-</u>	_	_	_	ND ND	ND	ND	ND	ND ND
38 39			G631C L641P		_	_	= -	_	= -	=	ND ND	ND ND	ND ND	ND ND	ND ND
40			S644F		_	_	_	_	_	-	ND	ND	ND	ND	ND
41			L649I A669V		=	+	= +	<b>=</b>	=	=	ND ND	ND ND	ND ND	ND ND	ND ND
43			F681L		_	_	_	_	_	_	ND	ND ND	ND ND	ND	ND ND
44			K716N	Inton 4 0 4 5 0 0	=	<u>=</u>	<u>=</u>	<b>=</b>	_	=	ND —	ND _	ND —	ND	ND ND
45 46		NP (2)	G287S T371M	Inter-1,3,4,5,6,8		=	= =	=	= =	=	= ND	= ND	= ND	ND ND	ND ND
47			D195N/R299K		=	+	+	+	=	=	ND	ND	ND	ND	ND
48		PB2(8)	D195N/R299K/N456D	Inter-2,7	<u>+</u>	+	++	++	=	=	=	=	=	ND ND	ND ND
50			M473V/D195N M473V/D195N/T683I	inter-2,1	_	_	_ _	=	_	=	T ND	T ND	ND	ND ND	ND ND
51			M473V/N456D		=	=	+	+	=	=	ND	ND	ND	ND	ND
52 53			M570I/K80R M570I/K80R/T129N	Inter-1,4,6	+ 	=	+ -	+ -	= -	=	= ND	= ND	= ND	ND ND	ND ND
54			M570I/K80R/T129N/I292M		_	_	_	_	_	-	ND	ND	ND	ND	ND
55 56 57			T182I/K214R T182I/K214R/L384S	Inter-2,7 Inter-6,8	+	_ =	++	+	=	=	=	= =	= =	ND ND	ND ND
			K198R/D175N	Inter-1	+	=	+++	++		=	++	++	+	+++	+++
58	Intra-segment combinations		A374T/K54R		<u>-</u>	_ +++	<del>-</del>	=	_	-	ND	ND	ND ND	ND	ND
60			P756S/P627L P756S/P627L/E618D		<del>+</del>   +	+++	+	+++	= =	=	ND ND	ND ND	ND ND	ND ND	ND ND
61			P756S/P627L/E618D/K386R	–	++	+++	++	++	=	=	_	_	=	ND	ND
62 63			N321S/K716N N321S/K716N/S400T	Inter-7 Inter-8	= -	= -	_ _	<b>=</b> <b>-</b>	_ _	_	– ND	– ND	= ND	ND ND	ND ND
64			N321S/K716N/S400T/K615R	inter 0	-	_	_	_	_	-	ND	ND	ND	ND	ND
65 66		PA(7)	G631C/E629H G631C/E629H/A669V		_	_	_	_	_	_	ND ND	ND ND	ND ND	ND ND	ND ND
67			G631C/E629H/A669V A669V/L649I	Inter-2	=	=	_ +	=	=	_	ND —	— Н	ND —	ND	ND ND
68		NID(A)	A669V/E382D	Inter-5		<u>=</u>	<u>=</u>	<u>=</u>	_			<u>_</u>	_	ND	ND ND
69 70		NP(1)	G287S/T371M Inter-1	Inter-2,7	= +	<u>=</u> +	= +++	<u>=</u> +++	= +	= +	++	+++	+	ND ND	ND ND
71	Inter-segment combination of mutations	Intersegment(16)	Inter-2		<u>+</u>	÷	+++	+++	<u> </u>	Ė	++	+++	=	+++	++
72			Inter-3		+ +	+	+++	+++	+ +	=	=	++	=	ND	ND
74			Inter-4 Inter-5		+	++	+++ +++	+++	<del>+</del>   +	+	++	+++	= =	+++ ND	+++ ND
75			Inter-6		+	+	+++	+++	++	+	_	_	=	ND	ND
76 77			Inter-7 Inter-8		<del>+</del>   +	= +	++	+++	++	=	_ _	=	= =	ND ND	ND ND
78			Inter-6			<u>-</u>	=	_ _	_	<u> </u>	ND	ND	ND	ND	ND
79			Inter-10		<u> </u>	=	++	++	<u>+</u>	=	ND	ND	ND	ND	ND ND
80 81			Inter-11 Inter-12		+	=	++	++	+ -	+	ND ND	ND ND	ND ND	ND ND	ND ND
<b>B2</b>			Inter-13		+	+	++	++	=	=	ND	ND	ND	ND	ND
83			Inter-14 Inter-15			_	+	=		=	ND ND	ND ND	ND ND	ND ND	ND ND
K /I			mer-15		_	_	_		<u> </u>		שוו	IND	עוו	שוו	IND

<sup>†</sup>Relative comparison with wild-type polymerase activity: +, 1.5-3-fold increase; ++, 3-5-fold increase; ++, >5-fold increase; =, similar to wild-type virus (<1.5-fold change); -, lower than wild-type.

<sup>&</sup>lt;sup>‡</sup>Relative comparison with wild-type virus yield: +, 2-5-fold increase; ++, >5-10-fold increase; +++, >10-fold increase; =, similar to wild-type virus (<2-fold change); -, lower that wild-type.

<sup>\*</sup>Relative comparison with wild-type virus titer: +, 2-10-fold increase; ++, >10-100-fold increase; +++, >100-fold increase; =, similar to wild-type virus (<2-fold change); -, lower that wild-type.

<sup>\*</sup>ND: not determined.