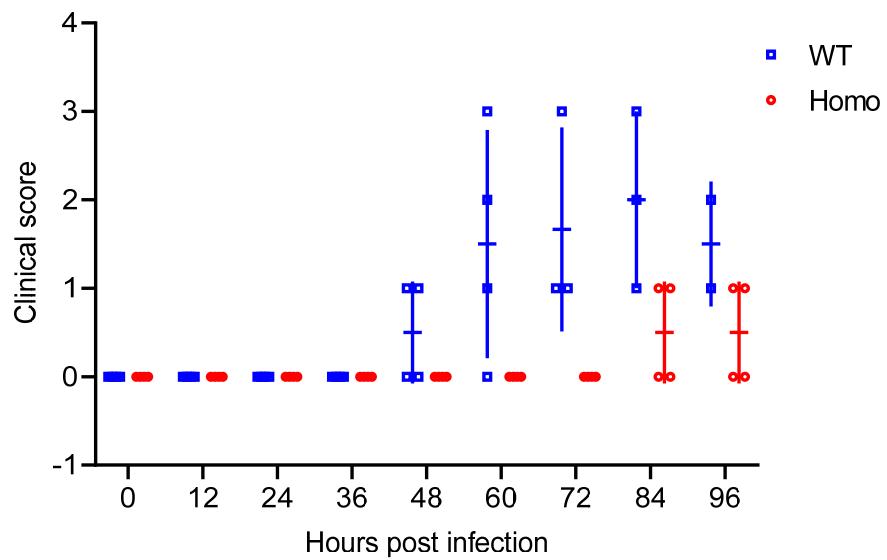


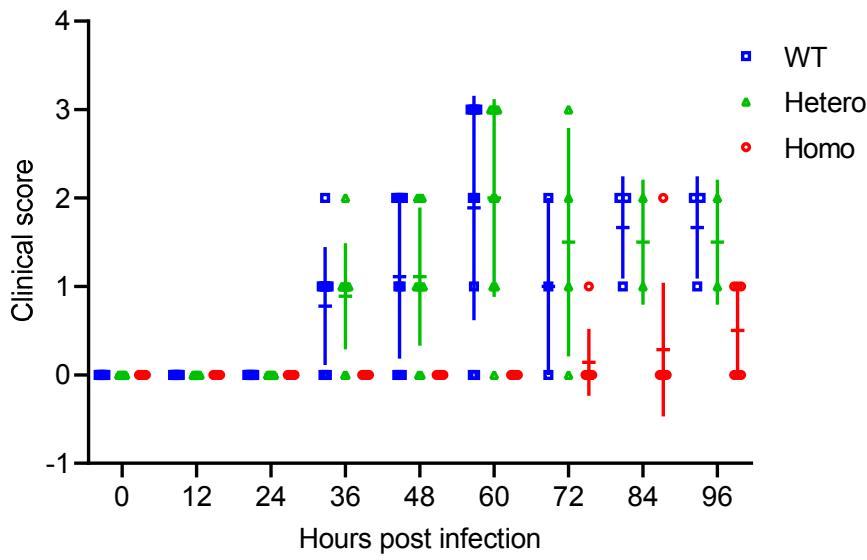
## Supplementary figures

Pig	1	MARMGLAGAAGRWWGLALGLTAFFLPGAHTQVVQVNDSMYGFIGTDVVLHCSFANPLPGV	60
Cattle	1	MARMGLAGAAGRWWGLALGLTAFFLPGAQAQMVKVNDSMYGFIGTDVVLHCSFANPLPGV	60
Human	1	MARMGLAGAAGRWWGLALGLTAFFLPGVHSQVVQVNDSMYGFIGTDVVLHCSFANPLPSV	60
Mouse	1	MARMGLAGAAGRWWGLALGLTAFFLPGTHTQVVQVNDSMYGFIGTDVVLHCSFANPLPSV	60
		*****.:::*****.*****.*	
Pig	61	KITQVTWQKATNGSKQNVAIYNPAMGVSVLAPYRERVEFLRPSFTDGTIRLSRLEDEG	120
Cattle	61	KITQVTWQKATNGSKQNVAIYNPAMGVSVLAPYRERVEFLRPSFTDGTIRLSRLEDEG	120
Human	61	KITQVTWQKSTNGSKQNVAIYNPSMGVSVLAPYRERVEFLRPSFTDGTIRLSRLEDEG	120
Mouse	61	KITQVTWQKASNGSKQNMAYNPTMGVSVLPPYEKRVEFLRPSFIDGTIRLSGLEDEG	120
		*****;:*****;*****;*****.**.;***** * *****	
Pig	121	VYICEFATFPAGNRESQLNLTVMAKPTNWIEGTQAVLRAKKGKDDKVLVATCTSANGKPP	180
Cattle	121	VYICEFATFPAGNRESQLNLTVMAKPTNWIEGTHAVLRARKGQDEKVLVATCTSANGKPP	180
Human	121	VYICEFATFPAGNRESQLNLTVMAKPTNWIEGTQAVLRAKKGQDDKVLVATCTSANGKPP	180
Mouse	121	MYICEFATFPAGNRESQLNLTVMAKPTNWIEGTRAVLRARKGQDDKVLVATCTSANGKPP	180
		:*****;*****;*****;*****;*****;*:*****	
Pig	181	SVVSWETHLKGEAEYQEIRNPNGTVTVisRYRLVPSREDHRQSLACIVNYHMDRFRESLT	240
Cattle	181	SVVSWETRLKGEAEYQEIRNPNGTVTVisRYRLVPSREAHRQSLACIVNYHMDRFWESLT	240
Human	181	SVVSWETRLKGEAEYQEIRNPNGTVTVisRYRLVPSREAHQQSLACIVNYHMDRFKESLT	240
Mouse	181	SAVSWETRLKGEAEYQEIRNPNGTVTVisRYRLVPSREAHRQSLACIVNYHLDRFRESLT	240
		*.*****:***** * :*****:*** ***	

**Figure S1.** Conservation of Nectin1 F129 residues across different mammalian species. Sequences were aligned with CLUSTALW. F129 residues are shown in red letters. Identical residues in all aligned sequences are denoted by asterisks.



**Figure S2.** Clinical score for WT and homozygous (Homo) nectin1 F129A mutant mice following PRV infection in experiment 1.



**Figure S3.** Clinical score for WT, heterozygous (Hetero), and homozygous (Homo) nectin1 F129A mutant mice following PRV infection in experiment 2.