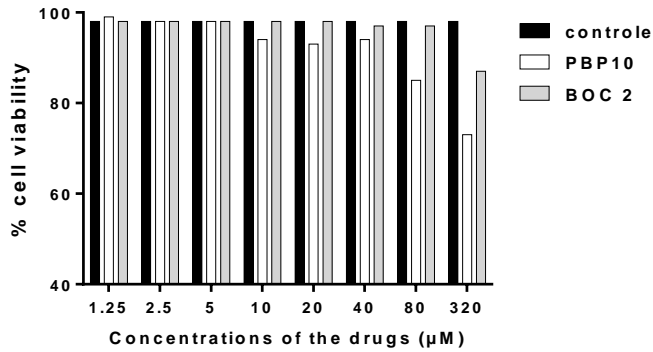
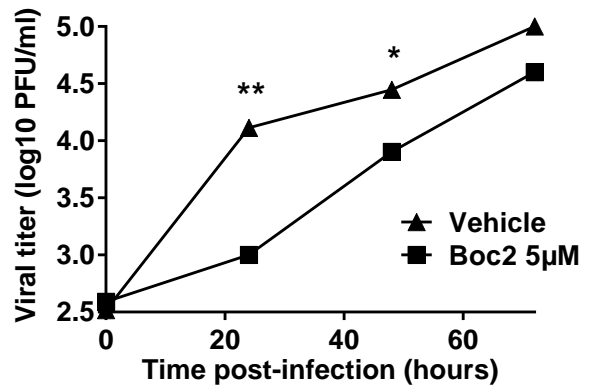
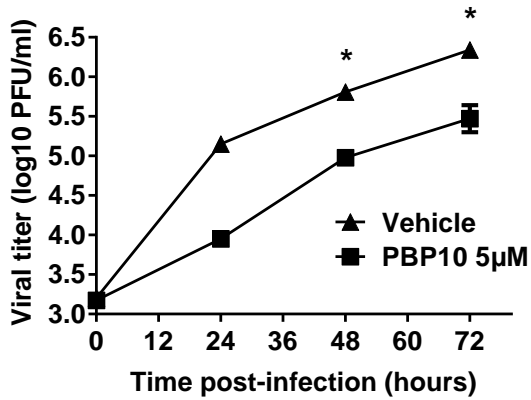


Fig 1

A



B



C

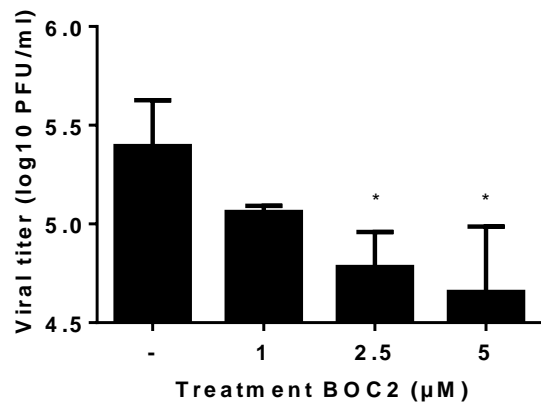
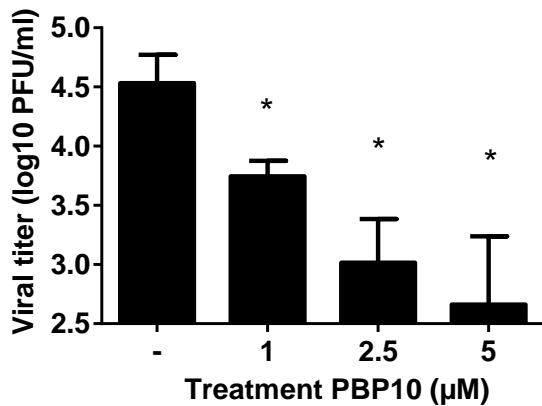
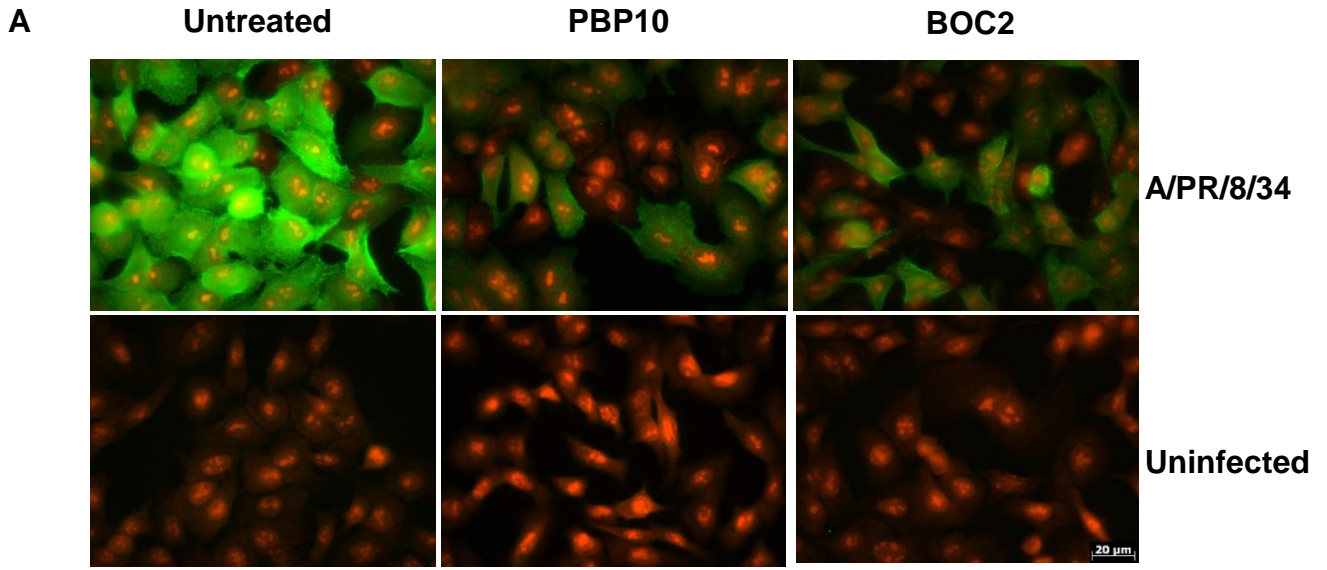


Fig 2



B

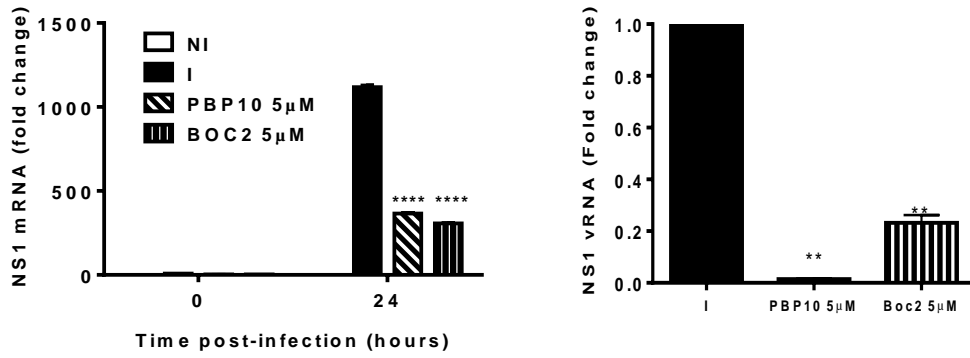


Fig 3

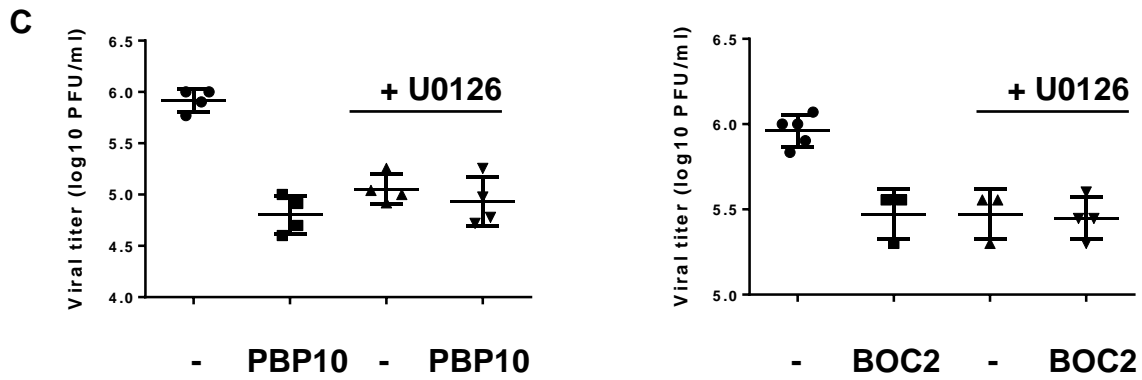
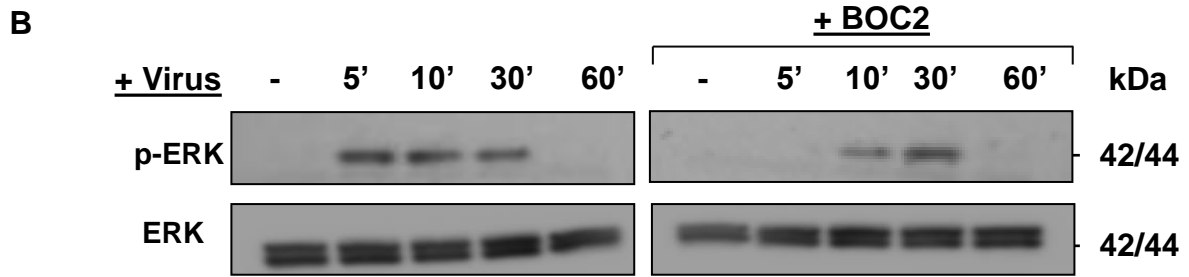
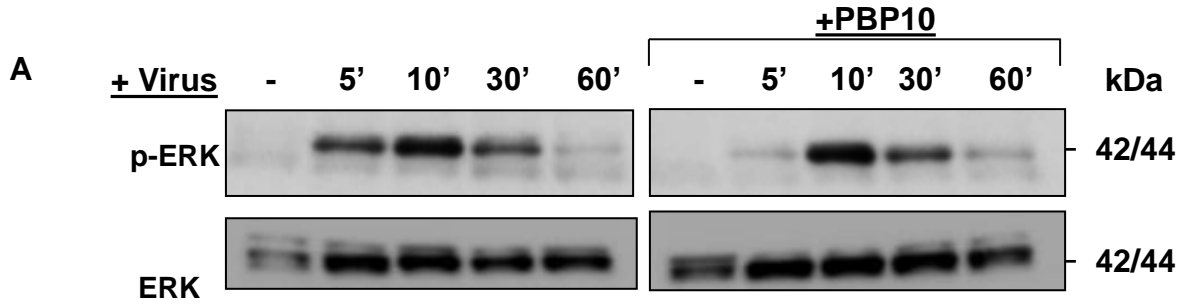


Fig 4

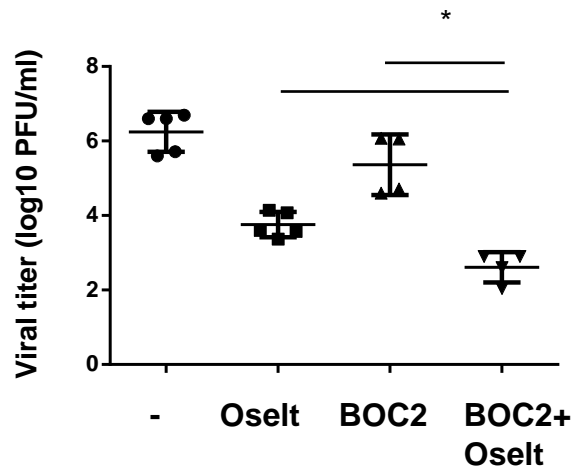
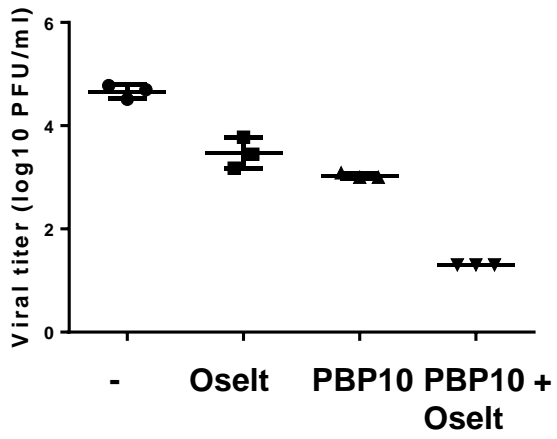


Fig 5

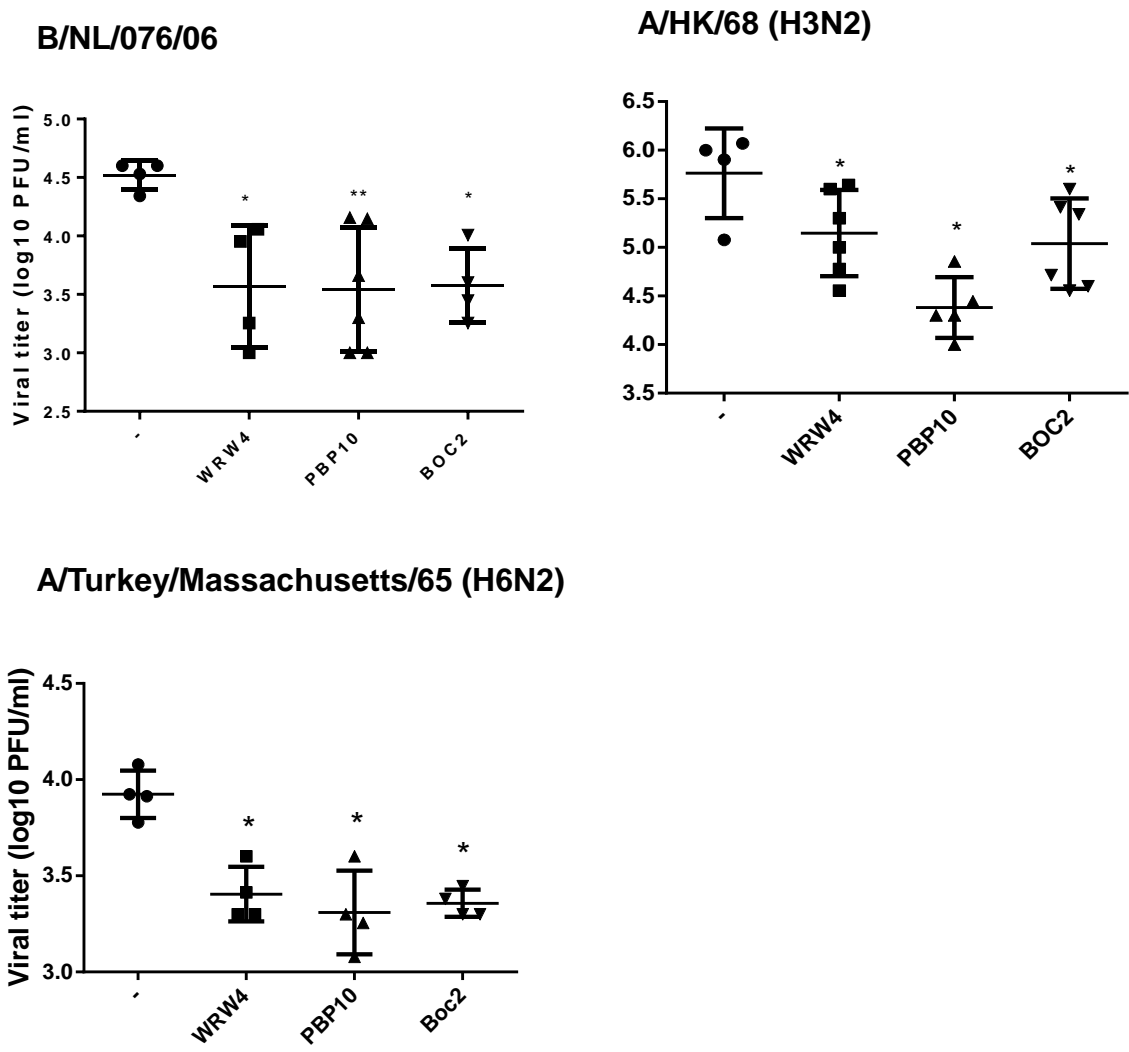
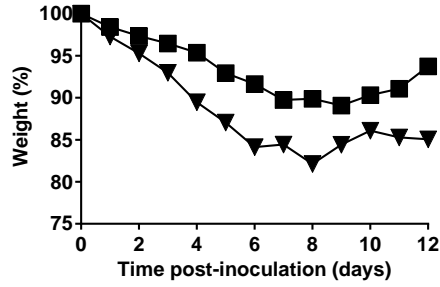
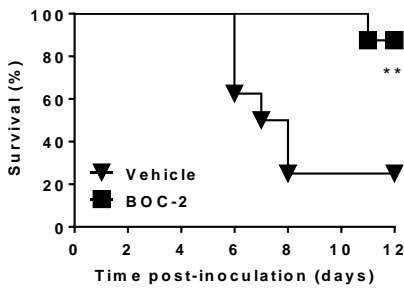


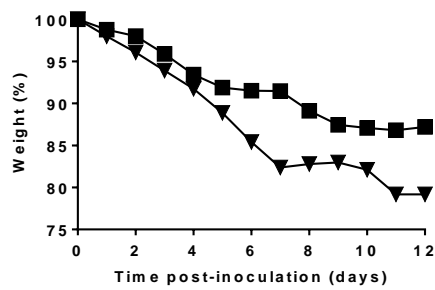
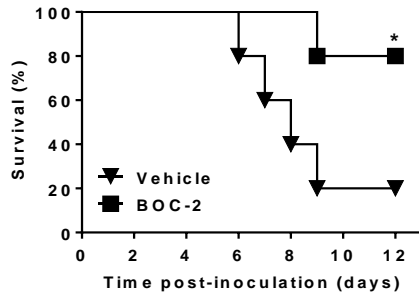
Fig 6

A

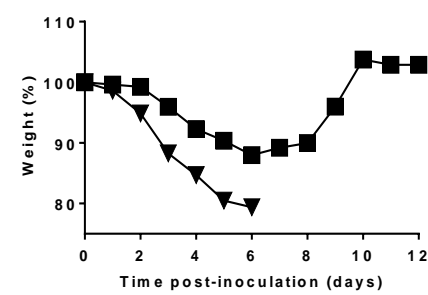
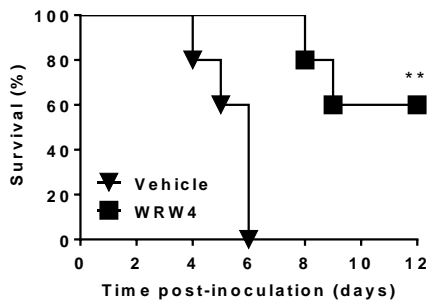


B

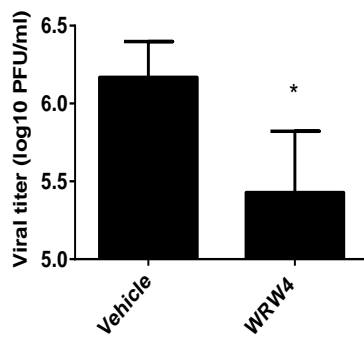
Prophylactic treatment



C



D



E

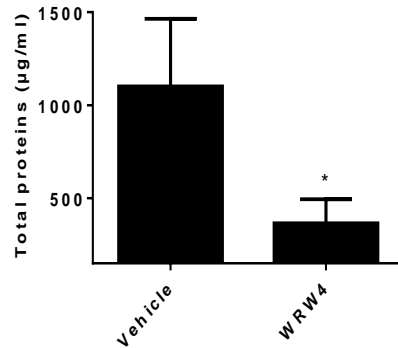


Fig 7

FPR2 protein BLAST

Query = Homo sapiens
Sbjct = Mus musculus

		Identities 267/350(76%)	Positives 300/350(85%)	Gaps 0/350(0%)	
Query	1	METNFS	TPLNEYEEVSYESAGY	TVLRILPLVVLGVTFV	LVGLGNGLVIWVAGFRMTRTVT 60
Sbjct	1	ME+N+S LN E V Y+S	VL IL +VV+ +TF	LGVLGNGLVIWVAGFRM TVT	
Query	61	TICYLNLALADFSFTATLPFLIVSMAMGEKWPFGWFLCKLIHIVVDINLFGSVFLIGFIA	TI YLNLALADFSFTATLPFL+V MAM EKWPFGWFLCKL+HIVVD+NLFGSVFLI IA		120
Sbjct	61	TIWYLNALADFSFTATLPFLLVEMAMKEKWPFGWFLCKLVHIVVDVNLFGSVFLIALIA			120
Query	121	LDRICICVLHPVWAQNHR	TVSLAMKVIVGPWILALVLTLPVFLFLLT	TVTIPNGD	TYCTFNF 180
Sbjct	121	LDRICICVLHPVWAQNHR	TVSLA KV+VGPWI AL+LTLF+P+FLTTV IP GD YCTFNF		180
Query	181	ASWGGTPEERLKVAITMLTARGIIRFVIGFSLPMSIVAICYGLIAAKIHKKGMIKSSRPL	SW T EE+L AIT +T RGIIRF+IGFS+PMSIVA+CYGLIA KI+++ ++ SSRPL		240
Sbjct	181	GSWAQDDEEKLN	TAITFVTTTRGIIRFLIGFSMPMSIVAVCYGLIAVKINRRNLVNSSRPL		240
Query	241	RVLTAVVASFFICWFPFQLVALLGTVWLKEMLFYGYKIIDILVNPTSSLAFFNSCLNPM	RVLTAVVASFFICWFPFQLVALLGTVW KE L G YKI+D+ VNPTSSLA+FNSCLNPM		300
Sbjct	241	RVLTAVVASFFICWFPFQLVALLGTVWFKETLLSGSYKILDMFVNPTSSLAYFNSCLNPM			300
Query	301	LYVFGQDFRERLIHSLPTS	SLERALSEDSAPTNDTAANSASPPAETELQA 350		
Sbjct	301	LYVFMGQDFRERFIHSLPYS	LERALSEDSGQTSDSSTSSPADI	ELKA 350	

Sbjct = Macaca mulatta

		Identities 336/348(97%)	Positives 343/348(98%)	Gaps 0/348(0%)	
Query	4	NFSTPLNEYEEVSYESAGYTVLRILPLVVLGVTFV	LVGLGNGLVIWVAGFRMTRTVTTIC 63		
Sbjct	1	NFSTPL+EYEEVSYESAGYTVL+ILPLVVLGVTFV	LVGLGNGLVIWVAGFRMTRTVTTIC 60		
Query	64	YLNALADFSFTATLPFLIVSMAMGEKWPFGWFLCKLIHIVVDINLFGSVFLIGFIALDR	YLNALADFSFTATLPFLIVSMAMGEKWPFGWFLCKLIHIVVDINLFGSVFLIGFIALDR		123
Sbjct	61	YLNALADFSFTATLPFLIVSMAMGEKWPFGWFLCKLIHIVVDINLFGSVFLIGFIALDR			120
Query	124	CICVLHPVWAQNHR	TVSLAMKVIVGPWILALVLTLPVFLFLLT	TVTIPNGD	TYCTFNFASW 183
Sbjct	121	CICVLHPVWAQNHR	TVSLAMKVIVGPWILALVLTLPVFLFLLT	TVTIPNGD	TYCTFNFASW 180
Query	184	GGTPEERLKVAITMLTARGIIRFVIGFSLPMSIVAICYGLIAAKIHKKGMIKSSRPLRVL	GGTPE+RLKVAITMLTARGIIRFVIGFS+PMSIVA CYGLIAAKIHKKGMIKSSRPLRVL		243
Sbjct	181	GGTPEKRLKVAITMLTARGIIRFVIGFSMPMSIVATCYGLIAAKIHKKGMIKSSRPLRVL			240
Query	244	TAVVASFFICWFPFQLVALLGTVWLKEMLFYGYKIIDILVNPTSSLAFFNSCLNPMPLYV	TAVVASFFICWFPFQLVALL TVWLKE+L GYKII+ILVNPTSSLAFFNSCLNPMPLYV		303
Sbjct	241	TAVVASFFICWFPFQLVALLSTVWLKEILVDGKYKIIINILVNPTSSLAFFNSCLNPMPLYV			300
Query	304	FVGQDFRERLIHSLPTS	SLERALSEDSAPTNDTAANSASPPAETELQAM 351		
Sbjct	301	FVGQDFRERLIHSLPTS	LERALSEDSAPTNDTAA+ ASPPAETELQAM 348		