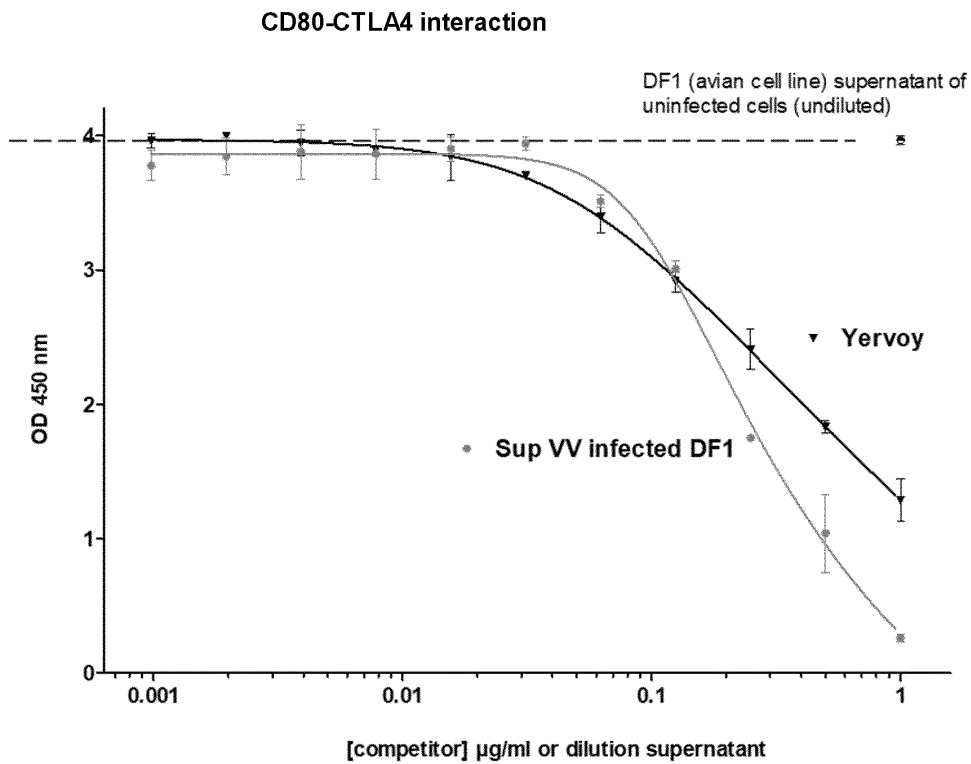
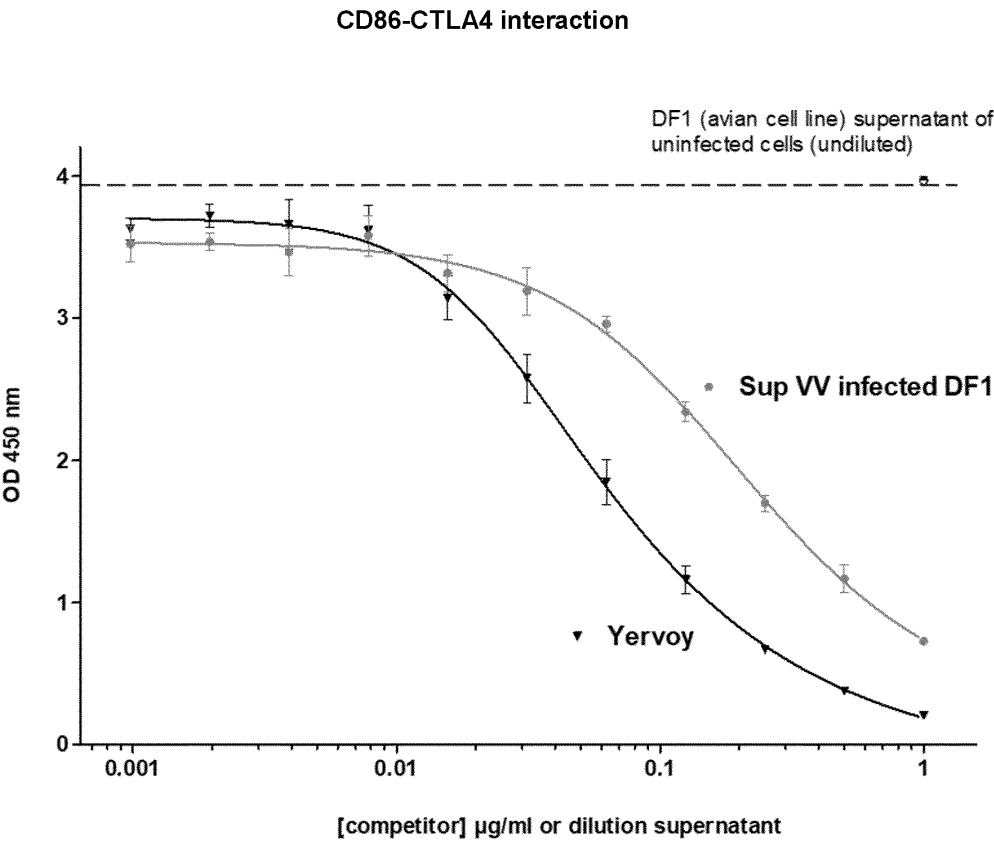


Fig.1A



2/15
Figure 1

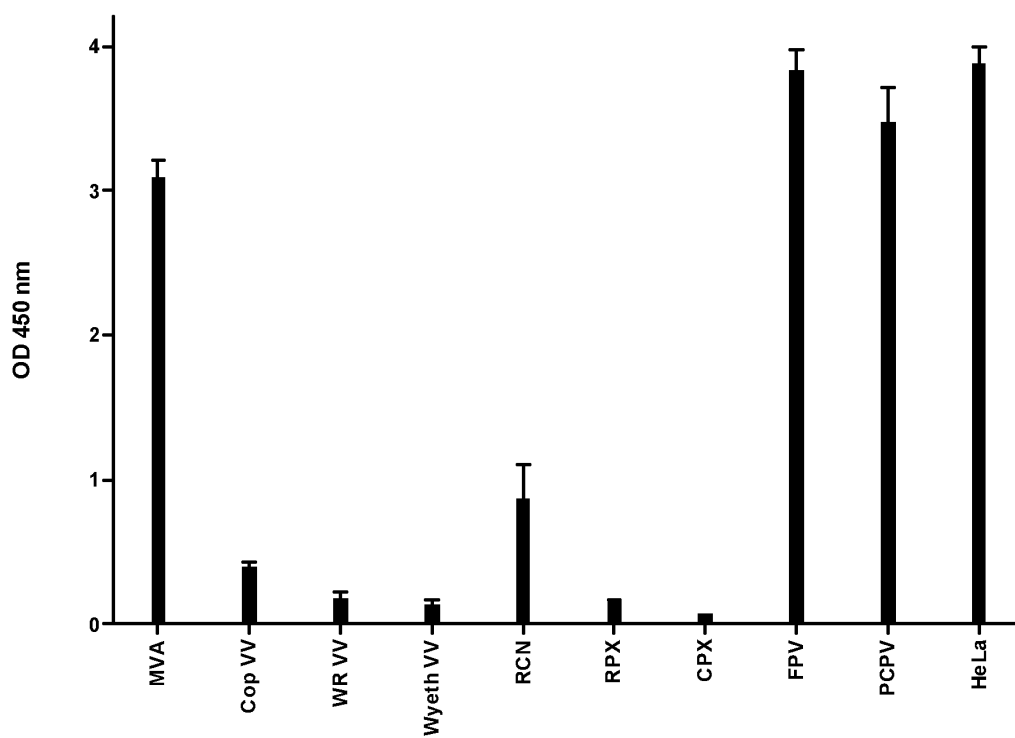
Fig. 1B



3/15

Figure 2

competition CD80/CTLA4



4/15

Figure 3

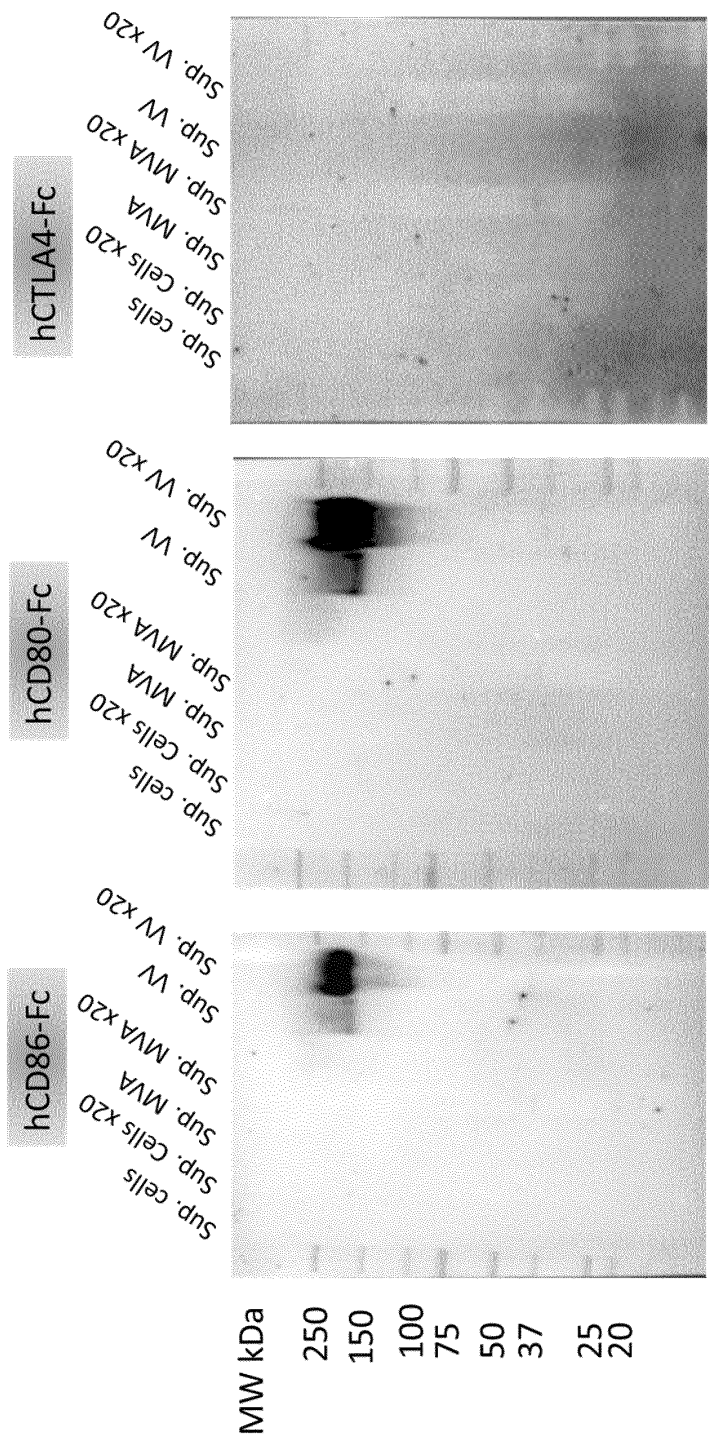
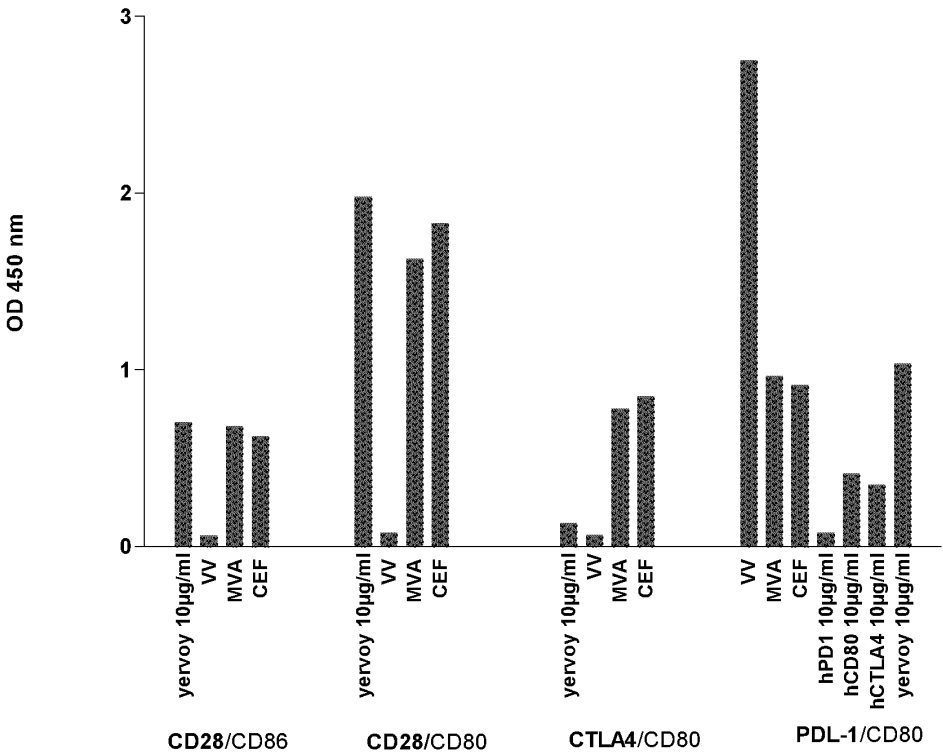


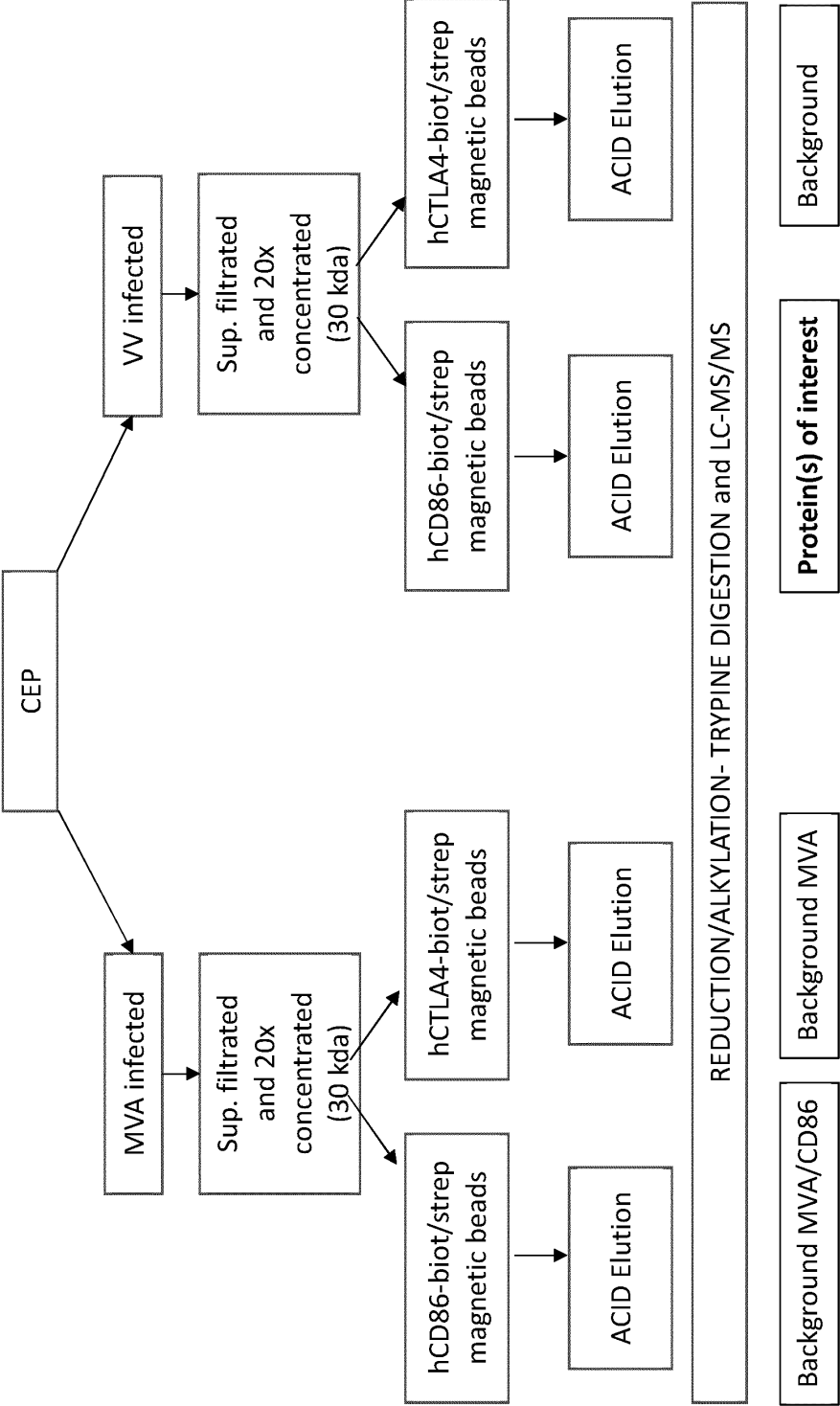
Figure 4



6/15

Figure 5

Fig. 5A



7/15

Figure 5

Fig. 5B

Score: 1741
Nominal mass (M_r): 25091
Calculated pI: 5.11
MS data file: D:\Partage\MGF_files\Q-Exactive\QE20317MS.mgf
Enzyme: Trypsin: cuts C-term side of KR unless next residue is P.
Variable modifications: Carbamidomethyl (C), Oxidation (C)
Protein sequence coverage: 75%
Matched peptides shown in **bold** .

1	MVYKLVLLFC	IASLGYSVEY	KNTICPPRQD	YRYWYFAAEL	TIGVNYDINS
51	TIIGECHMSE	SYIDR NANIV	LTGYGLEINM	TIMDTDQRFV	AAAEVGVKDN
101	KLSVLLFTTQ	RLDKVHHNIS	VTITCMEMNC	GTTYDSDLP	ESIAKSSSCD
151	ITINGSCVTC	VNLETDPTKI	NPHYLHPKDK	YLYHNSEYGM	RGSYGVTFID
201	ELNQCLLDIK	ELSYDICYRE			

Figure 6

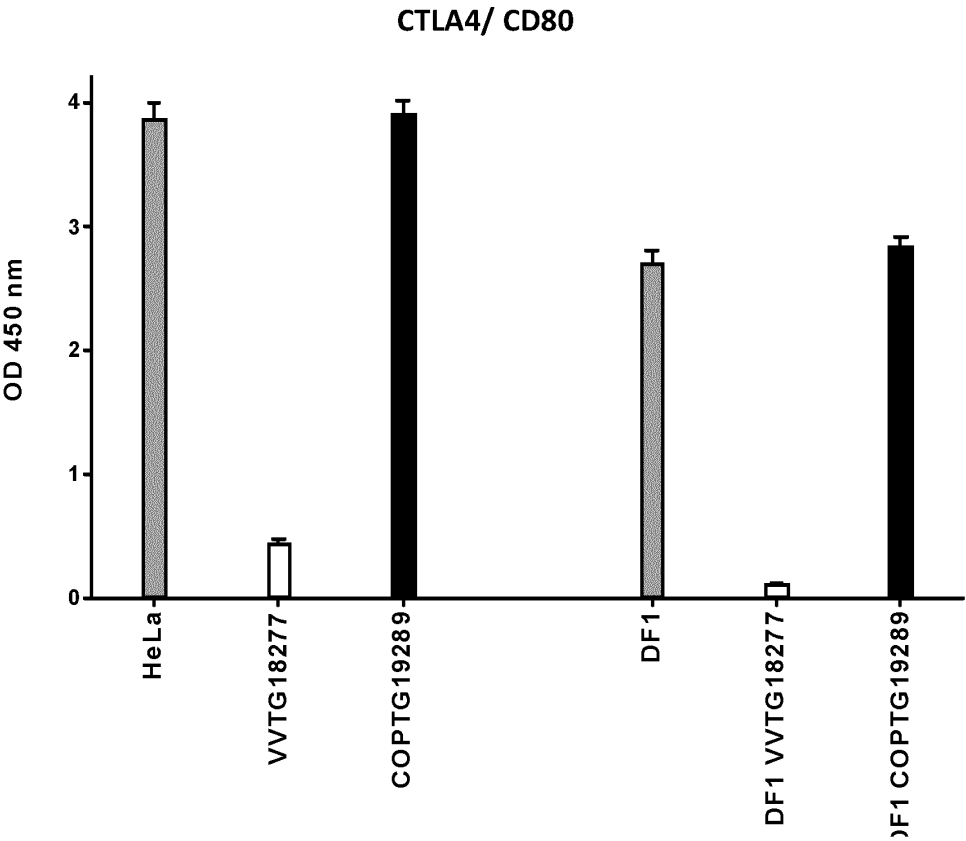
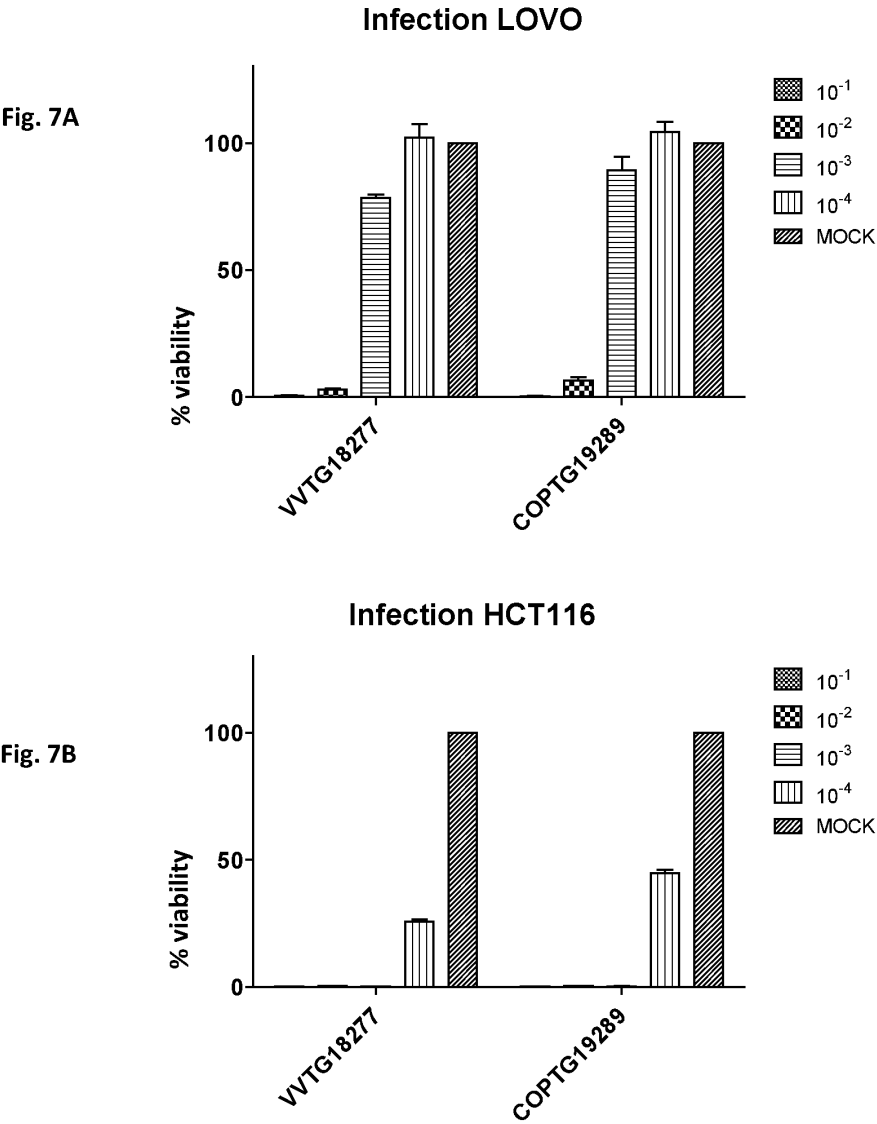
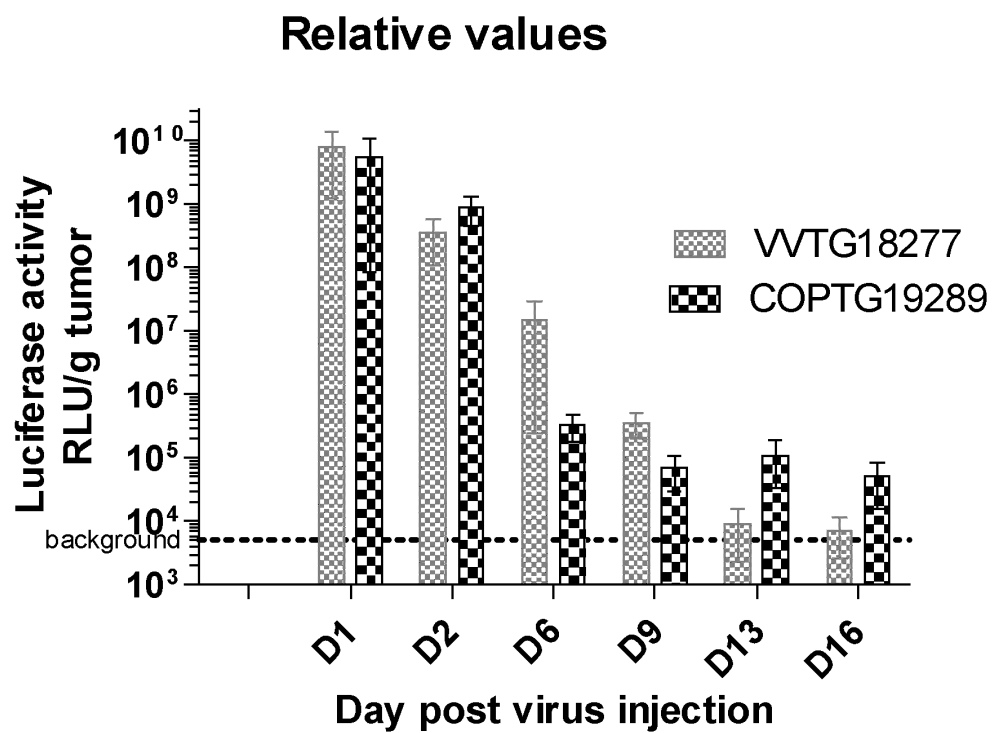


Figure 7



10/15

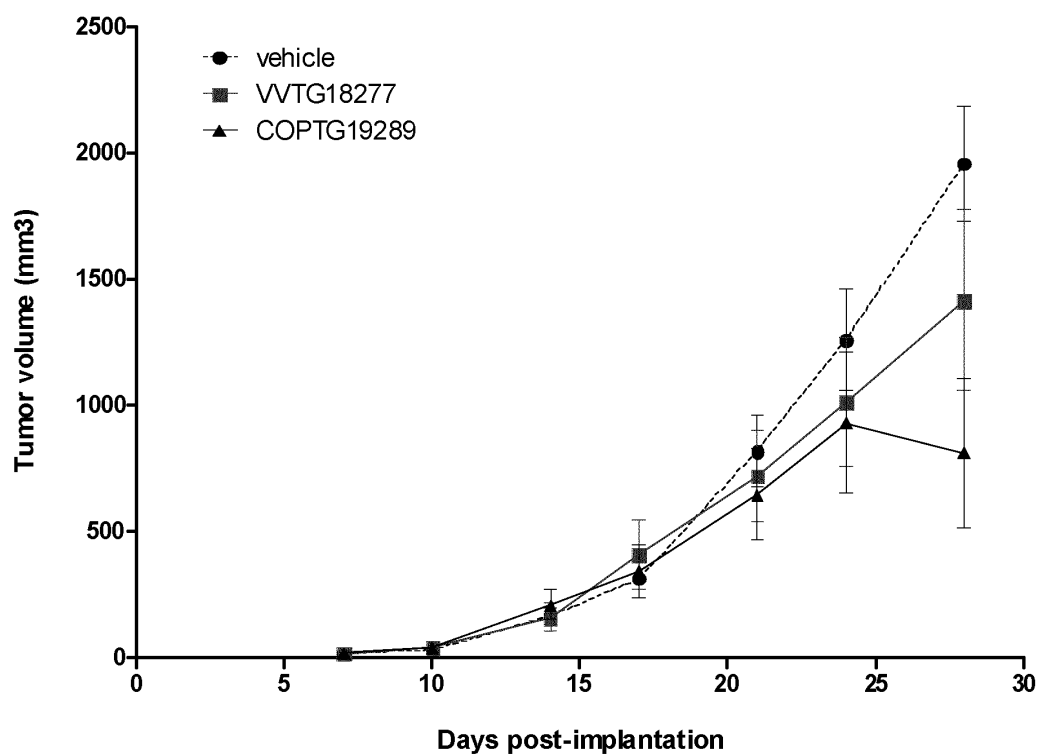
Figure 8



11/15

Figure 9

CT26 mean



12/15

Figure 10

Fig. 10A

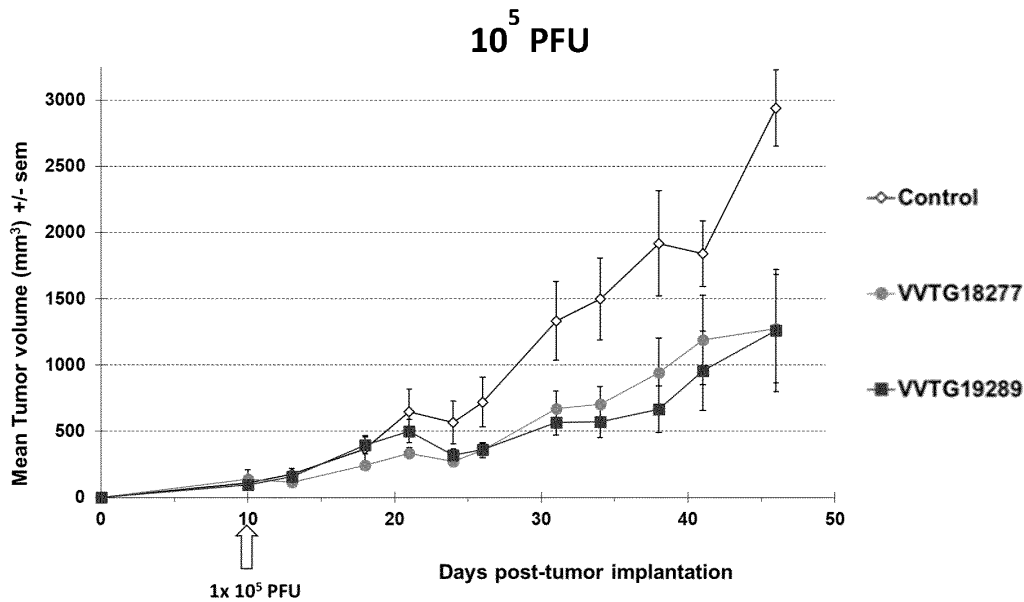


Fig. 10B

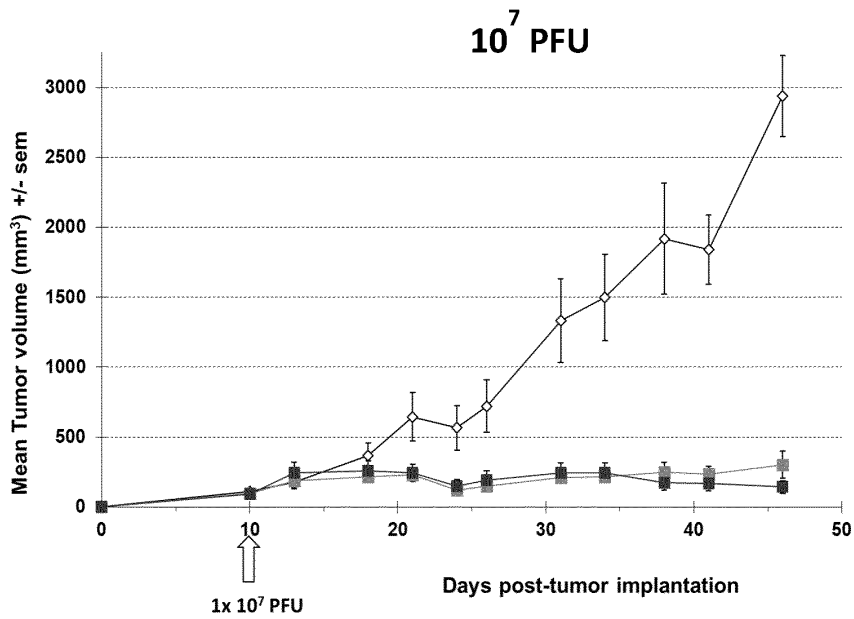


Figure 11

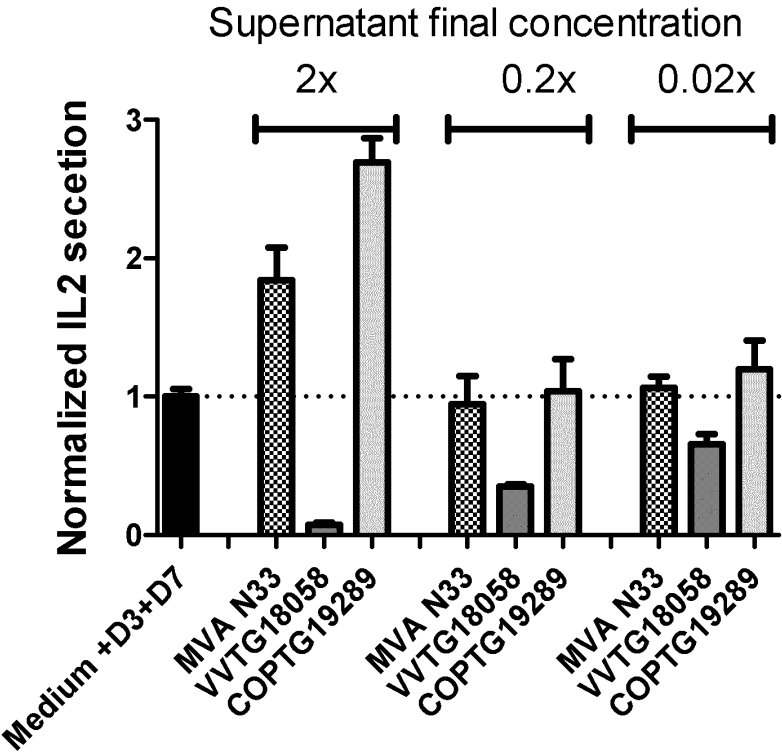


Figure 12

Fig. 12A 10⁶pfu

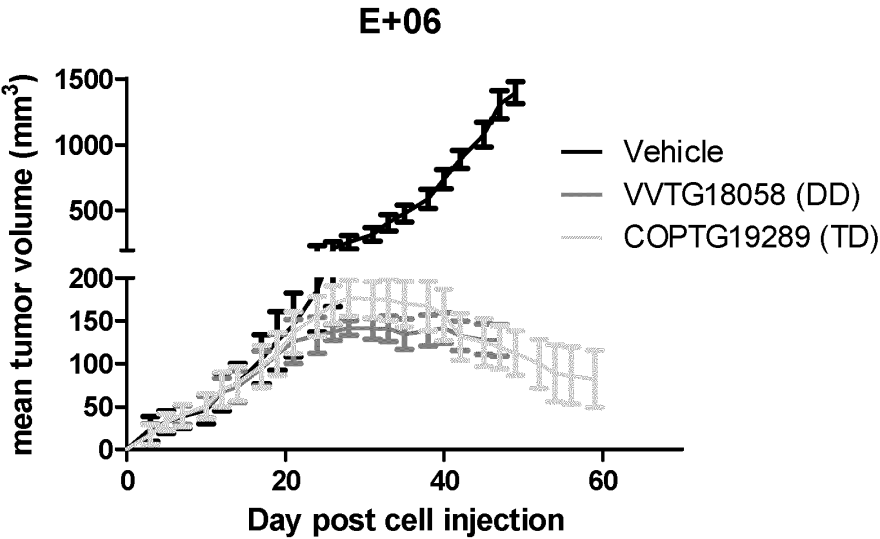
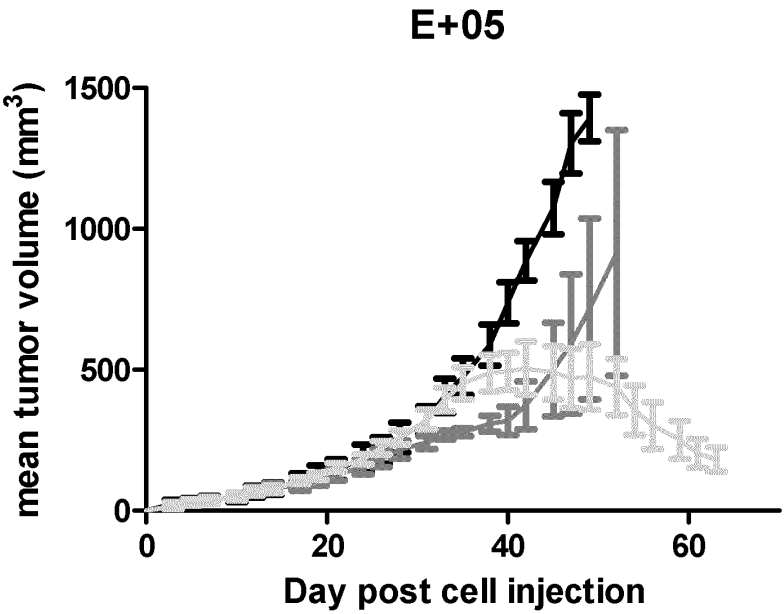


Fig. 12B 10⁵pfu



15/15

Figure 13

Fig. 13A 10⁶pfu

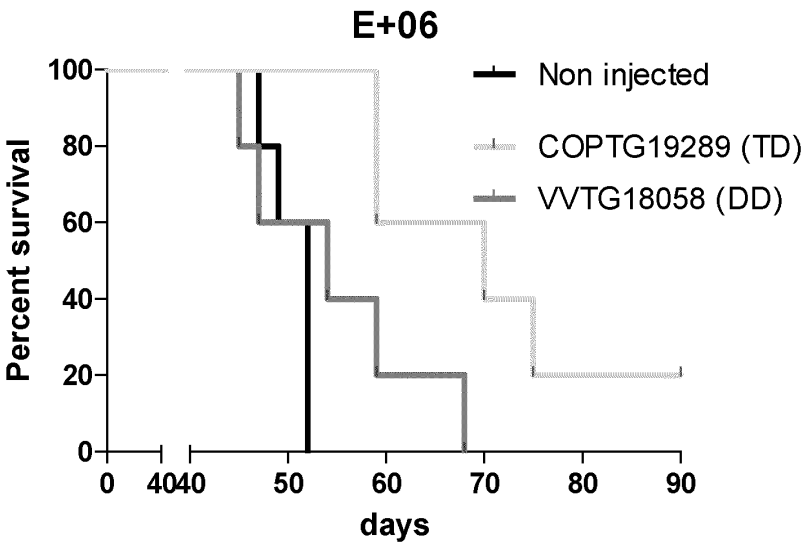


Fig. 13B 10⁵pfu

