

B

EGFR_human	SFLQRYSSDPTGALTEDSIDDTFLPVPE	1092	YINQSVPKRPAGS--VQN	1110	NPVYHNQPLNPA--PS	1120
EGFR_pig	SFLQRYSSDPTGALTEDSLDDTFLPAPE		YVNQSVPKRPAGS--VQN		NPVYHNQPLSAA--PG	1078
EGFR_mouse	AFLQRYSSDPTGAVTEDNIDDAFLPVPE		YVNQSVPKRPAGS--VQN		NPVYHNQPLHPA--PG	1120
EGFR_frog	SFVQRYSTDPTV-ILEDSDIDFLPVPE		YVNQSVMLKPL-VETQTN		SVYQNLI	1117
EGFR_chicken	SFVQRYSSDPTGNFLEESIDDGFLPAPE		YVNQLMPKKPSTAMVQNI		YNNISLTAISKLP	1134
EGFR_zebrafish	SMVLRYIPDPTERFQE----		GDFQPAPGYNEYMNQNE----		SSMINPVYQQPHGPPRTLL-	1110
	
EGFR_human	RDPHYQDPHSTAVGNPEY	1138	LNTVQPTCVNSTF----		DSPAHWAQKGSHQISLD	1172
EGFR_pig	RDPHYQNSHNSAVGNPEY		LNTPRPACINGGL----		DGPAFWAQTGSHQINLD	1135
EGFR_mouse	RDLHYQNPHNSAVGNPEY		LNTAQPTCLSSGF----		NSPALWIQKGSHQMSLD	1177
EGFR_frog	KENLYQNSHNSGLNPEY		LNADQTLVTKTDF----		DYSSIWDQKANLQINLD	1174
EGFR_chicken	MDSRYQNSHSTAVDNPEY		LNTNQSPKLVTF----		ESSPYWIQSGNHQINLD	1191
EGFR_zebrafish	-----HSSPALDETEEY		LNCFKSPAPASVVEYLNTSHTQLLSTKPF		FSSMDNPDYQQDFC	1165
	
EGFR_human	PKEAKPNGIFKGSTAENAEY	1197	LRVAPQSSEFIGA--			1210
EGFR_pig	PKEAKSNGICKGPAAENAEY		LRAAPASSDLTGA--			1168
EGFR_mouse	PKETKPNGIFKGPTAENAEY		LRVAPPSSEFIGA--			1210
EGFR_frog	PKEHKTNGHFKIPAAQNPEY		FGLAPVT-DYVEASA			1208
EGFR_chicken	PNETKPNGLLKVPAAENPEY		LRVAAPKSEYIEASA			1226
EGFR_zebrafish	PLELKTHTNGHLPAAQNQEY		MGLEVH-----			1191
	

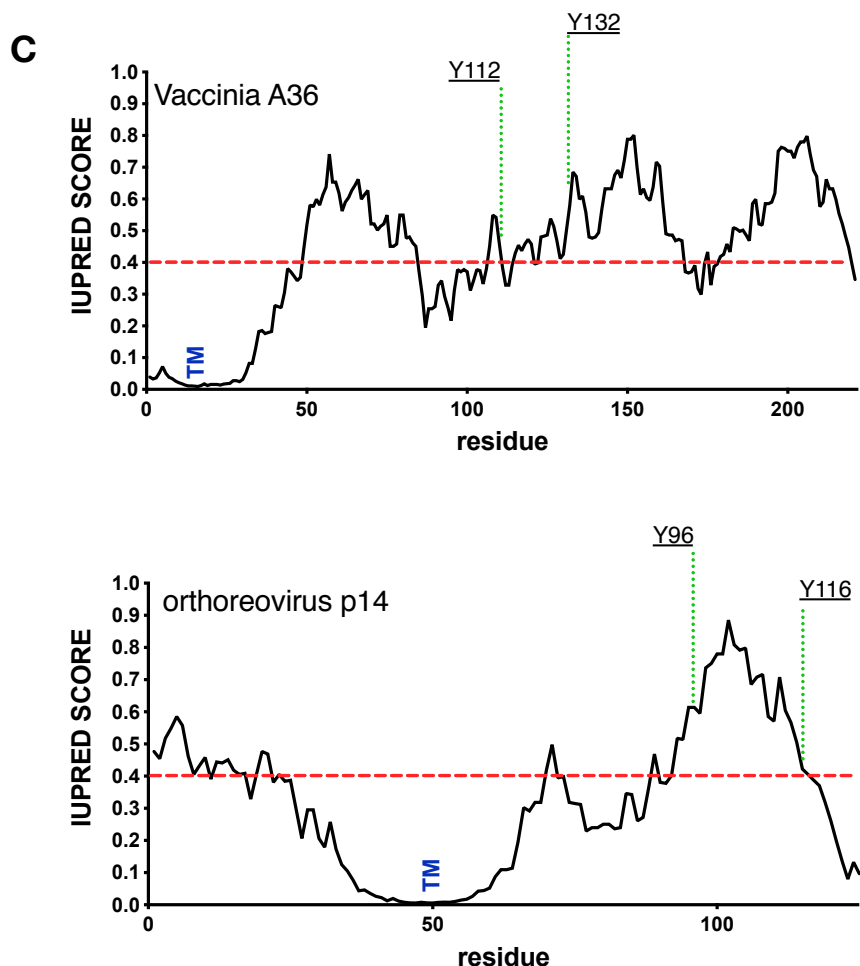


Figure 1 - suppl 1