

Supplementary Figures

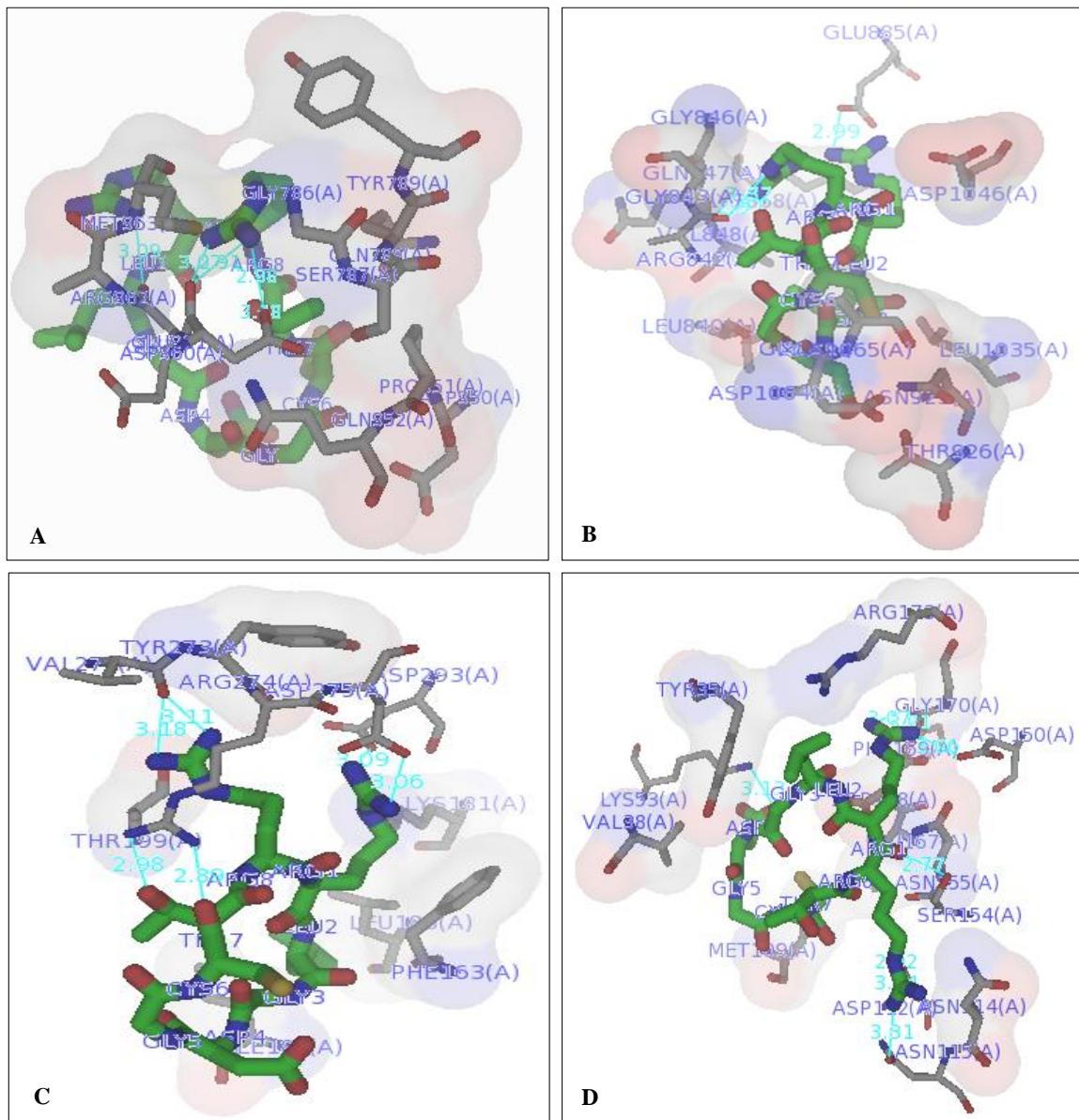


Fig. SB.1 Interaction of cyclosporin with various cancer-related proteins using LigPlot (3D representation). A) EGFR Kinase B) VEGFR2 Kinase C) PKB D) p38.

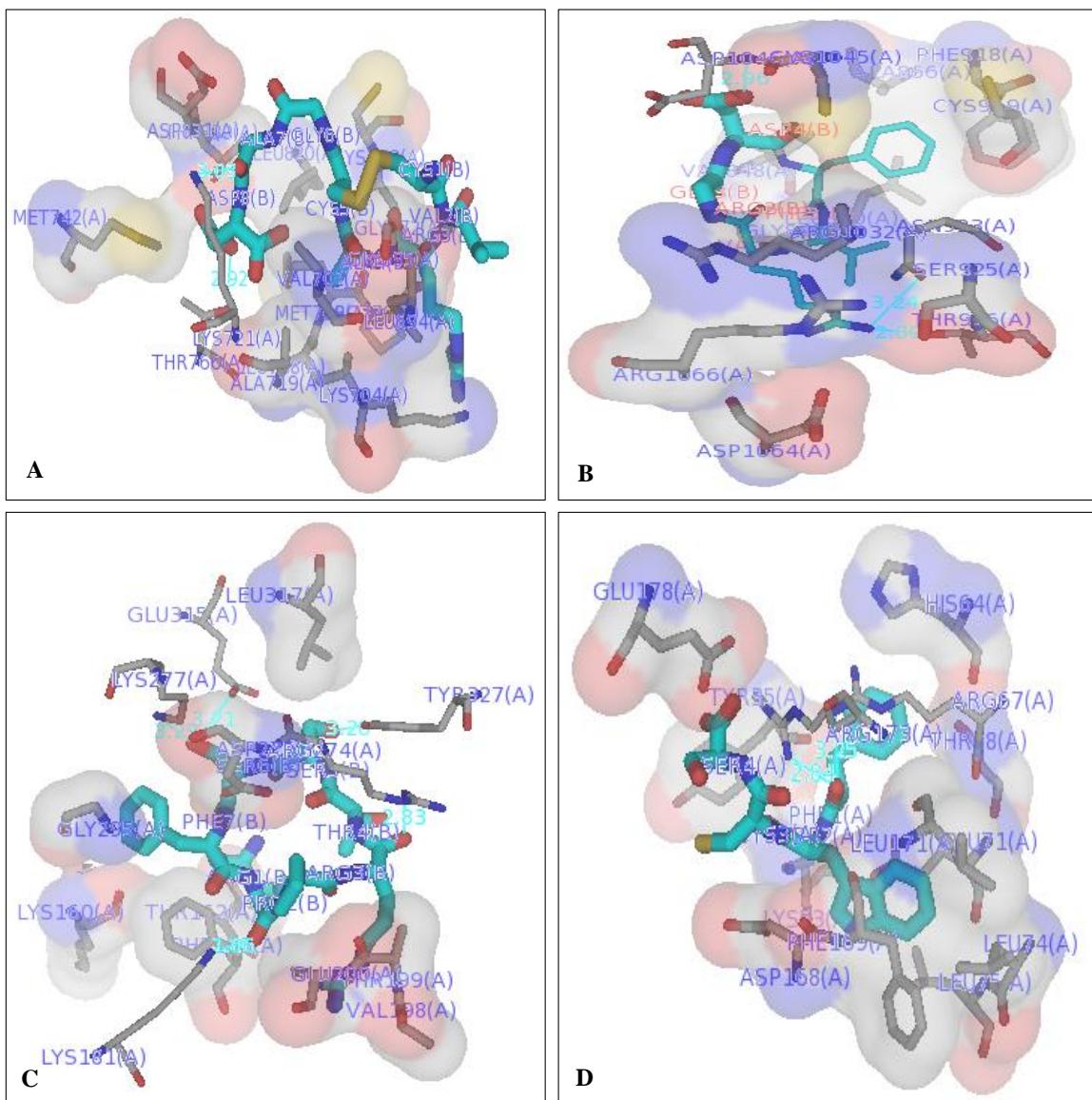


Fig. SB.2 Interaction of a few peptide-based ligands with various cancer-related proteins using LigPlot (3D representation). A) CVRACGAD and EGFR kinase B) Cilengitide and VEGFR2 kinase C) RPRTSSF and PKB D) FWCS and p38.

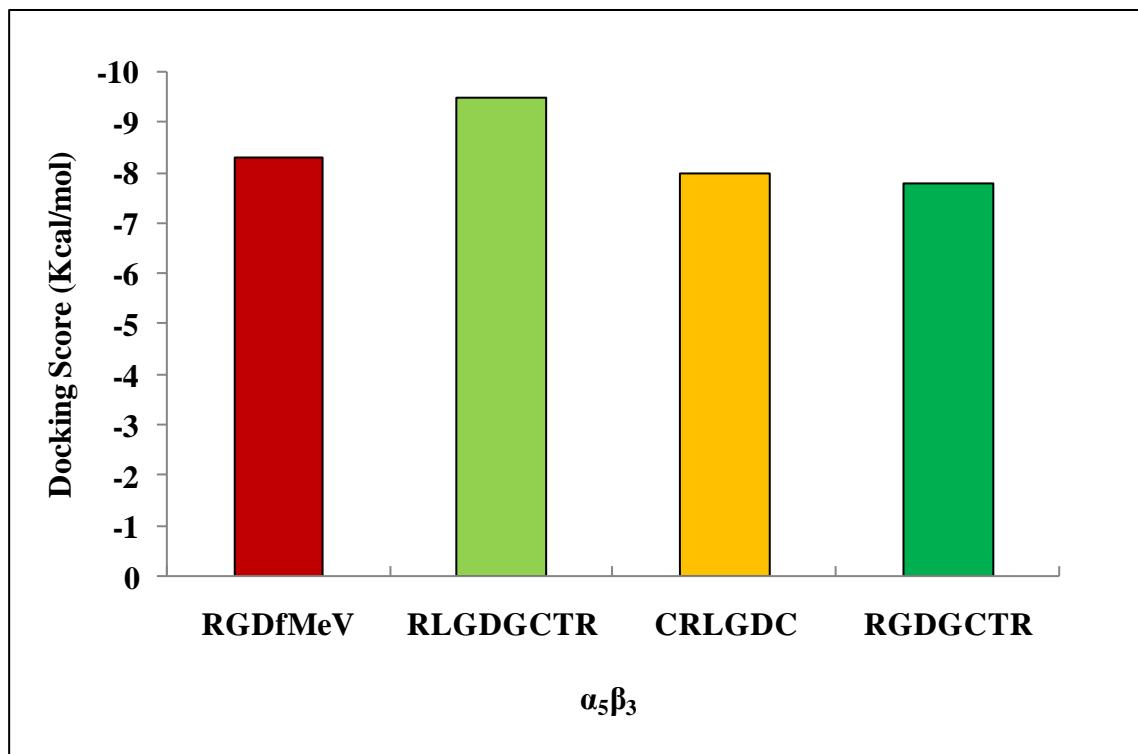


Fig. SB.3 Relative binding affinities (Kcal/mol) of RGD-like peptides towards integrin $\alpha_5\beta_3$

Fig. SB.4 The binding affinities of cyclosaplin along with other RGD-like peptides

Peptide	Binding affinities (Kcal/mol)
RGDFMev (Cilengitide)	-8.3
RLGDGCTR (Cyclosaplin)	-9.5
CRLGDC (Analogue I)	-8.0
RGDGCTR (Analogue II)	-7.8