

Figure S1. Venn diagram represents the modulated protein targets involved in HBV-induced HCC by the phytochemicals from *Andrographis paniculata* and *Thespesia populnea*.

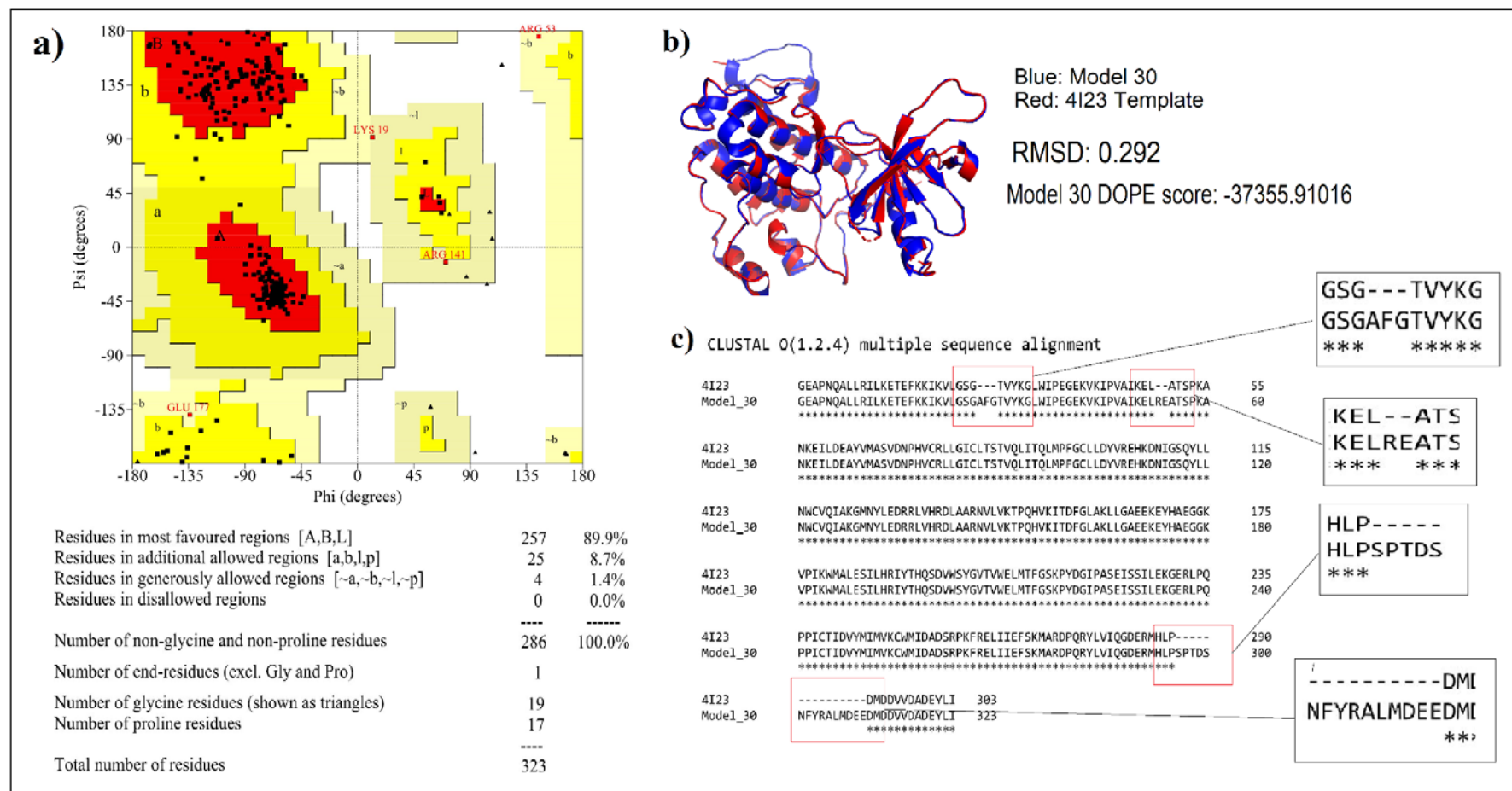


Figure S2. WT- *EGFR* model 30. a) Ramachandran plot for amino acid distribution b) PyMOL superimpose (Red: Template (PDB file 4I23) and Blue: Model 30), c) Clustal W template and model 30 alignment.

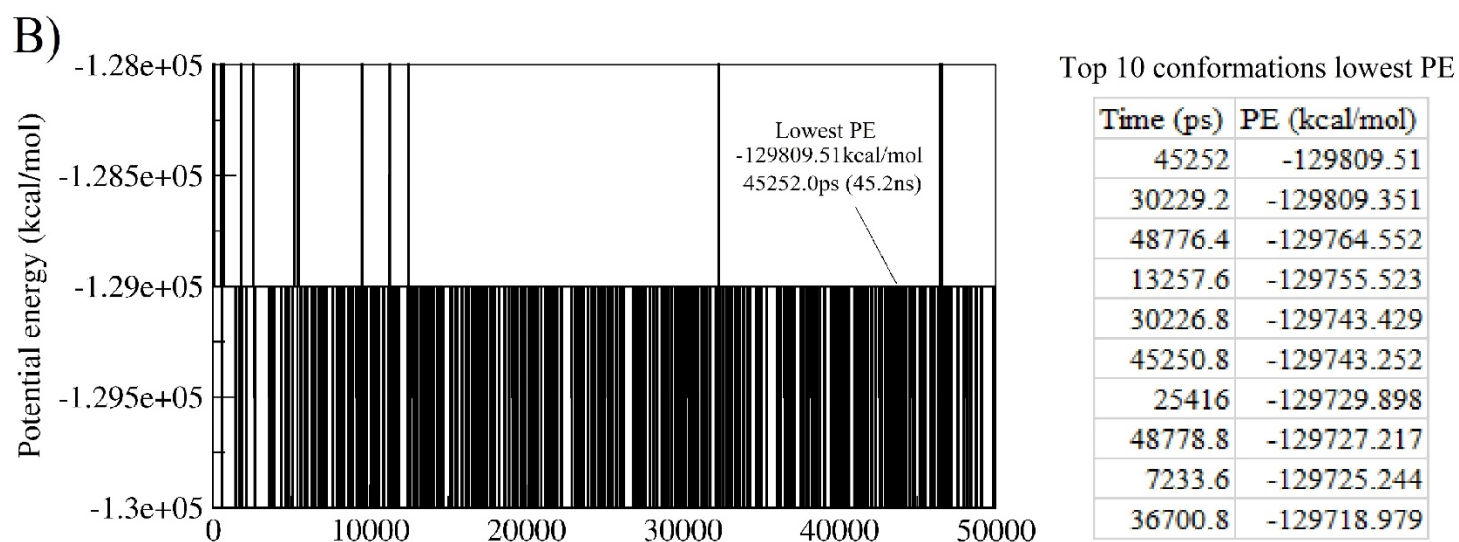
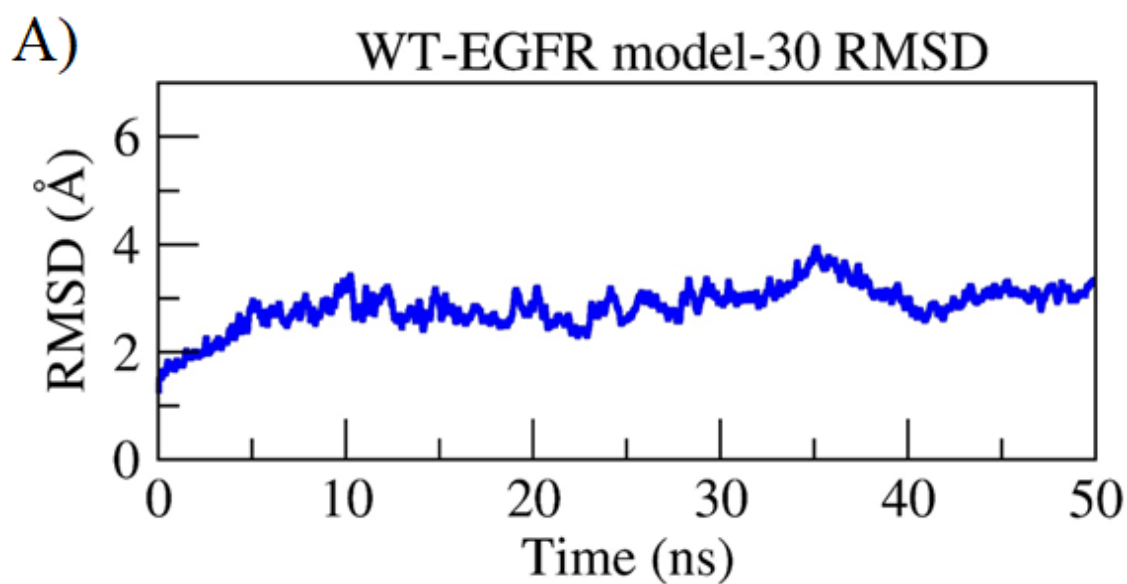


Figure S3. WT-EGFR model-30 RMSD plot showing the deviation (A) and minimum energy conformation observed at 45252.0thps (-129809.51kcal/mol) (B).