



Supplementary Fig. 3. Dynamical cross-correlated map (DCCM) of (A) wild type, (B) Mutant corresponding to c.904 [G>T]; p.E302* and (C) Mutation corresponding to c.453_454 [ins CGTT]; p.Tyr153Serfs*43 mutation. Deeper colour indicates more correlated or anti-correlated. Both x and y axis are residue indices of proteins.