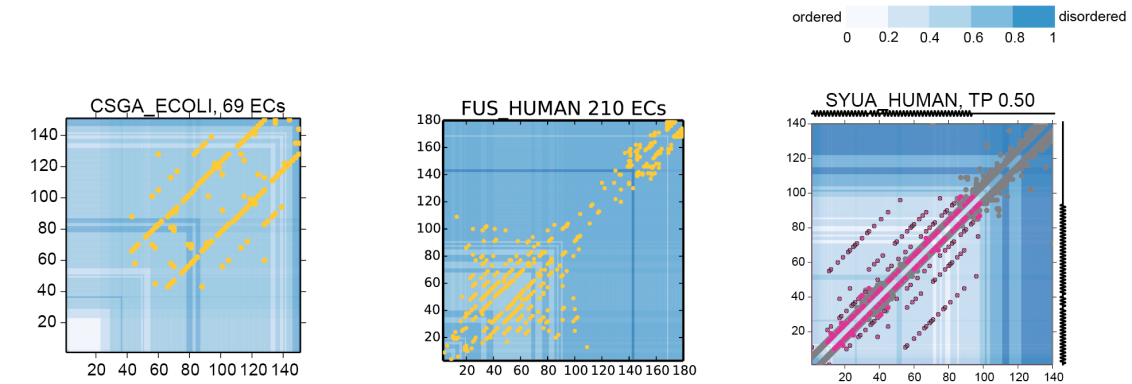
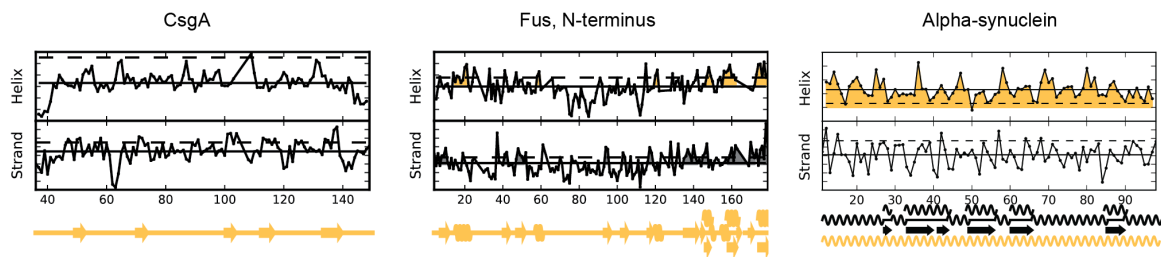


Figure S6. EC predictions of proteins that may form amyloids.

A. Evolutionary couplings



B. Secondary structure from ECs



**Figure S6. ECs predictions of proteins that may form amyloids.**

(A) Predicted contact maps of the regions with unknown structures in Csga (CSGA\_ECOLI), Fus N-terminal low-complexity domain (FUS\_HUMAN) and alpha-synuclein (SYUA\_HUMAN, 2kkw\_A). (B) Secondary structure inference from local ECs for CsgA, Fus prion domain, and alpha-synuclein predict malleable secondary structure (Experimental Procedures).