

Supplementary Figure 2. The SH3 domain aligns with the OB-fold of SSB. PDB file 2KXC containing an SH3 domain bound to a PXXP containing ligand was aligned onto *E.coli* SSB, PDB file 1EYG. The alignment was done using TM-align. Only one monomer of the tetramer of SSB is shown for clarity.

