Figure S1. Angle rotation graphs of unbound trajectory 1 (black), trajectory 2 (blue) and bound (red) forms for (A) Skp2, (C) Fbw7, (E) β-TrCP1, (G) Cdc4, (I) Fbs1, and (K) TIR1. The graphs of changes in the distances between hydrophobic residues from box domain and linker are shown for the unbound form trajectory 1 of (B) Skp2, (D) Fbw7, (F) β-TrCP1, (H) Cdc4, (J) Fbs1 and (L) TIR1.