



Figure S7 Dbf4 T105 residue was critical for the Dbf4-FHA1 domain interaction. (A) The *dbf4*-Δ100-109, *dbf4*-T105A and *dbf4*-NΔ109 mutants caused a loss of FHA1 domain binding in two-hybrid assays. The *dbf4*-S84A, -S92A, and -T95A mutants did not show any effect on FHA1 domain binding. (B) Substitution of T105A within various Dbf4 truncations consistently caused a loss of interaction with the FHA1 domain.