**Supplementary Data**



**Figure S1. Peak analyses of Twist1 ChIP-Seq Data Sets.** (A) Heat map view of signals observed in the presence of Twist1. (B) Density graph of the Twist1 peak locations in relation to the nearest transcriptional start sites (TSS). The x-axis represents the distance from the peak center to the nearest TSS, where zero is the position of the TSS. The y-axis represents the proportion of detected bindings that are located within each distance interval. (C) Twist1 ChIP-seq peak density on the Cul2.