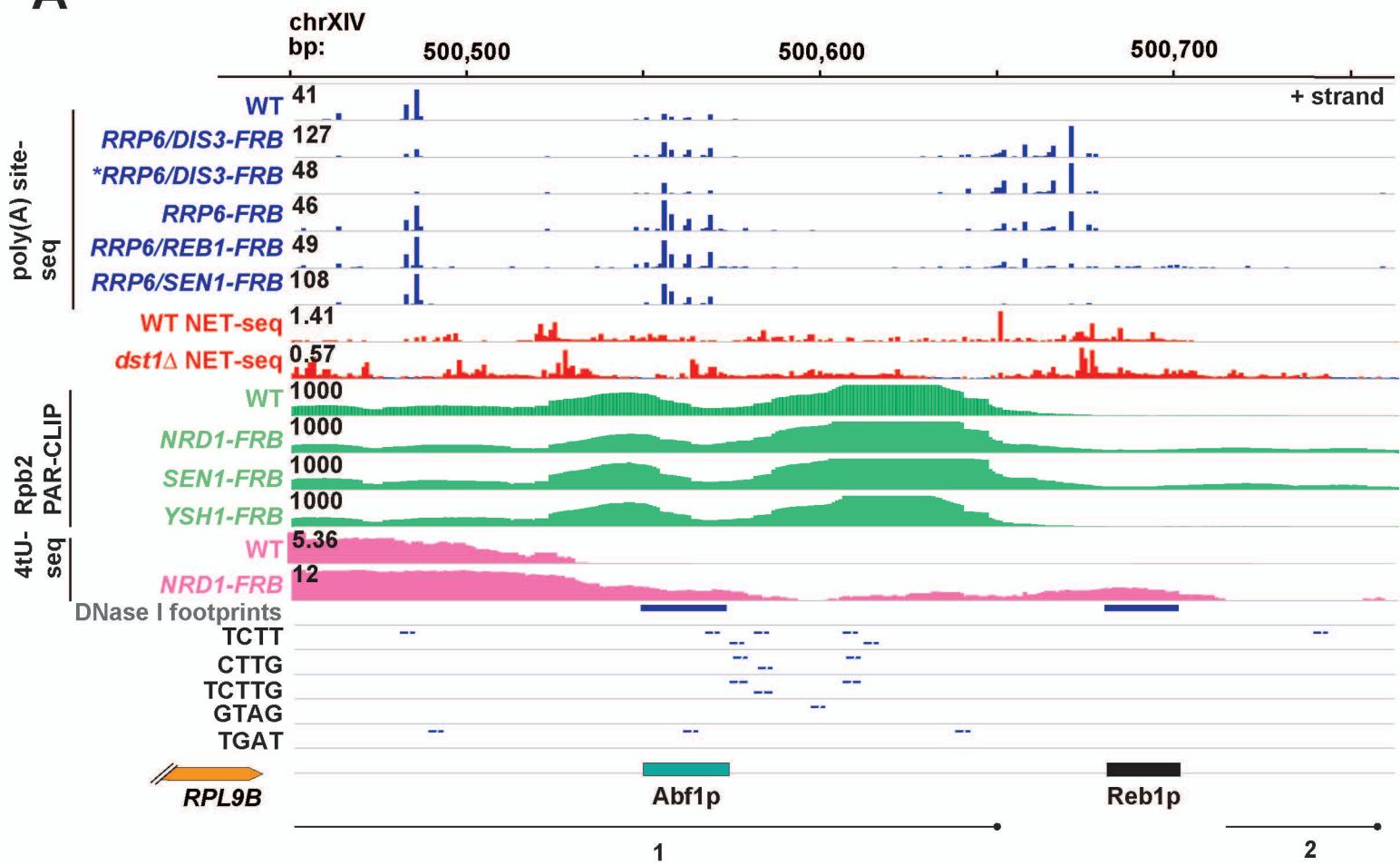
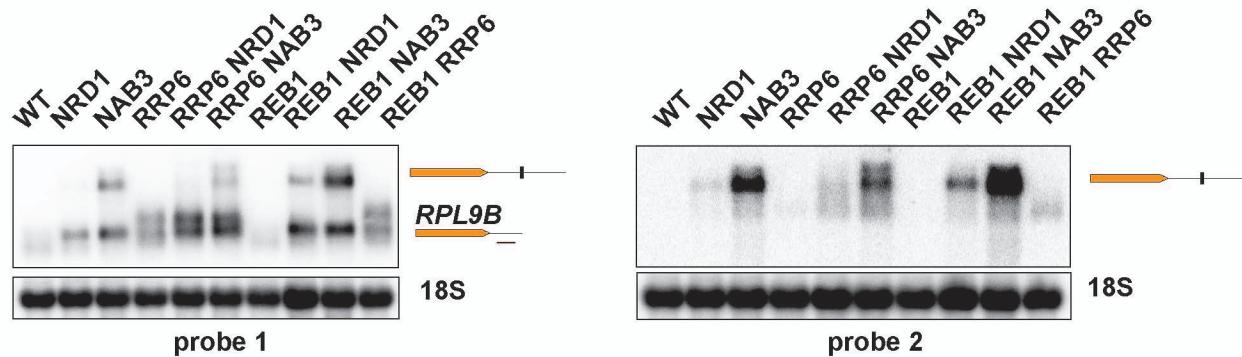


# Roy\_et\_al\_Supplemental\_Fig.S8

**A**



**B**



**Supplemental Figure S8. Control of RPL9B transcription termination by the NNS pathway and Reb1p, Related to Figure 4**

(A) Genome browser view of the *RPL9B* termination region. Tracks are shown as in **Supplemental Fig. S2**. 4-thio-uracil sequencing (4tU-seq) is from previously published data (Schulz et al. 2013). The relative positions of the Nrd1p and Nab3 motifs are indicated in the bottom tracks. The y-axis indicates reads per million for each dataset. (B) Northern blot analysis of *RPL9B* upon nuclear depletion of Nrd1p, Nab3p, Rrp6p and/or Reb1p by anchor away. *RPL9B* transcripts were detected with a 3' UTR riboprobe upstream of the Reb1p binding site (left panel), or downstream of the Reb1p site (right panel). The regions targeted by the probe are shown in (A).