



Supplementary Figure 12. Nascent RNA shows little change upon ROD1 depletion by Gro-seq. **(a)** The alignment of Gro-seq reads to TSSs in both the sense (upper) and antisense (lower) directions. The Gro-seq replicates in the presence of ROD1 are shown as black solid or dash lines. The Gro-seq replicates in the absence of ROD1 are shown as orange solid or dash lines. **(b)** The reproducibility of AID CLIP-seq analyzed by Pearson's correlation. Scatter plots show log₂-normalized read numbers between two replicates. R represents Pearson's correlation coefficient. **(c)** Snapshot of the AID CLIP-seq replicates at the *IgH* and *Ebf1* loci. Of note, AID binding sites at *Ebf1* were localized in a super-enhancer region³³. The signal is shown in red or blue to indicate its localization on the Watson or Crick strand, respectively.