SAGA occupancy by RNA levels (RNA-seq) wda saf6 2.5 -2.0 -2 · 1.5 1.0 0.5 rnaseq_quantile 0.0 log2 ratio (ChIP/Input) 10 spt3 ada2b 9 3 2 2 6 5 0 0 10 9 8 6 5 3 4 3 pol2 2 3 2 0 10 9 8 6 5 Distance to TSS (bp)