SAGA occupancy by RNA levels (RNA-seq) adab2 saf6 7.5 6 -5.0 2.5 2 · 0.0 -0 spt3 tbp rnaseq_quantile 8 -7.5 -6 RPM (log2) 5.0 2.5 2 0 -0.0 trf2 wda 8 -10.0 6 -7.5 4 5.0 2 2.5 0.0 0 5 10 9 10 9 8 6 Distance to TSS (bp)

10 9

3