



Figure S14. Top 15 TF modules for ChIP+KO+ATAC TRN. TFs were clustered into modules based on shared positive target genes between TFs (see [Methods](#)). The left panel shows enrichment of positive TF target interactions with up-regulated Th17 genes (“Th17” for potential Th17-promoting TFs) or down-regulated Th17 genes (“Non-Th17”, for potential Th17-repressing TFs). The lower panel shows relative TF gene expression over the Th17 and other T Helper cell time points.