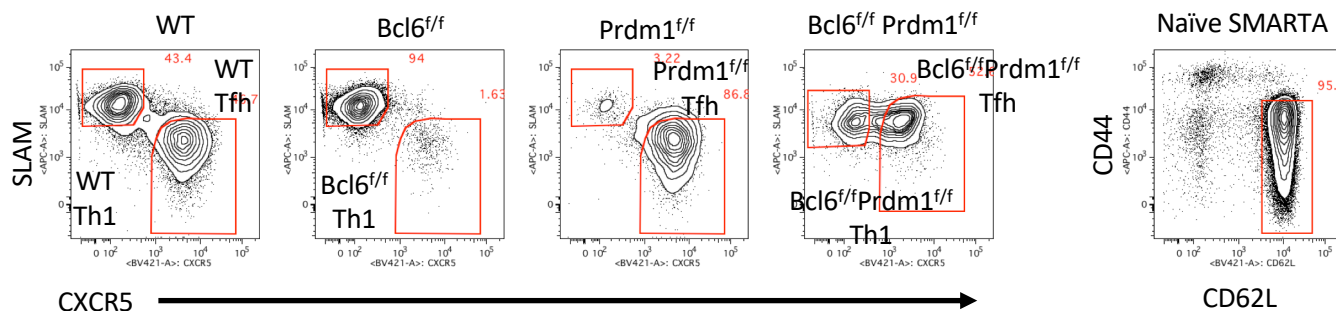


SMARTA CD45.1⁺

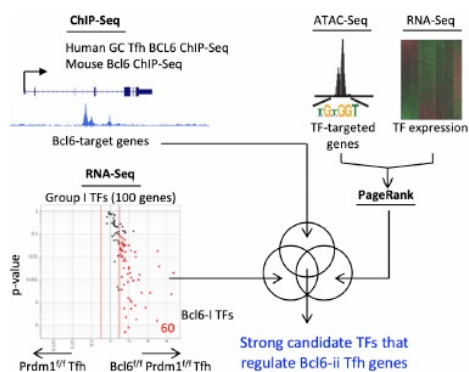
1. WT
2. Bcl6^{f/f}
3. Prdm1^{f/f}
4. Bcl6^{f/f}Prdm1^{f/f}



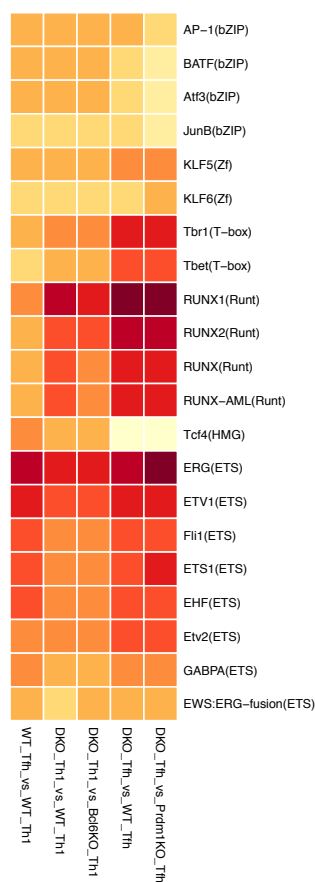
Experimental timeline diagram showing the sequence of events: cells, LCMV, Armstrong, Sort, and ATAC-Seq. Time points d-1, d0, and d7 are marked along a horizontal timeline. Arrows indicate the timing of each event: cells at d-1, LCMV at d0, Sort at d7, and ATAC-Seq following Sort.



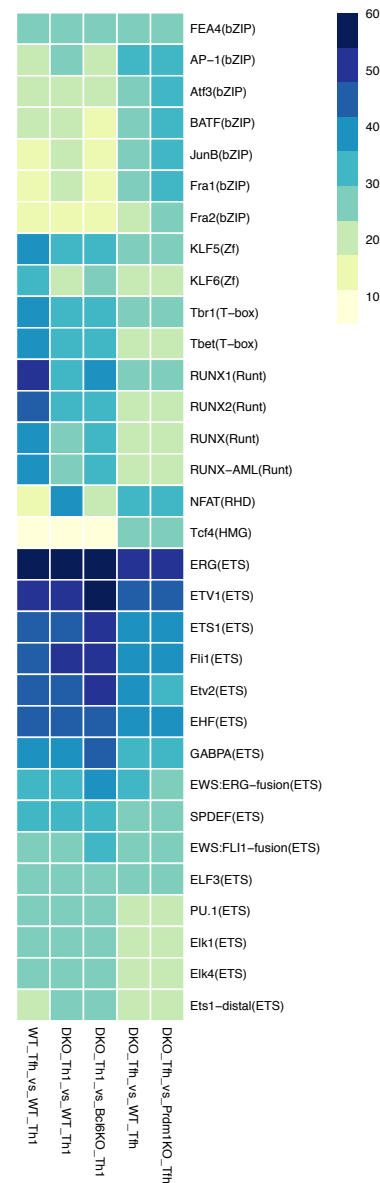
Jinyong will modify this



Open



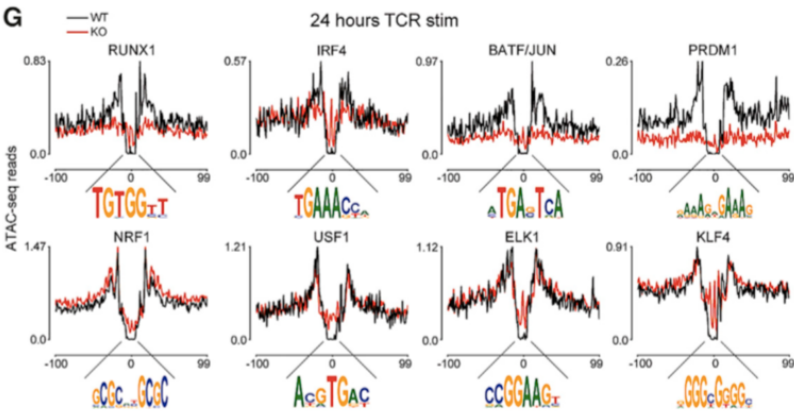
Closed



Huitian:
Can you please modify this
with more curated version
including Klf2, Gata3, and
Maf.

D Huitian:
Can you please generate this TF footprint analysis for Runx2, Runx3, Klf2, Gata3, and Maf.

TF footprint analysis:
WT Tfh vs. DKO Tfh



E Huitian:
Heatmap Can you please generate this figure with curated version including Runx2, Runx3, Klf2, Gata3, and Maf.

Group I Genes Group II Genes Group III Genes Group IV Genes

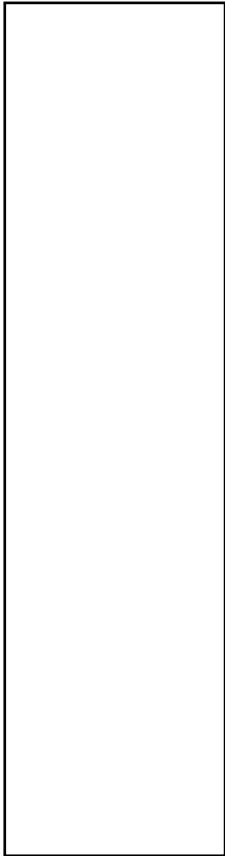
WT_TFH vs. WT TH1
DKO_TH1 vs. Bcl6KO TH1
DKO_TFH vs. Prdm1KO TFH

WT_TFH vs. WT TH1
DKO_TH1 vs. Bcl6KO TH1
DKO_TFH vs. Prdm1KO TFH

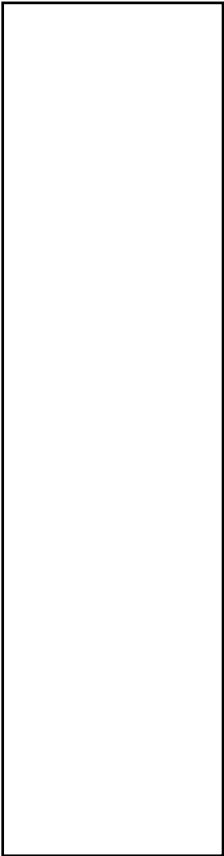
WT_TFH vs. WT TH1
DKO_TH1 vs. Bcl6KO TH1
DKO_TFH vs. Prdm1KO TFH

WT_TFH vs. WT TH1
DKO_TH1 vs. Bcl6KO TH1
DKO_TFH vs. Prdm1KO TFH

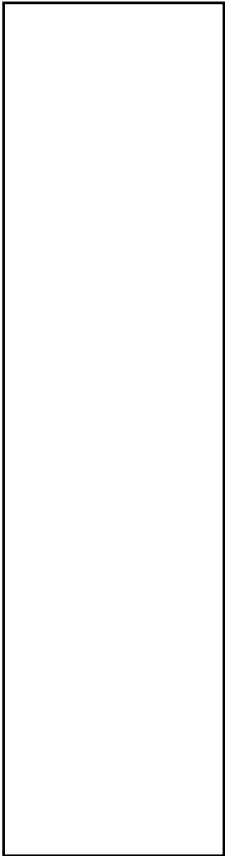
TFs w/ motif enrichment



TFs w/ motif enrichment



TFs w/ motif enrichment



TFs w/ motif enrichment

