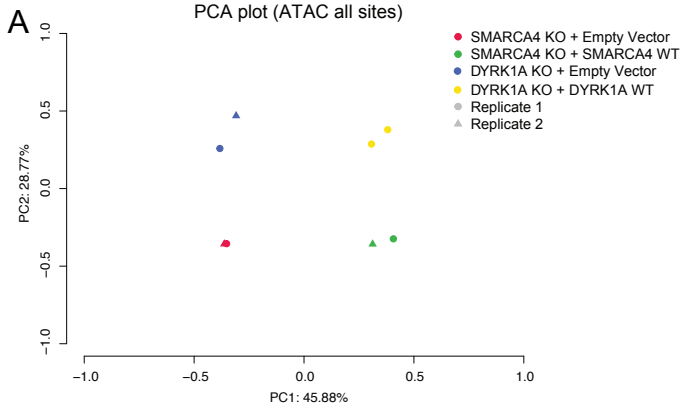
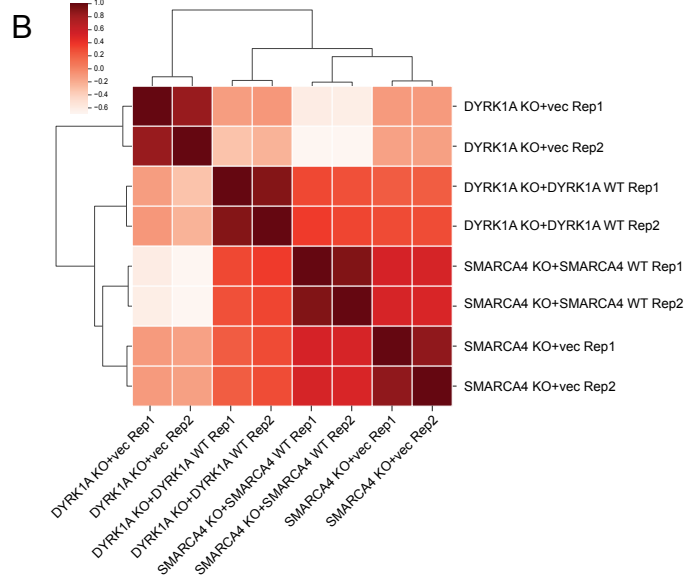


Supplemental Figure 4

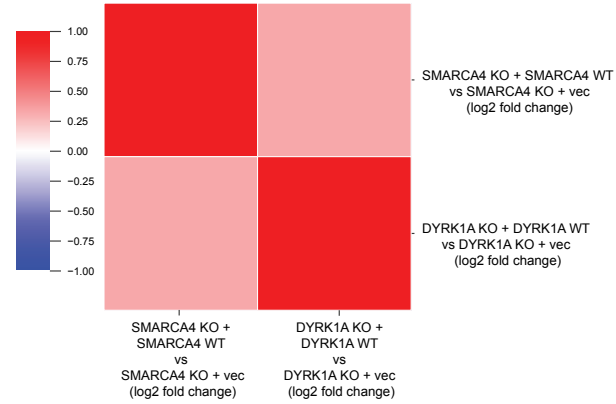
PCA plot (ATAC all sites)



Correlation heatmap of all ATAC sites



Correlation heatmap (ATAC log2 fold change)
Correlation coefficient = 0.33



D
Gained: SMARCA4 KO+WT vs. SMARCA4 KO+vec
Gained: DYRK1A KO+WT vs. DYRK1A KO+vec



Correlation heatmap (RNA-Seq log2 fold change)
Correlation coefficient = 0.08

