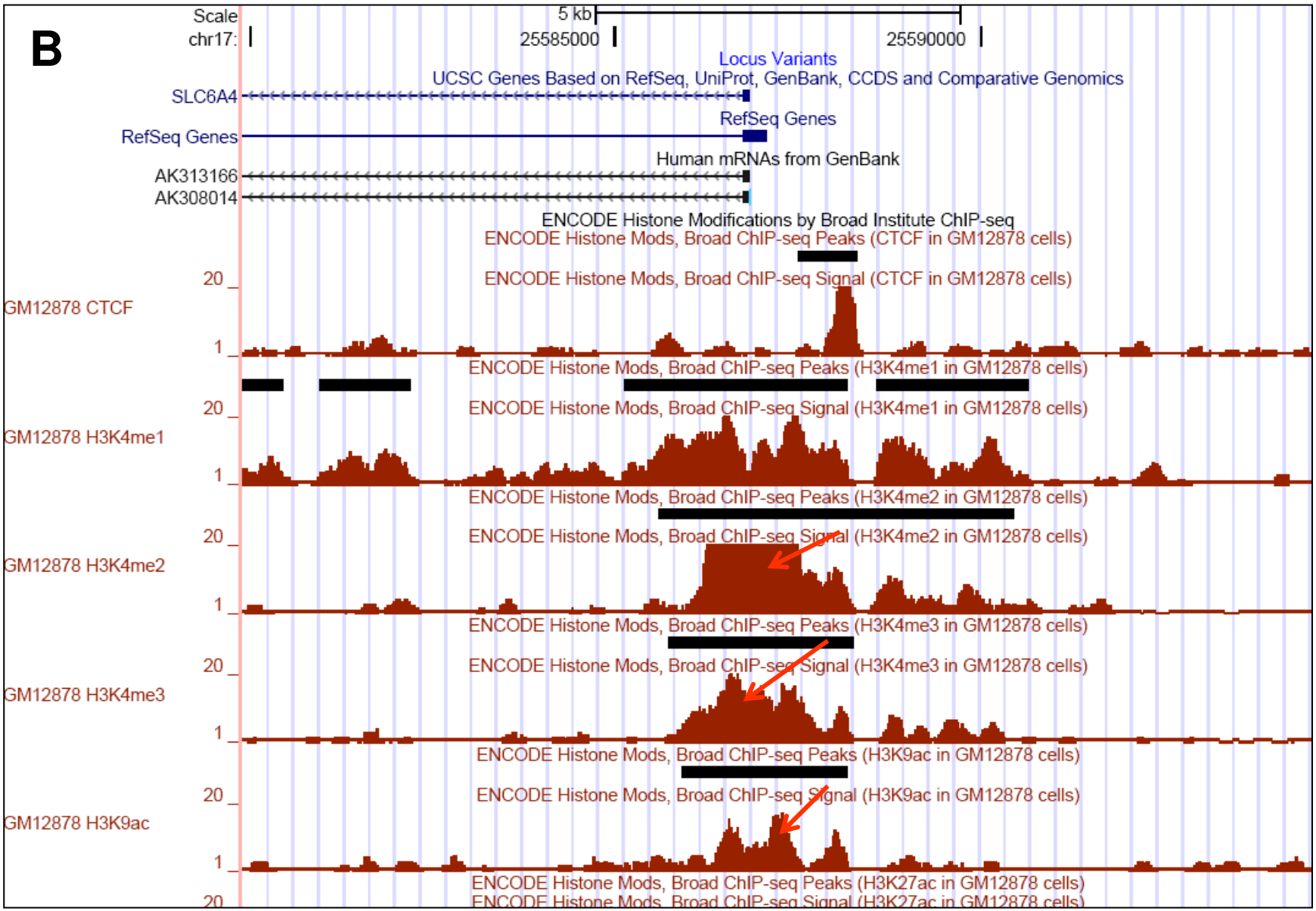
Figure S5B


B
Scale
chr17:
5 kb
25585000
25590000
Locus Variants
UCSC Genes Based on RefSeq, UniProt, GenBank, CCDS and Comparative Genomics
SLC6A4
RefSeq Genes
RefSeq Genes
Human mRNAs from GenBank
AK313166
AK308014
ENCODE Histone Modifications by Broad Institute ChIP-seq
ENCODE Histone Mods, Broad ChIP-seq Peaks (CTCF in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (CTCF in GM12878 cells)
GM12878 CTCF
20
1
ENCODE Histone Mods, Broad ChIP-seq Peaks (H3K4me1 in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (H3K4me1 in GM12878 cells)
GM12878 H3K4me1
20
1
ENCODE Histone Mods, Broad ChIP-seq Peaks (H3K4me2 in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (H3K4me2 in GM12878 cells)
GM12878 H3K4me2
20
1
ENCODE Histone Mods, Broad ChIP-seq Peaks (H3K4me3 in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (H3K4me3 in GM12878 cells)
GM12878 H3K4me3
20
1
ENCODE Histone Mods, Broad ChIP-seq Peaks (H3K9ac in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (H3K9ac in GM12878 cells)
GM12878 H3K9ac
20
1
ENCODE Histone Mods, Broad ChIP-seq Peaks (H3K27ac in GM12878 cells)
ENCODE Histone Mods, Broad ChIP-seq Signal (H3K27ac in GM12878 cells)
20