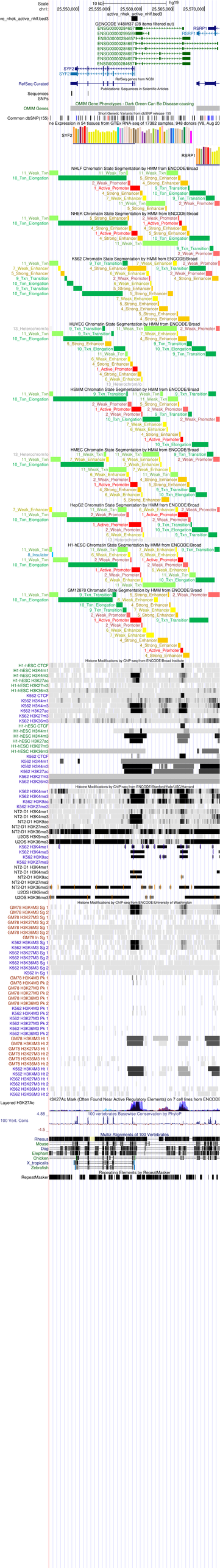
Scale
chr1:
10 kb
hg19
25,550,000
25,555,000
25,560,000
25,565,000
25,570,000
active_nhek_active_nhif.bed3
ve_nhek_active_nhif.bed3
GENCODE V48lift37 (28 items filtered out)
ENSG00000284657
RSRP1
SYF2
SYF2
RefSeq Curated
RefSeq genes from NCBI
Publications: Sequences in Scientific Articles
Sequences
SNPs
OMIM Genes
OMIM Gene Phenotypes - Dark Green Can Be Disease-causing
Short Genetic Variants from dbSNP release 155
Common dbSNP(155)
ne Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 20
SYF2
RSRP1
NHLF Chromatin State Segmentation by HMM from ENCODE/Broad
NHEK Chromatin State Segmentation by HMM from ENCODE/Broad
K562 Chromatin State Segmentation by HMM from ENCODE/Broad
HUVEC Chromatin State Segmentation by HMM from ENCODE/Broad
HSMM Chromatin State Segmentation by HMM from ENCODE/Broad
HMEC Chromatin State Segmentation by HMM from ENCODE/Broad
HepG2 Chromatin State Segmentation by HMM from ENCODE/Broad
H1-hESC Chromatin State Segmentation by HMM from ENCODE/Broad
GM12878 Chromatin State Segmentation by HMM from ENCODE/Broad
Histone Modifications by ChIP-seq from ENCODE/Broad Institute
Histone Modifications by ChIP-seq from ENCODE/Stanford/Yale/USC/Harvard
Histone Modifications by ChIP-seq from ENCODE/University of Washington
Layered H3K27Ac
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE
100 Vert. Cons
100 vertebrates Basewise Conservation by PhyloP
Multiz Alignments of 100 Vertebrates
Rhesus
Mouse
Dog
Elephant
Chicken
X_tropicalis
Zebrafish
Repeating Elements by RepeatMasker
RepeatMasker