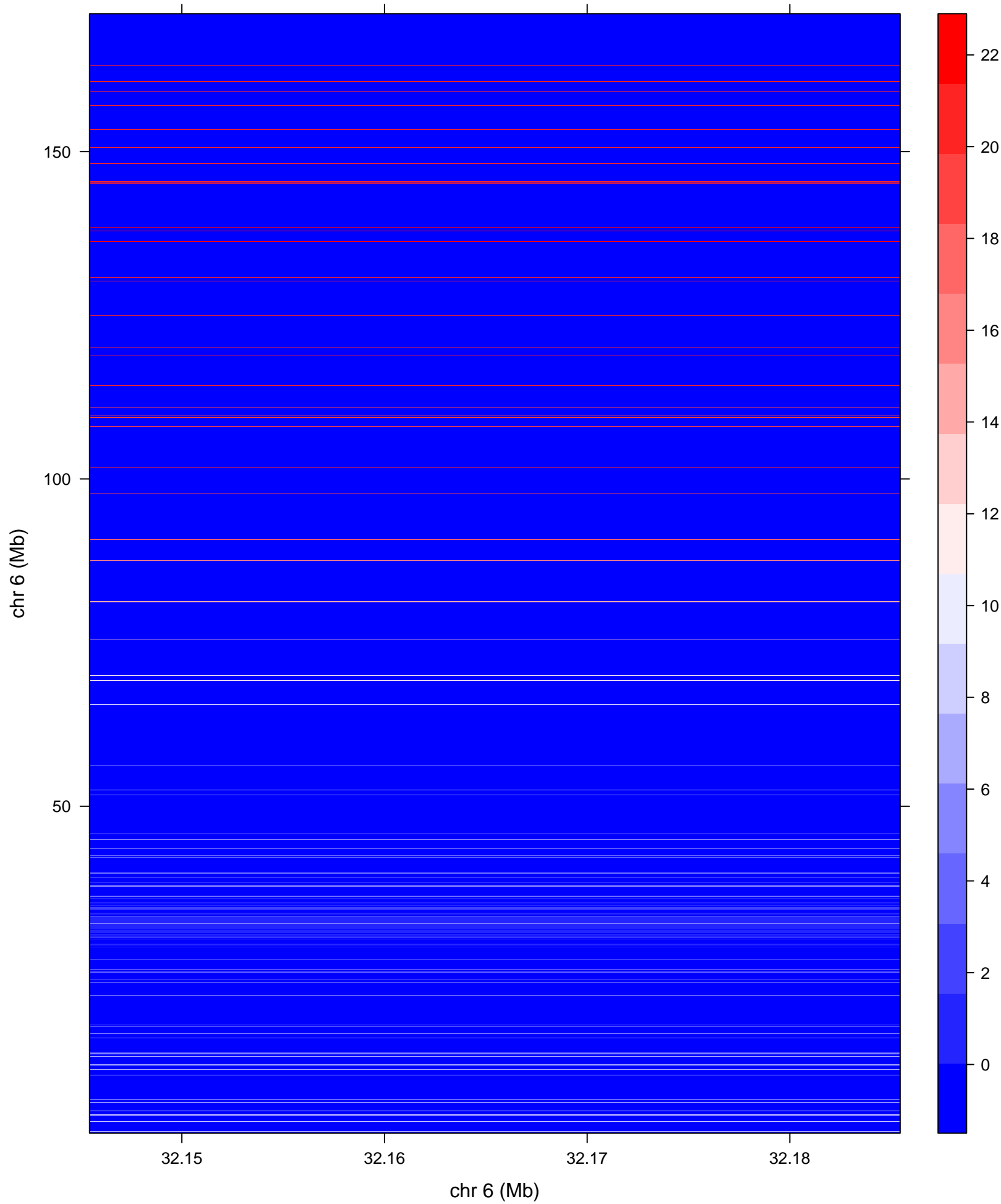












chr 6 chr 6 interaction  
cell\_line:k562

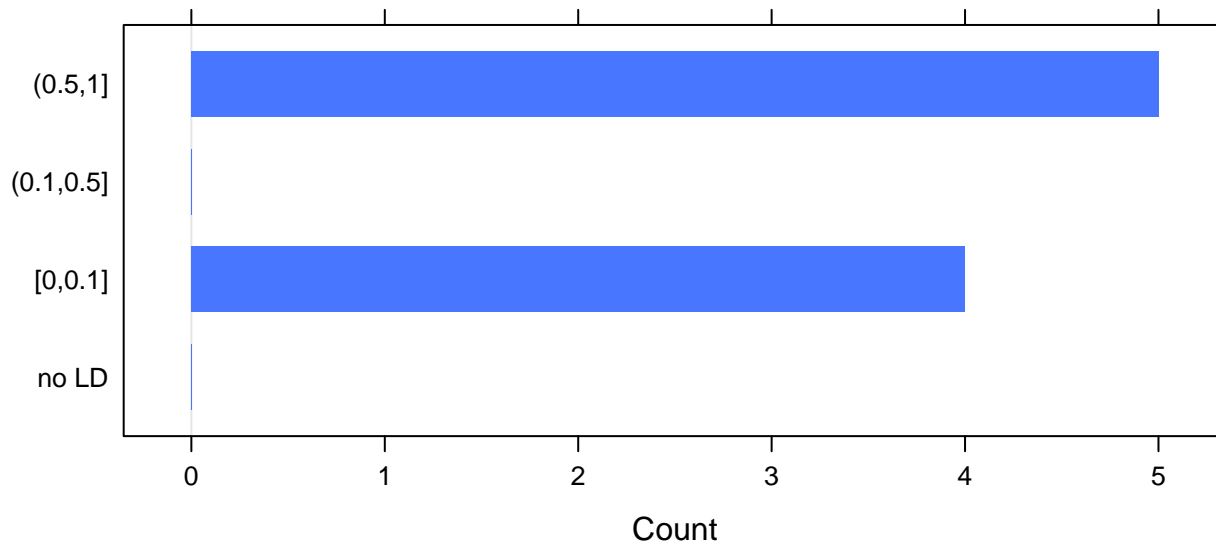


date: Mon Aug 25 17:05:36 2014  
build: hg19  
display range: chr6:32145444–32185444  
reference SNP: chr6:32165444  
number of SNPs plotted: 9  
max P:  $3.93\text{E}-12$  [chr6:32182759]  
min P:  $1\text{E}-1$  [chr6:32179896]

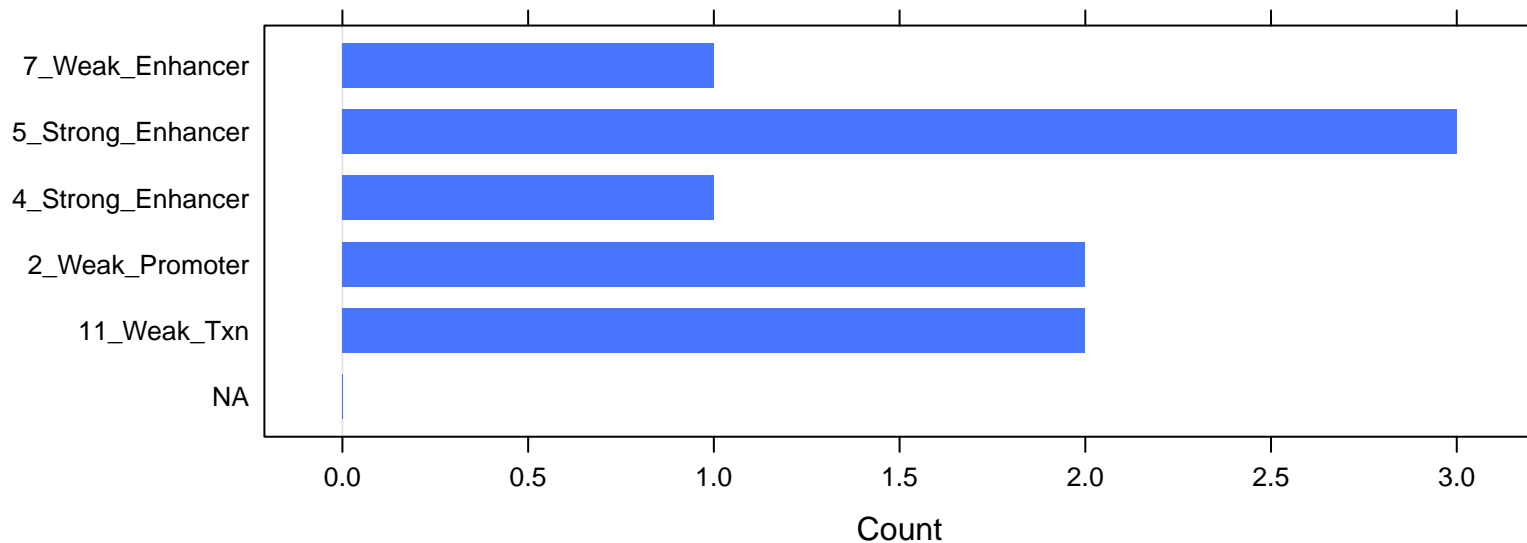
### Category Plot Legend

-  Active Promoter
-  Weak Promoter
-  Inactive/poised Promoter
-  Strong enhancer
-  Weak/poised enhancer
-  Insulator
-  Transcriptional transition/elongation
-  Weak transcribed
-  Polycomb–repressed
-  Heterochromatin; low signal; Repetitive/Copy Number Variation

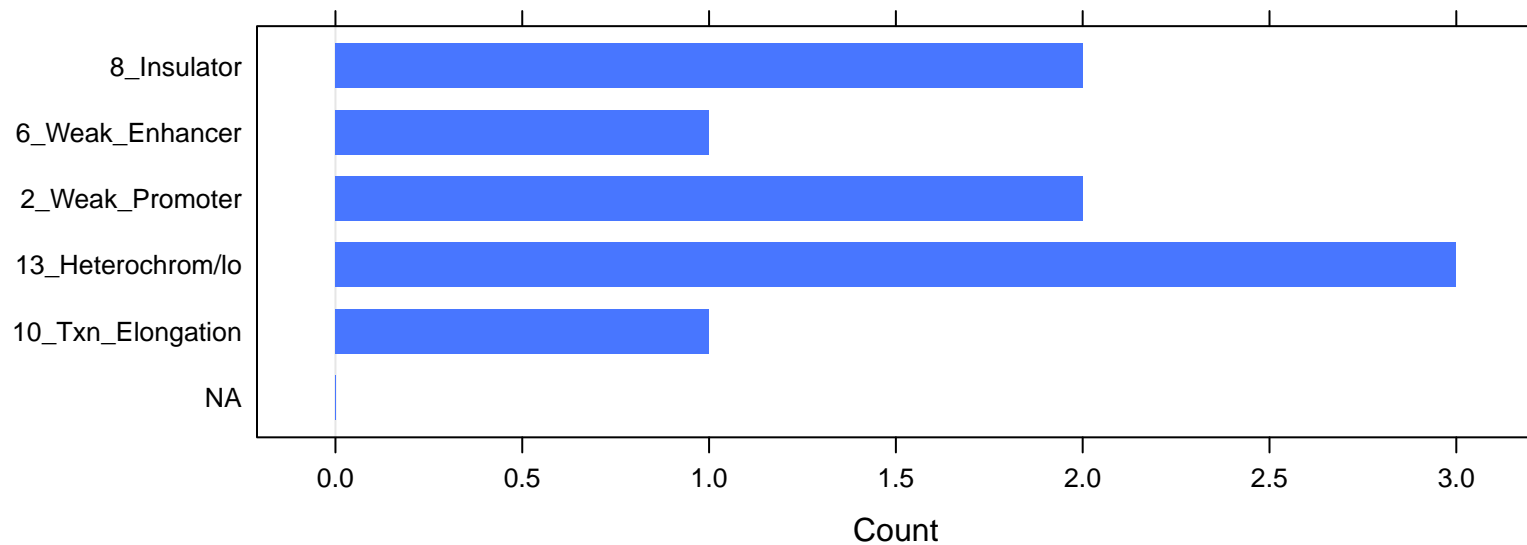
LD category



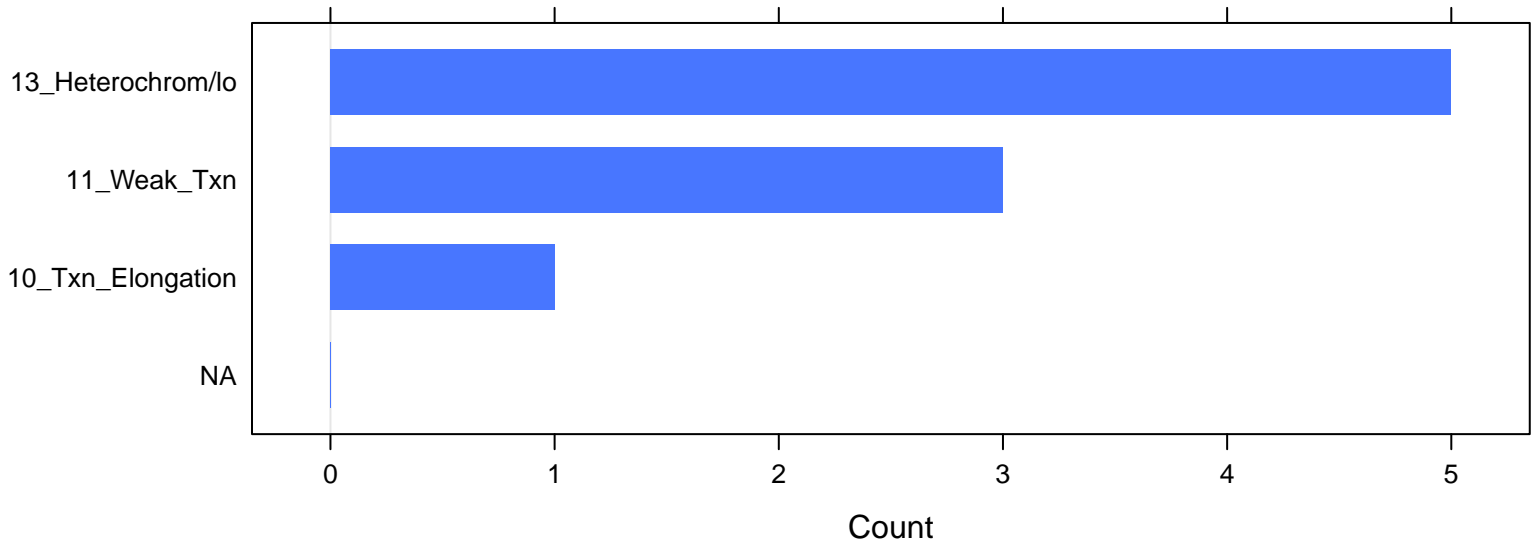
wgEncodeBroadHmmK562HMM category



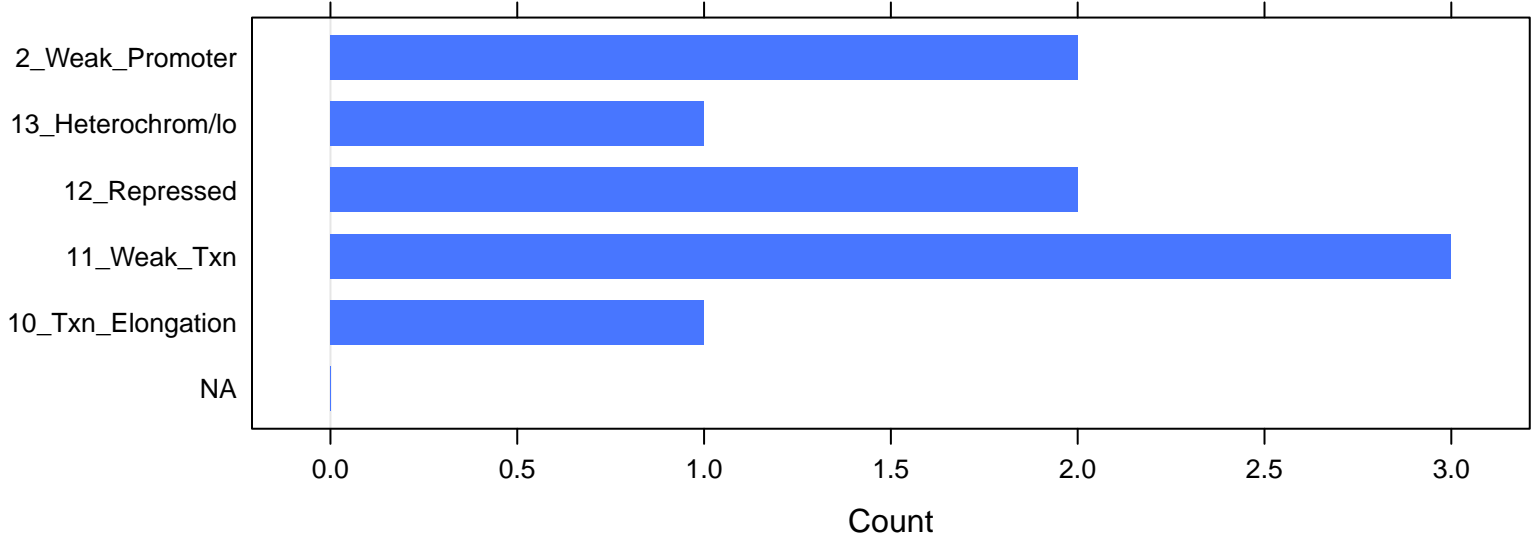
wgEncodeBroadHmmGm12878HMM category



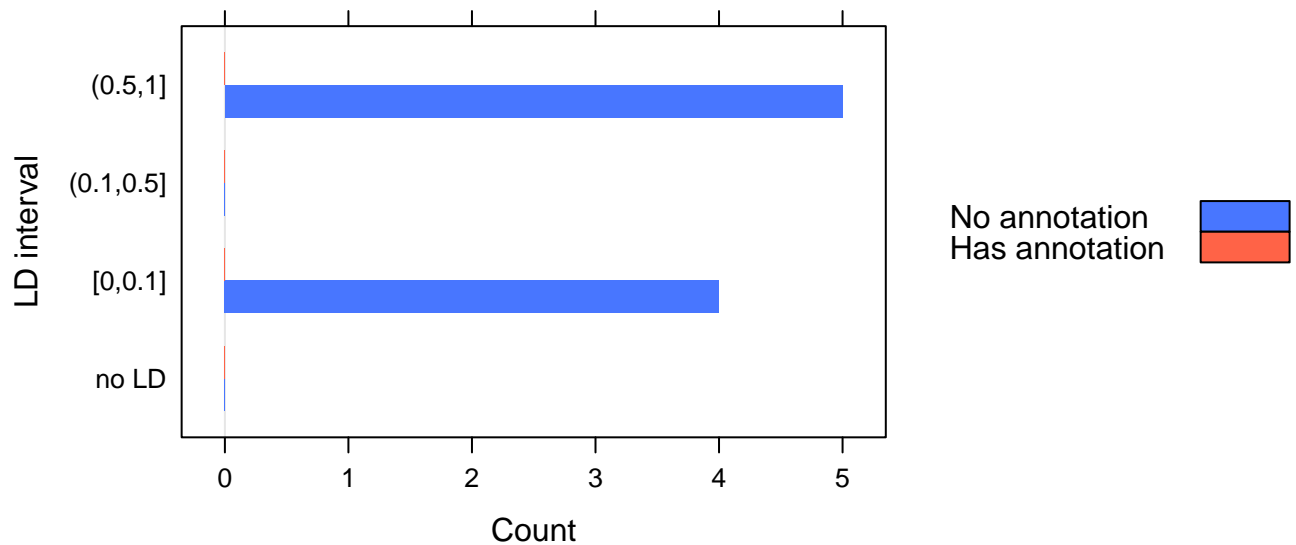
wgEncodeBroadHmmHsmmHMM category



wgEncodeBroadHmmHepg2HMM category



tfbsConsSites category



# wgEncodeRegTfbsClusteredV3 category

