

date: Thu Apr 17 18:46:44 2014

build: hg19

display range: chr15:32975979–33075979

reference SNP: chr15:33025979

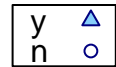
number of SNPs plotted: 394

max p: 1.67E–3 [chr15:33046012]

min p: 9.95E–1 [chr15:33018087]

GTEX_eQTL_11162013 summary: 0 'y' 394 'n'

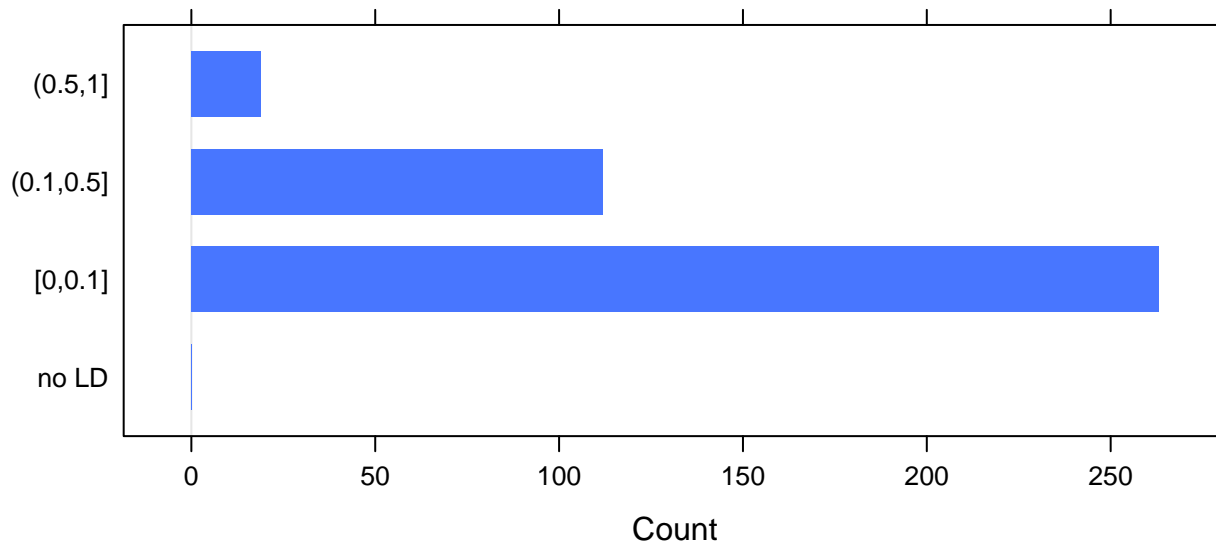
GTEX_eQTL_11162013



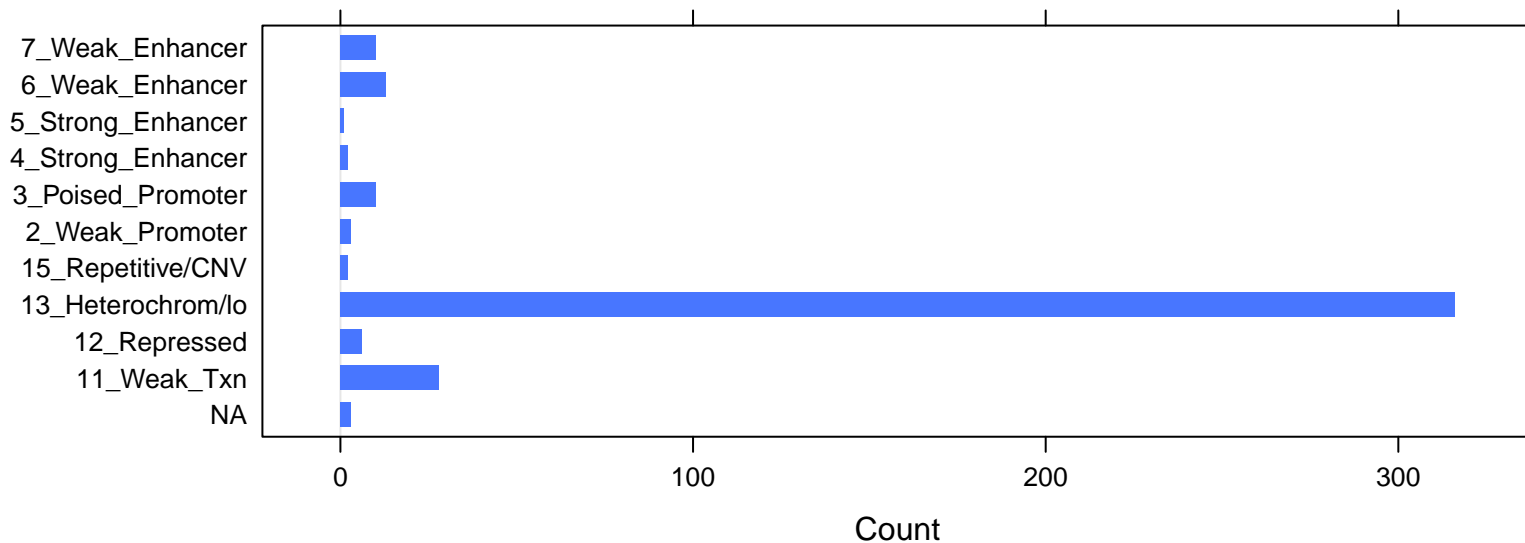
Category Plot Legend

- 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

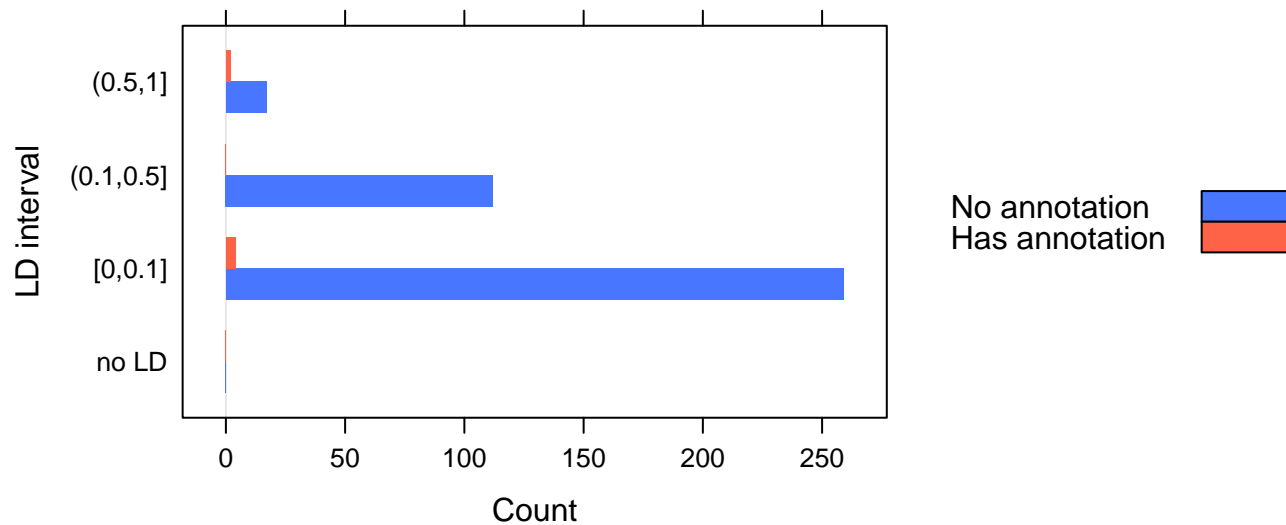
LD category



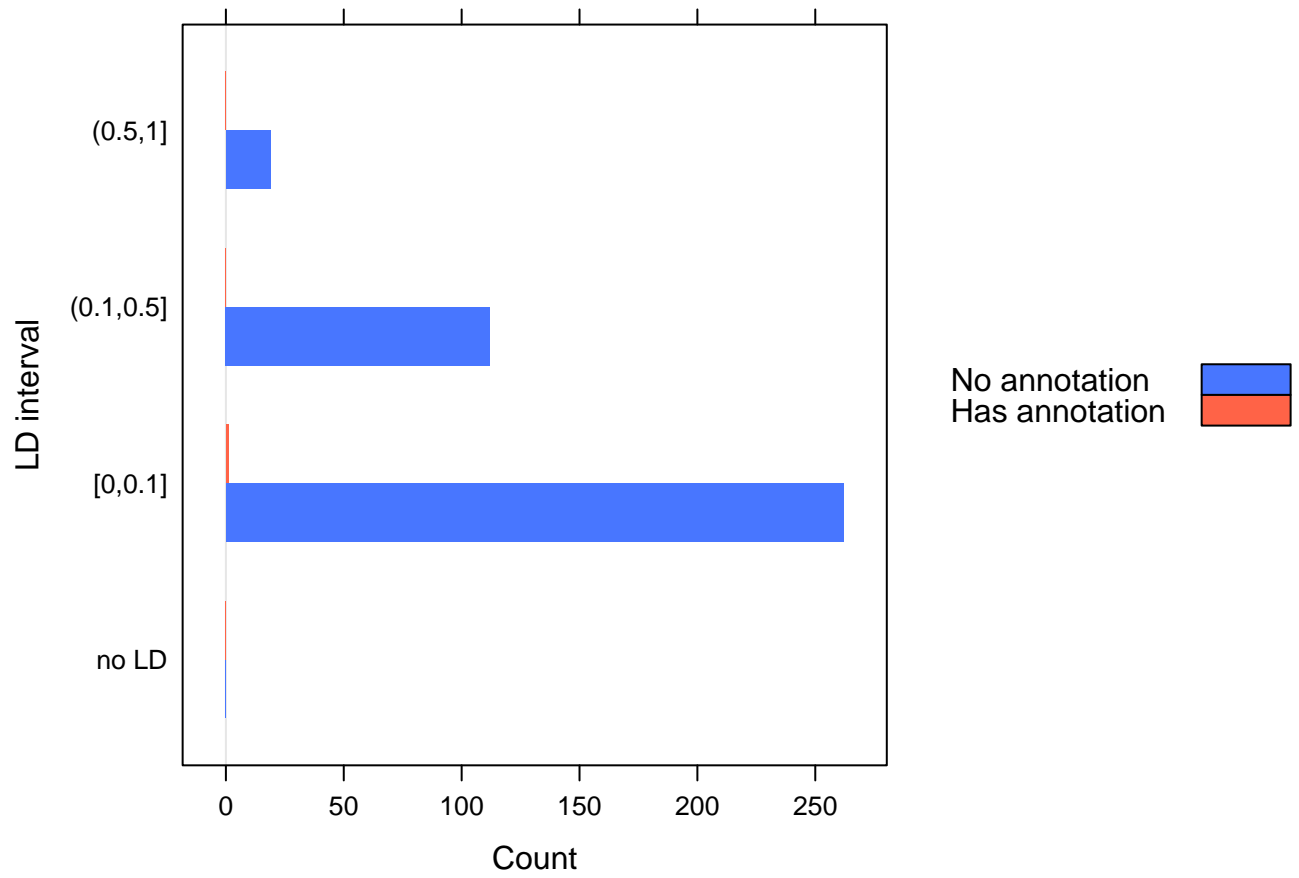
wgEncodeBroadHmmHepg2HMM category



wgEncodeOpenChromDnaseHepg2Pk category



wgEncodeUwDnaseCaco2PkRep1 category



wgEncodeUwDnaseHct116PkRep1 category

