A
puQTL region
GG
GC
CC
FPM
naive
txrevise alternative promoters for HDLBP
Distance from region start (bp)
B
rs62187434 (puQTL lead)
rs12624195 (caQTL lead)
−log10 p−value
HDLBP puQTL
R2
pos
C
caQTL p-values
rs12624195: G/C
−log10 p−value
ATAC−seq
RNA−seq
affected
first exon
accessible
regions
P2
P1
alternative
first exons
Chromosome 2 position (bp)