Week 2 Notes

Look at ExAC

Rprof()

\*Function here\*

Rprof(NULL)

prof = summaryRprof()

prof$by.self[1:5,]

mysql --user=genome --host=genome-mysql.cse.ucsc.edu -A

<https://genome.ucsc.edu/cgi-bin/hgTracks?db=hg38>

&lastVirtModeType=default&lastVirtModeExtraState=&virtModeType=default&virtMode=0&nonVirtPosition=&

position=chr11%3A47331397%2D47352702&

hgsid=499789521\_WALc2CraZ5dKcVbJSKK30ruCKCkC

/Users/jamesdiao/Documents/Kohane\_Lab/week\_2/twoBitToFa http://hgdownload.cse.ucsc.edu/gbdb/hg19/hg19.2bit stdout -seq=chr11 -start=47331396 -end=47352702

Coding regions don’t match

./twoBitToFa http://hgdownload.cse.ucsc.edu/gbdb/hg19/hg19.2bit stdout -seq=chr11 -start=47353421 -end=47353430

>chr11:47353421-47353430

TCACTGAGG

./twoBitToFa http://hgdownload.cse.ucsc.edu/gbdb/hg19/hg19.2bit stdout -seq=chr11 -start=47357128 -end=47357135

>chr11:47357128-47357135

tggcctc