## Exercise 7

## Your Name

22 Dec 2016

```
evals <- read.delim("gene.expression.txt",stringsAsFactors = FALSE)</pre>
evals[1:5,1:5]
                   NKI_4 NKI_6 NKI_7 NKI_8 NKI_9
##
## Contig56678_RC -0.261 0.346 0.047 -1.140 -0.110
## AF026004
                  -0.064 0.040 -0.165 -0.031 0.330
## AB033049
                  -0.307 0.046 -0.139 0.036 -0.154
                   0.582 0.216 0.091 -0.186 -0.156
## AB033050
## AB033086
                  -2.000 0.102 -0.016 -0.358 0.153
dim(evals)
## [1] 498 337
genes <- read.delim("gene.description.txt",stringsAsFactors = FALSE)</pre>
head(genes)
                            probe HUGO.gene.symbol Chromosome
                                                                   Start
## Contig56678_RC Contig56678_RC
                                             THSD4
                                                         chr15 71433788
## AF026004
                        AF026004
                                             CLCN2
                                                          chr3 184063973
## AB033049
                                           ANKRD50
                        AB033049
                                                          chr4 125585207
## AB033050
                         AB033050
                                             ZMIZ1
                                                         chr10 80828792
## AB033086
                        AB033086
                                                                 5808083
                                            NLGN4X
                                                          chrX
## NM_003008
                       NM_003008
                                             SEMG2
                                                         chr20 43850010
chr8Genes <- genes[genes$Chromosome=="chr8",]</pre>
head(chr8Genes)
                            probe HUGO.gene.symbol Chromosome
                                                                   Start
## Contig29827_RC Contig29827_RC
                                             FUT10
                                                          chr8 33228344
## NM_003046
                       NM_003046
                                            SLC7A2
                                                          chr8 17396286
## Contig55940_RC Contig55940_RC
                                             CYHR1
                                                          chr8 145675315
## NM_004133
                                             HNF4G
                                                          chr8 76452203
                       NM_004133
## NM 004374
                       NM 004374
                                             COX6C
                                                          chr8 100890223
## AF052142
                        AF052142
                                                          chr8 102698770
                                             NCALD
chr8GenesOrd <-chr8Genes[order(chr8Genes$Start),]</pre>
head(chr8GenesOrd)
                            probe HUGO.gene.symbol Chromosome
##
## NM_004745
                                            DLGAP2
                       NM_004745
                                                          chr8 1449569
## NM 018941
                       NM 018941
                                              CLN8
                                                          chr8 1711870
## AL117604
                        AL117604
                                              DLC1
                                                          chr8 12940872
## NM 003046
                       NM 003046
                                            SLC7A2
                                                          chr8 17396286
## Contig58301 RC Contig58301 RC
                                            SLC7A2
                                                          chr8 17396286
## NM 000662
                       NM 000662
                                              NAT1
                                                          chr8 18067618
write.table(chr8GenesOrd, "chromosome8.gene.info.txt", sep="\t")
```