Exercise 7

Your Name

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```
evals <- read.delim("gene.expression.txt", stringsAsFactors = FALSE)
evals[1:5,1:5]
dim(evals)
##
                   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
                  -0.307 0.046 -0.139 0.036 -0.154
## AB033049
## AB033050
                  0.582 0.216 0.091 -0.186 -0.156
## AB033086
                  -2.000 0.102 -0.016 -0.358 0.153
## [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
                                            NLGN4X
                                                          {\tt chrX}
                                                                 5808083
## NM_003008
                       NM_003008
                                             SEMG2
                                                        chr20 43850010
dim(genes)
## [1] 498
chr8Genes <- genes[genes$Chromosome=="chr8",]</pre>
head(chr8Genes)
##
                           probe HUGO.gene.symbol Chromosome
                                                                   Start
                                                         chr8 33228344
## Contig29827_RC Contig29827_RC
                                             FUT10
## NM_003046
                       NM_003046
                                            SLC7A2
                                                          chr8 17396286
## Contig55940_RC Contig55940_RC
                                             CYHR1
                                                         chr8 145675315
## NM_004133
                       NM_004133
                                             HNF4G
                                                          chr8 76452203
## NM_004374
                       NM_004374
                                             COX6C
                                                          chr8 100890223
## AF052142
                        AF052142
                                             NCALD
                                                          chr8 102698770
chr8GenesOrd <-chr8Genes[order(chr8Genes$Start),]</pre>
head(chr8GenesOrd)
```

##		probe	<pre>HUGO.gene.symbol</pre>	${\tt Chromosome}$	Start
##	NM_004745	NM_004745	DLGAP2	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")