STAT-S675

Homework 6

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Problem 1

Problem 2

Problem 3

Parts 1 and 2

```
import::from(readr, read_table)
import::from(magrittr, `%>%`, `%<>%`)
source('http://pages.iu.edu/~mtrosset/Courses/675/stress.r')
library(ggplot2)
theme_set(ggthemes::theme_base())
set.seed(6756)
# load the data
# note that it's a lower triangular matrix--fill in the rest
data.url <- 'http://pages.iu.edu/~mtrosset/Courses/675/colors.dat'</pre>
Delta <- read table(data.url, col names = FALSE) %>%
  as.matrix() %>%
  \{. + t(.)\}
# generate random 2 column matrix
rows <- nrow(Delta)</pre>
cols <- 2
stdev <- 40
X.rand <- matrix(rnorm(rows * cols, sd = stdev), nrow = rows, ncol = cols)</pre>
# compute raw stress of the random matrix
mds.stress.raw.eq(X.rand, Delta)
```

[1] 160697

```
# center X.rand
X.rand %<>% apply(2, function(x) x - mean(x))
apply(X.rand, 2, sum)
```

```
[1] -2.309264e-14 6.217249e-15

# and check its stress criterion
mds.stress.raw.eq(X.rand, Delta)
```

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Centering the data matrix did not change the stress criterion. Looking at mds.stress.raw.eq more closely, this is expected, since it just compares the distance matrix of X.rand to Delta, and shifting X.rand doesn't

do anything to its distance matrix.

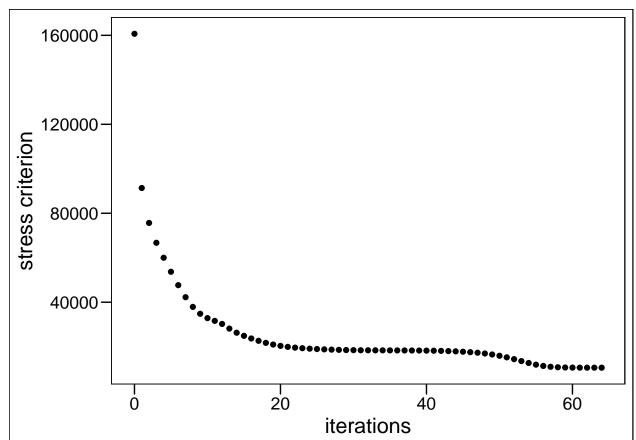
Part 3

```
# iterations
K <- 64
k.vector <- seq(0, K)

# initialize stress vector
stress.vector <- rep(NA, K + 1)
stress.vector[1] <- mds.stress.raw.eq(X.rand, Delta)

# iterate
for (k in k.vector[-1]) {
    X.rand <- mds.guttman.eq(X.rand, Delta)
    stress.vector[k + 1] <- mds.stress.raw.eq(X.rand, Delta)
}

mds.df <- dplyr::data_frame(k = k.vector, random.stress = stress.vector)
ggplot(mds.df) +
    geom_point(aes(x = k, y = random.stress)) +
    labs(x = 'iterations', y = 'stress criterion')</pre>
```



Part 4

```
X.cmds <- cmdscale(Delta)
mds.stress.raw.eq(X.cmds, Delta)
[1] 25880.08</pre>
```

Part 5

```
# initialize
cmds.stress.vector <- rep(NA, K + 1)</pre>
cmds.stress.vector <- mds.stress.raw.eq(X.cmds, Delta)</pre>
# iterate
for (k in k.vector[-1]) {
 X.cmds <- mds.guttman.eq(X.cmds, Delta)</pre>
  cmds.stress.vector[k + 1] <- mds.stress.raw.eq(X.cmds, Delta)</pre>
}
mds.df$cmds.stress <- cmds.stress.vector</pre>
ggplot(mds.df) +
  geom_point(aes(x = k,
                 y = random.stress,
                  colour = 'GMA with random initialization')) +
  geom_point(aes(x = k,
                 y = cmds.stress,
                 colour = 'GMA with CMDS initialization')) +
  scale_colour_brewer(palette = 'Set1') +
  theme(legend.position = 'bottom') +
  labs(x = 'iterations', y = 'stress criterion', colour = NULL)
```

