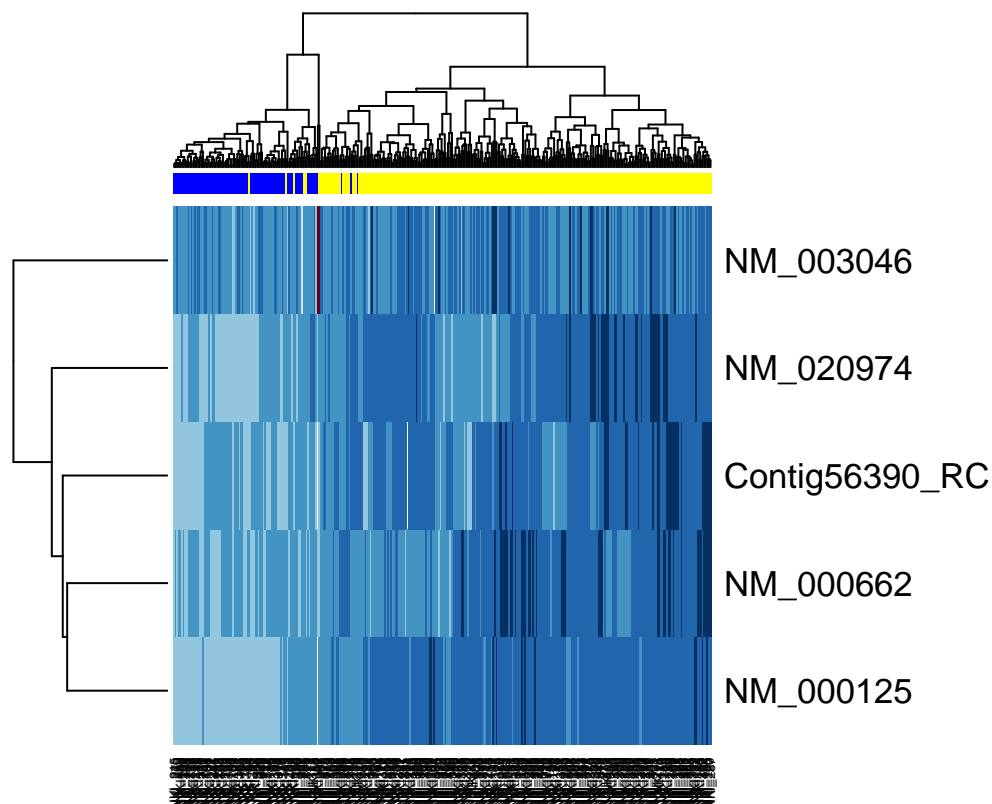# Exercise 8

Your Name

11 Dec 2015

- Modifiy the code chunk so this is run once only

```
genes <- read.delim("gene.description.txt")
subjects <- read.delim("cancer.patients.txt")
evals <- read.delim("gene.expression.txt",stringsAsFactors = FALSE)
```

The number of subjects in the study is 337. The number of genes being measured was 498

- Create a list of R packages and versions

```
sessionInfo()
```

```
## R version 3.2.2 (2015-08-14)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Ubuntu 14.04.3 LTS
##
## locale:
##  [1] LC_CTYPE=en_GB.UTF-8       LC_NUMERIC=C
##  [3] LC_TIME=en_GB.UTF-8        LC_COLLATE=en_GB.UTF-8
##  [5] LC_MONETARY=en_GB.UTF-8    LC_MESSAGES=en_GB.UTF-8
```

```
##  [7] LC_PAPER=en_GB.UTF-8       LC_NAME=C
##  [9] LC_ADDRESS=C               LC_TELEPHONE=C
## [11] LC_MEASUREMENT=en_GB.UTF-8 LC_IDENTIFICATION=C
##
## attached base packages:
## [1] stats     graphics  grDevices utils     datasets  methods   base
##
## other attached packages:
## [1] RColorBrewer_1.1-2
##
## loaded via a namespace (and not attached):
## [1] magrittr_1.5    formatR_1.2.1   tools_3.2.2     htmltools_0.2.6
## [5] yaml_2.1.13     stringi_0.5-5   rmarkdown_0.8   knitr_1.11
## [9] stringr_1.0.0   digest_0.6.8    evaluate_0.8
```