

# 3770 HW6

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## Problem 6.1.6

The sample mean of our data set can be calculated by adding all our values and dividing by the sample size:\

$$\bar{x} = \sum_{i=1}^n x_i \approx 14.35895$$

On the other hand, our standard deviance is simply the squareroot of our variance  $s^2$ :

$$s^2 = \frac{\sum_{i=1}^n (x_i - \mu)^2}{n-1} \approx 356.4716$$

$$s = \sqrt{s^2} = \sqrt{356.4716} \approx 18.88045$$

Thus, we get a mean of approx. 14.35895 and a standard deviation of approx. 18.88045.

```
times = read.table("6-8.txt", header = TRUE)
times = times$time
```

```
mean(times)
```

```
## [1] 14.35895
```

```
var(times)
```

```
## [1] 356.4716
```

```
sqrt(var(times))
```

```
## [1] 18.88045
```

## Problem 6.2.4

The two middle numbers in our data set are both 90.4. Since  $\frac{90.4+90.4}{2} = 90.4$ , our median is thus, 90.4.

```
rating = read.table("6-30.txt", header = TRUE)
rating = rating$Rating
```

```
median(rating)
```

```
## [1] 90.4
```

Our Quartiles:

```
quantiles = quantile(rating,type=6)
quantiles
```

```
##      0%      25%      50%      75%     100%
## 83.400  88.575  90.400  92.200 100.300
```

Our Stem and Leaf Plot:

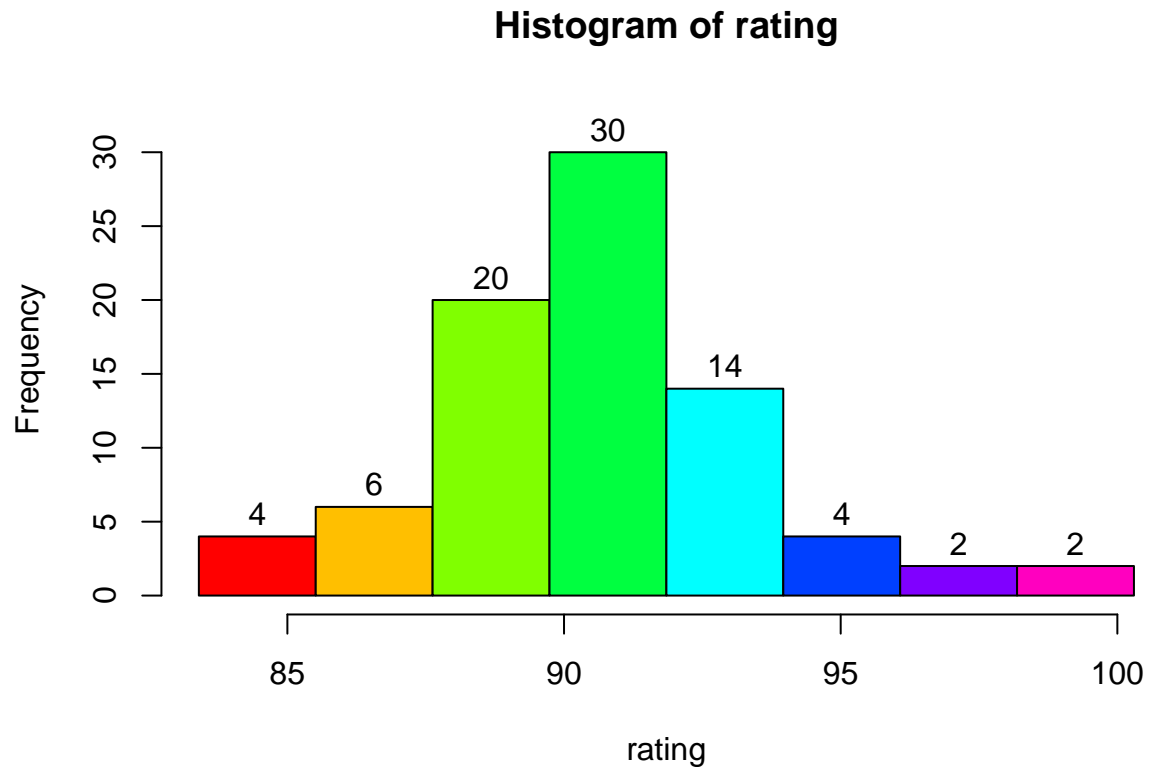
```
stem(rating, scale = 2)
```

```
##
## The decimal point is at the |
##
##      83 | 4
##      84 | 33
##      85 | 3
##      86 | 777
##      87 | 456789
##      88 | 23334556679
##      89 | 0233678899
##      90 | 0111344456789
##      91 | 0001112256688
##      92 | 22236777
##      93 | 023347
##      94 | 2247
##      95 |
##      96 | 15
##      97 |
##      98 | 8
##      99 |
##     100 | 3
```

### Problem 6.3.2

Our Histogram:

```
bin=seq(min(rating),max(rating),by=(max(rating)-min(rating))/8)
freqs = hist(rating, breaks=bin, label=TRUE, right=FALSE, col=rainbow(8), ylim=c(0, 32))
```



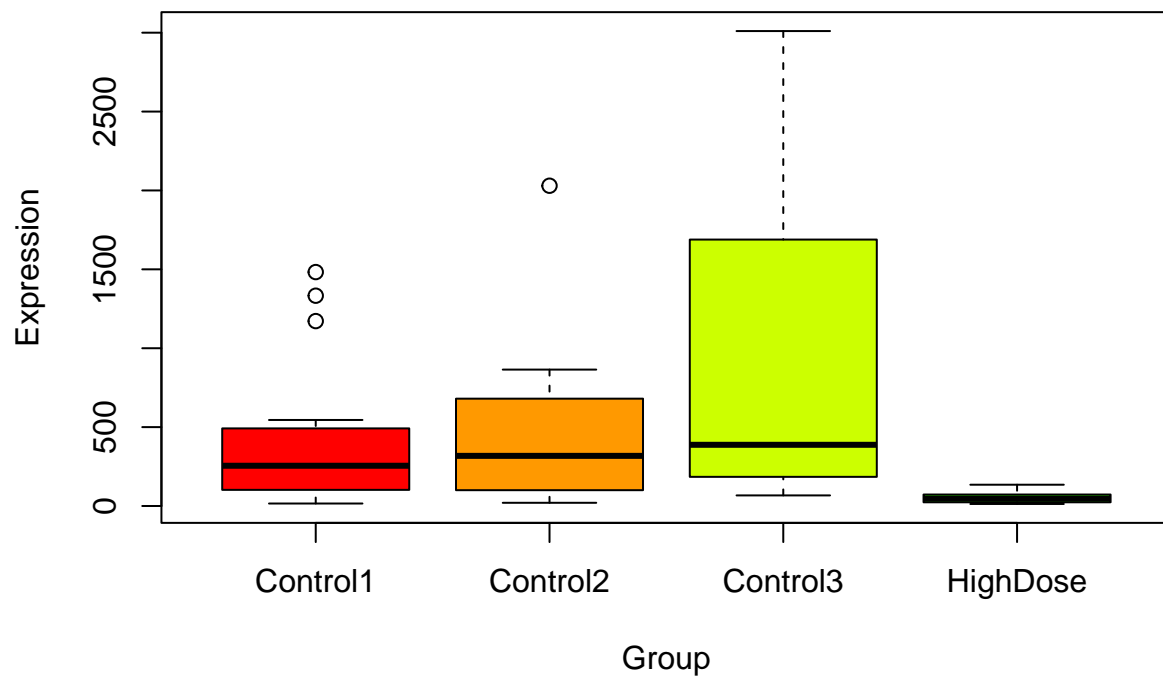
```
freqs
## $breaks
## [1] 83.4000 85.5125 87.6250 89.7375 91.8500 93.9625 96.0750 98.1875
## [9] 100.3000
##
## $counts
## [1] 4 6 20 30 14 4 2 2
##
## $density
## [1] 0.02309136 0.03463703 0.11545678 0.17318516 0.08081974 0.02309136 0.01154568
## [8] 0.01154568
##
## $mids
## [1] 84.45625 86.56875 88.68125 90.79375 92.90625 95.01875 97.13125 99.24375
##
## $xname
## [1] "rating"
##
## $equidist
## [1] TRUE
```

```
##  
## attr("class")  
## [1] "histogram"
```

### Problem 6.4.9

Based on my boxplot below, I think it's hard to definitively tell if the treatment is effective or not in gene expression. Because while it does have the smallest range - and practically no outliers - there's still a ton of difference in variance between the control groups. However, the variance and outliers in the High Dosage group itself definitely displayed a much smaller range compared to the other groups. So if I had to choose a definitive answer, I'd say that the high dosage treatment definitely helped in minimizing gene expression.

```
treatmentData = read.table("6-81.txt", header=TRUE)
attach(treatmentData)
boxplot(Expression~Group, col=rainbow(10))
```



### Problem 6.7.2

Based on the normal distribution line below, it would be reasonable to assume that the octane rating follows a normal distribution. This is because most of our ratings/points follow the expected normal distribution line.

```
qqnorm(rating)
qqline(rating, col="red")
```

