

R Module 6 Rubric

Question 1: Boxplot

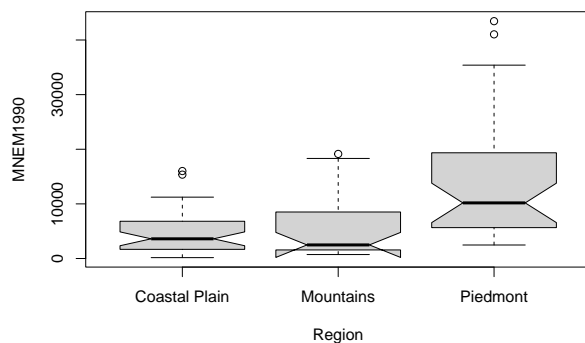
Notched boxplot of your regions

Base R:

```
library(sf)
NC <- read_sf("data/NC_REGION.shp")

boxplot(MNEM1990 ~ Region, data = NC, notch = TRUE)
```

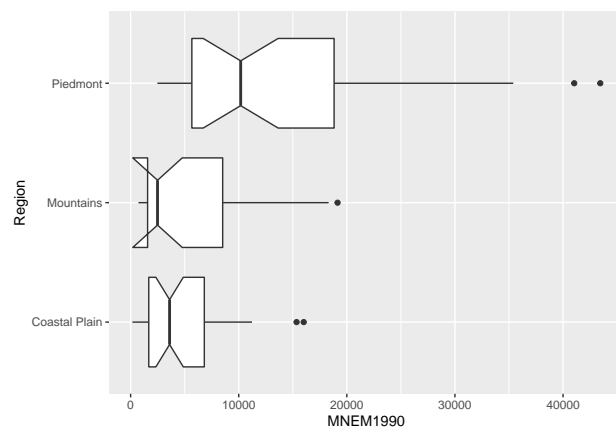
```
## Warning in (function (z, notch = FALSE, width = NULL, varwidth = FALSE, : some notches went outside
## hinges ('box'): maybe set notch=FALSE
```



ggplot2:

```
library(ggplot2)

ggplot(NC, aes(x = MNEM1990, y = Region)) +
  geom_boxplot(notch = TRUE)
```



Question 2: ANOVA

Report your F-Statistic and if you should reject or fail to reject the null hypothesis

```
anova <- aov(MNEM1990 ~ Region, data = NC)
```

```
print(anova)
```

```
## Call:
## aov(formula = MNEM1990 ~ Region, data = NC)
##
## Terms:
##              Region  Residuals
## Sum of Squares 2127131896 6155559536
## Deg. of Freedom      2      97
##
## Residual standard error: 7966.139
## Estimated effects may be unbalanced
```

```
summary(anova)
```

```
##              Df      Sum Sq   Mean Sq F value   Pr(>F)
## Region         2 2.127e+09 1.064e+09   16.76 5.6e-07 ***
## Residuals     97 6.156e+09 6.346e+07
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Question 3: oneway.test, Equal Variance

```
oneway.test(MNEM1990 ~ Region, data = NC, var.equal = TRUE)
```

```
##
## One-way analysis of means
##
## data: MNEM1990 and Region
## F = 16.76, num df = 2, denom df = 97, p-value = 5.6e-07
```

Question 4: Bartlett K-Squared

```
bartlett.test(MNEM1990 ~ Region, data = NC)
```

```
##
## Bartlett test of homogeneity of variances
##
## data: MNEM1990 by Region
## Bartlett's K-squared = 38.195, df = 2, p-value = 5.083e-09
```

Question 5: oneway.test, Unequal Variance

```
oneway.test(MNEM1990 ~ Region, data = NC, var.equal = FALSE)
```

```
##
## One-way analysis of means (not assuming equal variances)
##
## data: MNEM1990 and Region
## F = 11.66, num df = 2.000, denom df = 47.748, p-value = 7.526e-05
```

Question 6: Shapiro-Wilk Normality Test

Student's don't need to use dplyr or even transform the data, as long as the distribution is normal.

```
library(dplyr)
NC <- NC %>%
  mutate(
    sqrt_MNEM1990 = sqrt(MNEM1990)
  )
```

```
shapiro.test(NC$sqrt_MNEM1990)
```

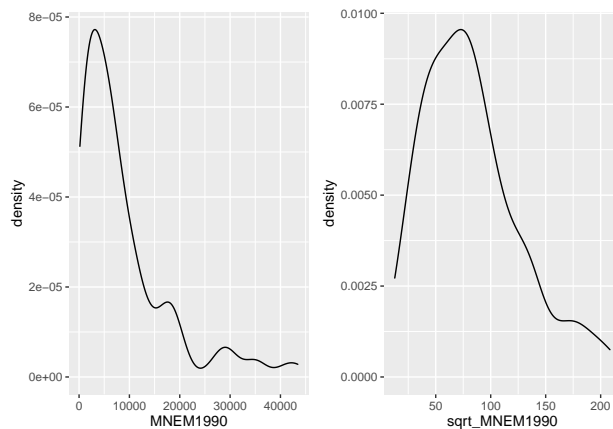
```
##
##  Shapiro-Wilk normality test
##
## data:  NC$sqrt_MNEM1990
## W = 0.94203, p-value = 0.0002569
```

Illustrate data using density plot for both un-transformed and transformed data

```
unt <- ggplot(NC, aes(x = MNEM1990)) +
  geom_density()
```

```
tra <- ggplot(NC, aes(x = sqrt_MNEM1990)) +
  geom_density()
```

This is just to get two plots side-by-side
cowplot::plot_grid(unt, tra)



Question 7: Tukey Test

```
aov <- aov(sqrt_MNEM1990 ~ Region, data = NC)
```

```
tukey <- TukeyHSD(aov)
tukey
```

```
##  Tukey multiple comparisons of means
##    95% family-wise confidence level
##
```

```
## Fit: aov(formula = sqrt_MNEM1990 ~ Region, data = NC)
##
## $Region
```

	diff	lwr	upr	p adj
Mountains-Coastal Plain	2.87440	-20.29441	26.04321	0.9530922
Piedmont-Coastal Plain	49.85867	29.54578	70.17157	0.0000002
Piedmont-Mountains	46.98427	23.24429	70.72426	0.0000244

```
plot(tukey)
```

