MANOVA

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attach data:

```
Skull_data <- read.csv(file.choose(),header = T)
attach(Skull_data)
names(Skull_data)</pre>
```

```
## [1] "X" "epoch" "mb" "bh" "bl" "nh"
```

check normality:

```
library(moments)
skewness(Skull_data[,3:6])
```

```
## mb bh bl nh
## -0.02837297 -0.17603322 0.13948131 0.08255358
```

accepted range from -1 to +1

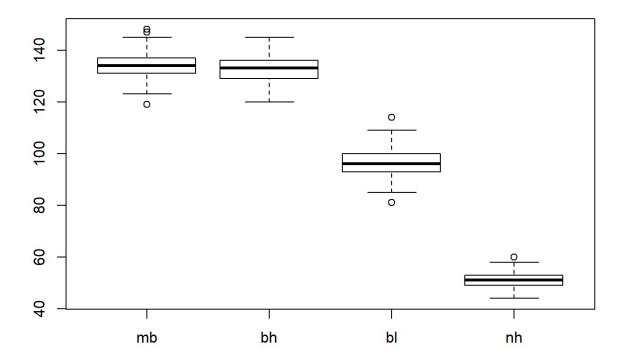
```
kurtosis(Skull_data[,3:6])
```

```
## mb bh bl nh
## 3.259550 2.889839 3.185601 2.830727
```

accepted range from -2 to +2, may to +3

```
boxplot(Skull_data[,3:6],main= "Diff. in Skull dimensions")
```

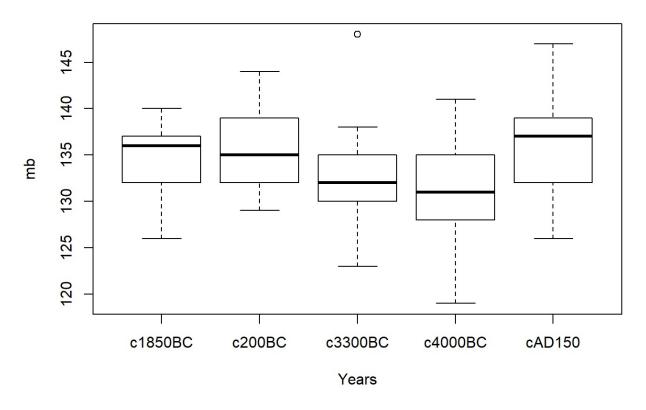
Diff. in Skull dimensions



data is normally distributed

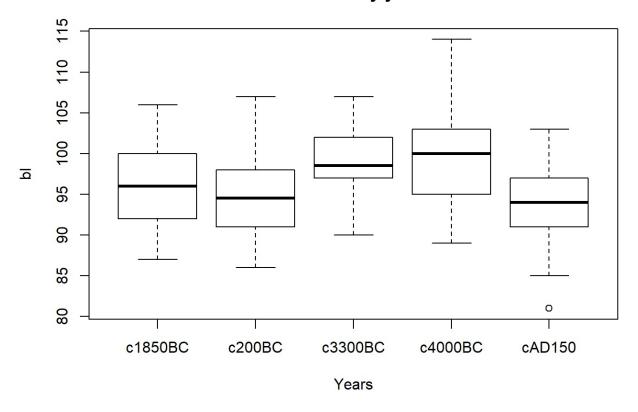
```
boxplot(mb~epoch,main="Diff. in mb by years",xlab = "Years")
```

Diff. in mb by years



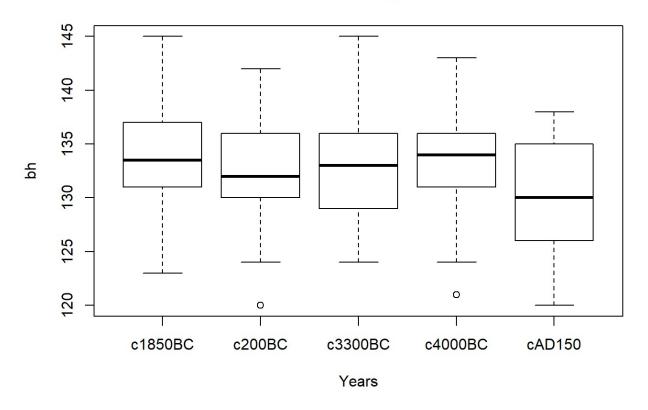
boxplot(bl~epoch,main="Diff. in bl by years",xlab = "Years")

Diff. in bl by years



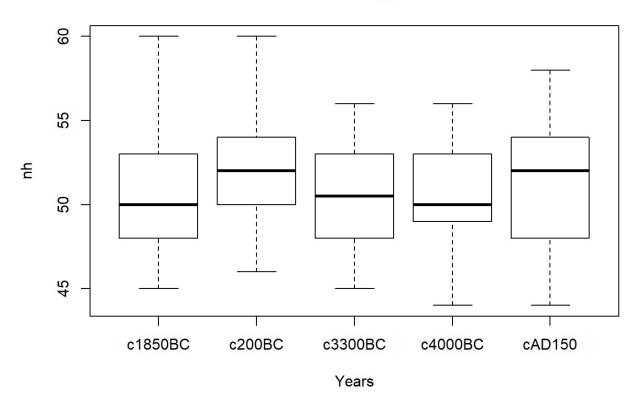
boxplot(bh~epoch,main="Diff. in bh by years",xlab = "Years")

Diff. in bh by years



boxplot(nh~epoch,main="Diff. in nh by years",xlab = "Years")

Diff. in nh by years



test: H0: diffrence in mean is the same in all years:

```
MANOVA1 <- manova(formula=(cbind(mb,bh,bl,nh))~as.factor(epoch) , data=Skull_data)
summary(MANOVA1)</pre>
```

```
## Df Pillai approx F num Df den Df Pr(>F)

## as.factor(epoch) 4 0.35331 3.512 16 580 4.675e-06 ***

## Residuals 145

## ---

## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

p-value < 0.05, reject H0, Skull dimensions differ by years

to know which is affected separetly:

```
summary.aov(MANOVA1)
```

```
## Response mb:
                  Df Sum Sq Mean Sq F value Pr(>F)
## as.factor(epoch) 4 502.83 125.707 5.9546 0.0001826 ***
## Residuals 145 3061.07 21.111
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
  Response bh :
                 Df Sum Sq Mean Sq F value Pr(>F)
## as.factor(epoch) 4 229.9 57.477 2.4474 0.04897 *
## Residuals 145 3405.3 23.485
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
  Response bl :
                 Df Sum Sq Mean Sq F value
## as.factor(epoch) 4 803.3 200.823 8.3057 4.636e-06 ***
## Residuals 145 3506.0 24.179
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Response nh:
                  Df Sum Sq Mean Sq F value Pr(>F)
## as.factor(epoch) 4 61.2 15.300
                                    1.507 0.2032
## Residuals 145 1472.1 10.153
```

mb & bl are highly significant bh is marginally significant nh is not significant