

# ISYE 6414 - HW4

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## Antler Length Data

In [1]:

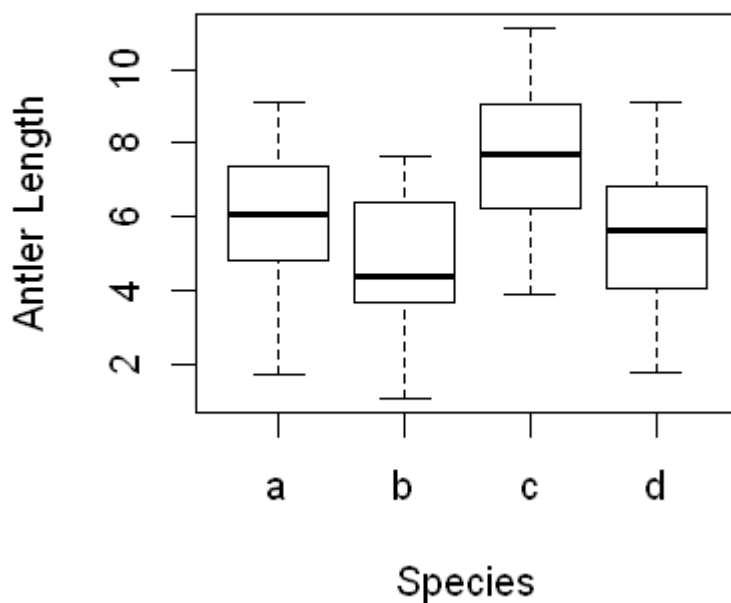
```
data = read.csv('homework04data01.csv')
```

Q1.

In [5]:

```
options(repr.plot.width = 3.5, repr.plot.height = 3.5)
boxplot(Length~Species,
data=data,
main="Different boxplots for each species",
xlab="Species",
ylab="Antler Length")
```

## Different boxplots for each specie



As we can see from the box plots, there is overlap between all 4 of the species, and it is hard to tell if the antler lengths of the four species are different. Although their means are clearly different, their variability is high.

**Q2.**

In [3]:

```
model = aov(Length ~ Species, data=data)
```

In [4]:

```
summary(model)
```

```

              Df Sum Sq Mean Sq F value    Pr(>F)
Species        3   90.0   30.001    8.121 8.58e-05 ***
Residuals     81  299.2    3.694
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

In [10]:

```
model.tables(model, type = 'means')
```

Tables of means

Grand mean

6.014706

```

Species
      a      b      c      d
6.063 4.697 7.594 5.445
rep 20.000 15.000 22.000 28.000

```

$$H_o : \mu_1 = \mu_2 = \dots = \mu_k$$

 $H_a$  = some means are different

$$F_0 = \frac{MST_R}{MSE} = \frac{30.001}{3.694} = 8.1215$$

From tables,  $F_{\alpha, k-1, N-k} = F_{0.05, 3, 81} = 2.7173$

$F_o > \text{critical } F \implies H_o$  can be rejected. Hence, some means are different.

**Q3.**

From the model summary:

SSE = 299.2

SSTr = 90.0

MSE = 3.694

MSTr = 30.001

**Q4.**

In [5]:

```
TukeyHSD(model)
```

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

```
Fit: aov(formula = Length ~ Species, data = data)
```

```
$Species
      diff      lwr      upr    p adj
b-a -1.365667 -3.08777369  0.3564404 0.1683081
c-a  1.5306364 -0.02706805  3.0883408 0.0559481
d-a -0.6176429 -2.09373459  0.8584489 0.6920003
c-b  2.8963030  1.20807683  4.5845292 0.0001303
d-b  0.7480238 -0.86520632  2.3612539 0.6183736
d-c -2.1482792 -3.58469904 -0.7118594 0.0010291
```

c-b and d-c imply a difference in average antler length between species.

**Hospital Data**

In [6]:

```
data1 = read.csv('homework04Hospital.csv')
```

**Q5.**

In [8]:

```
model1 = lm(Hours~.,data = data1)
```

In [9]:

summary(model1)

Call:

lm(formula = Hours ~ ., data = data1)

Residuals:

	Min	1Q	Median	3Q	Max
	-677.23	-270.19	60.93	228.32	517.70

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	1946.80204	504.18193	3.861	0.00226	**
Xray	0.03858	0.01304	2.958	0.01197	*
BedDays	1.03939	0.06756	15.386	2.91e-09	***
Length	-413.75780	98.59828	-4.196	0.00124	**

---

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

Residual standard error: 387.2 on 12 degrees of freedom

Multiple R-squared: 0.9961, Adjusted R-squared: 0.9952

F-statistic: 1028 on 3 and 12 DF, p-value: 9.919e-15

$$\beta_O = 1946.80204$$

$$\beta_{Xray} = 0.03858$$

$$\beta_{BedDays} = 1.03939$$

$$\beta_{Length} = -413.75780$$

**Q6.**

When BedDays and Length are held constant, for every unit increase in Xray, Hours increase by 0.03858.

When Xray and Length are held constant, for every unit increase in BedDays, Hours increase by 1.03939.

When BedDays and Xray are held constant, for every unit increase in Length, Hours decrease by 413.75780.

 $B_o = 1946.80204$  is the mean value of Hours when all the other values are 0.**Q7.**For BedDays,  $p\text{-value} = 2.91e-09 \ll \alpha = 0.05 \implies$  BedDays is statistically significant

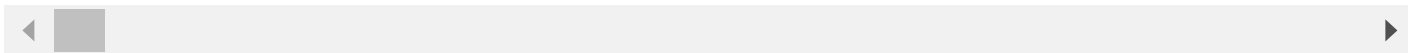
$$H_0 : \beta_{BedDays} = 0$$

$$H_a : \beta_{BedDays} \neq 0$$

$$t\text{-value} = 15.386$$

$$\text{critical } t = t_{0.025, 12} = 2.179$$

 $|t\text{-value}| > \text{critical } t \implies H_0$  can be rejected  $\implies$  BedDays is statistically significant.

**Q8.**

For Xray,  $p - value = 0.01197$

With  $\alpha = 0.05$ , Xray is statistically significant as  $p - value < \alpha$ .

But with  $\alpha = 0.01$ , Xray is not statistically significant as  $p - value > \alpha$ .

**Q9.**

For Length,  $p - value = 0.00124$

With  $\alpha = 0.05$ , Length is statistically significant as  $p - value < \alpha$ .

With  $\alpha = 0.10$ , Length is statistically significant as  $p - value < \alpha$ .