

# Stat 306: Finding Relationships in Data.

## Lecture 9

Section 3.3 + Section 3.6 (recap) + Section 3.7  
+ Section 3.8 + Section 3.10

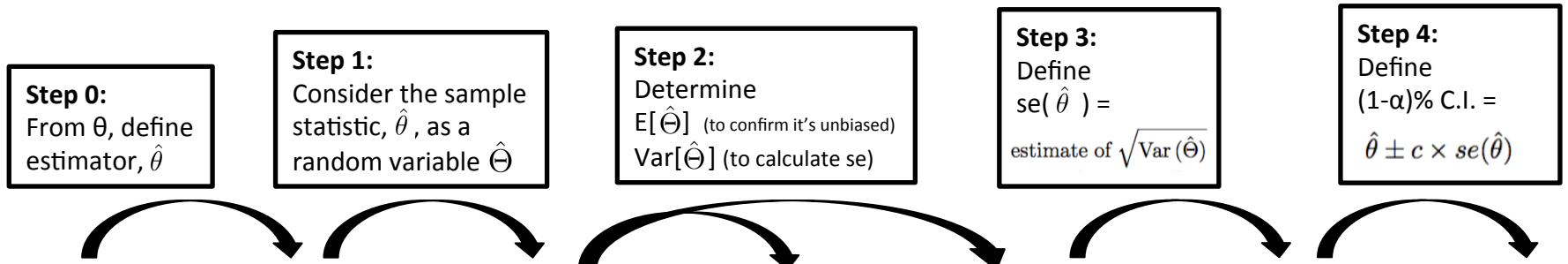
## 3.3 Statistical model for multiple regression

The main assumption of linear regression is that the outcomes,  $Y_i$ , (for  $i = 1, \dots, n$ ), are independently normally distributed. This follows directly from our model:

$$Y_i = \beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip} + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2) \text{ independently.}$$

(3.36)

# 3.6 Interval estimates and standard errors



| Population parameter or "something we would like to estimate" | Sample statistic ("estimator")   | Estimator as a Random Variable   | Expected Value of the estimator   | Variance of the estimator   | Standard Error of estimator   | Confidence Interval        |
|---|--|--|-----------------------------------|---|---|----------------------------|
| $\beta$   | <b>b</b> <b>1.</b><br>$= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$ | <b>B</b> ~ <b>2.</b><br>$\mathbf{N}(\beta, \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1})$ | $E[\mathbf{b}] = \beta$ <b>3.</b> | $\text{Var}[\mathbf{B}]$ <b>4.</b><br>$= \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1}$ | $se(\mathbf{b})$ <b>5.</b><br>$= \hat{\sigma} \sqrt{[(\mathbf{X}^T \mathbf{X})^{-1}]_{jj}}$ | C.I. for $\beta$ <b>6.</b> |
| $\sigma^2$  | $s^2$ or MS(Res)   | $S^2$  | $E[S^2]$                          | $\text{Var}[S^2]$   | $se(s^2)$   | C.I. for $\sigma^2$        |
| $\mu_Y(\mathbf{x})$   | $(\hat{\mu}_Y(x))$   | $(\hat{\mu}_Y(x))$   | $E(\hat{\mu}_Y(x))$               | $\text{Var}(\hat{\mu}_Y(x))$  | $se(\hat{\mu}_Y(x))$  | C.I. for $\mu_Y(x)$        |

# 1.

We obtain the estimator that minimize the Sum of squares by simple matrix calculus:

$$SS(\mathbf{b}) = (\mathbf{y} - \mathbf{X}\mathbf{b})^2 = (\mathbf{y} - \mathbf{X}\mathbf{b})^T(\mathbf{y} - \mathbf{X}\mathbf{b})$$

$$\frac{\delta SS(\mathbf{b})}{\delta \mathbf{b}} = 0$$

See 3.29 and 3.30

$\Rightarrow$

(3.10)

$$(\mathbf{X}^T \mathbf{X}) \hat{\mathbf{b}} = \mathbf{X}^T \mathbf{y}$$

(3.11)

$$\text{or } \hat{\mathbf{b}} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}.$$

Equation (3.11) assumes that  $\mathbf{X}^T \mathbf{X}$  is a non-singular matrix so that its inverse is defined. The discussion of a condition for non-singularity is given in Section 3.11.

# 2.

- **Thing 1:**

- Linear combinations of independent normal random variables also have normal distributions! (see Appendix B)



or stated explicitly in terms of vectors and matrices:

- If  $\mathbf{Y} \sim \mathcal{N}_n(\boldsymbol{\mu}, \boldsymbol{\Sigma})$ , then if we have a linear combination of  $\mathbf{Y}$ ,  $\mathbf{CY}$ , where  $\mathbf{C}$  is  $(q \times n)$ , then  $\mathbf{CY} \sim \mathcal{N}_q(\mathbf{C}\boldsymbol{\mu}, \mathbf{C}\boldsymbol{\Sigma}\mathbf{C}')$ .

Then we have that  $\mathbf{B}$  is normally distributed random vector, since:

From (3.11), with  $\hat{\mathbf{B}} = \hat{\boldsymbol{\beta}}$  as a random vector, and  $k = p + 1$  as the dimension of  $\hat{\boldsymbol{\beta}}$ ,

$$(3.66) \quad \hat{\mathbf{B}} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y} = \mathbf{A} \mathbf{Y},$$

$$(3.67) \quad \mathbf{A} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T = \begin{pmatrix} \mathbf{a}_0^T \\ \mathbf{a}_1^T \\ \vdots \\ \mathbf{a}_p^T \end{pmatrix},$$

$$(3.68) \quad (k \times n) \quad (k \times k) \quad (k \times n)$$

# 3. Is the least squares estimator for $\beta$ unbiased? Does $E[\hat{\mathbf{b}}] = \beta$ ?

**Proof that  $\hat{\beta}$  is an unbiased estimator of  $\beta$ .**

We know from earlier that  $\hat{\beta} = (X'X)^{-1}X'y$  and that  $y = X\beta + \epsilon$ . This means that

$$\begin{aligned}\hat{\beta} &= (X'X)^{-1}X'(X\beta + \epsilon) \\ \hat{\beta} &= \beta + (X'X)^{-1}X'\epsilon\end{aligned}\tag{24}$$

since  $(X'X)^{-1}X'X = I$ . This shows immediately that OLS is unbiased so long as either (i)  $X$  is fixed (non-stochastic) so that we have:

$$\begin{aligned}E[\hat{\beta}] &= E[\beta] + E[(X'X)^{-1}X'\epsilon] \\ &= \beta + (X'X)^{-1}X'E[\epsilon]\end{aligned}\tag{25}$$

where  $E[\epsilon] = 0$  by assumption or (ii)  $X$  is stochastic but independent of  $\epsilon$  so that we have:

$$\begin{aligned}E[\hat{\beta}] &= E[\beta] + E[(X'X)^{-1}X'\epsilon] \\ &= \beta + (X'X)^{-1}E[X'\epsilon]\end{aligned}\tag{26}$$

where  $E(X'\epsilon) = 0$ .

# 4.

We have that:

$$\text{Var}(B) = \text{Var}(AY)$$

$$\text{Var}(B) = A \text{Var}(Y) A^T$$

More information: [https://en.wikipedia.org/wiki/Covariance\\_matrix#Generalization\\_of\\_the\\_variance](https://en.wikipedia.org/wiki/Covariance_matrix#Generalization_of_the_variance)

We also have that the variance-covariance matrix of **Y (random vector of length n)** is  $\text{Var}(Y) = \sigma^2 I_n$ , where  $I_n$  is the n by n identity matrix.

Therefore, following equations 3.69 – 3.72, we have:

$$\text{Var}(B) =$$

$$(3.73) \quad \begin{pmatrix} \text{Var}(\hat{B}_0) & \text{Cov}(\hat{B}_0, \hat{B}_1) & \cdots & \text{Cov}(\hat{B}_0, \hat{B}_p) \\ \text{Cov}(\hat{B}_1, \hat{B}_0) & \text{Var}(\hat{B}_1) & \cdots & \text{Cov}(\hat{B}_1, \hat{B}_p) \\ \vdots & \vdots & \ddots & \vdots \\ \text{Cov}(\hat{B}_p, \hat{B}_0) & \text{Cov}(\hat{B}_p, \hat{B}_1) & \cdots & \text{Var}(\hat{B}_p) \end{pmatrix} = \sigma^2 \begin{pmatrix} \mathbf{a}_0^T \mathbf{a}_0 & \mathbf{a}_0^T \mathbf{a}_1 & \cdots & \mathbf{a}_0^T \mathbf{a}_p \\ \mathbf{a}_1^T \mathbf{a}_0 & \mathbf{a}_1^T \mathbf{a}_1 & \cdots & \mathbf{a}_1^T \mathbf{a}_p \\ \vdots & \vdots & \ddots & \vdots \\ \mathbf{a}_p^T \mathbf{a}_0 & \mathbf{a}_p^T \mathbf{a}_1 & \cdots & \mathbf{a}_p^T \mathbf{a}_p \end{pmatrix}$$

$$(3.74) \quad = \sigma^2 \begin{pmatrix} \mathbf{a}_0^T \\ \mathbf{a}_1^T \\ \vdots \\ \mathbf{a}_p^T \end{pmatrix} (\mathbf{a}_0 \quad \cdots \quad \mathbf{a}_p) = \sigma^2 \mathbf{A} \mathbf{A}^T$$

$$(3.75) \quad = \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{X} (\mathbf{X}^T \mathbf{X})^{-1}$$

$$(3.76) \quad = \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1} \stackrel{\text{def}}{=} \Sigma_{\hat{\beta}}$$

# 5.

Since a standard error is defined as an estimated square root of the variance of an estimator,

$$(3.77) \quad se(\hat{\beta}_j) = \hat{\sigma} \sqrt{[(\mathbf{X}^T \mathbf{X})^{-1}]_{jj}}, \quad j = 0, 1, \dots, p.$$



# 6.

For 95% confidence intervals for  $\beta$ 's or subpopulation means, or for 95% prediction intervals, the appropriate SE is multiplied by  $t_{n-k,0.975}$  to get the margin of error to add/subtract from the point estimate.

95% Confidence Interval:

$$\mathbf{b} \pm t_{n-k,0.975} \text{ se}(\mathbf{b})$$

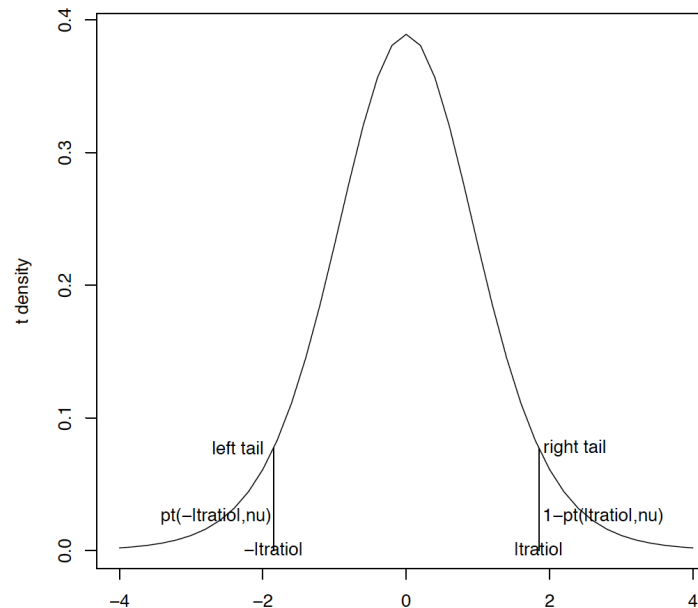
```
> betahat[1] - qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[1,1])
[1] 1.96994
> betahat[1] + qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[1,1])
[1] 44.56221
>
> betahat[2] - qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[2,2])
[1] 0.1792634
> betahat[2] + qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[2,2])
[1] 1.183658
>
> betahat[3] - qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[3,3])
[1] -0.6870086
> betahat[3] + qt(0.975, n-k)*sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[3,3])
[1] 0.1327291
>
> confint(lm(y~x1+x2))
              2.5 %      97.5 %
(Intercept) 1.9699396 44.5622139
x1           0.1792634  1.1836579
x2          -0.6870086  0.1327291
```

# 7. p-values...

Using the R function  $\text{pt}(\cdot, \nu)$  for the cdf of a  $t_\nu$  random variable, the 2-sided P-value is  $2[1 - \text{pt}(|\text{tratio}|, \nu)]$  (see Figure 3.2).

Common thresholds for P-values are 0.10, 0.05, 0.01 and 0.001. Let  $t_j = \hat{\beta}_j / \text{se}(\hat{\beta}_j)$  be the t-ratio for variable  $x_j$ .

Figure 3.2: P-value based on  $t_\nu$  density; by symmetry, the left-tail area and right-tail area are the same. The sum of the two tail areas is the P-value.



# 7. p-values...

```
> x1 <- c(82, 45, 71, 22, 29, 9, 12, 18, 24)
> x2 <- c(26, 49, 76, 37, 40, 0, 2, 10, 92)
> y <- c(71, 54, 43, 45, 21, 11, 30, 45, 10)
> n <- 9
> x1bar <- (1/n)*sum(x1)
> sx1 <- sqrt( sum((x1-x1bar)^2)/(n-1) )
>
> x2bar <- (1/n)*sum(x2)
> sx2 <- sqrt( sum((x2-x2bar)^2)/(n-1) )
>
> ybar <- (1/n)*sum(y)
> sy <- sqrt( sum((y-ybar)^2)/(n-1) )
>
> sx1y <- (1/(n-1))*sum((x1-x1bar)*(y-ybar))
> rx1y <- sx1y/(sx1*sy)
>
> sx2y <- (1/(n-1))*sum((x2-x2bar)*(y-ybar))
> rx2y <- sx2y/(sx2*sy)
>
> X <- cbind(1,x1,x2)
> y <- cbind(y)
.
```

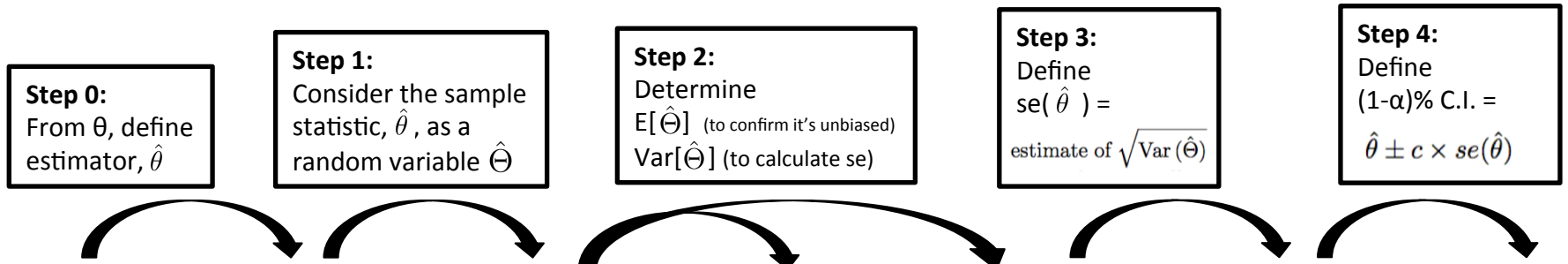
# 7. p-values...

```
> betahat <- solve(t(X) %*% X) %*% t(X) %*% y
> betahat
              y
23.2660767
x1  0.6814606
x2 -0.2771398
>
> yhat <- X%*%betahat
> k <- dim(X)[2]
> p <- k-1
> residuals <- yhat-y
> SS_Res <- sum(residuals^2)
> SS_Res
[1] 1159.452
> MS_Res <- SS_Res/(n-k)
```

# 7. p-values...

```
> se_betahat0<-sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[1,1])
> se_betahat1<-sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[2,2])
> se_betahat2<-sqrt(MS_Res)*sqrt(solve(t(X)%*%X)[3,3])
>
> tratio0 <- betahat[1]/se_betahat0
> tratio1 <- betahat[2]/se_betahat1
> tratio2 <- betahat[3]/se_betahat2
>
> # two-sided p-values:
> 2*(1-pt(abs(tratio0), n-k))
[1] 0.0368661
> 2*(1-pt(abs(tratio1), n-k))
[1] 0.01599726
> 2*(1-pt(abs(tratio2), n-k))
[1] 0.1491044
```

# 3.6 Interval estimates and standard errors



| Population parameter or "something we would like to estimate" | Sample statistic ("estimator")   | Estimator as a Random Variable   | Expected Value of the estimator   | Variance of the estimator   | Standard Error of estimator   | Confidence Interval        |
|---|--|--|-----------------------------------|---|---|----------------------------|
| $\beta$   | <b>b</b> <b>1.</b><br>$= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$ | <b>B</b> ~ <b>2.</b><br>$\mathbf{N}(\beta, \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1})$ | $E[\mathbf{b}] = \beta$ <b>3.</b> | $\text{Var}[\mathbf{B}]$ <b>4.</b><br>$= \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1}$ | $se(\mathbf{b})$ <b>5.</b><br>$= \hat{\sigma} \sqrt{[(\mathbf{X}^T \mathbf{X})^{-1}]_{jj}}$ | C.I. for $\beta$ <b>6.</b> |
| $\sigma^2$  | $s^2$ or MS(Res) <b>1.</b>   | $S^2$ <b>2.</b>  | $E[S^2]$ <b>3.</b>                | $\text{Var}[S^2]$   | $se(s^2)$   | C.I. for $\sigma^2$        |
| $\mu_Y(\mathbf{x})$   | $(\hat{\mu}_Y(x))$   | $(\hat{\mu}_Y(x))$   | $E(\hat{\mu}_Y(x))$               | $\text{Var}(\hat{\mu}_Y(x))$  | $se(\hat{\mu}_Y(x))$  | C.I. for $\mu_Y(x)$        |

# 1.

MS(Res) is our estimator for  $\sigma^2$

- Sum of squares of residuals

$$(3.41) \quad SS(Res) = \sum_{i=1}^n e_i^2 = \sum_{i=1}^n (y_i - \hat{y}_i)^2.$$

- Mean square of residuals or estimated  $\sigma^2$ :

$$(3.42) \quad \hat{\sigma}^2 = (n - k)^{-1} \sum_{i=1}^n e_i^2 = (n - k)^{-1} \sum_{i=1}^n (e_i - \bar{e})^2 = \frac{SS(Res)}{(n - k)} = MS(Res).$$

The residual standard deviation (called residual standard error in R output) is the sample standard deviation of the residuals with a denominator of  $n - k$  instead of  $n - 1$ . A mathematical explanation of this denominator is given in Section 3.7. A property of the residuals after a least squares fit is that

$$(3.43) \quad \bar{e} = n^{-1} \sum_{i=1}^n e_i = 0$$

—————  
↖  
**Typo: should be n-2**

# 3.

MS(Res) is an unbiased estimator for  $\sigma^2$ .

This is explained in Section 3.7

## 3.7 Denominator of the residual SD

The quantity  $\hat{\sigma}^2$  in (3.42), also known as the mean square of residuals, in matrix form is:

$$(3.92) \quad MS(Res) = \hat{\sigma}^2 = \frac{\mathbf{e}^T \mathbf{e}}{n - k} = \frac{SS(Res)}{n - k},$$

where  $e_i$  is defined in (3.40) and  $\mathbf{e} = (e_1, \dots, e_n)^T$  is the column vector of residuals. From (3.39) and (3.40),  $e_i = y_i - \hat{\beta}_0 - \hat{\beta}_1 x_{i1} - \dots - \hat{\beta}_p x_{ip}$  so that

$$(3.93) \quad \mathbf{e} = \mathbf{y} - \mathbf{X}\hat{\boldsymbol{\beta}}$$

$$(3.94) \quad = \mathbf{y} - \mathbf{X}(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y} \quad \underline{\hspace{10em}}$$

$$(3.95) \quad = (\mathbf{I}_n - \mathbf{X}(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T) \mathbf{y}. \quad \underline{\hspace{10em}}$$

Let

$$(3.96) \quad \mathbf{P} = \mathbf{X}(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T = (P_{ij})_{1 \leq i, j \leq n}.$$



# 3.

MS(Res) is an unbiased estimator for  $\sigma^2$ .

This is explained in Section 3.7

## 3.7 Denominator of the residual SD

This is called the *projection matrix* in Section 4.3. Note that  $\mathbf{P}^T = \mathbf{P}$  and

$$(3.97) \quad \mathbf{P}^2 = \mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T\mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T$$

$$(3.98) \quad = \mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T = \mathbf{P};$$

a matrix whose square is itself is said to be *idempotent*. Check that  $(\mathbf{I}_n - \mathbf{P})^T = \mathbf{I}_n - \mathbf{P}$  and that (3.98) implies  $(\mathbf{I}_n - \mathbf{P})^2 = \mathbf{I}_n - \mathbf{P}$ .

Now, we can compute the expected value of  $SS(Res)$  as a random variable, with  $\mathbf{E}$  as the random vector version of the residual vector  $\mathbf{e}$ :

$$(3.99) \quad E[SS(Res)] = E[\mathbf{E}^T\mathbf{E}] = E[\mathbf{Y}^T(\mathbf{I}_n - \mathbf{P})(\mathbf{I}_n - \mathbf{P})\mathbf{Y}] \quad \underline{\hspace{10em}}$$

$$(3.100) \quad = E[\mathbf{Y}^T(\mathbf{I}_n - \mathbf{P})\mathbf{Y}] \quad \underline{\hspace{10em}}$$

$$(3.101) \quad = E[(\mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon})^T(\mathbf{I}_n - \mathbf{P})(\mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon})] \quad \underline{\hspace{10em}}$$

$$(3.102) \quad = E[\boldsymbol{\beta}^T\mathbf{X}^T(\mathbf{I}_n - \mathbf{P})\mathbf{X}\boldsymbol{\beta} + 2\boldsymbol{\beta}^T\mathbf{X}^T(\mathbf{I}_n - \mathbf{P})\boldsymbol{\epsilon} + \boldsymbol{\epsilon}^T(\mathbf{I}_n - \mathbf{P})\boldsymbol{\epsilon}] \quad \underline{\hspace{10em}}$$

$$(3.103) \quad = E[\boldsymbol{\epsilon}^T(\mathbf{I}_n - \mathbf{P})\boldsymbol{\epsilon}] \quad \underline{\hspace{10em}}$$

# 3.

MS(Res) is an unbiased estimator for  $\sigma^2$ .

This is explained in Section 3.7

## 3.7 Denominator of the residual SD

$$(3.99) \quad E[SS(Res)] = E[\mathbf{E}^T \mathbf{E}] = E[\mathbf{Y}^T (\mathbf{I}_n - \mathbf{P})(\mathbf{I}_n - \mathbf{P}) \mathbf{Y}] \quad \underline{\hspace{10em}}$$

$$(3.100) \quad = E[\mathbf{Y}^T (\mathbf{I}_n - \mathbf{P}) \mathbf{Y}] \quad \underline{\hspace{10em}}$$

$$(3.101) \quad = E[(\mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon})^T (\mathbf{I}_n - \mathbf{P})(\mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon})] \quad \underline{\hspace{10em}}$$

$$(3.102) \quad = E[\boldsymbol{\beta}^T \mathbf{X}^T (\mathbf{I}_n - \mathbf{P}) \mathbf{X} \boldsymbol{\beta} + 2\boldsymbol{\beta}^T \mathbf{X}^T (\mathbf{I}_n - \mathbf{P}) \boldsymbol{\epsilon} + \boldsymbol{\epsilon}^T (\mathbf{I}_n - \mathbf{P}) \boldsymbol{\epsilon}] \quad \underline{\hspace{10em}}$$

$$(3.103) \quad = E[\boldsymbol{\epsilon}^T (\mathbf{I}_n - \mathbf{P}) \boldsymbol{\epsilon}] \quad \underline{\hspace{10em}}$$

$$(3.104) \quad E[\boldsymbol{\epsilon}^T (\mathbf{I}_n - \mathbf{P}) \boldsymbol{\epsilon}] = E\left[\sum_{i=1}^n (1 - P_{ii}) \epsilon_i^2 + 2 \sum_{i < j} (0 - P_{ij}) \epsilon_i \epsilon_j\right] \quad \underline{\hspace{10em}}$$

$$(3.105) \quad = \sum_{i=1}^n (1 - P_{ii}) \sigma^2 + 0 = \sigma^2 \sum_{i=1}^n (1 - P_{ii})$$

$$(3.106) \quad = \sigma^2 \text{trace}(\mathbf{I}_n - \mathbf{P}) \quad \underline{\hspace{10em}}$$

$$(3.107) \quad = \sigma^2 [\text{trace}(\mathbf{I}_n) - \text{trace}(\mathbf{P})] \quad \underline{\hspace{10em}}$$

$$(3.108) \quad = \sigma^2 [n - k], \quad \underline{\hspace{10em}}$$

because  $\text{trace}(\mathbf{P}) = \text{trace}[(\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{X}] = \text{trace}(\mathbf{I}_k) = k$ .

Hence  $E[MS(Res)] = (n - k)^{-1} E[SS(Res)] = \sigma^2$ , or  $MS(Res)$  is an *unbiased* estimator of  $\sigma^2$ .

# 3.

MS(Res) is an unbiased estimator for  $\sigma^2$ .

This is explained in Section 3.7

## 3.7 Denominator of the residual SD

In different textbooks notation can be a bit different. For example, from “Introduction to Linear Regression Analysis” by Douglas C. Montgomery, we have:

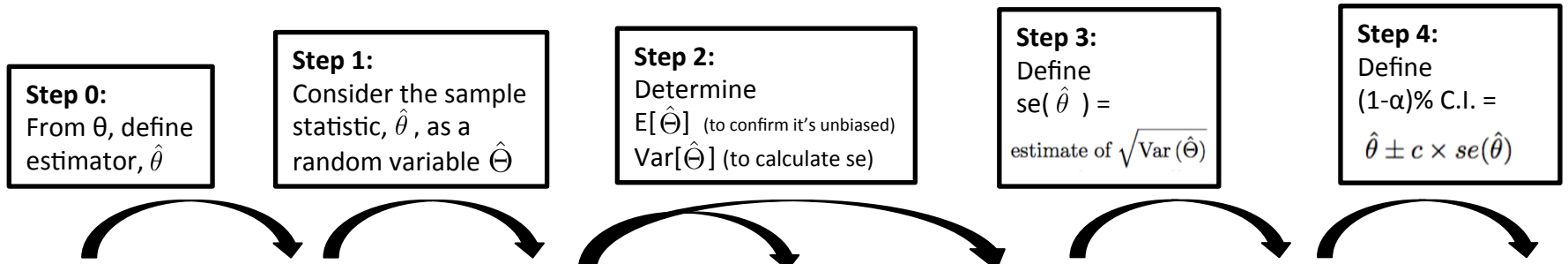
The expected value for  $SS_{\text{Res}}$  is

$$\begin{aligned} E(SS_{\text{Res}}) &= E(\mathbf{y}'[\mathbf{I} - \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}']\mathbf{y}) \\ &= \text{trace}([\mathbf{I} - \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}']\sigma^2\mathbf{I}) + E(\mathbf{y})'[\mathbf{I} - \mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}']E(\mathbf{y}) \\ &= (n - k)\sigma^2 \end{aligned}$$

As a result,

$$E(MS_{\text{Res}}) = E\left(\frac{SS_{\text{Res}}}{n - k}\right) = \sigma^2$$

# 3.6 Interval estimates and standard errors



| Population parameter or "something we would like to estimate" | Sample statistic ("estimator")   | Estimator as a Random Variable   | Expected Value of the estimator   | Variance of the estimator   | Standard Error of estimator   | Confidence Interval           |
|---|--|--|-----------------------------------|---|---|-------------------------------|
| $\beta$   | <b>b</b> <b>1.</b><br>$= (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$ | <b>B</b> ~ <b>2.</b><br>$\mathbf{N}(\beta, \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1})$ | $E[\mathbf{b}] = \beta$ <b>3.</b> | $\text{Var}[\mathbf{B}]$ <b>4.</b><br>$= \sigma^2 (\mathbf{X}^T \mathbf{X})^{-1}$ | $se(\mathbf{b})$ <b>5.</b><br>$= \hat{\sigma} \sqrt{[(\mathbf{X}^T \mathbf{X})^{-1}]_{jj}}$ | C.I. for $\beta$ <b>6.</b>    |
| $\sigma^2$  | $s^2$ or <b>MS(Res)</b> <b>1.</b>  | $S^2$ <b>2.</b>  | $E[S^2]$ <b>3.</b>                | $\text{Var}[S^2]$   | $se(s^2)$   | C.I. for $\sigma^2$           |
| $\mu_Y(\mathbf{x})$   | $(\hat{\mu}_Y(x))$ <b>1.</b>   | $(\hat{\mu}_Y(x))$ <b>2.</b>   | $E(\hat{\mu}_Y(x))$ <b>3.</b>     | $\text{Var}(\hat{\mu}_Y(x))$ <b>4.</b>  | $se(\hat{\mu}_Y(x))$ <b>5.</b>  | C.I. for $\mu_Y(x)$ <b>6.</b> |

# 1. 2. 3. 4. 5. 6.

## 3.6 Interval estimates and standard errors

Next consider the subpopulation mean  $\mu_Y(\mathbf{x}^*)$ , where  $\mathbf{x}^* = (1, x_1^*, \dots, x_p^*)^T$ . The point estimate is

$$(3.78) \quad \hat{\mu}_Y(\mathbf{x}^*) = \hat{\beta}_0 + \hat{\beta}_1 x_1^* + \dots + \hat{\beta}_p x_p^* = \mathbf{x}^{*T} \hat{\boldsymbol{\beta}}.$$

**Step 0:**  
From  $\theta$ , define estimator,

As a random variable, the variance is

$$(3.79) \quad \text{Var}[\hat{\mu}_Y(\mathbf{x}^*)] = \text{Var}[\mathbf{x}^{*T} \hat{\mathbf{B}}] = \mathbf{x}^{*T} \boldsymbol{\Sigma} \hat{\boldsymbol{\beta}} \mathbf{x}^*$$
$$(3.80) \quad = \sigma^2 \mathbf{x}^{*T} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{x}^*$$

**Step 2:**  
Determine  $E[\hat{\Theta}]$  (to confirm it's unbiased)  
 $\text{Var}[\hat{\Theta}]$  (to calculate se)

**Step 1:**  
Consider the sample statistic,  $\hat{\theta}$ , as a random variable

For the special case of  $p = 1$ , check that this is the same as (2.66). With the definition of the standard error,

$$(3.81) \quad se[\hat{\mu}_Y(\mathbf{x}^*)] = \hat{\sigma} \sqrt{\mathbf{x}^{*T} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{x}^*}.$$

**Step 3:**  
Define  $se(\hat{\theta}) =$   
estimate of  $\sqrt{\text{Var}(\hat{\Theta})}$

95% Confidence Interval:

**Step 4:**  
Define  $(1-\alpha)\%$  C.I. =  
 $\hat{\theta} \pm c \times se(\hat{\theta})$

$$\hat{\mu}_Y(\mathbf{x}^*) \pm t_{n-k, 0.975} se[\hat{\mu}_Y(\mathbf{x}^*)]$$

# Bonus: Prediction intervals

Next consider the prediction  $\hat{Y}(\mathbf{x}^*)$  for a future value at  $\mathbf{x}^*$ . Then

$$(3.82) \quad \hat{Y}(\mathbf{x}^*) = \hat{\beta}_0 + \hat{\beta}_1 x_1^* + \cdots + \hat{\beta}_p x_p^* = \mathbf{x}^{*T} \hat{\boldsymbol{\beta}}$$

is the same as the subpopulation mean in (3.78), which could be considered as the average of many observations at  $\mathbf{x}^*$ . Assuming the model is correct, the prediction error is

$$(3.83) \quad E = \hat{Y}(\mathbf{x}^*) - [\beta_0 + \beta_1 x_1^* + \cdots + \beta_p x_p^* + \epsilon(\mathbf{x}^*)],$$

where  $\epsilon(\mathbf{x}^*) \sim N(0, \sigma^2)$  independent of the previous data. Hence

$$(3.84) \quad \text{Var}(E) = \text{Var}[\mathbf{x}^{*T} \hat{\mathbf{B}}] + \text{Var}[\epsilon(\mathbf{x}^*)]$$

$$(3.85) \quad = \sigma^2 \mathbf{x}^{*T} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{x}^* + \sigma^2$$

and for the prediction error  $E$ ,

$$(3.86) \quad se[E] = \hat{\sigma} \sqrt{1 + \mathbf{x}^{*T} (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{x}^*}.$$

95% Prediction Interval:

$$\hat{Y}(\mathbf{x}^*) : +/- t_{n-k, 0.975} se[E]$$

# 3.6 Interval estimates and standard errors

## 95% Prediction Interval:

$$\hat{Y}(\mathbf{x}^*) : +/- t_{n-k,0.975} se[E]$$

$$se[E] = \hat{\sigma} \sqrt{1 + \mathbf{x}^{*T}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{x}^*}.$$

```
> # 50 year old who make 30K$ in income
> xstar <- c(1,50,30)
> xstar%%betahat
      y
[1,] 49.02491
>
> se_E <- sqrt(MS_Res)*sqrt(1 + t(xstar)%%solve(t(X)%%X)%%xstar)
> se_E
      [,1]
[1,] 15.12614
> mu_xstar -qt(0.975, n-k)*se_E
      y
[1,] 12.01257
> mu_xstar +qt(0.975, n-k)*se_E
      y
[1,] 86.03726
>
> predict(lm(y~x1+x2), newdata=data.frame(1,x1=50,x2=30), se.fit=TRUE,interval="prediction")
$fit
      fit      lwr      upr
1 49.02491 12.01257 86.03726
```

# Chapter 3

3.1 Least squares with two or more explanatory variables

3.4 Statistical software output for multiple regression

- $R^2$  and  $\text{adj}R^2$  and 3.4.1 Properties of  $R^2$  and  $\sigma^2$

- Sum of squares decomposition

3.5 Important explanatory variables

3.6 Interval estimates and standard errors

3.7 Denominator of the residual SD

**3.8 Residual plots**

3.9 Categorical explanatory variables

3.10 Partial correlation



# 3.8 Residual plots

Residual plots include the following.

- (i) Check for homoscedasticity versus heteroscedasticity and possible structural deviations from model (plot of residuals versus predicted values, plots of residuals versus each explanatory variable).
- (ii) Check for normality (normal quantile plot of residuals) if the plots from (i) look OK.

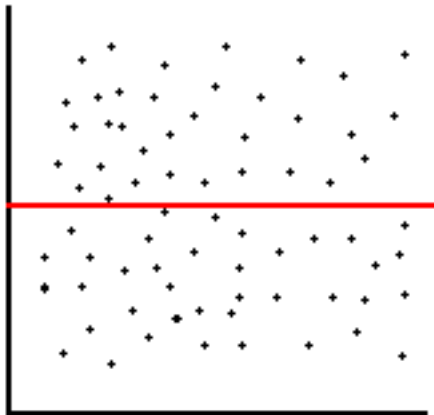
**We are looking at the residuals to verify that our model is correct:**

$$Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2) \text{ independently.}$$

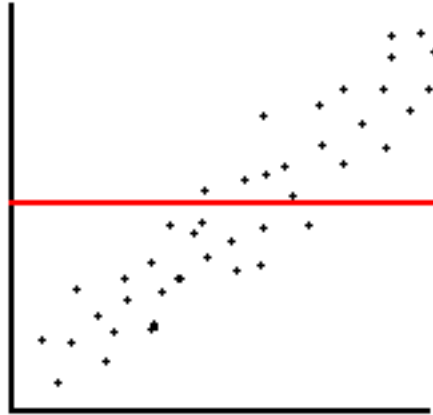
**This model has the following assumptions:**

1. residuals are independent
2. residuals are normally distributed
3. residuals all have common variance  $\sigma^2$  (homoscedasticity)  
-i.e. variance does not depend on any X or combination of X
4. Residuals have constant mean of 0 (i.e. no trends).

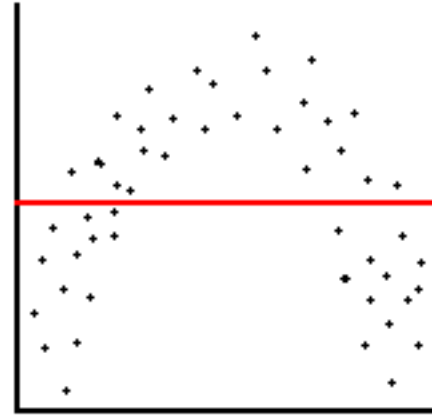
# 3.8 Residual plots



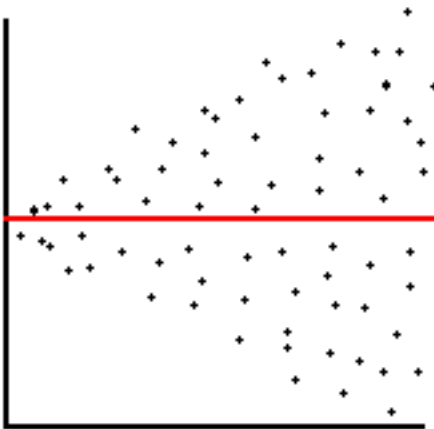
(a) Unbiased and Homoscedastic



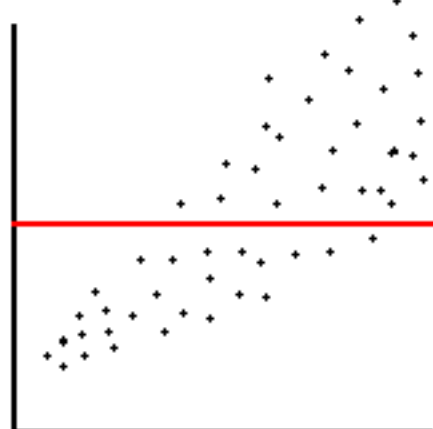
(b) Biased and Homoscedastic



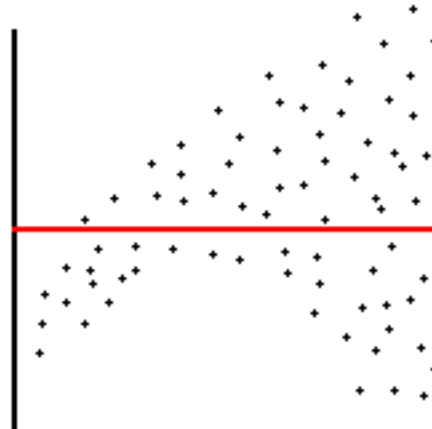
(c) Biased and Homoscedastic



(d) Unbiased and Heteroscedastic

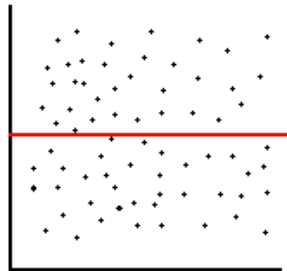


(e) Biased and Heteroscedastic

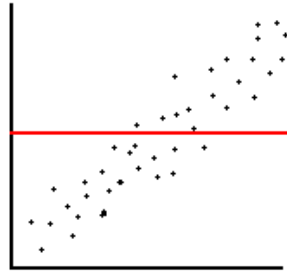


(f) Biased and Heteroscedastic

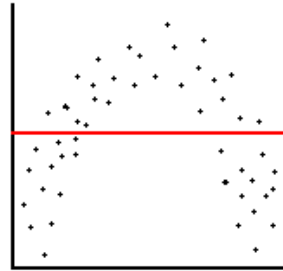
# 3.8 Residual plots



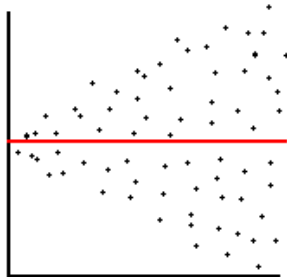
(a) Unbiased and Homoscedastic



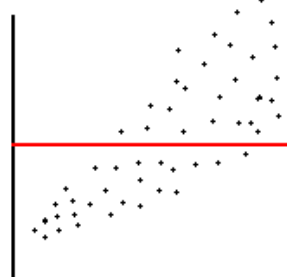
(b) Biased and Homoscedastic



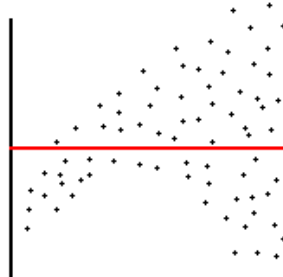
(c) Biased and Homoscedastic



(d) Unbiased and Heteroscedastic



(e) Biased and Heteroscedastic



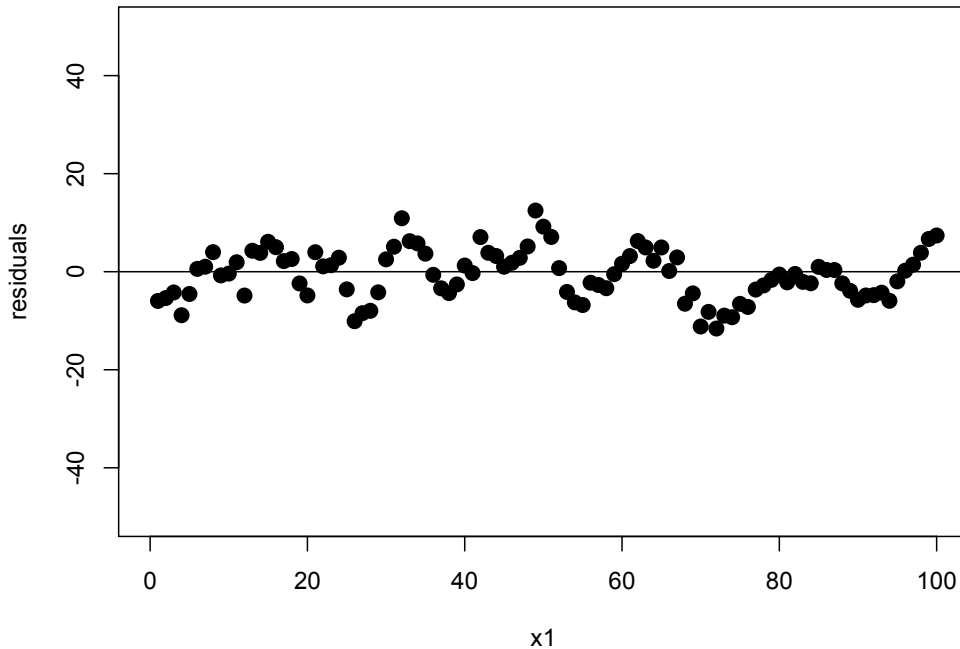
(f) Biased and Heteroscedastic

$$Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2) \text{ independently.}$$

**This model has the following assumptions:**

1. residuals are independent
2. residuals are normally distributed
- 3. residuals all have common variance  $\sigma^2$  (homoscedasticity)**  
-i.e. variance does not depend on any X or combination of X
- 4. Residuals have constant mean of 0 (i.e. no trends).**

# 3.8 Residual plots



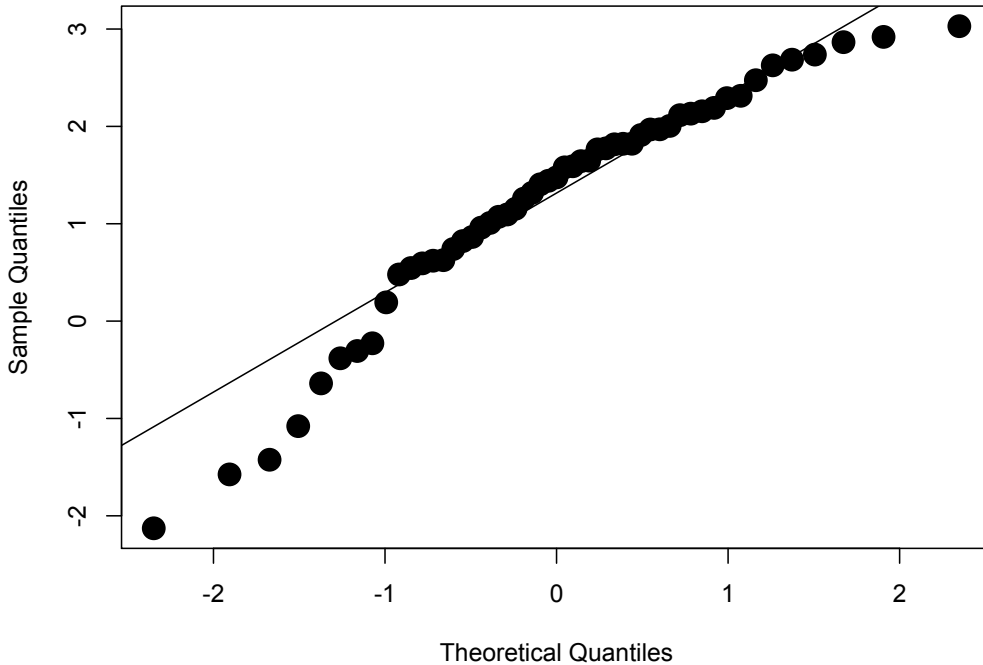
$$Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2) \text{ independently.}$$

**This model has the following assumptions:**

- 1. residuals are independent**
2. residuals are normally distributed
3. residuals all have common variance  $\sigma^2$  (homoscedasticity)  
-i.e. variance does not depend on any X or combination of X
4. Residuals have constant mean of 0 (i.e. no trends).

# 3.8 Residual plots

Normal Q-Q Plot



$$Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} + \epsilon_i, \quad i = 1, \dots, n, \quad \epsilon_i \sim N(0, \sigma^2) \text{ independently.}$$

**This model has the following assumptions:**

1. residuals are independent
- 2. residuals are normally distributed**
3. residuals all have common variance  $\sigma^2$  (homoscedasticity)  
-i.e. variance does not depend on any X or combination of X
4. Residuals have constant mean of 0 (i.e. no trends).

# *The art of linear regression*

- Categorical predictors
- Quadratic (polynomial) relationships
- Outliers
- How to fix heterogeneity
- Regression to the mean
- Simpsons Paradox
- Unobserved Confounding



# Chapter 3

3.1 Least squares with two or more explanatory variables

3.4 Statistical software output for multiple regression

- $R^2$  and  $\text{adj}R^2$  and 3.4.1 Properties of  $R^2$  and  $\sigma^2$

- Sum of squares decomposition

3.5 Important explanatory variables

3.6 Interval estimates and standard errors

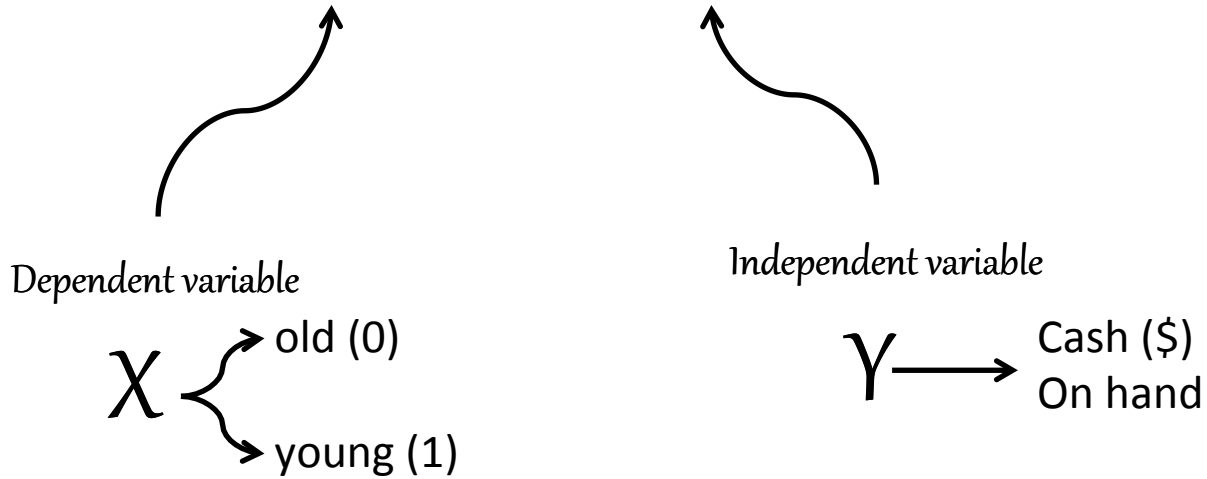
3.7 Denominator of the residual SD

3.8 Residual plots

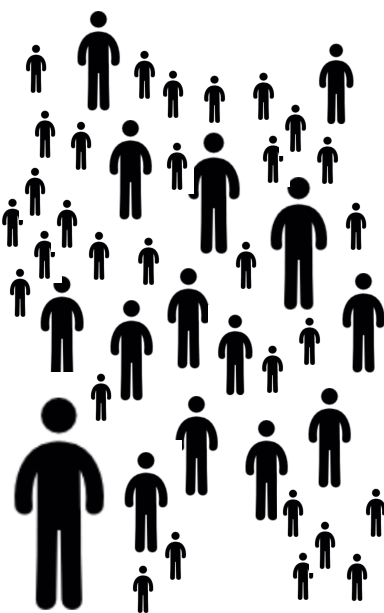
**3.9 Categorical explanatory variables**

3.10 Partial correlation

# Age vs. Money



## Population



Population parameters

$$\mu_0, \mu_1, \sigma^2$$

Hypothesis Test

$$H_0: \mu_0 = \mu_1$$

$$H_1: \mu_0 \neq \mu_1$$

Sample statistics

$$\bar{y}_0 = 56$$

$$\bar{y}_1 = 27$$

$$\bar{y}_0 - \bar{y}_1 = 29$$










$$s_p = 10.81$$

$$t = 2.68, df = 7$$

$$p\text{-value} = 0.03$$

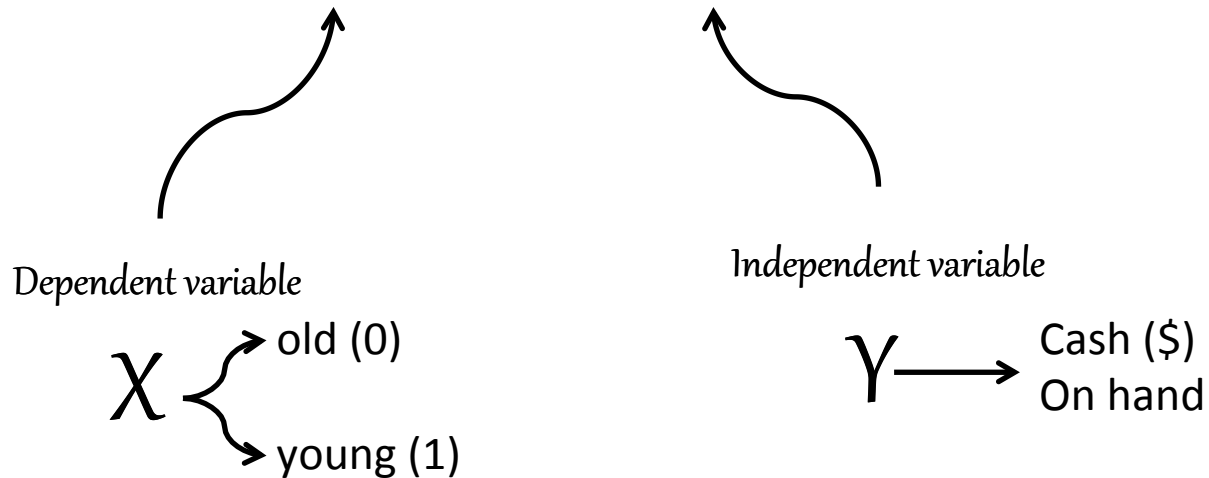
$$95\% \text{ C.I.} = [3.4, 54.6]$$

## Sample, n=9

|   | $X$   | $y$ |
|---|-------|-----|
|    | old   | 71  |
|   | old   | 54  |
|   | old   | 43  |
|  | young | 45  |
|  | young | 21  |
|  | young | 11  |
|  | young | 30  |
|  | young | 45  |
|  | young | 10  |



# Age vs. Money



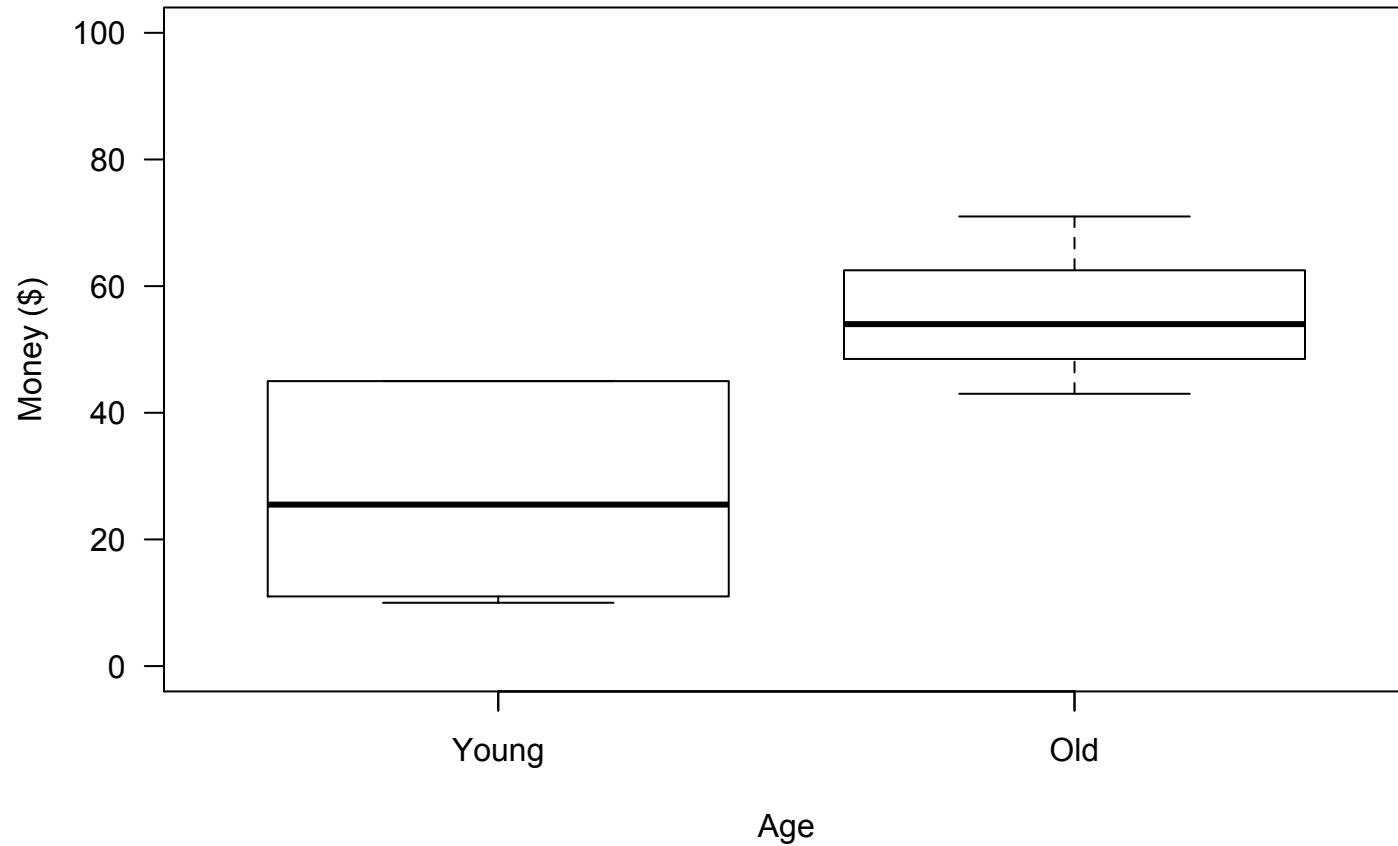
```
> old=c(71,54,43); young=c(45,21,11,30,45,10)  
> t.test(x=x2, y=x1, var.equal=TRUE)
```

Two Sample t-test

```
data: x2 and x1  
t = 2.6827, df = 7, p-value = 0.03142  
alternative hypothesis: true difference in means is not equal to 0  
95 percent confidence interval:  
 3.438298 54.561702  
sample estimates:  
mean of x mean of y  
    56      27
```

# Age vs. Money

Boxplot



# Age vs. Money

Dependent variable

$X$   $\left\{ \begin{array}{l} \text{old (0)} \\ \text{young (1)} \end{array} \right.$

Independent variable

$Y$   $\longrightarrow$  Cash (\$)  
On hand

```
> x <- c(0, 0, 0, 1, 1, 1, 1, 1, 1)
> y <- c(71, 54, 43, 45, 21, 11, 30, 45, 10)
> summary(lm(y~x))
```

# Age vs. Money



```
> x <- c(0, 0, 0, 1, 1, 1, 1, 1, 1)
> y <- c(71, 54, 43, 45, 21, 11, 30, 45, 10)
> summary(lm(y~x))
```

Call:  
lm(formula = y ~ x)

Residuals:

| Min | 1Q  | Median | 3Q | Max |
|-----|-----|--------|----|-----|
| -17 | -13 | -2     | 15 | 18  |

Coefficients:

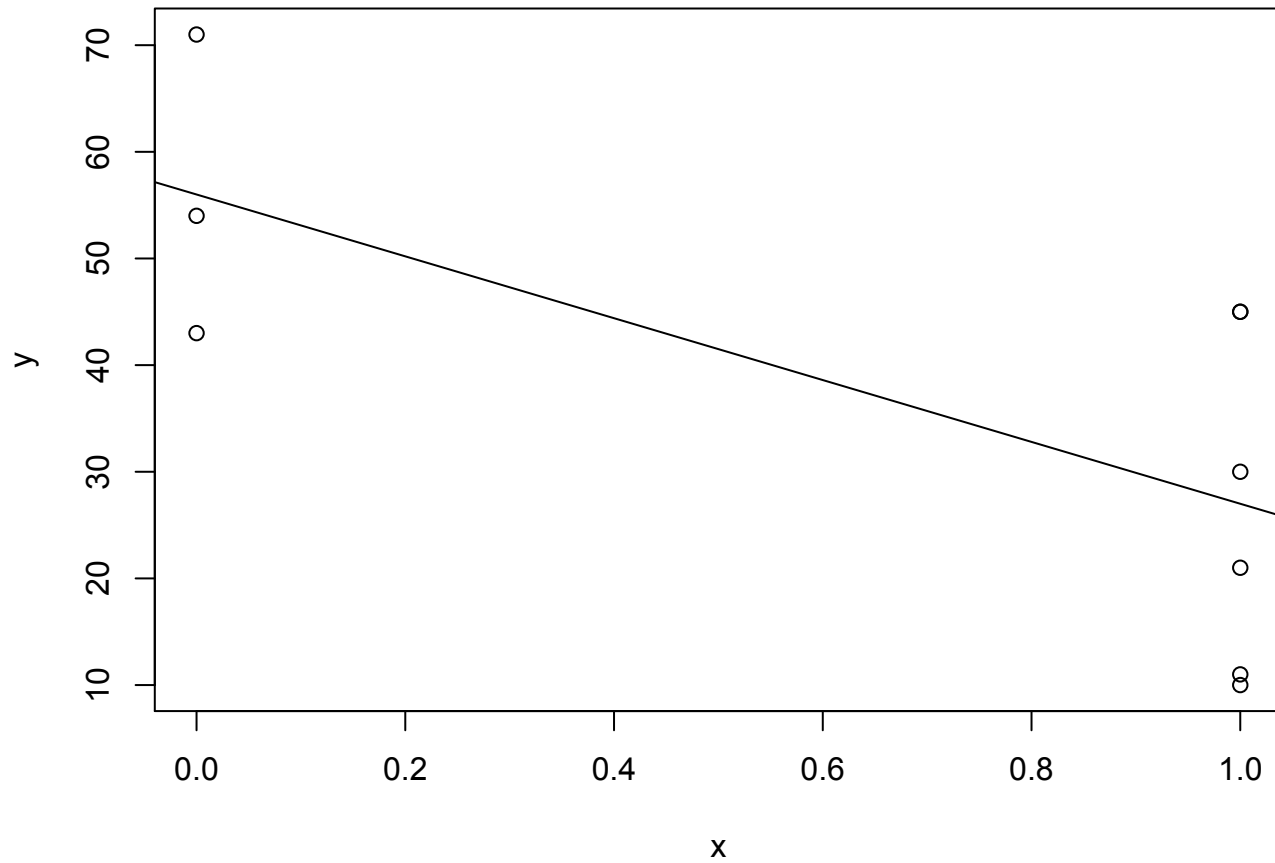
|             | Estimate | Std. Error | t value | Pr(> t ) |     |
|-------------|----------|------------|---------|----------|-----|
| (Intercept) | 56.000   | 8.826      | 6.345   | 0.000387 | *** |
| x           | -29.000  | 10.810     | -2.683  | 0.031417 | *   |

---

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

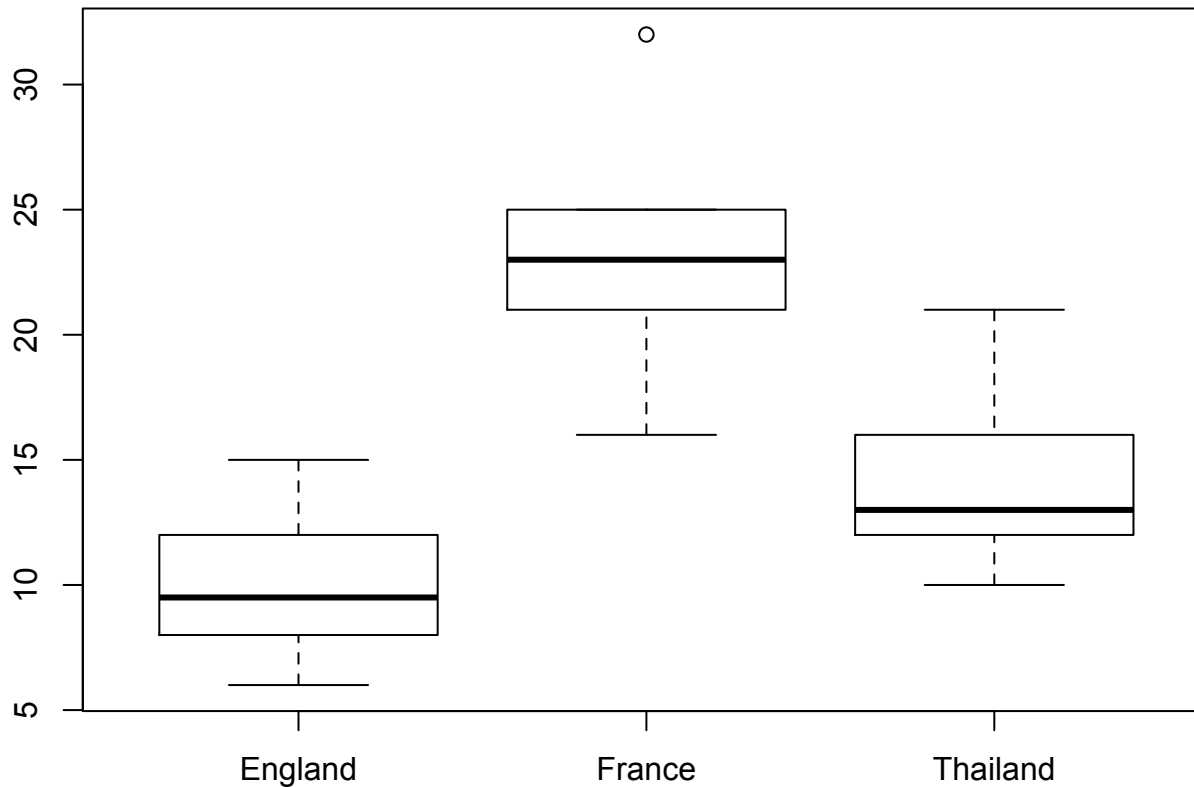
Residual standard error: 15.29 on 7 degrees of freedom  
Multiple R-squared: 0.5069, Adjusted R-squared: 0.4365  
F-statistic: 7.197 on 1 and 7 DF, p-value: 0.03142

# Age vs. Money



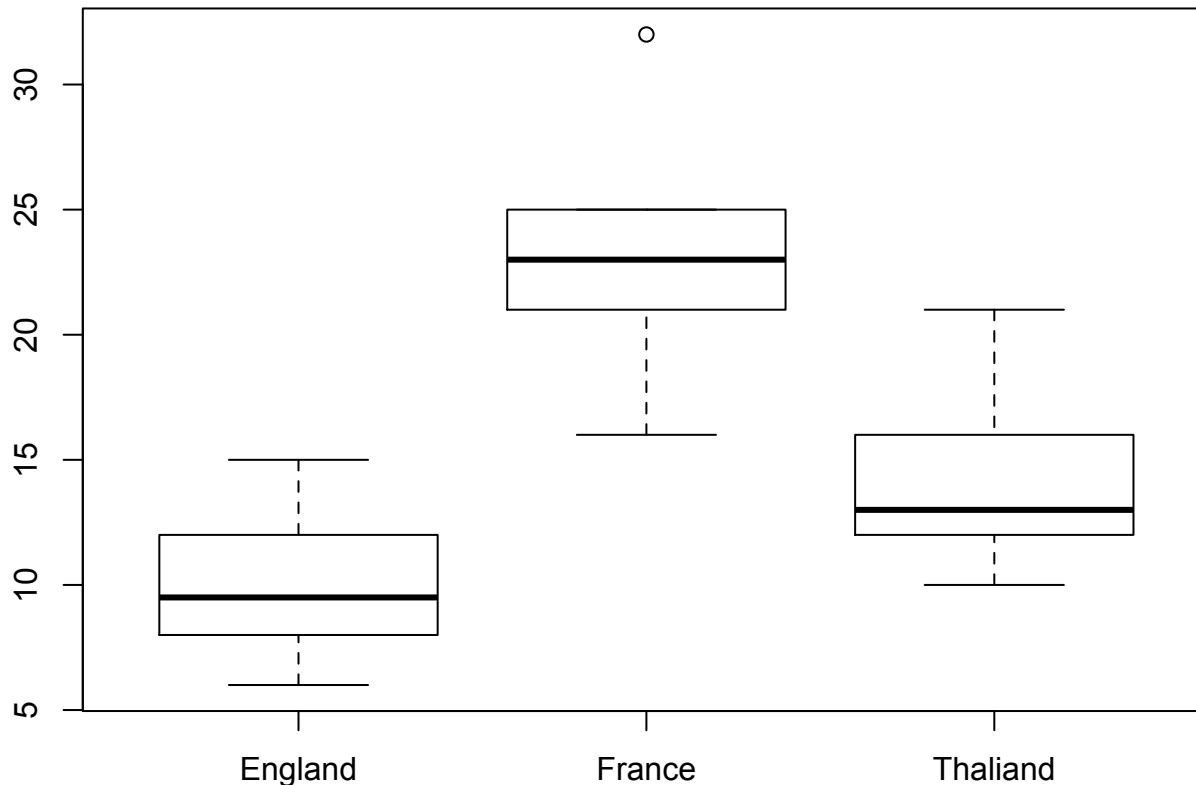
```
> plot(y~x)
> abline(lm(y~x))
```

# 3.9 Categorical explanatory variables



```
> country<-c(rep("France",6),rep("England",6),rep("Thailand",6))
> y<-c(23, 25, 21, 32, 16, 23, 15, 10, 8, 9, 6, 12, 13, 13, 12, 21, 16, 10)
> boxplot(y~country)
> |
```

# 3.9 Categorical explanatory variables



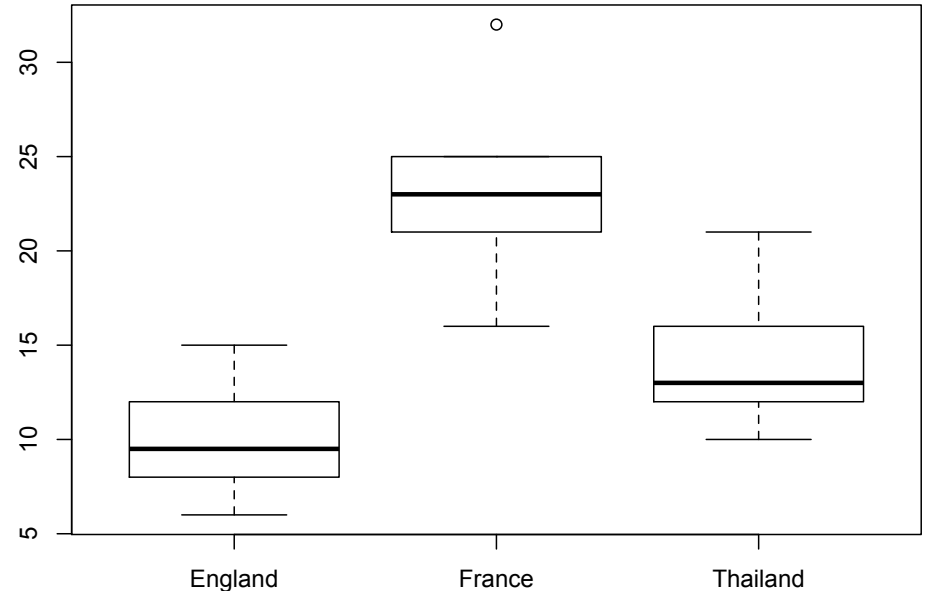
```
> summary(aov(y ~ country, data = mydata))
      Df Sum Sq Mean Sq F value    Pr(>F)
country  2  558.3   279.17   15.97 0.000192 ***
Residuals 15  262.2    17.48
```

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

# 3.9 Categorical explanatory variables

```
> data.frame(y, country)
```

|    | y  | country  |
|----|----|----------|
| 1  | 23 | France   |
| 2  | 25 | France   |
| 3  | 21 | France   |
| 4  | 32 | France   |
| 5  | 16 | France   |
| 6  | 23 | France   |
| 7  | 15 | England  |
| 8  | 10 | England  |
| 9  | 8  | England  |
| 10 | 9  | England  |
| 11 | 6  | England  |
| 12 | 12 | England  |
| 13 | 13 | Thailand |
| 14 | 13 | Thailand |
| 15 | 12 | Thailand |
| 16 | 21 | Thailand |
| 17 | 16 | Thailand |
| 18 | 10 | Thailand |

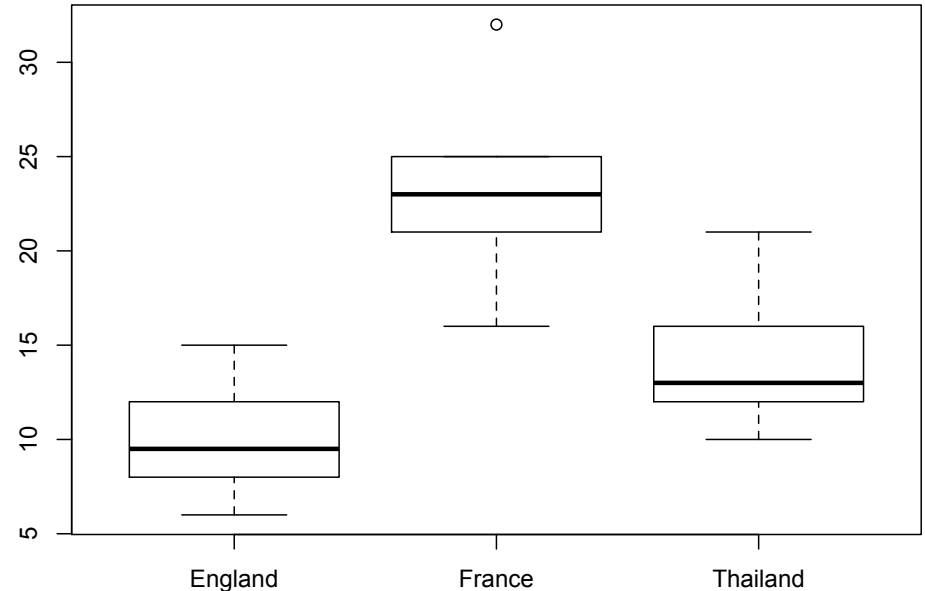




# 3.9 Categorical explanatory variables

```
> data.frame(y, country, x=as.numeric(as.factor(mydata$country))-1)
```

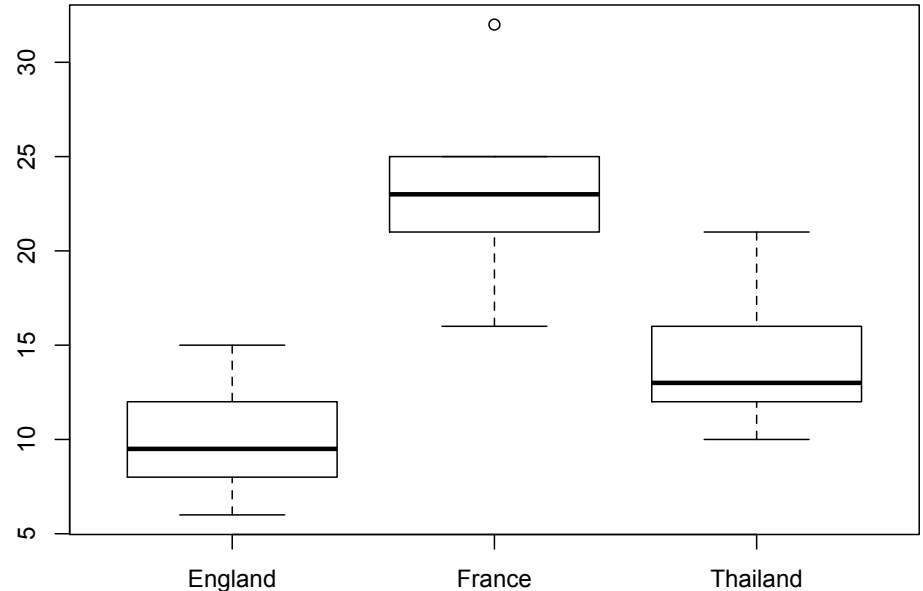
|    | y  | country  | x |
|----|----|----------|---|
| 1  | 23 | France   | 1 |
| 2  | 25 | France   | 1 |
| 3  | 21 | France   | 1 |
| 4  | 32 | France   | 1 |
| 5  | 16 | France   | 1 |
| 6  | 23 | France   | 1 |
| 7  | 15 | England  | 0 |
| 8  | 10 | England  | 0 |
| 9  | 8  | England  | 0 |
| 10 | 9  | England  | 0 |
| 11 | 6  | England  | 0 |
| 12 | 12 | England  | 0 |
| 13 | 13 | Thailand | 2 |
| 14 | 13 | Thailand | 2 |
| 15 | 12 | Thailand | 2 |
| 16 | 21 | Thailand | 2 |
| 17 | 16 | Thailand | 2 |
| 18 | 10 | Thailand | 2 |



# 3.9 Categorical explanatory variables

```
> data.frame(y, country, x=as.numeric(as.factor(mydata$country))-1)
```

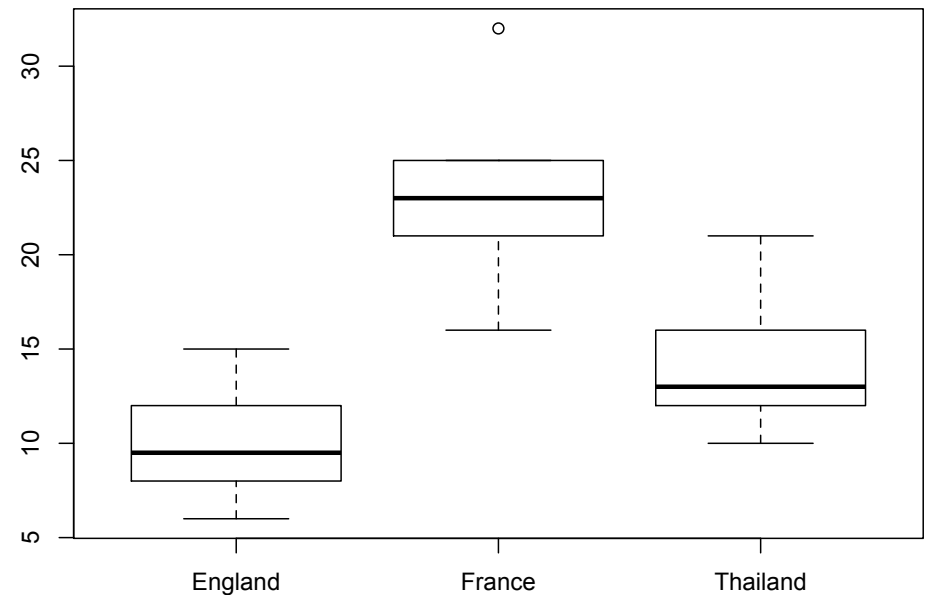
|    | y  | country  | x |
|----|----|----------|---|
| 1  | 23 | France   | 1 |
| 2  | 25 | France   | 1 |
| 3  | 21 | France   | 1 |
| 4  | 32 | France   | 1 |
| 5  | 16 | France   | 1 |
| 6  | 23 | France   | 1 |
| 7  | 15 | England  | 0 |
| 8  | 10 | England  | 0 |
| 9  | 8  | England  | 0 |
| 10 | 9  | England  | 0 |
| 11 | 6  | England  | 0 |
| 12 | 12 | England  | 0 |
| 13 | 15 | England  | 0 |
| 14 | 13 | Thailand | 2 |
| 15 | 12 | Thailand | 2 |
| 16 | 21 | Thailand | 2 |
| 17 | 16 | Thailand | 2 |
| 18 | 10 | Thailand | 2 |



# 3.9 Categorical explanatory variables

```
> data.frame(y, country, x1, x2)
```

|    | y  | country  | x1 | x2 |
|----|----|----------|----|----|
| 1  | 23 | France   | 1  | 0  |
| 2  | 25 | France   | 1  | 0  |
| 3  | 21 | France   | 1  | 0  |
| 4  | 32 | France   | 1  | 0  |
| 5  | 16 | France   | 1  | 0  |
| 6  | 23 | France   | 1  | 0  |
| 7  | 15 | England  | 0  | 0  |
| 8  | 10 | England  | 0  | 0  |
| 9  | 8  | England  | 0  | 0  |
| 10 | 9  | England  | 0  | 0  |
| 11 | 6  | England  | 0  | 0  |
| 12 | 12 | England  | 0  | 0  |
| 13 | 13 | Thailand | 0  | 1  |
| 14 | 13 | Thailand | 0  | 1  |
| 15 | 12 | Thailand | 0  | 1  |
| 16 | 21 | Thailand | 0  | 1  |
| 17 | 16 | Thailand | 0  | 1  |
| 18 | 10 | Thailand | 0  | 1  |



# 3.9 Categorical explanatory variables

```
> mydata<-data.frame(y, country, x1, x2)
> summary(lm(y ~ x1 + x2, data=mydata))
```

```
Call:
lm(formula = y ~ x1 + x2, data = mydata)
```

Residuals:

| Min     | 1Q      | Median  | 3Q     | Max    |
|---------|---------|---------|--------|--------|
| -7.3333 | -2.1250 | -0.6667 | 1.7917 | 8.6667 |

Coefficients:

|             | Estimate | Std. Error | t value | Pr(> t )     |
|-------------|----------|------------|---------|--------------|
| (Intercept) | 10.000   | 1.707      | 5.859   | 3.14e-05 *** |
| x1          | 13.333   | 2.414      | 5.524   | 5.84e-05 *** |
| x2          | 4.167    | 2.414      | 1.726   | 0.105        |

---

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

Residual standard error: 4.181 on 15 degrees of freedom  
Multiple R-squared: 0.6805, Adjusted R-squared: 0.6379  
F-statistic: 15.97 on 2 and 15 DF, p-value: 0.0001922

# 3.9 Categorical explanatory variables

```
> mydata<-data.frame(y, country, x1, x2)
> summary(lm(y ~ x1 + x2, data=mydata))
```

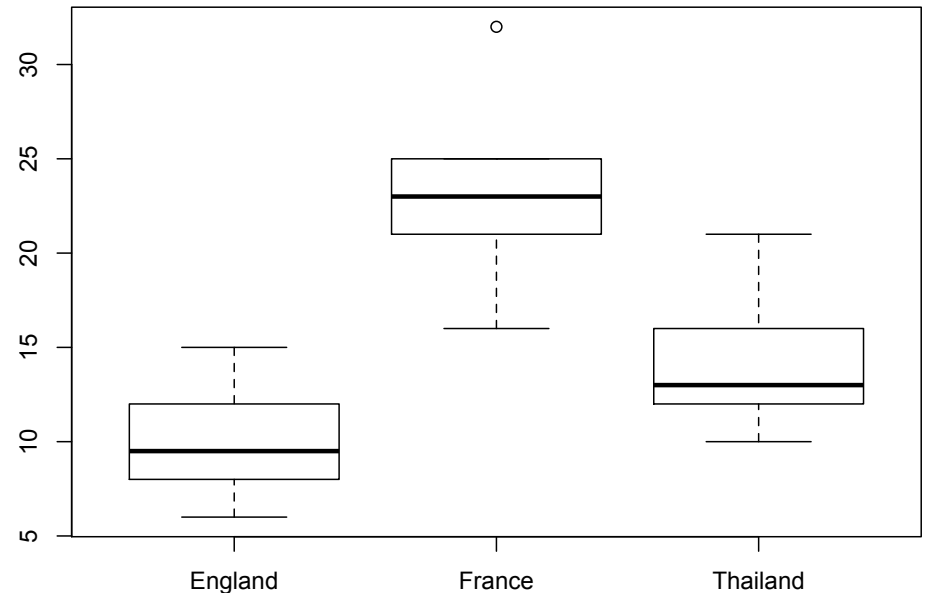
```
Call:
lm(formula = y ~ x1 + x2, data = mydata)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-7.3333 -2.1250 -0.6667  1.7917  8.6667
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept)  10.000      1.707    5.859 3.14e-05 ***
x1           13.333      2.414    5.524 5.84e-05 ***
x2            4.167      2.414    1.726  0.105
---
```

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

```
Residual standard error: 4.181 on 15 degrees of freedom
Multiple R-squared:  0.6805, Adjusted R-squared:  0.6379
F-statistic: 15.97 on 2 and 15 DF, p-value: 0.0001922
```



# 3.9 Categorical explanatory variables

```
> summary(lm(y ~ as.factor(country), data=mydata))
```

```
Call:  
lm(formula = y ~ as.factor(country), data = mydata)
```

Residuals:

| Min     | 1Q      | Median  | 3Q     | Max    |
|---------|---------|---------|--------|--------|
| -7.3333 | -2.1250 | -0.6667 | 1.7917 | 8.6667 |

Coefficients:

|                            | Estimate | Std. Error | t value | Pr(> t )     |
|----------------------------|----------|------------|---------|--------------|
| (Intercept)                | 10.000   | 1.707      | 5.859   | 3.14e-05 *** |
| as.factor(country)France   | 13.333   | 2.414      | 5.524   | 5.84e-05 *** |
| as.factor(country)Thailand | 4.167    | 2.414      | 1.726   | 0.105        |

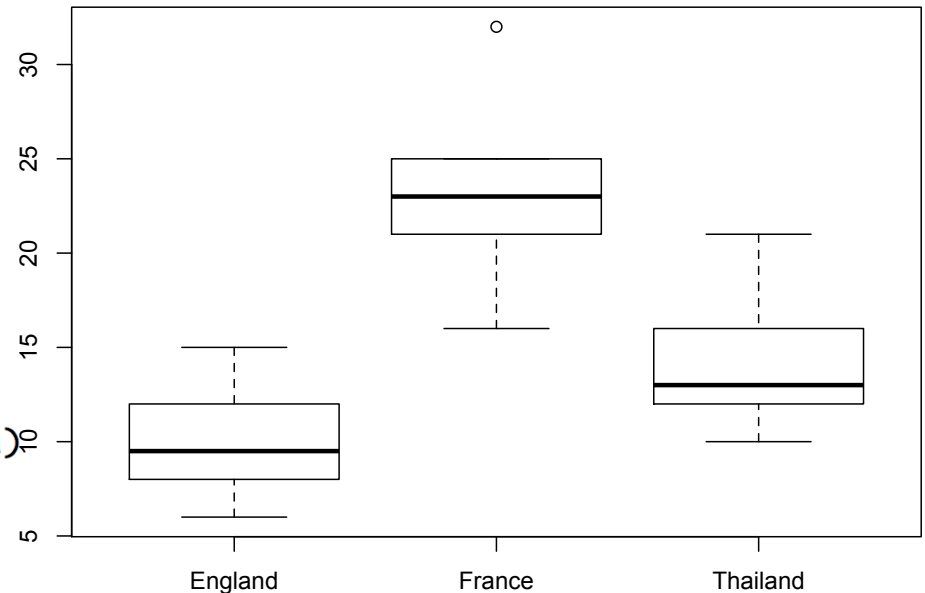
---

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

Residual standard error: 4.181 on 15 degrees of freedom

Multiple R-squared: 0.6805, Adjusted R-squared: 0.6379

F-statistic: 15.97 on 2 and 15 DF, p-value: 0.0001922



# 3.9 Categorical explanatory variables

## How do we interpret this model?

```
> x3<-c(34, 39, 32, 44, 22, 39, 41, 33, 37, 37, 27, 36, 67, 65, 56, 68, 60, 59)
> x3
[1] 34 39 32 44 22 39 41 33 37 37 27 36 67 65 56 68 60 59
> mydata<-data.frame(y, country, x1, x2, x3)
> summary(lm(y ~ x1 + x2 +x3, data=mydata))
```

```
Call:
lm(formula = y ~ x1 + x2 + x3, data = mydata)
```

```
Residuals:
    Min     1Q   Median     3Q      Max
-3.7297 -2.1410  0.1536  1.5329  3.7008
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -10.0295     4.0969  -2.448 0.028149 *
x1           13.4283     1.4859   9.037 3.23e-07 ***
x2          -11.4013     3.4177  -3.336 0.004899 **
x3           0.5696     0.1126   5.058 0.000175 ***
```

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

```
Residual standard error: 2.574 on 14 degrees of freedom
Multiple R-squared:  0.887, Adjusted R-squared:  0.8628
F-statistic: 36.63 on 3 and 14 DF, p-value: 7.02e-07
```

# 3.9 Categorical explanatory variables

## How do we make predictions from this model?

```
> x3<-c(34, 39, 32, 44, 22, 39, 41, 33, 37, 37, 27, 36, 67, 65, 56, 68, 60, 59)
> x3
[1] 34 39 32 44 22 39 41 33 37 37 27 36 67 65 56 68 60 59
> mydata<-data.frame(y, country, x1, x2, x3)
> summary(lm(y ~ x1 + x2 +x3, data=mydata))
```

```
Call:
lm(formula = y ~ x1 + x2 + x3, data = mydata)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-3.7297 -2.1410  0.1536  1.5329  3.7008
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -10.0295     4.0969  -2.448 0.028149 *
x1           13.4283     1.4859   9.037 3.23e-07 ***
x2          -11.4013     3.4177  -3.336 0.004899 **
x3           0.5696     0.1126   5.058 0.000175 ***
```

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

```
Residual standard error: 2.574 on 14 degrees of freedom
Multiple R-squared:  0.887, Adjusted R-squared:  0.8628
F-statistic: 36.63 on 3 and 14 DF,  p-value: 7.02e-07
```