

$$R^2 = 1 - \frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\sum_{i=1}^n (y_i - \bar{y})^2}$$

Proporción
de variabilidad
Debida a
los residuos

Respecto de
la variabilidad
Total

\approx chico

≈ 1

cuando tengo
un buen
ajuste

$$\hat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x}) Y_i}{\sum_{i=1}^n (x_i - \bar{x})^2},$$

$$\hat{\beta}_1 | X = x \sim N \left(\beta_1, \frac{\sigma^2}{\sum_{i=1}^n (x_i - \bar{x})^2} \right)$$

$$\hat{\beta}_0 = \bar{Y} - \hat{\beta}_1 \bar{x}.$$

$$\hat{\beta}_0 | X = x \sim N \left(\beta_0, \sigma^2 \left(\frac{1}{n} + \frac{\bar{x}^2}{\sum_{i=1}^n (x_i - \bar{x})^2} \right) \right)$$

$$T = \frac{\hat{\beta}_0 - \beta_0}{S_r \sqrt{\frac{1}{n} + \frac{\bar{x}^2}{S_{xx}}}} = \frac{\hat{\beta}_0 - \beta_0}{\text{se}(\hat{\beta}_0)} \sim t_{n-2}$$

29.6376

$$S_r^2 = \frac{\sum_{i=1}^n (Y_i - \bar{Y})^2}{n-2}, = \sigma^2$$

$$S_{xx} = \sum_{i=1}^n (x_i - \bar{x})^2.$$

$$T = \frac{\hat{\beta}_1 - \beta_1}{\frac{S_r}{\sqrt{S_{xx}}}} = \frac{\hat{\beta}_1 - \beta_1}{\text{se}(\hat{\beta}_1)} \sim t_{n-2}$$

0.7243

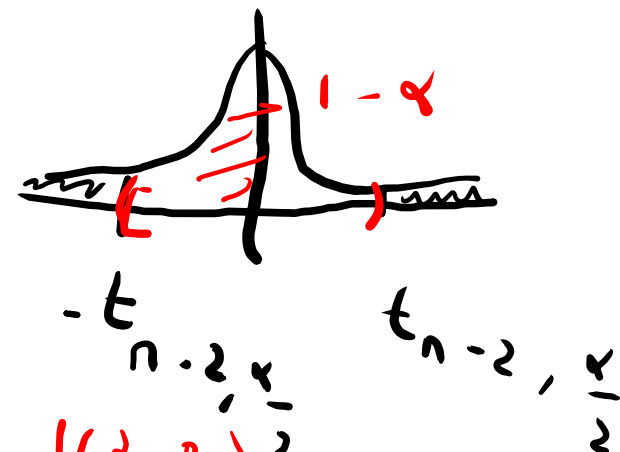
$$\left[\hat{\beta}_0 + t_{n-2, \alpha/2} S_r \sqrt{\frac{1}{n} + \frac{\bar{x}^2}{S_{xx}}}; \hat{\beta}_0 - t_{n-2, \alpha/2} S_r \sqrt{\frac{1}{n} + \frac{\bar{x}^2}{S_{xx}}} \right]$$

$$\left[\hat{\beta}_1 - t_{n-2, \alpha/2} \frac{S_r}{S_{xx}}; \hat{\beta}_1 + t_{n-2, \alpha/2} \frac{S_r}{S_{xx}} \right]$$

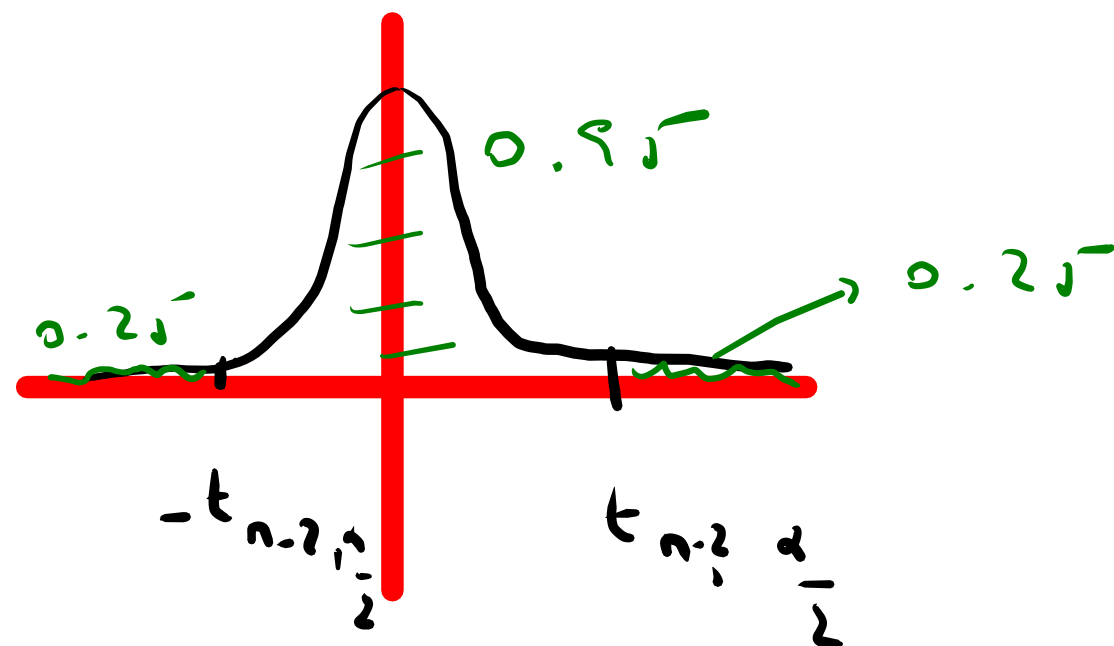
`confint(modelo)`

Nivel de confiança por default
é 95%.

##	2.5 %	97.5 %
## (Intercept)	41.265155	163.885130
## edad	3.822367	6.818986



IC Para β_0 de nível $1-\alpha$ é $[41.26, 163.88]$
 β_1 $[3.82, 6.81]$



Intervalo de confiança para σ^2

$$\left[\frac{(n-2)S_r^2}{\chi_{n-2,\alpha/2}^2}, \frac{(n-2)S_r^2}{\chi_{n-2,1-\alpha/2}^2} \right]$$

$\hat{\sigma}^2$

Test de hipótesis para β_0 y β_1 .

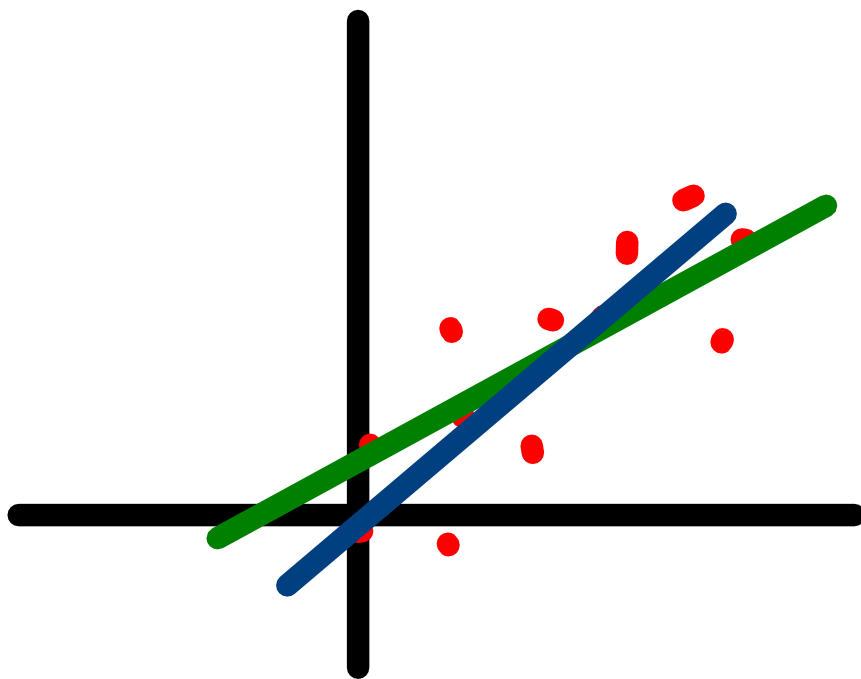
Recta de mejor ajuste

$$Y = \hat{\beta}_0 + \hat{\beta}_1 x$$

Test

$$H_0 \quad \beta_0 = 0 \quad \sim \quad H_1 \quad \beta_0 \neq 0$$

$$\beta_1 = 0 \quad \sim \quad \beta_1 \neq 0$$



$$y = \beta_0 + \beta_1 x$$

$$y = \beta x$$

$$\beta_0 \approx 0 \quad \text{vs} \quad \beta_0 \neq 0$$

No Intercept

Passo 2 es tudo um
modelos $y = \beta x$

$$y = \beta_0 +$$

$$\beta_1 = 0 \quad \sim \quad \beta_1 \neq 0$$

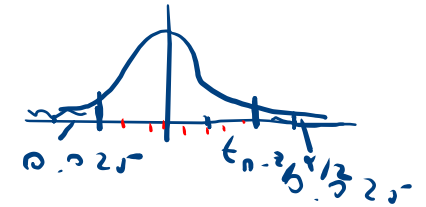
$$T = \frac{\hat{\beta}_1 - \beta_1}{\frac{S_r}{\sqrt{S_{xx}}}} = \frac{\hat{\beta}_1 - \beta_1}{\text{se}(\hat{\beta}_1)} \sim t_{n-2} \rightarrow \frac{\hat{\beta}_1}{\text{se}(\hat{\beta}_1)} \sim t_{n-2} \text{ Bajo } H_0$$

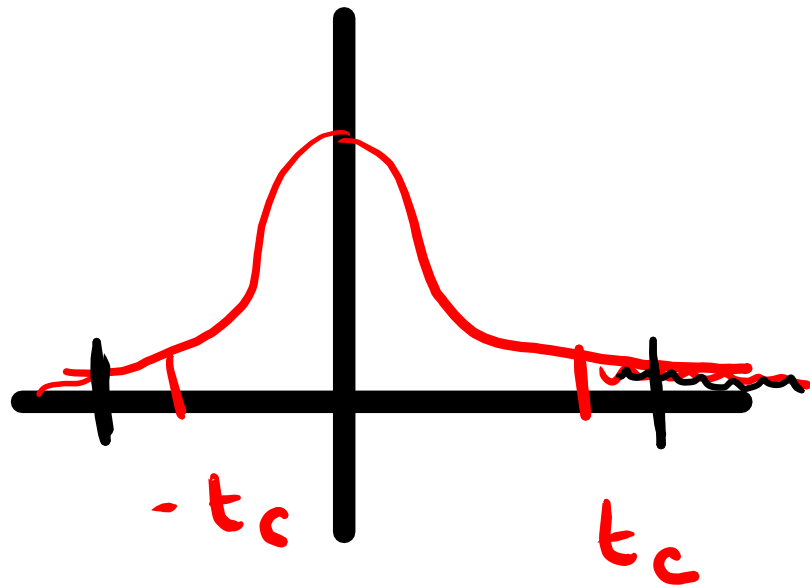
con los datos de las observaciones

$$\frac{\hat{\beta}_1}{\text{se}(\hat{\beta}_1)} \Rightarrow t_c = \frac{5.3207}{0.2243} = 17.3459$$

$\alpha = 0.05$

$$t_{n-2, \frac{\alpha}{2}}$$



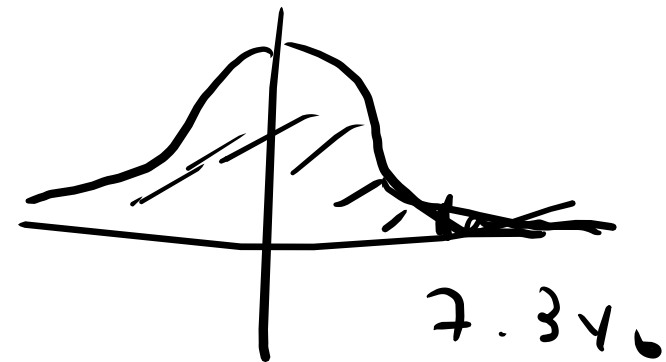


$$P(|T| > |t_c|) = p_{\text{value}}$$

$$P(T > 7.346) +$$

$$P(T < -7.346)$$

P_t

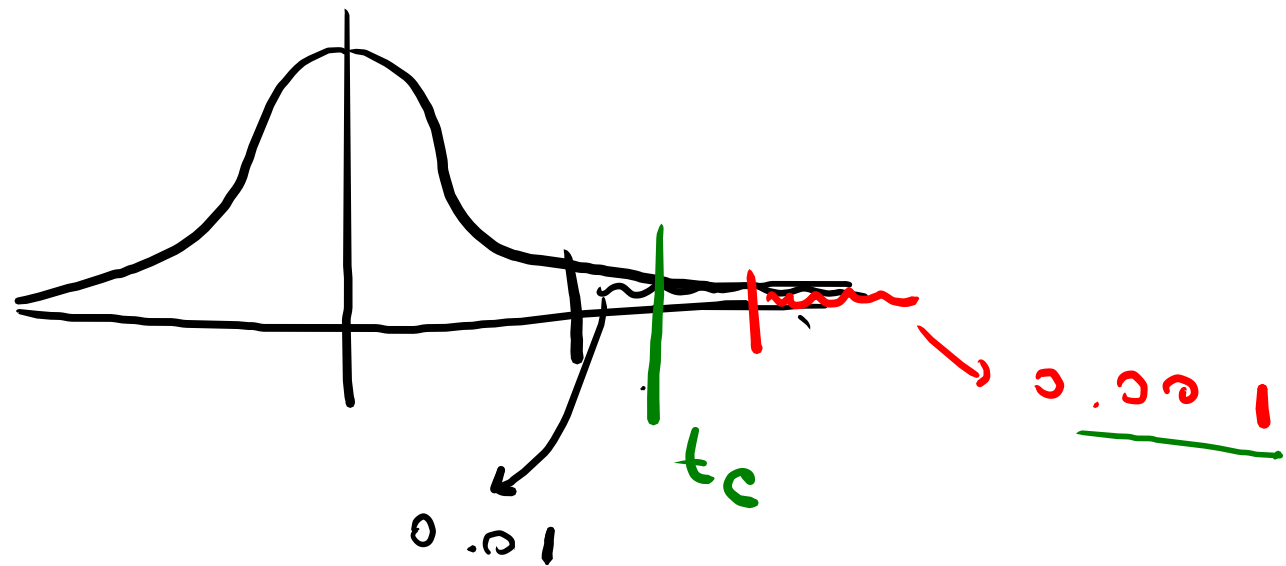


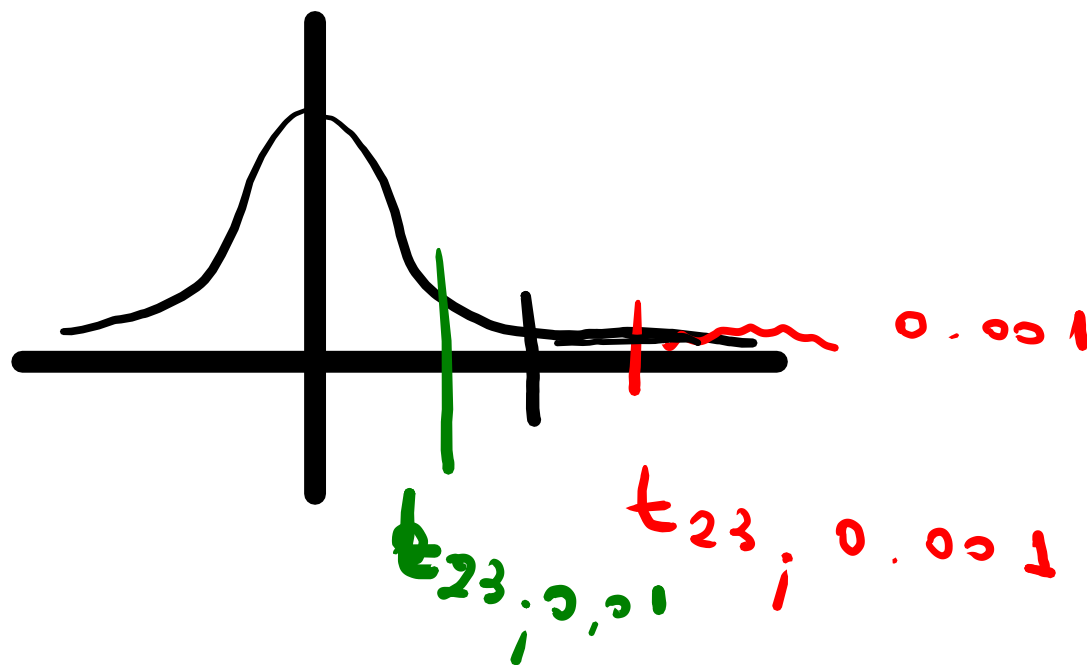
Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	102.5751	29.6376	3.461	0.00212	**
edad	5.3207	0.7243	7.346	1.79e-07	***

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

0.001 < 0.00212 < 0.01





P value ≈ 0.002

Rechazo H_0 a un nivel del 1%.

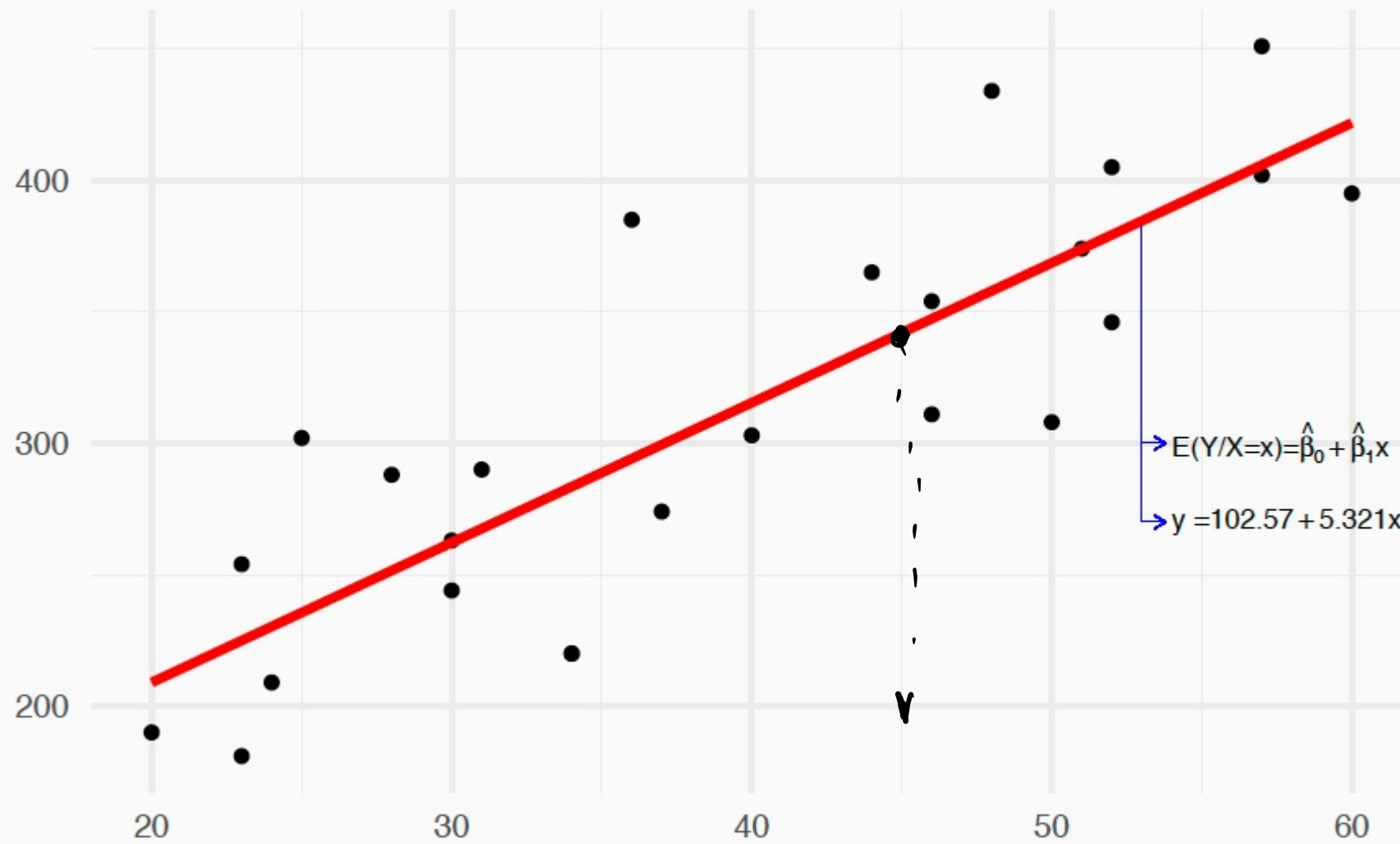
Pero no rechazo H_0 a un nivel del 1 por mil

El p value correspondiente a β_1 es muy chico. Entonces rechazamos H_0 para este test, a cualquier nivel

Conclusión

Para este problema, ambos parámetros son significativamente $\neq 0$

Es decir, hay suficiente evidencia para afirmar que ambos parámetros son $\neq 0$

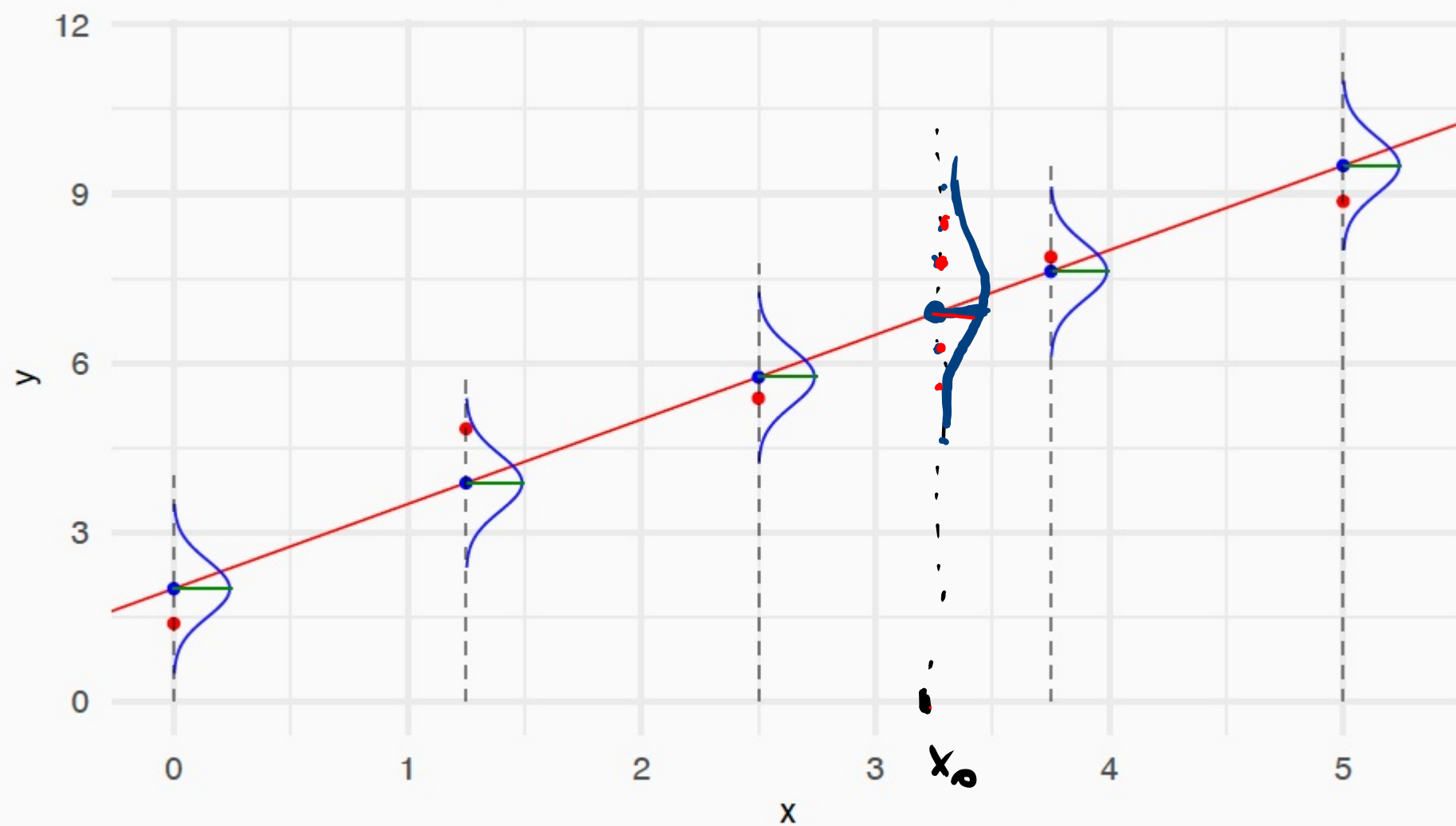


```
nuevas.edades <- data.frame(edad = c(30,45))  
nuevas.edades
```

```
##      edad  
## 1      30  
## 2      45
```

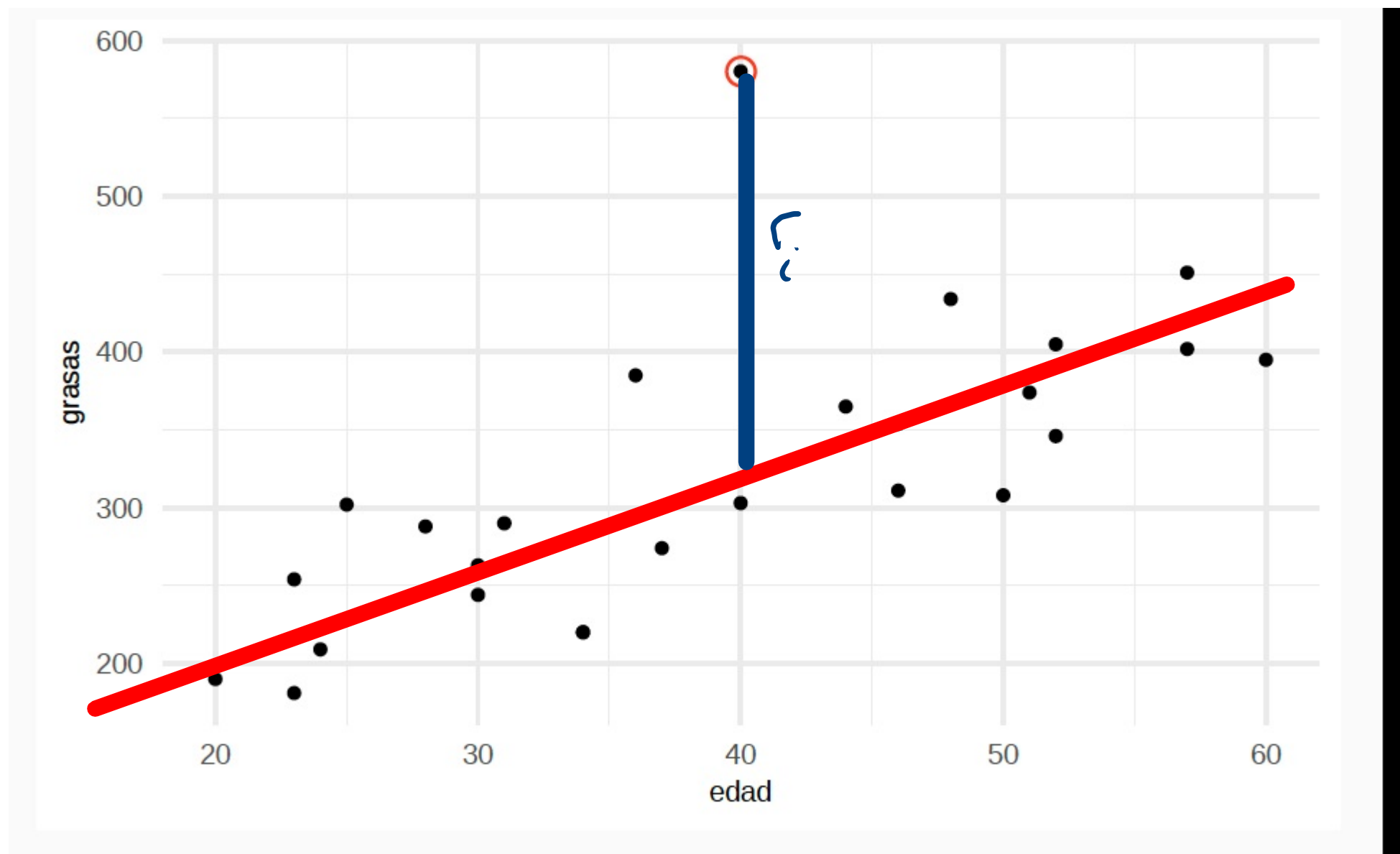
```
predict(modelo, nuevas.edades)
```

Regresión lineal simple con normales en cada x_i




```
predict(modelo, newdata=nuevas.edades, interval="confidence")
```

```
predict(modelo, newdata=nuevas.edades, interval="prediction")
```



```
> summary(modelo)
```

```
Call:
```

```
lm(formula = grasas ~ edad, data = grasas)
```

```
Residuals:
```

Min	1Q	Median	3Q	Max
-63.478	-26.816	-3.854	28.315	90.881

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	<u>102.5751</u>	29.6376	3.461	0.00212 **
edad	<u>5.3207</u>	0.7243	7.346	1.79e-07 ***

```
---
```

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

```
Residual standard error: 13.46 on 23 degrees of freedom
```

```
Multiple R-squared:  0.7012,    Adjusted R-squared:  0.6882
```

```
F-statistic: 53.96 on 1 and 23 DF,  p-value: 1.794e-07
```

```
> summary(regresion)
```

```
Call:
```

```
lm(formula = grasas ~ edad, data = datos)
```

```
Residuals:
```

Min	1Q	Median	3Q	Max
-73.34	-37.60	-13.00	19.36	254.59

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	110.324	46.304	2.383	<u>0.0255 *</u>
edad	<u>5.383</u>	1.133	4.753	7.78e-05 ***

```
---
```

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

```
Residual standard error: 67.97 on 24 degrees of freedom
```

```
Multiple R-squared:  0.4849,    Adjusted R-squared:  0.4634
```

```
F-statistic: 22.59 on 1 and 24 DF,  p-value: 7.784e-05
```

0.002 es chis

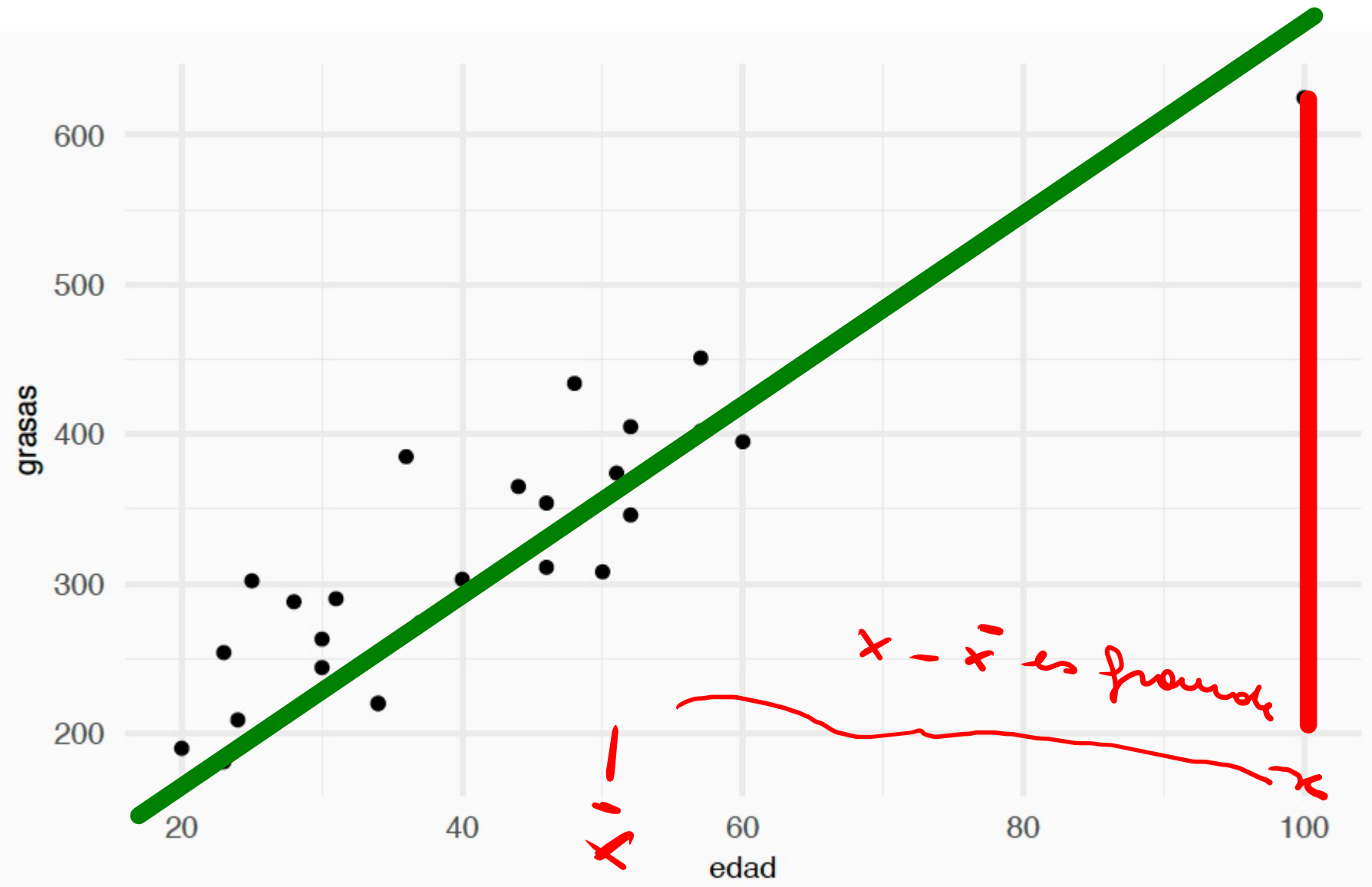
0.01 < 0.02 < 0.05

20; 25; 30; (31); 42; 50; 510

$$\frac{\sum x_i}{n}$$

$$\min_{\rho, \beta} \sum r_i^2$$

nombre [nombre >]



$$\text{Var}(r_i) = \sigma^2 (1 - h_{ii})$$

$$h_{ii} = \frac{1}{n} + \frac{(x_i - \bar{x})^2}{\sum (x_i - \bar{x})^2}$$

```
> summary(regresion.original) #sin punto de alto leverage
```

```
Call:
lm(formula = grasas ~ edad, data = datos.originales)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-63.478 -26.816  -3.854   28.315   90.881
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 102.5751    29.6376   3.461  0.00212 **
edad         5.3207     0.7243   7.346  1.79e-07 ***
```

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

```
Residual standard error: 43.46 on 23 degrees of freedom
Multiple R-squared:  0.7012,    Adjusted R-squared:  0.6882
F-statistic: 53.96 on 1 and 23 DF, p-value: 1.794e-07
```

```
> summary(regresion.ConLevBueno) #Con punto de alto leverage
```

```
Call:
lm(formula = grasas ~ edad, data = datos.ConLevBueno)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-63.695 -25.312  -3.459   27.712   90.821
```

```
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 105.4709    22.4607   4.696  9.00e-05 ***
edad         5.2419     0.5029  10.423  2.18e-10 ***
```

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

```
Residual standard error: 42.57 on 24 degrees of freedom
Multiple R-squared:  0.8191,    Adjusted R-squared:  0.8115
F-statistic: 108.6 on 1 and 24 DF, p-value: 2.175e-10
```

$$\bar{h} = \frac{\sum_{i=1}^n h_{ii}}{n}$$

en R → hat values

criterio que vamos a considerar es

$$h_{ii} > \bar{h} = 2 \times \frac{2}{25} = \frac{4}{25}$$

```
> summary(regresion.original) #sin punto de alto leverage
```

```
Call:
```

```
lm(formula = grasas ~ edad, data = datos originales)
```

```
Residuals:
```

Min	1Q	Median	3Q	Max
-63.478	-26.816	-3.854	28.315	90.881

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	<u>102.5751</u>	29.6376	3.461	<u>0.00212</u> **
edad	<u>5.3207</u>	0.7243	7.346	<u>1.79e-07</u> ***

```
---
```

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

```
Residual standard error: 43.46 on 23 degrees of freedom
```

```
Multiple R-squared: 0.7012, Adjusted R-squared: 0.6882
```

```
F-statistic: 53.96 on 1 and 23 DF, p-value: 1.794e-07
```

```
> summary(regresion) # Con punto de alto leverage
```

```
Call:
```

```
lm(formula = grasas ~ edad, data = datos)
```

```
Residuals:
```

Min	1Q	Median	3Q	Max
-87.163	-40.453	-4.734	29.165	118.566

```
Coefficients:
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	28.8926	29.0073	0.996	<u>0.329</u>
edad	<u>7.3254</u>	0.6495	11.279	<u>4.46e-11</u> ***

```
---
```

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

```
Residual standard error: 54.98 on 24 degrees of freedom
```

```
Multiple R-squared: 0.8413, Adjusted R-squared: 0.8347
```

```
F-statistic: 127.2 on 1 and 24 DF, p-value: 4.461e-11
```


Distancia de Cook

$$D_i = \frac{r_{si}^2}{2} \frac{h_{ii}}{1 - h_{ii}}$$

```
cooks.o <- cooks.distance(modelo.o)  
p <- length(coef(modelo.o))  
n <- dim(grasas.o)[1]  
cooks_umbral.o <- qf(0.5, p, n-p) &L
```

F distribución de Fisher

$F_{p, n-p}$

$F_{2, n-2}$