→ A4-Regresión Poisson

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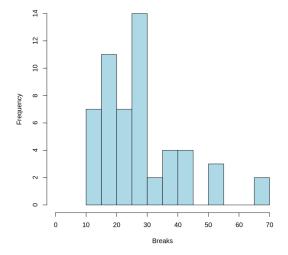
- 1 data<-warpbreaks
 2 head(data,10)</pre>
 - A data.frame: 10 × 3

	Additional to x o		
	breaks	wool	tension
	<dbl></dbl>	<fct></fct>	<fct></fct>
1	26	Α	L
2	30	Α	L
3	54	Α	L
4	25	Α	L
5	70	Α	L
6	52	Α	L
7	51	Α	L
8	26	Α	L
9	67	Α	L
10	18	Α	М

▼ 1. Histograma de numero de ropturas

```
1 hist(data$breaks,
      main = "Histogram of Breaks",
                                    # Title for the plot
      xlab = "Breaks",
                                      # Label for the x-axis
      ylab = "Frequency",
4
                                      # Label for the y-axis
      col = "lightblue",
5
                                      # Color of the bars
      border = "black",
                                      # Border color of the bars
      xlim = c(0, max(data\$breaks)),
                                      # Set the x-axis limits if needed
      breaks = 20)
                                     # Number of bins
```





▼ 2. Obtén la media y la varianza

```
1 mean_breaks <- mean(data$breaks)
2 variance_breaks <- var(data$breaks)
3
4 # Print the results</pre>
```

1

```
5 cat("Media de breaks", mean_breaks, "\n")
6 cat("Varianza de breaks", variance_breaks, "\n")
    Media de breaks 28.14815
    Varianza de breaks 174.2041
```

▼ 3. Ajusta el modelo de regresión Poisson

```
1 poisson.model<-glm(breaks ~ wool + tension, data, family = poisson(link = "log"))</pre>
2 summary(poisson.model)
   Call:
   glm(formula = breaks ~ wool + tension, family = poisson(link = "log"),
   Coefficients:
              Estimate Std. Error z value Pr(>|z|)
   (Intercept) 3.69196 0.04541 81.302 < 2e-16 ***
                          0.05157 -3.994 6.49e-05 ***
              -0.20599
   woolB
                         0.06027 -5.332 9.73e-08 ***
   tensionM
             -0.32132
   tensionH
             -0.51849
                          0.06396 -8.107 5.21e-16 ***
   Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
   (Dispersion parameter for poisson family taken to be 1)
       Null deviance: 297.37 on 53 degrees of freedom
   Residual deviance: 210.39 on 50 degrees of freedom
   ATC: 493.06
   Number of Fisher Scoring iterations: 4
```

Si bien la desviación residual es mayor que los grados de libertad, y esto es algo que se busca, esta es muy grande por lo que se puede interpretar que el modelo tiene estimaciones correctas pero los errores estandar son incorrectos. Con esto en cuenta se implementará un modelo cuasi Poisson para mejorar los resultados."

```
1 poisson.model2<-glm(breaks ~ wool + tension, data = data, family = quasipoisson(link = "log"))</pre>
2 summary(poisson.model2)
   Call:
   glm(formula = breaks ~ wool + tension, family = quasipoisson(link = "log"),
       data = data)
   Coefficients:
              Estimate Std. Error t value Pr(>|t|)
   (Intercept) 3.69196 0.09374 39.384 < 2e-16 ***
                          0.10646 -1.935 0.058673 .
   wool B
              -0.20599
   tensionM
              -0.32132
                         0.12441 -2.583 0.012775 *
   tensionH
              -0.51849
                          0.13203 -3.927 0.000264 ***
   Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
   (Dispersion parameter for quasipoisson family taken to be 4.261537)
       Null deviance: 297.37 on 53 degrees of freedom
   Residual deviance: 210.39 on 50 degrees of freedom
   AIC: NA
   Number of Fisher Scoring iterations: 4
```