Act 4: Regression de Poisson

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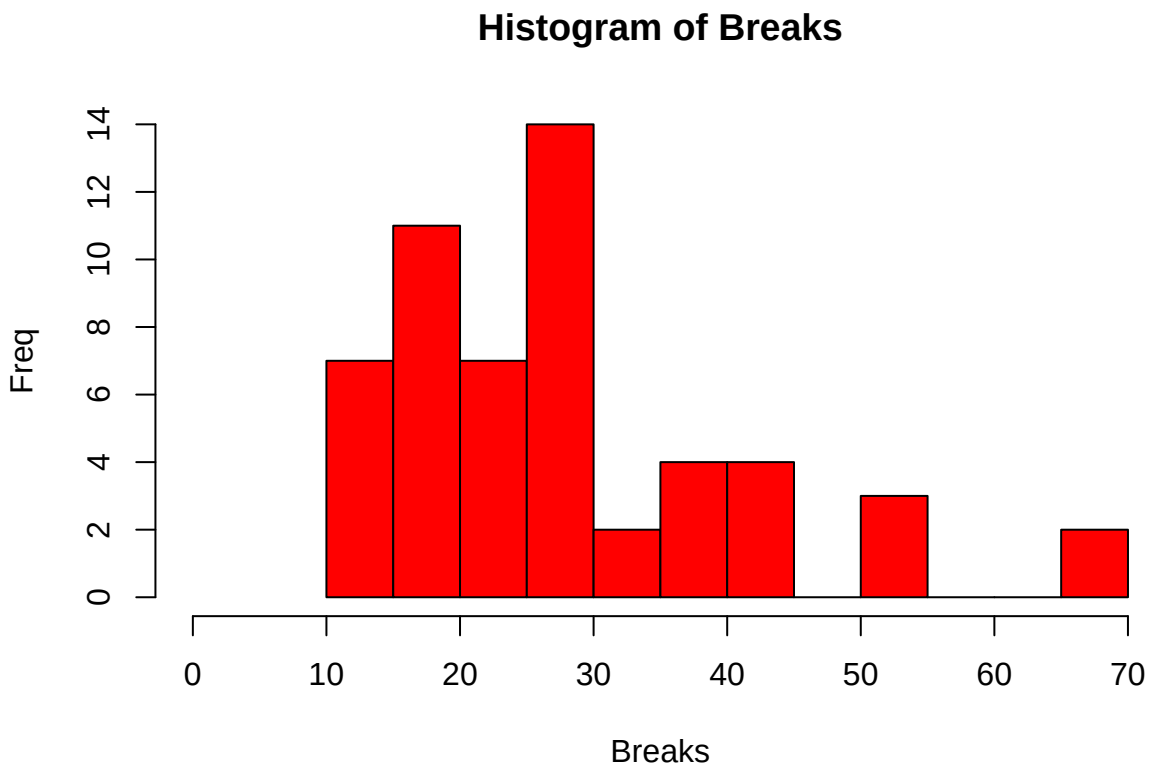
Importando los datos

```
data<-warpbreaks
head(data,10)
```

```
##    breaks wool tension
## 1      26    A       L
## 2      30    A       L
## 3      54    A       L
## 4      25    A       L
## 5      70    A       L
## 6      52    A       L
## 7      51    A       L
## 8      26    A       L
## 9      67    A       L
## 10     18    A       M
```

Histograma de rupturas

```
hist(data$breaks,
     main = "Histogram of Breaks",
     xlab = "Breaks",ylab = "Freq",
     col = "red",
     border = "black",
     xlim = c(0, max(data$breaks)),
     breaks = 20)
```



Media y varianza

```
mu <- mean(data$breaks)
var <- var(data$breaks)
```

Modelo de Poisson

```
poisson.model<-glm(breaks ~ wool + tension, data, family = poisson(link = "log"))
summary(poisson.model)
```

```
##
## Call:
## glm(formula = breaks ~ wool + tension, family = poisson(link = "log"),
##     data = data)
##
## Deviance Residuals:
##     Min       1Q   Median       3Q      Max
## -3.6871  -1.6503  -0.4269   1.1902   4.2616
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z|)
```

```
## (Intercept)  3.69196    0.04541  81.302  < 2e-16 ***
## woolB       -0.20599    0.05157  -3.994 6.49e-05 ***
## tensionM    -0.32132    0.06027  -5.332 9.73e-08 ***
## 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
## AIC: 493.06
##
## Number of Fisher Scoring iterations: 4
```

Lamentablemente vemos que la desviación residual es mayor que los grados de libertad asi que tendremos que hacer un modelo cuasipoisson.

```
poisson.model2<-glm(breaks ~ wool + tension, data = data, family = quasipoisson(link = "log"))
summary(poisson.model2)
```

```
##
## Call:
## glm(formula = breaks ~ wool + tension, family = quasipoisson(link = "log"),
##     data = data)
##
## Deviance Residuals:
##     Min       1Q   Median       3Q      Max
## -3.6871  -1.6503  -0.4269   1.1902   4.2616
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  3.69196    0.09374  39.384  < 2e-16 ***
## woolB       -0.20599    0.10646  -1.935 0.058673 .
## 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
```

Sin embargo, vemos que de nuevo la desviación es considerablemente alta.