

# Clase 4.0

## Análisis

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Estadística y Manejo de Datos con R (EMDR) — Virtual

# Modelos Lineales Generalizados (GLM)

# Modelos Lineales Generalizados

- Combinan variables predictoras discretas y continuas.
- Podemos especificar la naturaleza de la variable de respuesta mediante una función liga (por ejemplo: Gausiana, Poisson o Binomial).
- Cuando la variable de respuesta es binomial, también se conoce como *Regresión Logística*.

# Modelos Lineales Generalizados

- 1 variable continua.

```
?Puromycin  
head(Puromycin)
```

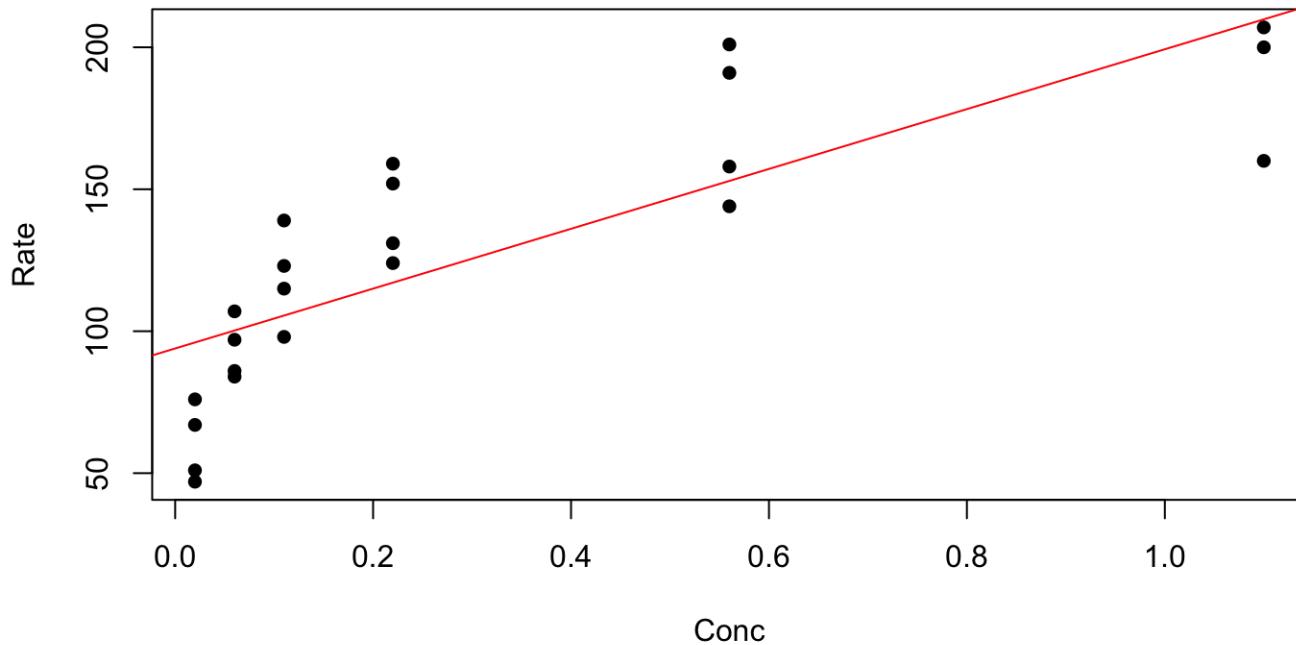
```
pmycin_glm <- glm(rate ~ conc, data = Puromycin)  
summary(pmycin_glm)
```

```
##  
## Call:  
## glm(formula = rate ~ conc, data = Puromycin)  
##  
## Deviance Residuals:  
##      Min        1Q    Median        3Q       Max  
## -49.861   -15.247    -2.861    15.686    48.054  
##  
## Coefficients:  
##             Estimate Std. Error t value Pr(>|t|)  
## (Intercept)  93.92      8.00  11.74 1.09e-10 ***  
## conc         105.40     16.92   6.23 3.53e-06 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## (Dispersion parameter for gaussian family taken to be 830.4261)  
##  
## Null deviance: 49665 on 22 degrees of freedom  
## Residual deviance: 17439 on 21 degrees of freedom  
## AIC: 223.78  
##  
## Number of Fisher Scoring iterations: 2
```

# Modelos Lineales Generalizados

- 1 variable continua.

```
plot(Puromycin$conc, Puromycin$rate, pch = 16, xlab = "Conc", ylab = "Rate")
abline(lm(rate ~ conc, data = Puromycin), col = "red")
```



# Modelos Lineales Generalizados

- 1 una variable discreta.

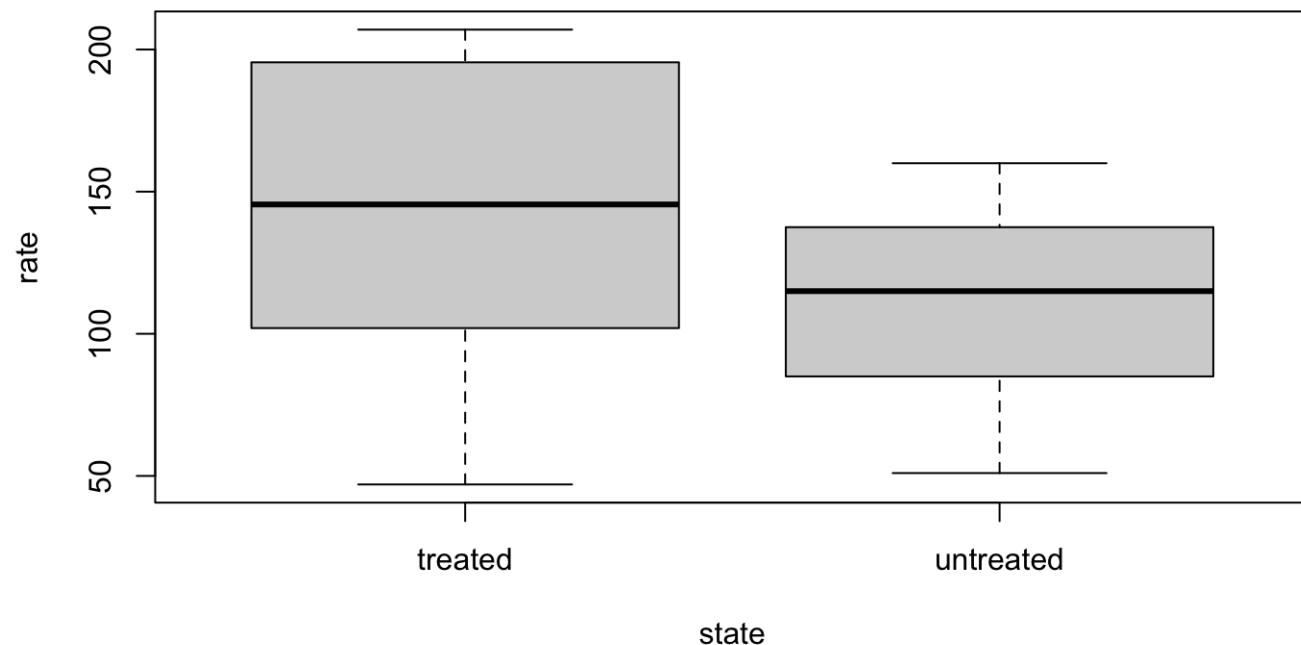
```
pmycin_glm <- glm(rate ~ state, data = Puromycin)
summary(pmycin_glm)
```

```
##
## Call:
## glm(formula = rate ~ state, data = Puromycin)
##
## Deviance Residuals:
##    Min      1Q  Median      3Q     Max
## -94.583 -30.655    4.273   40.273   65.417
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 141.58     13.24  10.690 5.94e-10 ***
## stateuntreated -30.86     19.15  -1.611    0.122
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 2104.814)
##
## Null deviance: 49665  on 22  degrees of freedom
## Residual deviance: 44201  on 21  degrees of freedom
## AIC: 245.17
##
## Number of Fisher Scoring iterations: 2
```

# Modelos Lineales Generalizados

- 1 una variable discreta.

```
boxplot(rate ~ state, data = Puromycin)
```



# Modelos Lineales Generalizados

- 2 variables e interacción.

```
pmycin_glm <- glm(conc ~ rate*state, data = Puromycin)
summary(pmycin_glm)
```

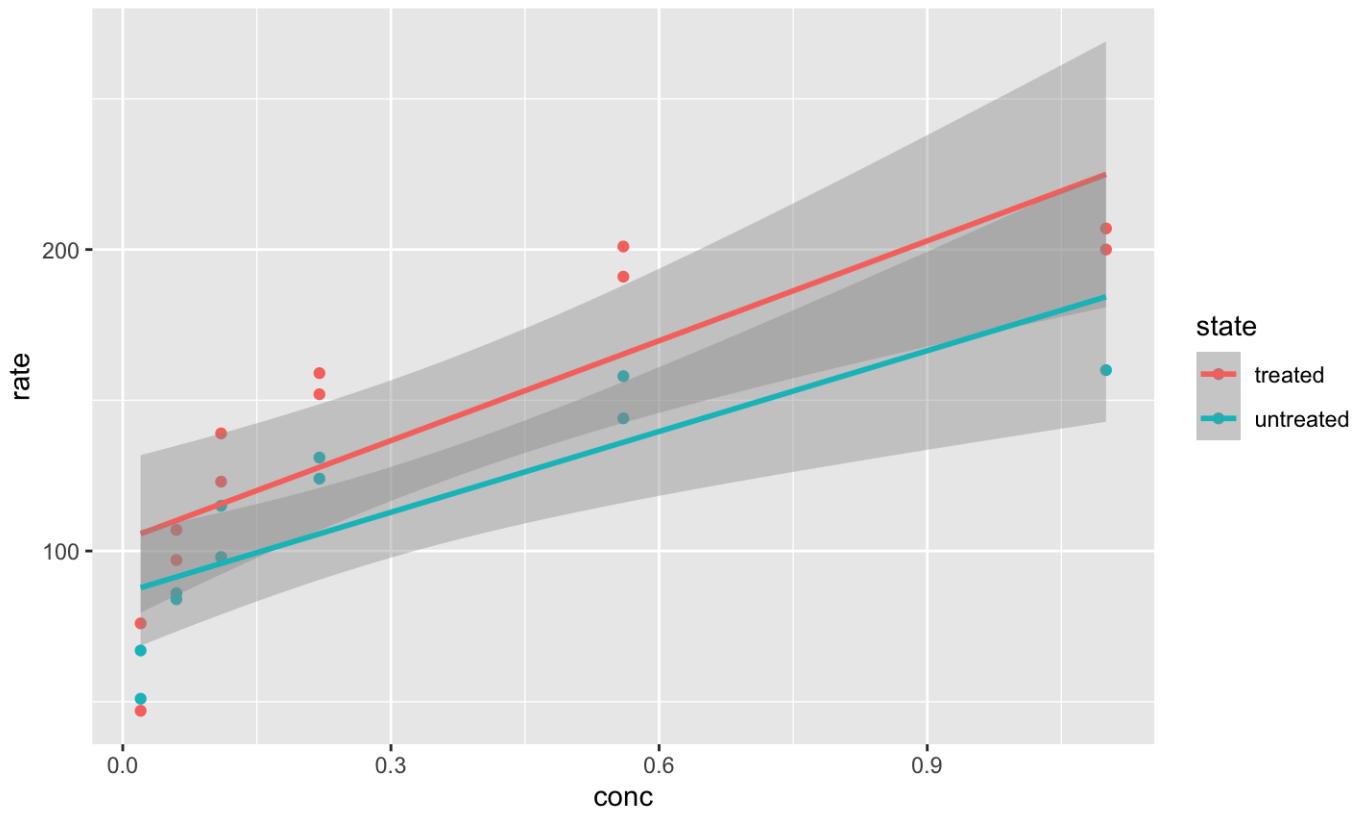
```
##
## Call:
## glm(formula = conc ~ rate * state, data = Puromycin)
##
## Deviance Residuals:
##      Min        1Q     Median        3Q       Max
## -0.23393  -0.15653  -0.06870   0.07926   0.45212
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)           -0.540523  0.187185 -2.888  0.00943 ** 
## rate                  0.006254  0.001245  5.025 7.52e-05 *** 
## stateuntreated       -0.017989  0.288642 -0.062  0.95096  
## rate:stateuntreated  0.001285  0.002265  0.567  0.57706  
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.04780568)
##
## Null deviance: 2.90099 on 22 degrees of freedom
## Residual deviance: 0.90831 on 19 degrees of freedom
## AIC: 0.94285
##
## Number of Fisher Scoring iterations: 2
```

# Modelos Lineales Generalizados

- 2 variables e interacción.

```
ggplot(Puromycin, aes(x=conc, y=rate, color=state)) + geom_point() + geom_smooth(method='lm')
```

```
## `geom_smooth()` using formula 'y ~ x'
```



# Modelos Lineales Generalizados

- Función liga.

```
?infert
```

```
head(infert)
```

```
##   education age parity induced case spontaneous stratum pooled.stratum
## 1    0-5yrs  26      6       1     1          2         1             3
## 2    0-5yrs  42      1       1     1          0         2             1
## 3    0-5yrs  39      6       2     1          0         3             4
## 4    0-5yrs  34      4       2     1          0         4             2
## 5   6-11yrs  35      3       1     1          1         5            32
## 6   6-11yrs  36      4       2     1          1         6            36
```

```
unique(infert$case)
```

```
## [1] 1 0
```

# Modelos Lineales Generalizados

- Función liga.

```
inf.bn <- glm(case ~ age + parity + education + spontaneous,
  data = infert, family = binomial())
summary(inf.bn)
```

```
##
## Call:
## glm(formula = case ~ age + parity + education + spontaneous,
##       family = binomial(), data = infert)
##
## Deviance Residuals:
##    Min      1Q  Median      3Q     Max
## -1.7219 -0.7698 -0.6353  0.9825  2.1823
##
## Coefficients:
##             Estimate Std. Error z value Pr(>|z| )
## (Intercept) -0.63833   1.33256 -0.479   0.6319
## age          0.01553   0.02920  0.532   0.5947
## parity       -0.30516   0.13729 -2.223   0.0262 *
## education6-11yrs -0.69194   0.76038 -0.910   0.3628
## education12+ yrs -0.84472   0.79294 -1.065   0.2867
## spontaneous   1.27783   0.22305  5.729 1.01e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
## Null deviance: 316.17 on 247 degrees of freedom
## Residual deviance: 278.26 on 242 degrees of freedom
## AIC: 290.26
```

# Modelos Lineales Generalizados

- Función liga.

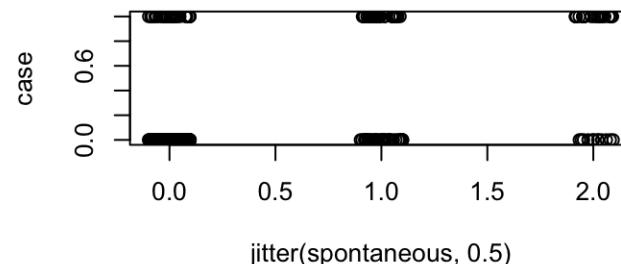
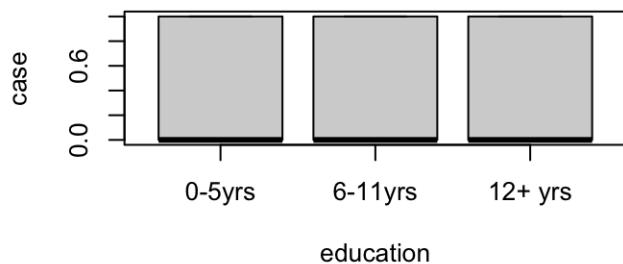
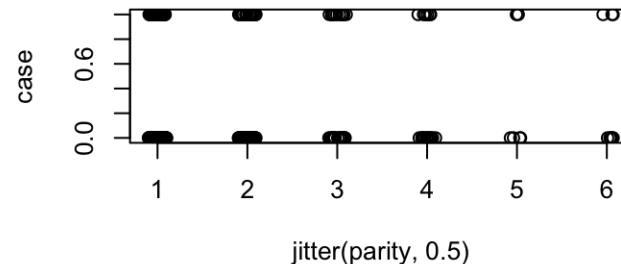
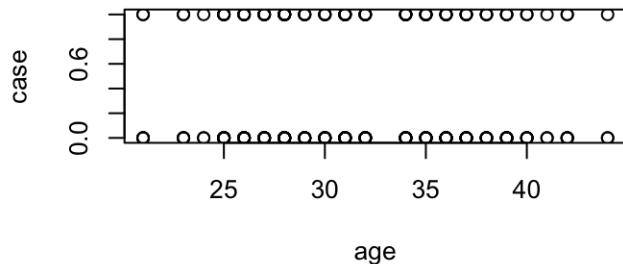
```
inf.no <- glm(case ~ age + parity + education + spontaneous,
  data = infert)
summary(inf.no)
```

```
##
## Call:
## glm(formula = case ~ age + parity + education + spontaneous,
##      data = infert)
##
## Deviance Residuals:
##    Min      1Q  Median      3Q     Max
## -0.7752 -0.2685 -0.1795  0.3808  0.9700
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 0.371789  0.258916   1.436   0.1523  
## age         0.002873  0.005664   0.507   0.6124  
## parity      -0.059536  0.026263  -2.267   0.0243 *  
## education6-11yrs -0.155096  0.146016  -1.062   0.2892  
## education12+ yrs -0.181238  0.151961  -1.193   0.2342  
## spontaneous   0.271910  0.041317   6.581 2.87e-10 *** 
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1935233)
##
## Null deviance: 55.222 on 247 degrees of freedom
## Residual deviance: 46.833 on 242 degrees of freedom
## AIC: 304.42
```

# Modelos Lineales Generalizados

- Comparación binomial.

```
par(mfrow=c(2,2))
with(infert, plot(case ~ age + jitter(parity,.5) + education + jitter(spontaneous,.5)))
```



# Modelos Lineales Generalizados

- Comparación *post hoc*.

```
library(multcomp)
```

```
summary(glht(inf.no, mcp(education="Tukey")))
```

```
##  
##   Simultaneous Tests for General Linear Hypotheses  
##  
##   Multiple Comparisons of Means: Tukey Contrasts  
##  
##  
## Fit: glm(formula = case ~ age + parity + education + spontaneous,  
##          data = infert)  
##  
## Linear Hypotheses:  
##                               Estimate Std. Error z value Pr(>|z|)  
## 6-11yrs - 0-5yrs == 0 -0.15510    0.14602 -1.062  0.519  
## 12+ yrs - 0-5yrs == 0 -0.18124    0.15196 -1.193  0.438  
## 12+ yrs - 6-11yrs == 0 -0.02614    0.06043 -0.433  0.896  
## (Adjusted p values reported -- single-step method)
```

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