Generalised linear models in R

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Researcher Development



Recap: Linear regression

Assumptions:

- A linear mean function is relevant.
- 2 Variances are equal across all predicted values of the response (homoscedatic).
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- Samples collected at random.
- 6 Errors are independent.

Generalised linear models (GLMs)

1 A linear mean (including any explanatory variables you want to) i.e $\beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$

2 A link function (like an "internal" transformation).

3 An error structure. So far we assumed normality $\epsilon \sim \mathcal{N}(0, \sigma^2)$

Link functions

Links your mean function to the scale of the observed data e.g.

$$E(Y) = g^{-1} \left(\beta_0 + \beta_1 X \right)$$

• $\mathbb{E}(Y)$ is the **expected value** (i.e. mean of Y).

• The function $g(\cdot)$ is known as the **link function**, and $g^{-1}(\cdot)$ denotes the **inverse** of $g(\cdot)$.

 $\textbf{1} \ \, \mathsf{A} \ \, \mathsf{linear} \ \, \mathsf{mean} \colon \ \, \beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$

② An error structure: $\epsilon \sim \mathcal{N}(0, \sigma^2)$

Solution Link function: identity $\mu = \beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$

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2 An error structure: $\epsilon \sim \mathcal{N}(0, \sigma^2)$

6 Link function: identity

$$\mu = \beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$$

 $\textbf{1} \ \, \mathsf{A} \ \, \mathsf{linear} \ \, \mathsf{mean} \colon \ \, \beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$

2 An error structure: $\epsilon \sim \mathcal{N}(0, \sigma^2)$

3 Link function: identity $\mu = \beta_0 + \beta_1 X_1 + \ldots + \beta_p X_p$

$$Y \sim \mathcal{N}(\mu, \sigma^2)$$
$$\mu = \beta_0 + \beta_1 X$$

GLMs in R

```
lm(height ~ weight, data=df)
```

Is equivalent to:

```
glm(height ~ weight, data=df, family=gaussian(link=identity))
```

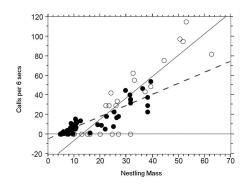
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family specifies the error structure and link function

Family	Link
gaussian	identity
binomial	logit, probit or cloglog
poisson	log, identity or sqrt
Gamma	inverse, identity or log
inverse.gaussian	1/mu^2
quasi	user-defined
quasibinomial	logit
quasipoisson	log

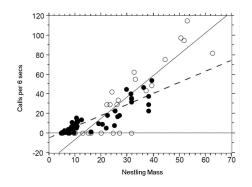
Poisson regression

Count data is discrete and non-negative



Poisson regression

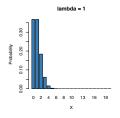
Count data is **discrete** and **non-negative**

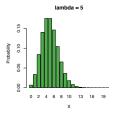


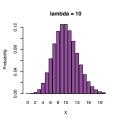
$$Y \sim \mathcal{N}(\mu, \sigma^2)$$
 $Y \sim \mathcal{P}ois(\mu)$
 $\mu = \beta_0 + \beta_1 X$ $\log \mu = \beta_0 + \beta_1 X$

Poisson distribution

- **Discrete** variable, defined on the range $0, 1, \ldots, \infty$.
- A single **rate** parameter λ , where $\lambda > 0$.
- Mean = λ
- Variance = λ







Poisson regression

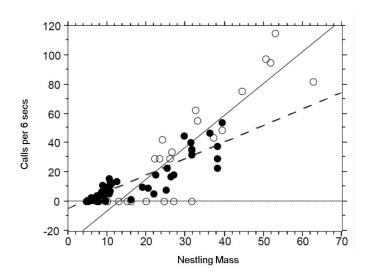
$$Y \sim \mathcal{P}ois(\lambda)$$
$$\log \lambda = \beta_0 + \beta_1 X$$

Using the rules of logarithm (i.e $\log \lambda = k$, then $\lambda = e^k$):

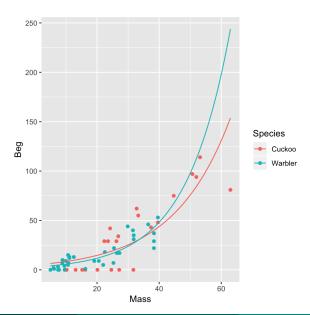
$$\log \lambda = \beta_0 + \beta_1 X$$
$$\lambda = e^{\beta_0 + \beta_1 X}$$

Thus we are effectively modelling the observed counts using an exponential distribution

glm(outcome ~ explanatory, data=df, family=poisson(link=log))



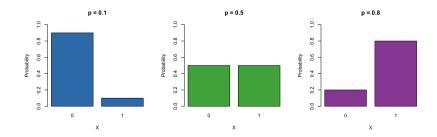
Cuckoo data



Consider a **categorical** response variable with two levels (e.g pass/fail). These type of **binary** data are assumed to be **Bernoulli** distributed:

$$Y \sim \mathcal{B}ern(p)$$

- A **probability** parameter p, where 0 .
- Mean = p
- Variance = p(1-p)



$$Y \sim \mathcal{N}(\mu, \sigma^2)$$
 $Y \sim \mathcal{P}ois(\lambda)$ $Y \sim \mathcal{B}ern(p)$
 $\mu = \beta_0 + \beta_1 X$ $\log \lambda = \beta_0 + \beta_1 X$ $?? = \beta_0 + \beta_1 X$

$$Y \sim \mathcal{N}(\mu, \sigma^2)$$
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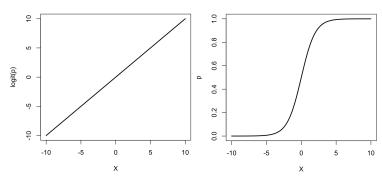
$$Y \sim \mathcal{B}ern(p)$$

$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 X$$

$$\log \operatorname{logit}(p) = \beta_0 + \beta_1 X$$

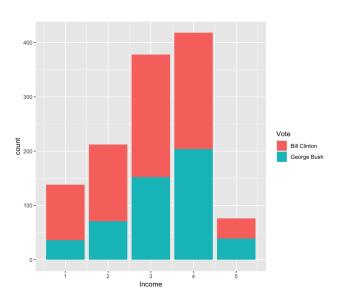
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$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 X$$
$$p = \frac{e^{\beta_0 + \beta_1 X}}{1 + e^{\beta_0 + \beta_1 X}}$$



glm(response ~ explanatory, data=df, family=binomial(link=logit))

1992 US election survey



```
fit <- glm(Vote ~ Income, data=USA, family=binomial(link=logit))
summary(fit)</pre>
```

```
##
## Call:
## glm(formula = Vote ~ Income, family = binomial(link = logit),
      data = USA)
## Deviance Residuals:
      Min
               1Q Median
                                         Max
## -1.2699 -1.0162 -0.8998 1.2152
                                     1 6199
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.3017 0.1828 -7.122 1.06e-12 ***
## Income
             0.3033 0.0551 5.505 3.69e-08 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1655.0 on 1221 degrees of freedom
## Residual deviance: 1623.5 on 1220 degrees of freedom
## ATC: 1627.5
##
## Number of Fisher Scoring iterations: 4
```

$$Y \sim \mathcal{B}ern(p)$$
$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 X$$

- '(Intercept)' = $\beta_0 = -1.3$
- 'Income' = $\beta_1 = 0.303$

It is common to interpret variables according to some **central tendency** e.g at the central income category (i.e X=3)

$$\begin{split} P(\text{Republican vote at } X=3) &= \mathsf{logit}^{-1} \left(-1.3 + 0.3 \times 3\right) \\ &= \frac{e^{-1.3 + 0.3 \times 3}}{1 + e^{-1.3 + 0.3 \times 3}} \\ &= 0.48. \end{split}$$

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