Generalised linear models in R

John Joseph Valletta

University of Exeter, Penryn Campus, UK

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

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

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

3 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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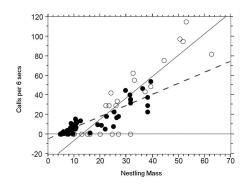
family specifies the error structure and link function

Default link functions

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

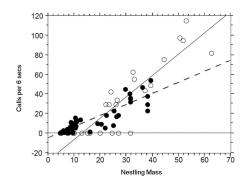
Poisson regression

Count data is discrete and non-negative



Poisson regression

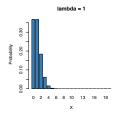
Count data is **discrete** and **non-negative**

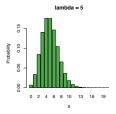


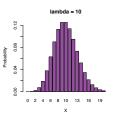
$$Y \sim \mathcal{N}(\mu, \sigma^2)$$
 $Y \sim \mathcal{P}ois(\mu)$
 $\mu = \beta_0 + \beta_1 X$ $\log \mu = \beta_0 + \beta_1 X$

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

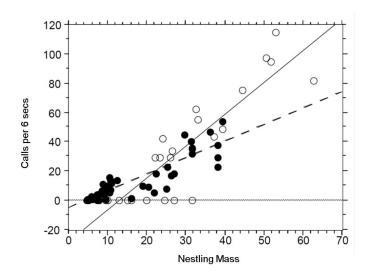
$$Y \sim \mathcal{P}ois(\lambda)$$
$$\log \lambda = \beta_0 + \beta_1 X$$

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

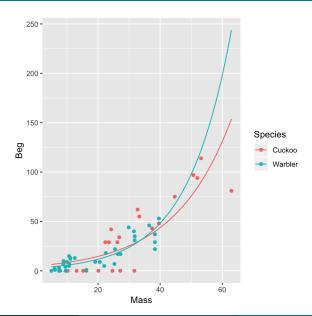
$$\log \lambda = \beta_0 + \beta_1 X$$
$$\lambda = e^{\beta_0 + \beta_1 X}$$

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

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



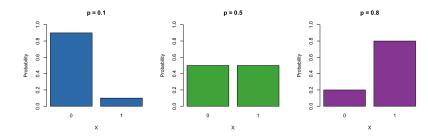
Cuckoo data



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)



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

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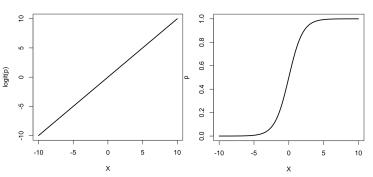
$$Y \sim \mathcal{B}ern(p)$$

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

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

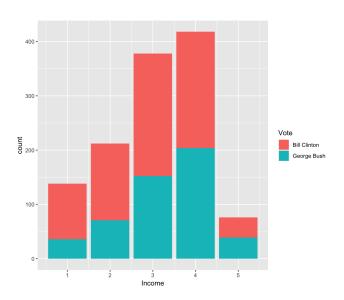
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$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 X$$
$$p = \frac{e^{\beta_0 + \beta_1 X}}{1 + e^{\beta_0 + \beta_1 X}}$$



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
```

1992 US election survey

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

- '(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)

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