

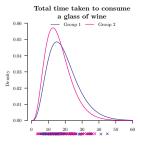


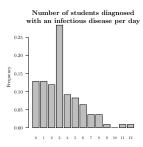
Generalised Linear Models (GLM)

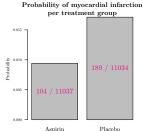
Cancer Research UK -19^{th} of July 2017

D.-L. Couturier / M. Dunning / R. Nicholls

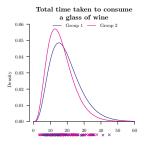
Examples of data with non-normal conditional distributions

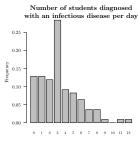


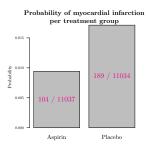




Examples of data with non-normal conditional distributions







Linear model not suitable:

Assumed model:

$$\begin{split} Y_i &= \mathbf{x}_i^T \boldsymbol{\beta} + \epsilon_i \text{ where } \epsilon_i \sim N(0, \sigma^2), \\ Y_i &| (\mathbf{x}_i, \boldsymbol{\beta}) \sim N(\mu_i, \sigma^2). \end{split}$$

- \triangleright theoretical range of $\epsilon_i = [-\infty, +\infty],$
- $\triangleright \mathbf{x}_i^T \boldsymbol{\beta}$ not bounded to $[0, \infty]$,
- $\triangleright Var[Y_i]$ independent of $E[Y_i]$.
- Solution:

$$Y_i|(\mathbf{x}_i,\boldsymbol{\beta}) \sim distribution(function(\mathbf{x}_i^T\boldsymbol{\beta}),\phi),$$

where distribution belongs to the exponential family and function is monotonically increasing.



GLM: conditional distributions

$$Y_i|(\mathbf{x}_i, \boldsymbol{\beta}) \sim distribution(function(\mathbf{x}_i^T \boldsymbol{\beta}), \phi),$$

- ► Some possible conditional *distributions*: statistical probability mass functions & density functions
 - ▶ Within the exponential family ['classical' GLM framework]

normal chi-squared Bernoulli Inverse Wishart exponential beta Poisson gamma Dirichlet Wishart ...

Outside the exponential family ['extended' GLM framework]

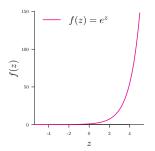
Box-Cox power exponential exponential Gaussian generalized beta generalized gamma generalized inverse Gaussian inverse Gaussian logistic power exponential reverse Gumbel skew power exponential Weibull Pareto type I, II, III Poisson inverse Gaussian



GLM: link functions

$$Y_i|(\mathbf{x}_i, \boldsymbol{\beta}) \sim distribution(function(\mathbf{x}_i^T \boldsymbol{\beta}), \phi),$$

- ► Most used link functions: connection between Y_i and $\mathbf{x}_i^T \boldsymbol{\beta}$



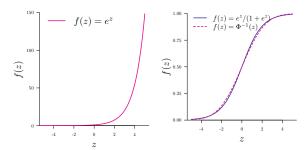


GLM: link functions

$$Y_i|(\mathbf{x}_i, \boldsymbol{\beta}) \sim distribution(function(\mathbf{x}_i^T \boldsymbol{\beta}), \phi),$$

- ► Most used link functions: connection between Y_i and $\mathbf{x}_i^T \boldsymbol{\beta}$
 - ▶ to restrict $f(\mathbf{x}_i^T\boldsymbol{\beta})$ to belong to $[0,\infty[:$ \triangleright log link: $f(z) = e^z$
 - ▶ to restrict $f(\mathbf{x}_i^T \boldsymbol{\beta})$ to belong to [0,1]:

 ▷ logit link: $f(z) = e^z/(1+e^z) = 1/(1+e^{-z})$ where z is positive
 ▷ probit link: $f(z) = \Phi^{-1}(z)$, where Φ denotes the N(0,1).





Distribution for dichotomous variates: Bernoulli

Example:

in Jones (Unpublished BSc dissertation, University of Southampton, 1975), the main outcome is the presence/absence of bronchitis:

lf

- n independent experiments,
- outcome of each experiment is dichotomous (success/failure),
- lacktriangle the probability of success π is the same for all experiments,

then, each dichotomous experiment, Y_i , follows a Bernoulli distribution with parameter π :

$$Y_i \sim Bernoulli(\pi)$$

$$P(Y_i = 1) = \pi$$

$$P(Y_i = 0) = 1 - \pi$$



Logistic regression: GLM for dichotomous variates

Example:

in Jones (Unpublished BSc dissertation, University of Southampton, 1975), the main outcome is the presence/absence of bronchitis as a function of the daily number of smoked cigarettes (X_1) and level of pollution (X_2) :

Sample of 212 men in Cardiff:
$$i = 1$$
 $i = 2$ $i = 3$ \cdots $i = 212$

lf

- ▶ n independent experiments,
- outcome of each experiment is dichotomous (success/failure),
- \blacktriangleright the probability of success π is the same for all experiments given the covariates,

then, each dichotomous experiment, Y_i , follows a Bernoulli distribution with parameter π_i :

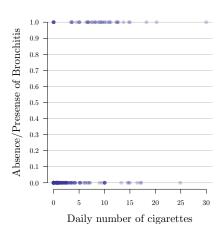
$$Y_i \sim Bernoulli(\pi_i)$$
 where $\pi_i = \frac{e^{\mathbf{x}_i^T \boldsymbol{\beta}}}{1 + e^{\mathbf{x}_i^T \boldsymbol{\beta}}}$
$$P(Y_i = 1) = \pi_i$$

$$P(Y_i = 0) = 1 - \pi_i$$



Example:

$$P(Y_i = 1) = \pi_i = \frac{e^{\beta_0 + \beta_1 x_{i1}}}{1 + e^{\beta_0 + \beta_1 x_{i1}}}$$

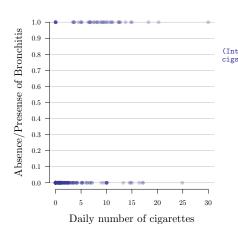


Example:

Model the probability of presence of bronchitis as a function of the daily number of smoked cigarettes (X_1) :

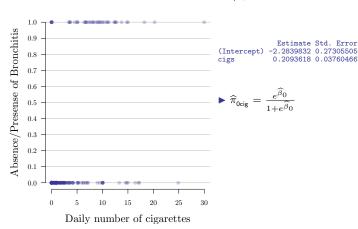
$$P(Y_i = 1) = \pi_i = \frac{e^{\beta_0 + \beta_1 x_{i1}}}{1 + e^{\beta_0 + \beta_1 x_{i1}}}$$

0.2093618 0.03760466



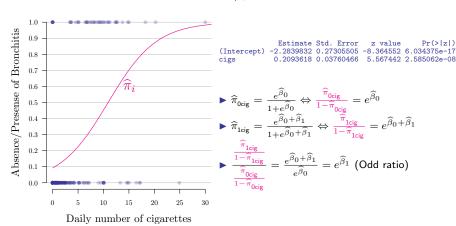
Example:

$$P(Y_i = 1) = \pi_i = \frac{e^{\beta_0 + \beta_1 x_{i1}}}{1 + e^{\beta_0 + \beta_1 x_{i1}}}$$



Example:

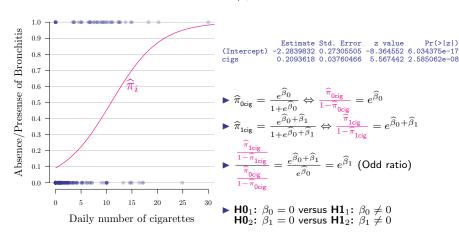
$$P(Y_i = 1) = \pi_i = \frac{e^{\beta_0 + \beta_1 x_{i1}}}{1 + e^{\beta_0 + \beta_1 x_{i1}}}$$





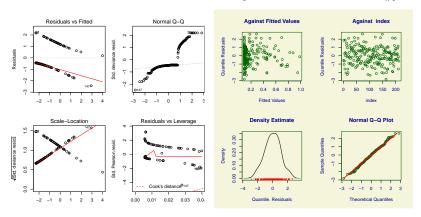
Example:

$$P(Y_i = 1) = \pi_i = \frac{e^{\beta_0 + \beta_1 x_{i1}}}{1 + e^{\beta_0 + \beta_1 x_{i1}}}$$



Logistic regression: model check

- ▶ pearson residuals $(y_i \widehat{\pi})/\sqrt{\mathsf{Var}(\widehat{\pi})}$,
- ▶ deviance residuals [Default in R],
- randomised normalised quantile residuals [Default in package gamlss()]





Distribution for count data: Poisson

Example:

Interest for the number of high school students diagnosed with an infectious disease

Sample of 115 days:
$$t=1$$
 $t=2$ $t=3$ \cdots $t=115$ y_i 6 8 12 \cdots 0

If, during a time interval or in a given area,

- events occur independently,
- ▶ at the same rate,
- and the probability of an event to occur in a small interval (area) is proportional to the length of the interval (size of the area),

then,

ightharpoonup a count occurring in a fixed time interval or in a given area, Y, may be modelled by means of a Poisson distribution with parameter μ :

$$Y \sim Poisson(\mu)$$
 where $\mu = E[Y] = Var[Y]$,

 \blacktriangleright the probability of observing x events during a fixed time interval or in a given area is given by

$$P(Y = y | \mu) = \frac{\mu^y e^{-\mu}}{y!}.$$



Poisson regression: GLM for count data

Example:

Interest for the number of high school students diagnosed with an infectious disease as a function of the number of days from the initial outbreak

Sample of 115 days:
$$t = 1$$
 $t = 2$ $t = 3$ \cdots $t = 115$

If, during a time interval or in a given area,

- events occur independently given the covariates,
- ▶ at the same rate given the covariates,
- and the probability of an event to occur in a small interval (area) is proportional to the length of the interval (size of the area) given the covariates,

then,

• each count occurring in a fixed time interval or in a given area, Y_t , may be modelled by means of a Poisson distribution with parameter μ_t :

$$Y_t \sim Poisson(\mu_t)$$
 where $\mu_t = \mathsf{E}[Y] = \mathsf{Var}[Y] = e^{\mathbf{x}_i^T \boldsymbol{\beta}}$,

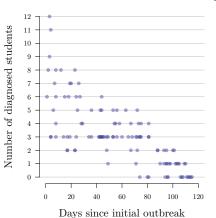
lacktriangle the probability of observing y during the fixed time interval or in the given area is given by

$$P(Y_t = y_t | \mu_t) = \frac{\mu_t^{y_t} e^{-\mu_t}}{y_t!}.$$



Example:

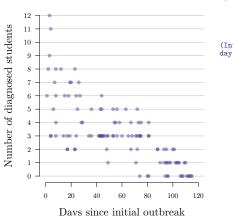
$$\mu_t = \frac{e^{\beta_0 + \beta_1 t}}{1 + e^{\beta_0 + \beta_1 t}}$$



Example:

Model the mean count of diagnosed students, μ_t , as a function of the number of days from the outbreak (T):

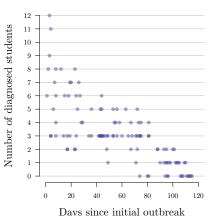
$$\mu_t = \frac{e^{\beta_0 + \beta_1 t}}{1 + e^{\beta_0 + \beta_1 t}}$$



| Estimate Std. Error z value | Pr(>|z | (Intercept) | 1.99023497 | 0.083935207 | 23.71156 | 2.739875e-1 | day | -0.01746317 | 0.001726709 | -10.11356 | 4.810392e-

Example:

$$\mu_t = \frac{e^{\beta_0 + \beta_1 t}}{1 + e^{\beta_0 + \beta_1 t}}$$

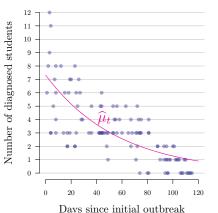




$$\blacktriangleright \ \widehat{\mu}_{\rm day0} = e^{\widehat{\beta}_0}$$

Example:

$$\mu_t = \frac{e^{\beta_0 + \beta_1 t}}{1 + e^{\beta_0 + \beta_1 t}}$$



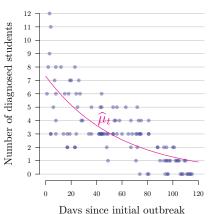
$$\widehat{\mu}_{\text{day0}} = e^{\widehat{\beta}_0}$$

$$\widehat{\mu}_{\text{dev}1} = e^{\widehat{\beta}_0 + \widehat{\beta}_1}$$

$$\frac{\widehat{\mu}_{\mathsf{day1}}}{\widehat{\mu}_{\mathsf{day0}}} = \frac{e^{\widehat{\beta}_0 + \widehat{\beta}_1}}{e^{\widehat{\beta}_0}} = e^{\widehat{\beta}_1}$$

Example:

$$\mu_t = \frac{e^{\beta_0 + \beta_1 t}}{1 + e^{\beta_0 + \beta_1 t}}$$



$$\blacktriangleright \ \widehat{\mu}_{\rm day0} = e^{\widehat{\beta}_0}$$

$$\mu_{\mathsf{day1}} = e^{\widehat{\mu}_{\mathsf{day1}}}$$

$$\frac{\widehat{\mu}_{\mathsf{day1}}}{\widehat{\mu}_{\mathsf{day1}}} = \frac{e^{\widehat{\beta}_0 + \widehat{\beta}_1}}{\widehat{\beta}_0} = e^{\widehat{\beta}_1}$$

▶
$$\mathbf{H0_1}$$
: $\beta_0 = 0$ versus $\mathbf{H1_1}$: $\beta_0 \neq 0$ $\mathbf{H0_2}$: $\beta_1 = 0$ versus $\mathbf{H1_2}$: $\beta_1 \neq 0$

Poisson regression: model check

- ▶ pearson residuals $(y_i \widehat{\pi})/\sqrt{\mathsf{Var}(\widehat{\pi})}$,
- ▶ deviance residuals [Default in R],
- randomised normalised quantile residuals [Default in package gamlss()]

