

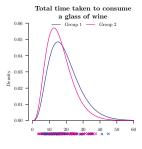


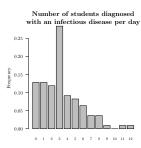
Generalised Linear Models (GLM)

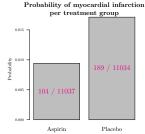
Cancer Research UK – 14th of May 2021

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

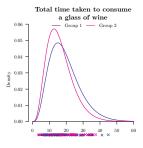
Examples of data with non-normal conditional distributions

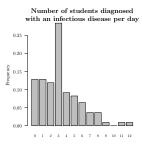


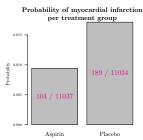




Examples of data with non-normal conditional distributions







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 $[0, \infty]$ or [0, 1],
- \triangleright Var[Y_i] independent of $E[Y_i]$.
- Solution:

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

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



GLM: conditional distributions

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

- ► 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 Gaussian inverse Gaussian exponential Gaussian logistic generalized beta generalized gamma generalized inverse skew power exponential skew power exponential

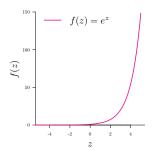
Weibull Pareto type I, II, III Poisson inverse Gaussian



GLM: link functions

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

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





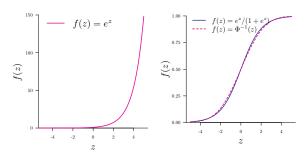
GLM: link functions

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

- 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 [0,1]:

 ▷ logit link: $f(z) = e^z/(1+e^z) = 1/(1+e^{-z})$ where z is positive
 ▷ probit link: $f(z) = \Phi(z)$, 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),
- ightharpoonup 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 :

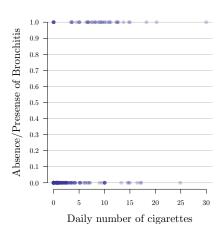
$$Y_i \sim Bernoulli(\pi_i)$$
 where $\pi_i = \frac{e^{\mathbf{x}_i^T \boldsymbol{\beta}}}{1 + e^{\mathbf{x}_i^T \boldsymbol{\beta}}}$
$$P(Y_i = 1) = \pi_i$$

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



Example:

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

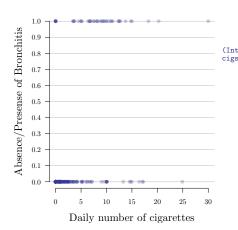


Example:

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

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

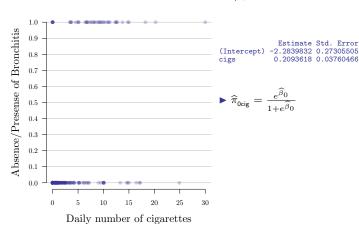
0.2093618 0.03760466



5.567442 2.585062e-08

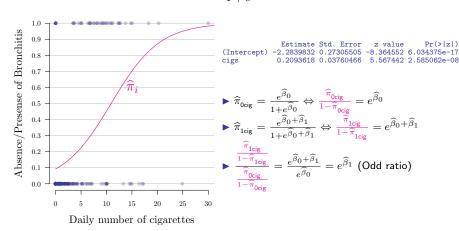
Example:

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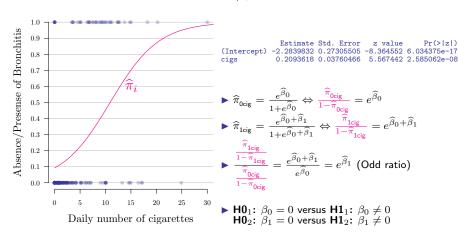
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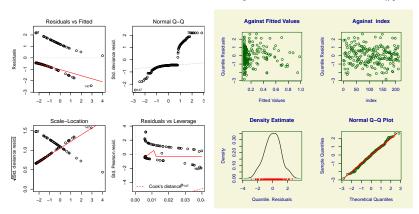
Example:

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



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 $\mu = E[Y] = Var[Y]$,

lacktriangle 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

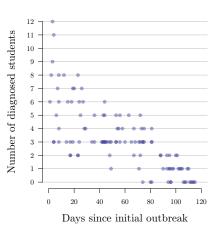
$$P(Y_t = y_t | \mu_t) = \frac{\mu_t^{y_t} e^{-\mu_t}}{y_t!}.$$



Example:

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

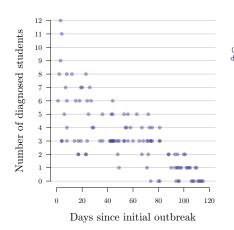
$$\mu_t = e^{\beta_0 + \beta_1 t}$$



Example:

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

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

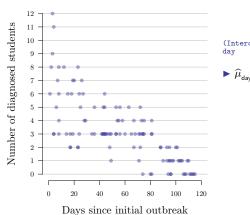


Estimate Std. Error z value Pr(>|z (Intercept) 1.99023497 0.083935207 23.71156 2.739875e-1: day -0.01746317 0.001726709 -10.11356 4.810392e-:

Example:

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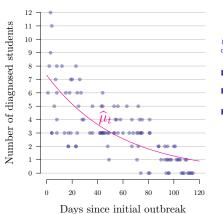




Example:

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

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| Estimate | Std. Error | z value | Pr(>|z| | (Intercept) | 1.99023497 | 0.083935207 | 23.71156 | 2.739875e-12 | day | -0.01746317 | 0.001726709 | -10.11356 | 4.810392e-2

$$\mathbf{\hat{\mu}}_{\cdot \cdot \cdot \cdot} = e^{\widehat{\beta}_0}$$

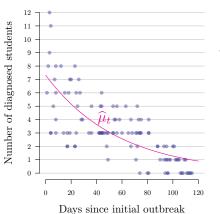
$$\widehat{\mu}_{1,1}=e^{\widehat{eta}_0+\widehat{eta}_1}$$

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

Example:

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

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



| Estimate Std. Error z value | Pr(>|z| (Intercept) 1.99023497 0.083935207 23.71156 2.739875e-12 | day -0.01746317 0.001726709 -10.11356 4.810392e-2

$$\mathbf{P} \widehat{\mu}_{\mathbf{A} = \mathbf{0}} = e^{\widehat{\beta}_0}$$

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

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

▶ **H0**₁:
$$\beta_0 = 0$$
 versus **H1**₁: $\beta_0 \neq 0$ **H0**₂: $\beta_1 = 0$ versus **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],
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