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What you have never been told about GLM

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Linear Model

Goal:

To find out if there exists a linear relationship between the random variable (rv) Y and the covariables $X_1, X_2, X_3, ..., X_n$.

Model:

$$\forall i \ Y_i = \beta_0 + x_{i1}\beta_1 + x_{i2}\beta_2 + \dots + x_{ip}\beta_p + \epsilon_i = \mu_i + \epsilon_i$$

Hipothesis:

- $\forall i \ \epsilon_i \sim N(0, \sigma_i^2)$.
- $\forall i \ \sigma_i^2 = \sigma^2$ (homoscedasticity).
- $\forall i, j, i \neq j \ \epsilon_i$ independent of ϵ_j .
- X values are fixed or if random, they are independent of errors.

Linear Model

In matrix form:

$$\begin{bmatrix} Y_1 \\ Y_2 \\ Y_3 \\ \vdots \\ Y_n \end{bmatrix} = \begin{bmatrix} 1 & x_{11} & x_{12} & x_{13} & \dots & x_{1p} \\ 1 & x_{21} & x_{22} & x_{23} & \dots & x_{2p} \\ 1 & x_{31} & x_{32} & x_{33} & \dots & x_{3p} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ 1 & x_{n1} & x_{n2} & x_{n3} & \dots & x_{np} \end{bmatrix} \cdot \begin{bmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \\ \vdots \\ \beta_p \end{bmatrix} + \begin{bmatrix} \epsilon_1 \\ \epsilon_2 \\ \epsilon_3 \\ \vdots \\ \epsilon_n \end{bmatrix}$$

So,
$$Y = X\beta + \epsilon$$

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Generalized Linear Model

From the previous expression, it is easy to see that $Y|X \sim N(X\beta, \sigma^2 \cdot Id_n)$ But what happens if Y is not normal distributed and we still want to find a linear relationship? In such situations we can use GLM, since they allow to assume that:

- Y follows a more general probability distribution, no necessarily Normal.
- The linearity is between a transformation of $\mu = E(Y|X)$ and the covariables (and not Y directly).

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Random component
Deterministic component
Link function

GLM components

A GLM has three components:

- Random component.
- Deterministic component.
- · Link function.

Random component

Given a random vector $Y = (Y_1, Y_2, Y_3, ..., Y_n)^t$ and given that $\mu = E(Y) = (\mu_1, \mu_2, \mu_3, ..., \mu_n)^t$, each Y_i should be written as:

Goodness of fit

$$Y_i \sim expigg(rac{y_i heta_i-b(heta_i)}{a(\phi)}+c(y_i,\phi)igg)$$

Comments:

- $h(\mu_i) = \theta_i$.
- ϕ is known as dispersion parameter (also detonted as σ^2).

Deterministic component

$X_{n \times p} \beta_{p \times 1}$ where p < n

$$\begin{bmatrix} x_{11} & x_{12} & x_{13} & \dots & x_{1p} \\ x_{21} & x_{22} & x_{23} & \dots & x_{2p} \\ x_{31} & x_{32} & x_{33} & \dots & x_{3p} \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ x_{n1} & x_{n2} & x_{n3} & \dots & x_{np} \end{bmatrix} \cdot \begin{bmatrix} \beta_1 \\ \beta_2 \\ \beta_3 \\ \vdots \\ \beta_p \end{bmatrix}$$

Random component Deterministic component Link function

Link function

Any monotone differentiable function η :

$$\eta = g(\mu) = X\beta$$

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$$f(y; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(y-\mu)^2}{2\sigma^2}\right)$$

This can be written as:

$$\exp\!\left(\frac{y\mu-\frac{1}{2}\mu^2}{\sigma^2}\!-\!\frac{y^2}{2\sigma^2}\!-\!\frac{1}{2}\log(2\pi\sigma^2)\right)$$

- $\theta = \mu$,
- h is the identity function.

$$\bullet \ a(\phi) = \phi = \sigma^2.$$

•
$$b(\theta) = \frac{\theta^2}{2}$$
.

•
$$c(y;\phi) = \frac{-y^2}{2\sigma^2} - log(\sqrt{2\pi\sigma^2})$$

$$Y_i \sim exp\left(\frac{y\theta - b(\theta)}{a(\phi)} + c(y,\phi)\right)$$

•
$$h(\mu) = \theta$$
.

$$f(y; m, p) = \binom{m}{y} p^{y} (1-p)^{m-y}$$

$$Y_i \sim exp\left(\frac{y\theta - b(\theta)}{a(\phi)} + c(y,\phi)\right)$$

This can be written as:

•
$$h(\mu) = \theta$$
.

$$\exp\left[ylog\left(\frac{\mu}{m-\mu}\right) - mlog\left(\frac{m}{m-\mu}\right) + log\left(\frac{m}{y}\right)\right]$$

•
$$\theta = \log\left(\frac{\mu}{m-\mu}\right) = h(\mu),$$

•
$$a(\phi) = 1$$
.

•
$$b(\theta) = log(1 + e^{\theta}).$$

•
$$c(y; \phi) = log\binom{m}{v}$$

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Link function

It is the link between E(Y) and the covariables $X_1, ..., X_p$. Remember:

•
$$g(\mu) = X\beta$$
.

Any monotone differentiable function.

Link function

- In the normal distribution, we have seen that $\theta=\mu.$ So h is the identity function.
- In the binomial distribution, we have seen that $\theta = log\left(\frac{\mu}{m-\mu}\right) = h(\mu)$.
- When g is choosen so that g=h, then it is said to use the canonical link function. In this case, $\theta=g(\mu)=X\beta$

Link function

In the normal case, when canonical link is used:

$$\bullet \ \mu = \beta_0 + X_1 \beta_1 + \cdots + X_p \beta_p$$

In the binomail case, when canonical link is used:

•
$$log\left(\frac{\mu}{m-\mu}\right) = \beta_0 + X_1\beta_1 + \cdots + X_p\beta_p$$
.

•
$$\mu = \frac{e^{X\beta}}{1+e^{X\beta}}$$
.

$$p = \frac{1}{m} \cdot \frac{e^{X\beta}}{1 + e^{x\beta}}.$$

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Variance function

Given that

$$Y \sim exp \left(rac{y heta - b(heta)}{a(\phi)} + c(y; \phi)
ight)$$

We can obtain the following expressions:

- $E(Y) = b'(\theta)$.
- $Var(Y) = a(\phi)b''(\theta)$.
- $V(\mu) = b''(\theta)$ is known as the **Variance function**.
- In case of binomial distribution, $V(\mu) = \frac{\mu(1-\mu)}{m}$.
- In case of normal distribution, $V(\mu)=1$. So, homoscedasticity must hold.

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Parameters estimation

To find the maximum likelihood estimator of β , the following iterative equation has to be solved:

$$X^{t}WXb^{m+1} = X^{t}WZ$$

where:

- W depends on the variance function and Z depends.
- Z depends on x, b^m , y, μ and η .

Parameters estimation

Given b^m , we proceed as follows:

- Get $\eta = X\beta^m$.
- Get $\mu = g^{-1}(X\beta^m)$.
- Get $\theta = h(\mu)$.
- Get Var(Y).
- Get W and Z.
- Get b^{m+1} from $X^t W X b^{m+1} = X^t W Z$.

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Predictions

Given $\hat{\beta}$ (estimation of β), the predicted mean value is equal to:

$$\hat{y} = \hat{\mu} = g^{-1}(X\beta)$$

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Goodness of fit

- The null model is defined as the model with just one variable (the intercept).
- The full model is defined as the model with as many variables as observations. It is the perfect model.
- We want a model as close as possible to the full model but with less variables. We will test our model with a hypothesis test.

- Let $I(\hat{\mu}, \phi, y)$ be the value of the log-likelihood corresponding to our model.
- Let $I(y, \phi, y)$ be the value of the log-likelihood corresponding to the perfect model.

The scaled deviance is defined as:

$$D^*(y, \hat{\mu}) = 2(I(y, \phi, y) - I(\hat{\mu}, \phi, y))$$

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Goodness of fit

In order to check if our model is as good as the perfect model, the following test is done:

 $\begin{cases} H_0 \colon \text{Our model is as good as the perfect one} \\ H_1 \colon \text{Our model is not as good the perfect one} \end{cases}$

Under H_0 , asymptotically $D^*(y, \hat{\mu}) \sim \chi^2_{n-p}$, where p is the number of parameters of our model.