Scoring

M2 D3S/EGR 2025-2026

Louis Olive

louis.olive@gmail.com / louis.olive@ut-capitole.fr

September 23, 2025

Outline

Outline

• The Logistic Regression model

Informal introductory example

We provide here some intuitions leading to the Logistic Regression model using a simulated data set from James et al. (2021) (the **Default** data set).

▼ Code

```
1 # Default data set (simulated) from ESLII/ISLR
2 default_data <- ISLR2::Default %>%
3     as_tibble()
4
5 glimpse(default_data)
```

This is a toy data set used for teaching purposes containing information on ten thousand customers.

The aim here is to assess which customers will **default** on their credit card debt (the target or response variable) based on the current credit card **balance** and other individual characteristics (the predictors or feature vector).

Informal introductory example

We can start to explore the Default data with a scatterplot (Figure 1) of the target variable (**default**) with respect to a predictor (**balance**):

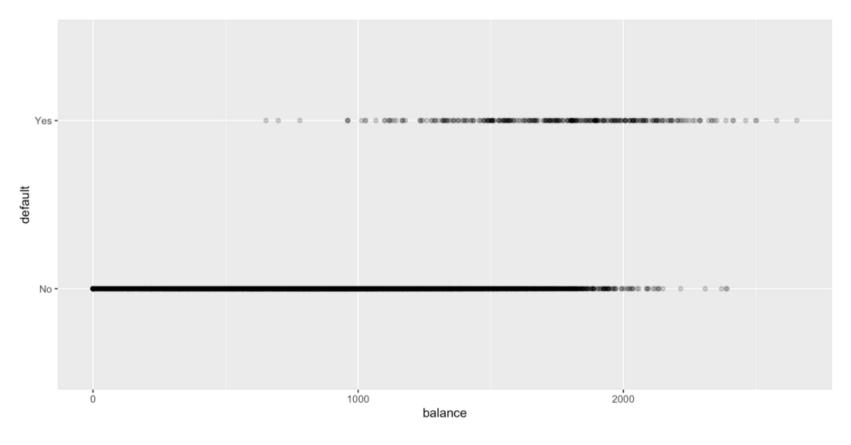
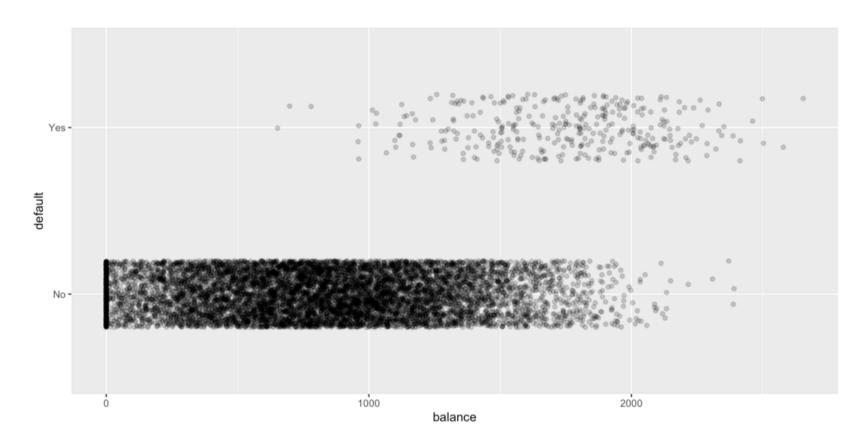


Figure 1: Scatterplot of variable *default* with respect to credit card *balance* for 10000 customers

Informal introductory example

In this scatterplot, all points fall on one of two parallel lines representing the absence (No) or occurrence (Yes) of **default**. We "jitter" the data vertically to avoid overplotting. The plot below shows that the response variable is imbalanced towards the absence of default:



Informal introductory example

We also show the boxplots of credit cards **balance** with respect to **default** status:

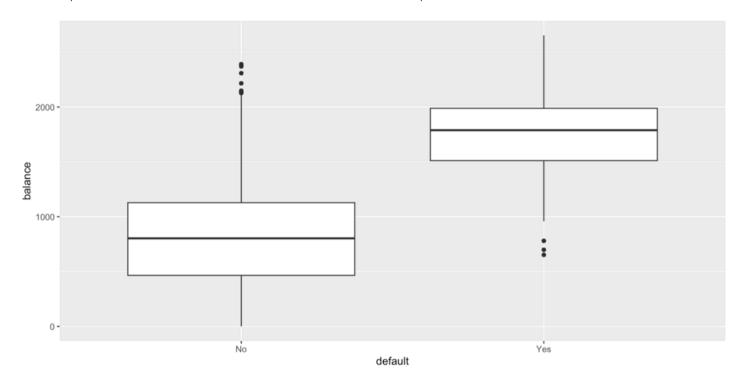


Figure 2: Variable **balance** with respect to **default** status

We can see from Figure 1 and Figure 2 that default tends to be more prevalent for accounts with a high balance. However it is difficult to guess a simple relationship between default and balance.

Informal introductory example

To investigate further we discretise the balance variables by classes of width 300\$ and compute the mean of response variable (**default** is Yes) within each balance class:

#	A tibble: 9	× 6				
	balance_bins	min	max	No	Yes	`Mean(default)`
	<fct></fct>	<dbl></dbl>	<dbl></dbl>	<int></int>	<int></int>	<dbl></dbl>
1	[0,300)	0	300	1497	0	0
2	[300,600)	300	600	1784	0	0
3	[600,900)	600	900	2305	3	0.0013
4	[900,1200)	900	1200	2098	19	0.009
5	[1200,1500)	1200	1500	1330	53	0.0383
6	[1500,1800)	1500	1800	527	96	0.154
7	[1800,2100)	1800	2100	115	114	0.498
8	[2100,2400)	2100	2400	11	41	0.788
9	[2400,2700)	2400	2700	0	7	1

Informal introductory example

Then we plot the mean of default (in red) within each balance class (of width 300\$):

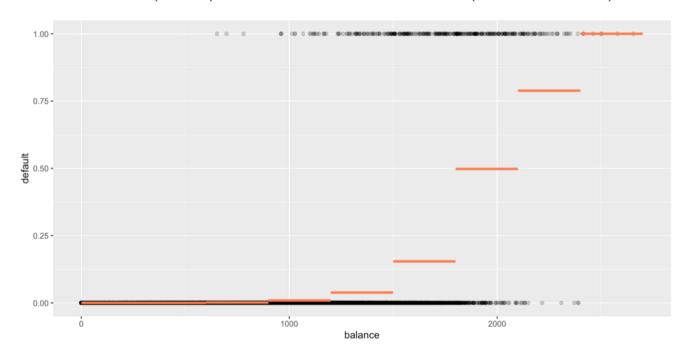
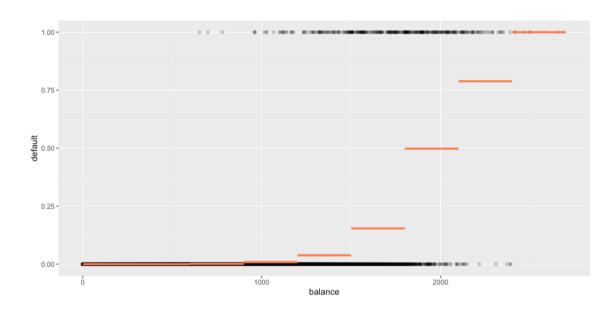


Figure 3: Mean occurrence of *default* within *balance* classes

The relationship between the mean occurrence of **default** and **balance** is easier to read.

Figure 3 clearly shows that as balance increases, the proportion of customers defaulting on their credit card increases.

Informal introductory example



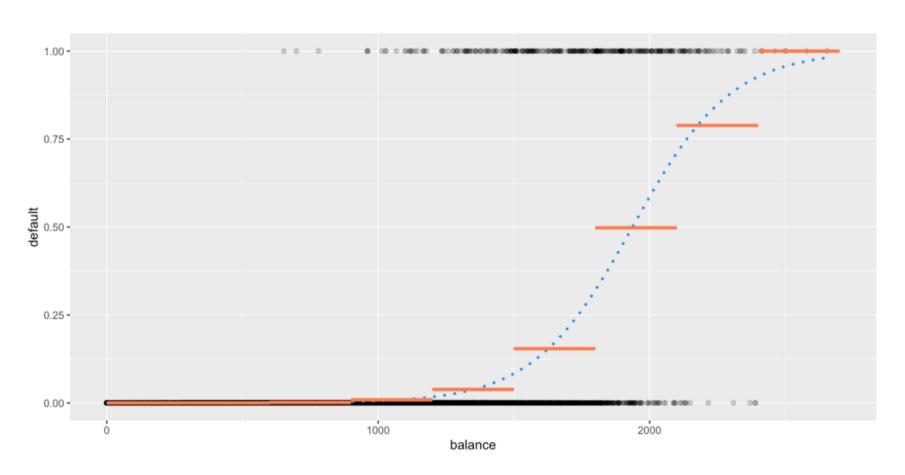
We also notice that the mean default occurrence with respect to balance classes follows a kind of "S"-shaped curve or **sigmoid** function. Going further and informally, considering that the mean of default occurrence is an estimate of $\mathbf{E}[Y|X=x]$ for each balance classes an idea would be to model:

$$\mathbb{E}[Y|X=x]=\mu_{eta}(x)$$

where μ_{β} is a **sigmoid** function in [0,1].

Informal introductory example

The Logistic Regression model uses the **sigmoid** function $\sigma: x \to \sigma(x) = \frac{e^x}{1+e^x}$ also known as the logistic function. Below a simple transform of this logistic function has been "fitted" (blue dots) to the **default** ~ **balance** data set:



A more formal definition - the discriminative approach

Reminding the statistical learning / scoring concepts introduced before. We try to predict the output default ($Y \in \{0,1\}$) using a training set of inputs X: this is a binary classification problem.

We remind that to estimate an optimal classifier for output $Y \in \{0,1\}$ using input $X = (X_1, \dots, X_p)$ one approach was to:

- ullet model the conditional distribution Y|X (the **discriminative** approach),
- estimate $\eta(x)$ with:

$$\eta(x)=\mathbb{P}[Y=1|X=x)]=\mathbb{E}[Y=1|X=x)]$$

- $\eta(x)$ can be used as a Scoring function
- ullet define the classifier f_s using a cutoff s and Scoring function $\eta(x)$ to predict output Y:

$$f_s(x) = egin{cases} 1 & ext{if } \eta(x) \geq s \ 0 & ext{otherwise} \end{cases}$$

A more formal definition

In the case of **Logistic Regression** classifier, we model:

$$|Y|X = x |\sim B(\eta(x))$$

with

$$\eta(x) = \sigma(x^Teta) = rac{exp(x^Teta)}{1 + exp(x^Teta)}$$

for some parameter $\beta=(\beta_1,\cdots,\beta_p)\in\mathbb{R}^p$, usually $x_1=1$ and β_1 is an intercept. σ is the sigmoid logistic function we have seen before.

In the literature is usual to denote $\eta(X)=p_{eta}(X)$ or $\eta(X)=\pi_{eta}(X)$.

From now, we will use the notation $p_{eta}(X)$.

A more formal definition

Defining $\log it: x \to \log\left(\frac{x}{1-x}\right)$, which is the inverse of σ the logistic function (show it as an exercise), we have:

$$\operatorname{logit}(p_{eta}(X)) = X^T eta$$

The Logistic Regression model:

We are given $(x_i, y_i) \in \mathbb{R}^p \times \{0, 1\}$, $i = 1, \dots, n$

The Logistic Regression model assumes that outputs y_i are independent Bernoulli with parameter $p_{\beta}(x_i)$ depending on x_i :

$$\operatorname{logit}(p_{eta}(x_i)) = x_i^T eta$$

How to fit with R

The syntax to fit the Logistic model in R using glm() is:

```
glm(y \sim x, data = dataframe, family = binomial(link = ''logit'')
```

The formula $y \sim x$ depicts the model (i.e. inputs are X, output is Y) and the **data**= argument points to the training set contained in a R dataframe (or tibble). This is quite similar to the **lm()** function.

We also need to specify the distribution for the conditional Y values (binomial) and the link function (logit) via the **family**= argument.

How to fit with R

For our default example:

The command **summary** produces result summaries of the fitted model:

```
Call:
glm(formula = default ~ ., family = "binomial", data = default data)
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.087e+01 4.923e-01 -22.080 < 2e-16 ***
studentYes -6.468e-01 2.363e-01 -2.738 0.00619 **
balance
            5.737e-03 2.319e-04 24.738 < 2e-16 ***
            3.033e-06 8.203e-06 0.370 0.71152
income
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 2920.6 on 9999 degrees of freedom
Residual deviance: 1571.5 on 9996 degrees of freedom
AIC: 1579.5
Number of Fisher Scoring iterations: 8
```

We will see in the next lesson, how this model is fitted in practice and how to interpret or understand what is printed by the **summary** function.

Estimation

We are given $(x_i, y_i) \in \mathbb{R}^p \times \{0, 1\}$, $i = 1, \dots, n$ where outputs y_i are independent Bernoulli with parameter $p_\beta(x_i)$ depending on x_i :

$$\operatorname{logit}(p_{eta}(x_i)) = x_i^T eta$$

The parameters β of the Logistic Regression model are usually determined using Maximum Likelihood Estimation (MLE). It consists on finding β for which the joint probability of the observed data is greatest.

As y_i are independent the likelihood function (joint probability) is the product of the probability mass functions:

$$L(Y,eta) = \prod_{i=1}^n p_eta(x_i)^{y_i} (1-p_eta(x_i))^{1-y_i}$$

with $Y=(y_1,\cdots,y_n)$ and $eta=(eta_1,\cdots,eta_p)$.

Estimation

We seek to maximize the likelihood function over β , it is equivalent but easier to maximize the log-likelihood:

$$egin{aligned} \ell(Y,eta) &= \log L(Y,eta) = \sum_{i=1}^n \left(y_i \log(p_eta(x_i)) + (1-y_i) \log(1-p_eta(x_i))
ight) \ &= \sum_{i=1}^n \left(y_i \log(rac{p_eta(x_i)}{1-p_eta(x_i)}) + \log(1-p_eta(x_i))
ight) \ &= \sum_{i=1}^n \left(y_i x_i^T eta - \log(1+\exp(x_i^T eta))
ight) \end{aligned}$$

If the MLE \hat{eta} exists, the gradient of log-likelihood satisfies (first order necessary condition):

$$abla \ell(Y,eta) = \left(rac{\partial \ell(Y,eta)}{\partial eta_1}, \cdots, rac{\partial \ell(Y,eta)}{\partial eta_p}
ight) = \mathbf{0}.$$

Estimation

We have for $j=1,\cdots,p$:

$$rac{\partial \ell(Y,eta)}{\partial eta_j} = \sum_{i=1}^n \left(y_i x_{ij} - x_{ij} rac{\exp(x_i^T eta)}{1 + \exp(x_i^T eta)}
ight) = \sum_{i=1}^n x_{ij} \left(y_i - p_eta(x_i)
ight)$$

In vector form:

$$abla \ell(Y,eta) = \sum_{i=1}^n x_i \left(y_i - p_eta(x_i)
ight) = X^T(Y-P_eta)$$

where:

$$X = egin{pmatrix} x_{11} & \cdots & x_{1p} \ x_{21} & \cdots & x_{2p} \ dots & dots & dots \ x_{n1} & \cdots & x_{np} \end{pmatrix} = egin{pmatrix} x_1^T \ x_2^T \ dots \ x_n^T \end{pmatrix} \in \mathbb{R}^{n imes (p)}, \quad Y = egin{pmatrix} y_1 \ y_2 \ dots \ y_n \end{pmatrix} \quad and \quad P_eta = egin{pmatrix} p_eta(x_1) \ p_eta(x_2) \ dots \ p_eta(x_n) \end{pmatrix}$$

Estimation

In the literature $abla \ell(Y, eta)$ is denoted as the Fisher's score function S(eta).

If the MLE $\hat{\beta}$ exists, we have:

$$S(\hat{eta}) =
abla \ell(Y,\hat{eta}) = X^T(Y-P_{\hat{eta}}) = 0$$

Solving this equation involves solving p non-linear equations in $eta=(eta_1,\cdots,eta_p)$:

$$y_1x_{1j}+\cdots+y_nx_{nj}=x_{1j}rac{\exp(x_1^Teta)}{1+\exp(x_1^Teta)}+\cdots+x_{nj}rac{\exp(x_n^Teta)}{1+\exp(x_n^Teta)},\quad j=1,\cdots,p$$

Numerical methods are used to solve these non-linear equations as no closed-form solution exist.

Estimation - MLE existence

If we assume that rank(X) = p, we have that $S(\beta)$ is concave in β hence if we find a local maximum it is a global maximum.

We have for $(k,l)\in (1,\cdots,p)^2$:

$$egin{aligned} rac{\partial \ell}{\partial eta_k \partial eta_l}(eta) = &rac{\partial}{\partial eta_k} \sum_{i=1}^n x_{il} (y_i - rac{\exp(x_i^T eta)}{1 + \exp(x_i^T eta)}) \ = &- \sum_{i=1}^n x_{il} x_{ik} rac{\exp(x_i^T eta)}{(1 + \exp(x_i^T eta))^2} \ = &- \sum_{i=1}^n x_{ik} p_eta(x_i) (1 - p_eta(x_i)) x_{il} \end{aligned}$$

Estimation - MLE existence

We obtain that in matrix form:

$$H(\beta) = \nabla^2 \ell(Y, \beta) = -X^T W_{\beta} X$$

where:

$$W_eta = egin{pmatrix} p_eta(x_1)(1-p_eta(x_1)) & \cdots & \cdots \ dots & \ddots & dots \ \cdots & \cdots & p_eta(x_n)(1-p_eta(x_n)) \end{pmatrix}$$

We have $p_{\beta}(x_i)(1-p_{\beta}(x_i))\geq 0$ hence $W(\beta)$ is semi-definite negative and since rank(X)=p, $H(\beta)$ is concave.

Estimation - MLE existence

It is shown in (Albert & Aanderson, 1984) that if additionally there is no complete separation in the training set:

Maximum likelihood estimates in logistic regression

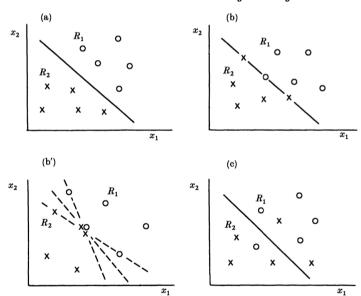


Fig. 1. Possible configurations of sample points in the case of two variables, x_1 and x_2 , and two groups, H_1 , shown by circles, and H_2 , shown by crosses. Regions R_1 and R_2 define corresponding allocation rule. (a) Complete separation. (b) Quasicomplete separation $\rho(X^q) = 2$. (b') Quasicomplete separation $\rho(X^q) = 1$; at the point of intersection of the lines, there are three observations, one from H_1 , and two from H_2 . (c) Overlap.

Estimation - The Newton-Raphson method

In practice a numerical/iterative method such as the Newton-Raphson method is used to solve the equation:

$$S(eta) =
abla \ell(Y,eta) = X^T(Y-P_eta) = 0$$

Using Taylor expansion of Score S(eta) around an initial guess $eta^{(0)}$ of \hat{eta} :

$$S(\hat{eta})pprox S(eta^{(0)}) + H(eta^{(0)})(\hat{eta}-eta^{(0)})$$

giving a first estimate of $\hat{\beta}$:

$$eta^{(1)} = eta^{(0)} - H^{-1}(eta^{(0)}) S(eta^{(0)})$$

Then until convergence, the Newton-Raphson method iterates:

$$eta^{(k+1)} = eta^{(k)} - H^{-1}(eta^{(k)}) S(eta^{(k)})$$

Estimation - The Newton-Raphson method

We show below a naive implementation of Newton-Raphson method to estimate β :

▼ Code

```
1 # We put the data frame in matrix form
 2 # also adding an intercept
 3 X <- cbind(rep(1, nrow(default data)),</pre>
                               as.matrix(default data %>% select(balance, income)))
 5 colnames(X) <- c("(Intercept)", "balance", "income")</pre>
     <- nrow(X)
     We extract the output as vector
 9 Y <- default data %>% mutate(default = if else(default=='Yes', 1, 0)) %>% pull(default)
10
11 # We set an initial guess for beta and criterion for stopping
12 beta <-c(0.01, 0.0, 0.0)
13 nb iter <- 25
14 tol <- 1e-4
15
16 lr solve <- function(X, Y, beta, nb iter, tol){</pre>
17
        for(i in 1:nb iter){
18
            # first compute p beta(X)
            p beta <-\exp(X %*% beta) / (1 + \exp(X %*% beta))
19
20
            # then the Score
21
            Score beta <- t(X) %*% (Y-p beta)
22
            # and the Hessian
23
            W beta <- matrix(0, n, n)</pre>
24
            diag(W beta) <- p beta*(1-p beta)</pre>
2.5
            Hessian beta <- -t(X) %*% W beta %*% X
26
            # we update beta
27
            new beta <- beta - solve(Hessian beta) %*% Score beta
28
            # we check for convergence
```

Estimation - The Newton-Raphson method

We verify that the R glm() function and our algorithm give close values for coefficients β :

• R glm():

```
(Intercept) balance income 
-11.540468 0.005647 0.000021
```

Newton-Raphson:

```
(Intercept) balance income [1,] -11.53791 0.005646 2.1e-05
```

Estimation - Machine Learning approach

We rewrite the log-likelihood equation stated before:

$$egin{aligned} \ell(Y,eta) &= \log L(Y,eta) = \sum_{i=1}^n \left(y_i \log(p_eta(x_i)) + (1-y_i) \log(1-p_eta(x_i))
ight) \ &= -\sum_{i=1}^n \ell_{logistic}\left(p_eta(x_i),y_i
ight) \ &= -n\hat{ ext{R}}(p_eta) \end{aligned}$$

where $\ell_{logistic}: \{0,1\} imes \{0,1\}
ightarrow \mathbb{R}^+$:

$$\ell_{logistic}(y,z) = -y\log(z) - (1-y)\log(1-z) = egin{cases} -\log(z) & ext{if } y=1 \ -\log(1-z) & ext{if } y=0 \end{cases}$$

and $\hat{\mathrm{R}}(p_{eta})$ is the empirical risk on the training set.

Estimating β by maximizing the log-likelihood is equivalent to minimizing with respect to β the empirical risk of p_{β} for the logistic loss.

Interpretation - the R glm() output

▼ Code

```
1 summary(glm_default)
```

```
Call:
glm(formula = default ~ ., family = "binomial", data = default data)
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.087e+01 \quad 4.923e-01 -22.080 \quad < 2e-16 ***
studentYes -6.468e-01 2.363e-01 -2.738 0.00619 **
balance
             5.737e-03 2.319e-04 24.738 < 2e-16 ***
             3.033e-06 8.203e-06 0.370 0.71152
income
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 2920.6 on 9999 degrees of freedom
Residual deviance: 1571.5 on 9996 degrees of freedom
AIC: 1579.5
Number of Fisher Scoring iterations: 8
```

Based on this output, the fitted model is (we have re-scaled balance and income for better readability):

$$\log\left(rac{p_{eta}(x_i)}{1-p_{eta}(x_i)}
ight) = -1.08 - 0.65(\mathbb{1}_{ ext{student}_i= ext{Yes}}) + 5.74(rac{ ext{balance}_i}{1000}) + 0.03(rac{ ext{income}_i}{10000})$$

Interpretation - logits

$$\log\left(rac{p_{eta}(x_i)}{1-p_{eta}(x_i)}
ight) = -1.08 - 0.65(\mathbb{1}_{ ext{student}_i= ext{Yes}}) + 5.74(rac{ ext{balance}_i}{1000}) + 0.03(rac{ ext{income}_i}{10000})$$

Coefficients interpretation

We cannot interpret the coefficients in the same manner as we interpret coefficients from a linear model, as the outcome is now expressed in "logits":

- ullet The predicted logit (or as we will see later log-odds) of defaulting for non-students with zero balance and income are -1.08.
- Each one-unit difference in $\frac{\text{balance}}{1000}$ is associated with a difference of 5.74 in the predicted logit of defaulting.
- Each one-unit difference in $\frac{\text{income}}{10000}$ is associated with a difference of 0.03 in the predicted logit of defaulting.

Interpretation - odds

We remind the following relationship:

$$\operatorname{logit}(p_{eta}(x)) = \operatorname{log}(rac{p_{eta}(x)}{1 - p_{eta}(x)}) = x^T eta$$

The ratio on which we take the logarithm is called odds:

$$odd_eta(x) = rac{p_eta(x)}{1-p_eta(x)} = exp(x^Teta)$$

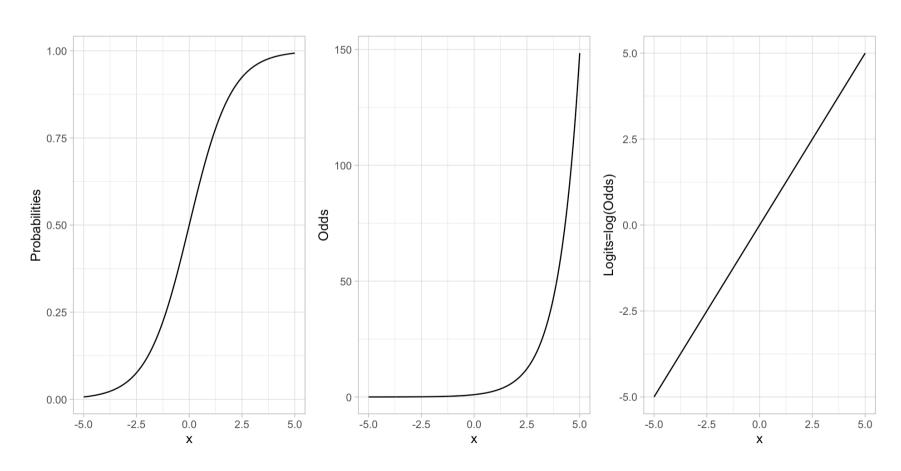
It represents the chance an event occurs $(p_{\beta}(x))$ versus the chance that same event does not occur $(1-p_{\beta}(x))$.

We can also rewrite:

$$p_{eta}(x) = rac{odds_{eta}(x)}{1 + odds_{eta}(x)}$$

Interpretation - odds

To set these ideas, for a variable x in [-5, 5], we plot the logistic curve (i.e. the probabilities) together with the odds and the logits (ie log(odds)):



Interpretation - odds ratio

For two observations x and \tilde{x} we define odds ratio as:

$$OR(x, ilde{x}) = rac{odds(x)}{odds(ilde{x})}$$

Odds ratio are used to compare probabilities between two observations:

- $OR(x, \tilde{x}) = 1 \Leftrightarrow p(x) = p(\tilde{x})$
- $OR > 1 \Leftrightarrow p(x) > p(\tilde{x})$
- $OR < 1 \Leftrightarrow p(x) < p(\tilde{x})$

They are also used to measure the impact of a predictor:

$$OR(x, ilde{x}) = \exp(eta_1(x_1 - ilde{x_1})) \cdots exp(eta_p(x_p - ilde{x_p}))$$

Choosing (x, \tilde{x}) differing by only one predictor x_i :

$$OR(x, \tilde{x}) = \exp(\beta_j (x_j - \tilde{x_j}))$$

Interpretation - odds ratio

 $exp(eta_j)$ is the odds ratio associated with a one-unit increase in the x_j .

Tip

Interpret the coefficients in terms of odds:

- The coefficient of balance / 1000 is 5.737. Hence an increase of balance by 1000 points increases odds for default by a factor of exp(5.737)=310.
- The coefficient of income / 10000 is 0.03. Hence an increase of income by 10000 points increases odds for default by a factor of exp(0.03)=1.03.
- The coefficient of "being a student" is -0.647. Hence being a student decreases odds for default by a factor of exp(-0.647)=0.52.

More on odds ratio interpretation can be found here.

Inference - asymptotic properties of MLE

Under certain assumptions (see for example Gourieroux & Monfort (1981) or Fahrmeir & Kaufmann (1986)), the Maximum Likelihood Estimator has the following asymptotic properties:

$$\hat{\beta} \xrightarrow{p} \beta$$
, as $n \to \infty$

and

$$\sqrt{n}(\hat{eta}-eta) \stackrel{\mathcal{L}}{
ightarrow} \mathcal{N}(0,\mathcal{I}(eta)^{-1}) ext{, as n}
ightarrow \infty$$

where:

$$\mathcal{I}(eta) = -\mathbb{E}[
abla^2 \ell(Y,eta)] = -rac{1}{n}
abla^2 \ell(Y,eta) = rac{1}{n}X^T W_eta X$$

where $\mathcal{I}(eta)$ is the Fisher information matrix. We recognize the Hessian matrix we have seen before.

Inference - asymptotic properties of MLE

The asymptotic property rewrites:

$$(\hat{eta} - eta)^T n \mathcal{I}(eta) (\hat{eta} - eta) \xrightarrow{\mathcal{L}} \chi_p^2$$

As $\mathcal{I}(\beta)$ is unknown we use instead $\mathcal{I}(\hat{\beta}) = \frac{1}{n} X^T W_{\hat{\beta}} X$. Since $\hat{\beta} \xrightarrow{p} \beta$ and p_{β} continuous in β it can be shown that:

$$(\hat{eta}-eta)^T X^T W_{\hat{eta}} X (\hat{eta}-eta) \stackrel{\mathcal{L}}{
ightarrow} \chi_p^2$$

Or equivalently:

$$\hat{eta} - eta \overset{\mathcal{L}}{
ightarrow} \mathcal{N}(0, (X^T W_{\hat{eta}} X)^{-1})$$

Inference - Wald statistics

Using the preceding asymptotic properties we can derive confidence interval and tests for the coefficients eta_j , $j=1,\cdots,p$ of the model:

$$rac{\hat{eta}_j - eta_j}{\hat{\sigma}_j} \stackrel{\mathcal{L}}{ o} \mathcal{N}(0,1)$$

where $\hat{\sigma}_j^2 = s.\,e.\,(\hat{eta}_j)^2$ denotes the j-th term of $(X^TW_{\hat{eta}}X)^{-1}$ diagonal.

The typical formula for a 1-lpha confidence interval is:

$$\hat{eta}_j \pm z_{1-lpha/2} \hat{\sigma}_j$$

where $z_{1-lpha/2}$ is the (1-lpha/2) quantile of the standard normal distribution.

Inference - Wald statistics

Going further, the asymptotic properties of MLE also allow to test the "statistical significance" of each coefficient in the model, the Wald test.

Denoting: \mathbf{H}_0 : $\mathbf{eta}_j = \mathbf{0}$ and \mathbf{H}_1 : $\mathbf{eta}_j
eq \mathbf{0}$ we have under \mathbf{H}_0 :

$$rac{\hat{eta}_j}{\hat{\sigma}_j} \stackrel{\mathcal{L}}{ o} \mathcal{N}(0,1)$$

We will reject H_0 at level α if the absolute of the observed value $\frac{\hat{\beta}_j}{\hat{\sigma}_j}$ (denoted in glm output as z value) is above the $(1-\alpha/2)$ quantile of the standard normal distribution.

Inference - Wald statistics

```
Coefficients: 

Estimate Std. Error z value Pr(>|z|) (Intercept) -1.154e+01 4.348e-01 -26.545 < 2e-16 *** balance 5.647e-03 2.274e-04 24.836 < 2e-16 *** income 2.081e-05 4.985e-06 4.174 2.99e-05 *** -- Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 (Dispersion parameter for binomial family taken to be 1) Null deviance: 2920.6 on 9999 degrees of freedom Residual deviance: 1579.0 on 9997 degrees of freedom AIC: 1585
```

Inference - Wald statistics

We reject H_0 at level lpha when $p=\mathbf{P}(|z|>|rac{\hat{eta}_j}{\hat{\sigma}_j}|)<lpha$.

 $oldsymbol{p}$ is called the p-value.

The output of **glm** in **R** shows:

- $\hat{\beta}_j$ as **Estimate**,
- $\hat{\sigma}_j$ as Std. Error,
- the observed test statistic $\frac{\hat{\beta}_j}{\hat{\sigma}_j}$ as **z** value,
- and the p-value as Pr(>|z|)

Inference - Wald statistics

We obtain the **z** value $\frac{\hat{\beta}_j}{\hat{\sigma}_j}$ using estimate and its standard deviation:

▼ Code

```
1 z <- glm_bal_inc$coefficients[3] / (summary(glm_bal_inc))$coefficients[3,2]
2 z</pre>
```

income 4.174178

and then the p-value:

▼ Code

```
1 (1-pnorm(abs(z)))*2
```

income 2.990638e-05

Inference - Wald statistics

Using the hessian matrix obtained before as a side product of the Newton-Raphson algorithm, we retrieve comparable values with ${\tt glm}$ outputs for $\hat{\sigma}_i$:

▼ Code

```
1 std_errors <- sqrt(diag(solve(-as.matrix(sol$hessian))))
2 std_errors

(Intercept) balance income
4.346349e-01 2.273110e-04 4.984579e-06</pre>
```

the **z** values and then the p-values:

```
1 z <- sol$beta / std_errors
2 t(z)

(Intercept) balance income
[1,] -26.54622 24.83714 4.173579

▼ Code

1 t((1-pnorm(abs(z)))*2)

(Intercept) balance income
[1,] 0 0 2.998516e-05
```

Inference - confidence intervals

The R command to get confidence interval of estimators based on Wald statistic is the following (by default lpha=5%)

▼ Code

```
1 confint.default(glm_bal_inc)

2.5 % 97.5 %

(Intercept) -1.239258e+01 -1.068836e+01
balance 5.201460e-03 6.092746e-03
income 1.103823e-05 3.057972e-05
```

We can retrieve it manually using coefficient estimate and standard deviation:


```
1 output_bal_inc = summary(glm_bal_inc)$coefficients
2 bal_std_estimate <- output_bal_inc[2,1]
3 bal_std_error <- output_bal_inc[2,2]
4
5 # upper bound for beta(balance) at 5%
6 upper <- bal_std_estimate + 1.96 * bal_std_error
7
8 # lower bound for beta(balance) at 5%
9 lower <- bal_std_estimate - 1.96 * bal_std_error
10 (bal_confint <- c(lower, upper))</pre>
```

Inference - confidence intervals

The following R command provides confidence interval of estimators using a more advanced profile likelihood method:

▼ Code

```
1 confint(glm_bal_inc)

2.5 % 97.5 %
```

```
(Intercept) -1.241910e+01 -1.071361e+01
balance 5.214030e-03 6.105971e-03
income 1.105359e-05 3.060844e-05
```

More details on profile likelihood method can be found here.

Inference - tests on model coefficients

Based on the same idea, it is possible to test for the nullity of a subset of the model coefficients.

Denoting: \mathbf{H}_0 : $\beta_1 = \cdots = \beta_q = 0$, \mathbf{H}_1 : $\exists j \in \{1, \dots, q\} \mid \beta_j \neq 0$, $\hat{\beta} = (\hat{\beta}_1, \dots, \hat{\beta}_p)$ the MLE and $\hat{\beta}_{1:q} = (\hat{\beta}_1, \dots, \hat{\beta}_q)$ the vector of first q parameters.

We have under \mathbf{H}_0 :

$$\hat{eta}_{1:q}^T (X^T W_{\hat{eta}} X)_{1:q}^{-1} \hat{eta}_{1:q} \stackrel{\mathcal{L}}{
ightarrow} \chi_q^2$$

where $(X^TW_{\hat{\beta}}X)_{1:q}^{-1}$ is the $q \times q$ upper left block matrix extracted from the inverse of hessian.

We will reject H_0 at level α if the observed value $\hat{\beta}_{1:q}^T(X^TW_{\hat{\beta}}X)_{1:q}^{-1}\hat{\beta}_{1:q}$ is above the $1-\alpha$ quantile of the χ_q^2 distribution.

Inference - tests on model coefficients

We show below the Wald tests for each coefficient in the model using **summary**:

```
1 summary(glm default)
Call:
qlm(formula = default ~ ., family = "binomial", data = default data)
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.087e+01 4.923e-01 -22.080 < 2e-16 ***
studentYes -6.468e-01 2.363e-01 -2.738 0.00619 **
balance
            5.737e-03 2.319e-04 24.738 < 2e-16 ***
income
            3.033e-06 8.203e-06 0.370 0.71152
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 2920.6 on 9999 degrees of freedom
Residual deviance: 1571.5 on 9996 degrees of freedom
AIC: 1579.5
Number of Fisher Scoring iterations: 8
```

Inference - tests on model coefficients

These tests can be also performed in **R** using **car::Anova** or **aod::wald.test** routines. In particular when categorical variables have more than two levels these functions allow to test each variables as a whole (vs coefficient by coefficient when using **summary**)

```
1 car::Anova(glm_default, type=3, test.statistic= "Wald")

Analysis of Deviance Table (Type III tests)

Response: default

Df Chisq Pr(>Chisq)
(Intercept) 1 487.5303 < 2.2e-16 ***
student 1 7.4947 0.006188 **
balance 1 611.9470 < 2.2e-16 ***
income 1 0.1368 0.711520

---
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Inference - tests on model coefficients

Inference - tests on model coefficients

We can retrieve these outputs manually:

▼ Code

```
1 sum_default <- summary(glm_default)
2 beta_income <- sum_default$coefficients[4,1]
3 stdev_income <- sum_default$coefficients[4,2]
4
5 wald <- beta_income ^ 2 / stdev_income ^ 2
6
7 l-pchisq(wald, df = 1)</pre>
```

[1] 0.7115203

▼ Code

```
1 z_val <- sum_default$coefficients[4,3]
2
3 2*(1-pnorm(abs(z_val)))</pre>
```

[1] 0.7115203

With all routines, the p-value for the income coefficient is 0.71 validating the null hypothesis.

Inference - tests on model coefficients

Using **Terms** or **L** parameters in **aod::wald.test** it is also feasible to test the null hypothesis for a subsets of parameters:

▼ Code

```
Chi-squared test:
X2 = 698.3, df = 3, P(> X2) = 0.0
```

The null hypothesis is rejected for the model with balance and student.

Likelihood ratio tests

It is possible to test for the nullity of a subset of the model coefficients using Likelihood Ratio statistics.

Denoting: H_0 : $\beta_1=\dots=\beta_q=0$, H_1 : $\exists j\in\{1,\dots,q\}\mid \beta_j\neq 0$, and $\hat{\beta}=(\hat{\beta}_1,\dots,\hat{\beta}_p)$ the MLE, we have under H_0 :

$$-2\left(\ell_{\mathrm{H}_0}(Y,\hat{eta}_{\mathrm{H}_0}) - \ell(Y,\hat{eta})
ight) \stackrel{\mathcal{L}}{
ightarrow} \chi_q^2$$

where $\ell_{\mathrm{H}_0}(Y,\hat{eta}_{\mathrm{H}_0})$ is the log-likelihood of:

$$\operatorname{logit}(p_{eta}(X)) = x_{q+1}\beta_{q+1} + \dots + x_n\beta_n$$

Likelihood ratio tests

Consider two models, a larger model with l parameters and likelihood L_L and a smaller model with sparameters and likelihood L_S , where the smaller model represents a subset of the larger model. Typically the smaller model is equivalent to the large model where we have imposed:

$$H_0: \ \beta_j = \ldots = \beta_{j+r} = 0$$

Likelihood Ratio tests on variables may be performed in R using car:: Anova:

```
1 car::Anova(glm default, type=3, test.statistic= "LR")
Analysis of Deviance Table (Type III tests)
Response: default
       LR Chisq Df Pr(>Chisq)
student
        7.42 1 0.006445 **
balance 1335.95 1 < 2.2e-16 ***
income
          0.14 1 0.711514
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Likelihood ratio tests

Using base **R anova** it is also possible to test subsets of variables and in particular individual variables within the "full" model, we have to fit **glm** for each sub model:

```
glm_wo_student <- glm(default ~ balance + income,
data = default_data,
family = "binomial")

glm_wo_balance <- glm(default ~ student + income,
data = default_data,
family = "binomial")

glm_wo_income <- glm(default ~ student + balance,
data = default_data,
family = "binomial")</pre>
```

Likelihood ratio tests

Testing the model without student predictor:

▼ Code

```
1 anova(glm_wo_student, glm_default, test = "LRT")

Analysis of Deviance Table

Model 1: default ~ balance + income
Model 2: default ~ student + balance + income
    Resid. Df Resid. Dev Df Deviance Pr(>Chi)
1 9997 1579.0
2 9996 1571.5 1 7.4214 0.006445 **
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

We can retrieve this result manually:

```
1 LRT <- 2 * (logLik(glm_default)-logLik(glm_wo_student))
2 1 - pchisq(LRT, df = 1)</pre>
```

^{&#}x27;log Lik.' 0.006445112 (df=4)

Likelihood ratio tests

Testing the other sub models:

```
▼ Code
```

```
1 anova(glm wo balance, glm default, test = "LRT")
Analysis of Deviance Table
Model 1: default ~ student + income
Model 2: default ~ student + balance + income
  Resid. Df Resid. Dev Df Deviance Pr(>Chi)
       9997
                2907.5
       9996
                1571.5 1
                             1336 < 2.2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
▼ Code
  1 anova(glm wo income, glm default, test = "LRT")
Analysis of Deviance Table
Model 1: default ~ student + balance
Model 2: default ~ student + balance + income
  Resid. Df Resid. Dev Df Deviance Pr(>Chi)
       9997
                1571.7
       9996
                1571.5 1 0.13677 0.7115
```

The deviance defined as $D=-2\ell$ is often reported by statistical software in place of log-likelihood. A large likelihood corresponding to a small deviance.

Likelihood ratio tests

A better and full coverage of tests in the context of Logistic Regression can be found here or in (Hosmer et al., 2013). See also here for a data analysis using **R** and see the question here for a very detailed description of the outputs of **glm()** (in particular this answer).

Goodness of Fit test / Calibration

Although it is generally not recommended by practitioners and theoreticians, the Hosmer & Lemeshow test (see here, here or here) allows to quickly assess the "goodness of fit" of a Logistic Regression.

But more than the test in itself, the underlying motivation is interesting: the Logistic Regression model provides an estimate of the probability of an outcome (success/failure, here the default is success or 1). The estimated probability of this outcome should be close to the true observed probability.

The Hosmer & Lemeshow test assess if observed event rates match expected event rates in subgroups of "similar" observations. Models for which expected and observed event rates agree on these subgroups are considered well calibrated.

Goodness of Fit test / Calibration

A first step of the test is to order the predicted probabilities of the outcome and divide it into Q groups (usually using deciles, Q=10). Then the average predicted probability for each group is computed and compared to the observed probability.

The Hosmer & Lemeshow test statistic H is compared to a χ^2_{Q-2} distribution:

$$H = \sum_{q=1}^Q rac{(o_q - m_q \mu_q)^2}{m_q \mu_q (1 - \mu_q)}$$

where:

- ullet o_q denotes the number of success (Y=1) observed in group q,
- ullet μ_q denotes the mean of $p_{\hat{eta}}(x_i)$ in group q,
- m_q denotes the number of observations in group q, so that $m_q \mu_q$ is the expected number of success in group q.

The null hypothesis is that observed/expected outcomes are close along all subgroups.

Goodness of Fit test / Calibration

▼ Code

```
1 library(glmtoolbox)
2 hltest(glm_default)
```

```
The Hosmer-Lemeshow goodness-of-fit test
Group Size Observed
                        Expected
    1 1000
                      0.02653992
    2 1000
                      0.10737240
    3 1000
                  0 0.29143249
    4 1000
                  1 0.67265778
    5 1000
                  2 1.39515666
    6 1000
                      2.87108745
    7 1000
                  7 5.98948667
    8 1000
                 16 13.74542953
                 45 39.52811751
    9 1000
   10 1000
                261 268.37271994
        Statistic = 3.68229
degrees of freedom = 8
```

p-value = 0.88459

Here the p-value for a chi-squared statistic of H=3.68 with df=Q-2=8 is p=0.885 which is well above the usual levels (eg 0.05), so that the null hypothesis is accepted, goodness of fit is acceptable.

However the Hosmer & Lemeshow test is dependent on the choice of Q and the binning performed on probabilities and is sometimes considered unreliable.

Goodness of Fit test / Calibration

Nonetheless it is usual to assess or diagnose the good calibration of a model probabilities using Calibration Plots or Probability Calibration Curves (see here for a recent R package from the tidyverse/tidymodel ecosystem and here for a scikit-learn version): they are used to visualize if predictions are consistent with the observed event rates (be it on the training set or a testing set, which is better).

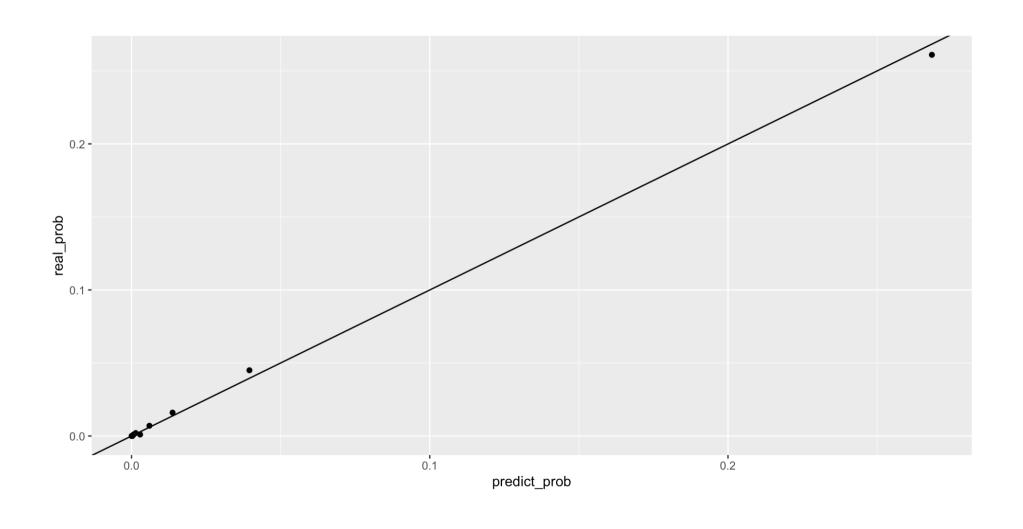
Goodness of Fit test / Calibration

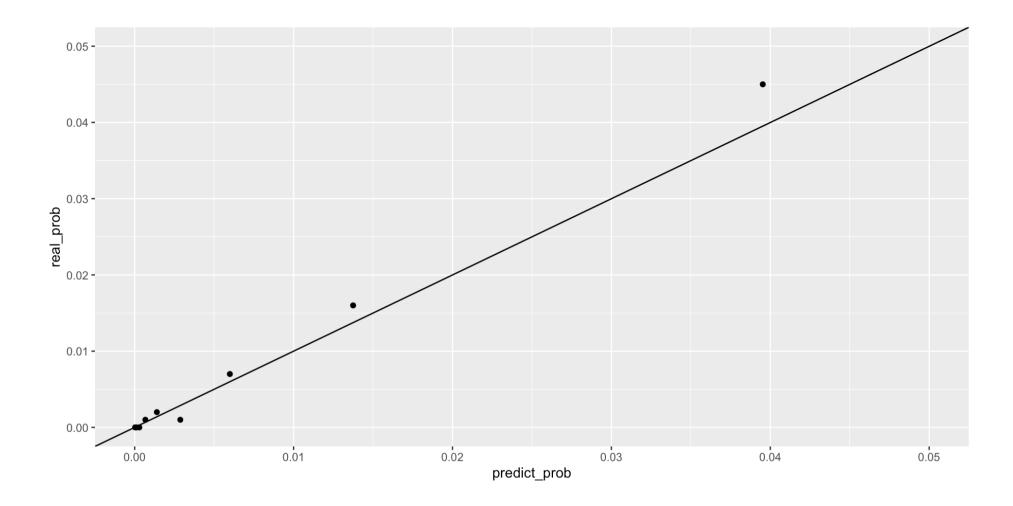
For example considering the **default** data set we have:

```
1 check default prob <- as tibble(cbind(fitted=glm default$fitted.values,</pre>
 2
                                           Y = default data %>%
 3
                                             mutate(default = if else(default == "Yes", 1, 0)) %>%
                                             pull(default)))
    (calibration data <- check default prob %>%
      mutate(bins prob = cut(fitted, breaks = quantile(fitted,seq(0,1,0.10)), include.lowest = TRUE)) %>%
      group by(bins prob) %>%
 8
      summarize(n = n(),
 9
                def = sum(Y),
10
                no def = n - def,
11
                predict prob = mean(fitted),
12
                real prob = def/n,
13
                forecast acc = def / sum(check default prob$Y)))
```

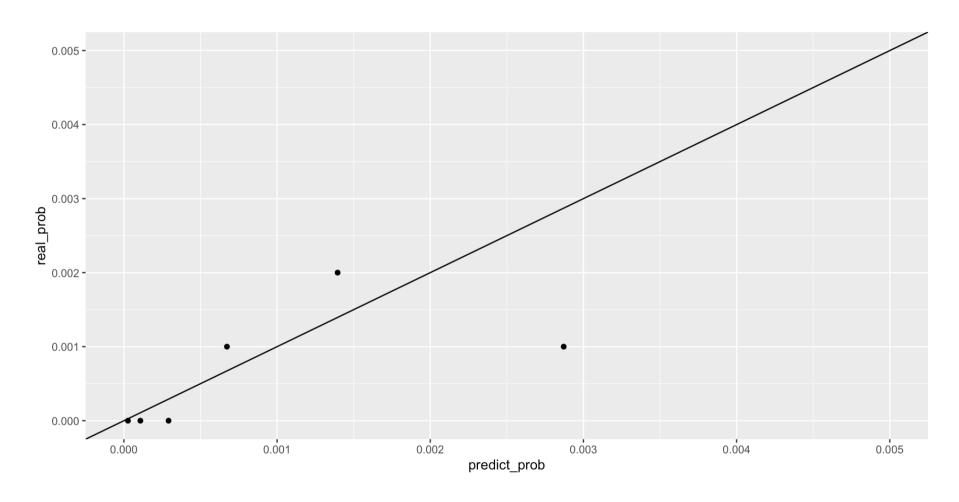
```
# A tibble: 10 × 7
  bins prob
                                def no def predict prob real prob forecast acc
   <fct>
                        <int> <dbl>
                                     <dbl>
                                                   <dbl>
                                                              <dbl>
                                                                            <dbl>
                                                              0
1 [1.03e-05,5.14e-05] 1000
                                       1000
                                               0.0000265
                                                                         0
 2 (5.14e-05,0.000176]
                        1000
                                       1000
                                               0.000107
                                                              0
                                                                         0
 3 (0.000176,0.000443]
                                       1000
                                               0.000291
                        1000
                                       999
                        1000
                                               0.000673
                                                              0.001
                                                                         0.00300
 4 (0.000443,0.0009451
                                       998
                                                              0.002
 5 (0.000945,0.00197]
                         1000
                                               0.00140
                                                                         0.00601
 6 (0.00197,0.00402]
                         1000
                                  1
                                       999
                                               0.00287
                                                              0.001
                                                                         0.00300
                                                              0.007
 7 (0.00402,0.00881
                         1000
                                       993
                                               0.00599
                                                                         0.0210
 8 (0.0088,0.021]
                         1000
                                 16
                                       984
                                               0.0137
                                                              0.016
                                                                         0.0480
 9 (0.021, 0.07091
                         1000
                                 45
                                       955
                                               0.0395
                                                              0.045
                                                                         0.135
10 (0.0709, 0.978)
                         1000
                                261
                                       739
                                               0.268
                                                              0.261
                                                                         0.784
```

Goodness of Fit test / Calibration





Goodness of Fit test / Calibration



The model tends to slightly overestimate/underestimates some deciles without a clear pattern.

References

- Albert, A., & Aanderson, J. A. (1984). On the existence of maximum likelihood estimates in logistic regression models. *Biometrika*, 71(1), 1–10. https://doi.org/10.1093/biomet/71.1.1
- Fahrmeir, L., & Kaufmann, H. &. (1986). Asymptotic inference in discrete response models. *Statistische Hefte*, 27(1), 179–205. https://doi.org/10.1007/bf02932567
- Gourieroux, C., & Monfort, A. (1981). Asymptotic properties of the maximum likelihood estimator in dichotomous logit models. *Journal of Econometrics*, *17*(1), 83–97. https://doi.org/10.1016/0304-4076(81)90060-9
- Hosmer, D. W., Lemeshow, S., & Sturdivant, R. X. (2013). Applied logistic regression. In *Wiley Series in Probability and Statistics*.
- James, G., Witten, D., Hastie, T., & Tibshirani, R. (2021). *An introduction to statistical learning: With applications in r.* Springer US. https://doi.org/10.1007/978-1-0716-1418