

# Introduction to Logistic Regression

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# Still Classification Problem

- Given a set of features (age, gender, BMI, BP, etc.), a person has diabetes (1) or not (0)?
- Given the temperature, the age of equipment, and other features, classify if the equipment will get failure (1) or not (0) in the coming working round?
- Given some features, a student gets admitted into NUS (1) or not (0)?

# Notations

- Assume we have a set of  $n$  observations used to build a model.
- We have  $p$  features,  $X_1, \dots, X_p$  in general.
- The outcome variable  $Y$  is binary with two values, 0 and 1.

Obs	$X_1$	$X_2$	$\dots$	$X_p$	$Y$
1	$x_{11}$	$x_{21}$	$\dots$	$x_{p1}$	$y_1$
2	$x_{21}$	$x_{22}$	$\dots$	$x_{p1}$	$y_2$
$\vdots$	$\vdots$	$\vdots$		$\vdots$	$\vdots$
$n$	$x_{n1}$	$x_{2n}$	$\dots$	$x_{pn}$	$y_n$

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# Binary Response

- Assume a point with known features, denote

$$P(Y = 1) = p.$$

- We may have a linear regression model for  $p$ :

$$p \sim \beta_0 + \beta_1 X_1 + \dots + \beta_p X_p.$$

- However, the fitted value might be negative or more than 1, which is not possible for a probability.
- Instead of forming a model for  $p$ , we can form a model for a function of  $p$ !

# Logistic Model

- If we assume  $Y = 1$  as a success, then  $P(Y = 1) = p$  is the success probability. The *odds of success* is then defined as

$$\frac{p}{1 - p}.$$

- We then consider model for the log-odds, or called “logit”:

$$\log\left(\frac{p}{1 - p}\right) = \beta_0 + \beta_1 X_1 + \dots + \beta_p X_p.$$

- Keeping other features constant, for each unit increased in  $X_1$ , the **log odds** increases by  $\beta_1$ .
- This is a type of **generalized linear model (GLM)**.



# Logistic Model

- From the logit equation, we can have the equivalent version:

$$p = \frac{e^{\beta_0 + \beta_1 X_1 + \dots + \beta_p X_p}}{1 + e^{\beta_0 + \beta_1 X_1 + \dots + \beta_p X_p}}.$$

- Regardless the values of the features, the range for  $p$  is always between 0 and 1.

# Logistic Model

- Just like linear regression, in logistic regression the parameters  $\beta_0, \beta_1, \dots, \beta_p$  need to be estimated based on the training data.
- Instead of the method of ordinary least squares (OLS), parameter estimation in logistic regression is based on the method called Maximum Likelihood Estimation (MLE).
- In our course, we'll not introduce the details of MLE.

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## Example: Customer Churn

- A wireless telecommunications company wants to predict whether a customer will churn (switch to a different company) in the next six months.
- With a reasonably accurate prediction of a person's churning, the sales and marketing groups can attempt to retain the customer by offering various incentives.
- Data on 8,000 current and prior customers was obtained. The variables collected for each customer follow:
  - (i) Age (years)
  - (ii) Married (true/false)
  - (iii) Duration as a customer (years)
  - (iv) Churned contacts—Number of the customer's contacts that have churned (count)
  - (v) Churned (true/false)—Whether the customer churned

## Example: Customer Churn

```
> churn = read.csv("C:/Data/churn.csv")
```

```
> churn[1:3,]
```

	ID	Churned	Age	Married	Cust_years	Churned_contacts
1	1	0	61	1	3	1
2	2	0	50	1	3	2
3	3	0	47	1	2	0

```
> churn$Churned = as.factor(churn$Churned)
```

```
> churn$Married = as.factor(churn$Married)
```

```
> churn= churn[,-1] # Remove ID column
```

```
> attach(churn)
```

```
> #table(Churned)
```

```
> prop.table(table(Churned))
```

Churned

0	1
---	---

0.782125	0.217875
----------	----------

- About 21.8% of the customers churned in the given data.

## Example: Customer Churn

- Logistic regression can be performed using the Generalized Linear Model function, **glm()** in R.
- Specify the family to be binomial, the logit link is set as the default.

```
> M1<- glm( Churned ~., data =churn,  
+          family = binomial(link ="logit"))
```

```
> summary(M1)
```

```
Call:
```

```
glm(formula = Churned ~ ., family = binomial(link = "logit"),  
     data = churn)
```

```
Deviance Residuals:
```

Min	1Q	Median	3Q	Max
-2.4542	-0.5206	-0.1971	-0.0728	3.3786

```
Coefficients:
```

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.415201	0.163734	20.858	<2e-16 ***
Age	-0.156643	0.004088	-38.320	<2e-16 ***
Married1	0.066432	0.068302	0.973	0.331
Cust_years	0.017857	0.030497	0.586	0.558
Churned_contacts	0.382324	0.027313	13.998	<2e-16 ***
---				

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(Dispersion parameter for binomial family taken to be 1)
```

```
Null deviance: 8387.3  on 7999  degrees of freedom
```

## p-value of a Coefficient

- In a linear regression model, the column ' $\Pr(>|Z|)$ ' indicate the p-value for a test to test the significance of the coefficient in the fitted model.
- Similarly, in a logistic model, we'll have p-value for each coefficient in the last column in the table 'Coefficients'.
- A large p-value means the contribution of the coefficient (equivalently, of the feature) to the model is not significant.
- It's optional to drop or to keep an insignificant feature in the model. However, dropping it will simplify the model.



## Example: Customer Churn

- From the initial model, 'Cust\_years' is most insignificant. We can drop it.
- Re-fit the logistic model without 'Cust\_years'. We have model M2.

```
> M2<- glm( Churned ~ Age + Married + Churned_contacts,  
+           data = churn, family = binomial)
```

```
> summary(M2)
```

```
Call:
```

```
glm(formula = Churned ~ Age + Married + Churned_contacts, family = b  
     data = churn)
```

```
Deviance Residuals:
```

Min	1Q	Median	3Q	Max
-2.4476	-0.5178	-0.1972	-0.0723	3.3776

```
Coefficients:
```

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.472062	0.132107	26.282	<2e-16 ***
Age	-0.156635	0.004088	-38.318	<2e-16 ***
Married1	0.066430	0.068299	0.973	0.331
Churned_contacts	0.381909	0.027302	13.988	<2e-16 ***

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(Dispersion parameter for binomial family taken to be 1)
```

```
Null deviance: 8387.3  on 7999  degrees of freedom
```

```
Residual deviance: 5358.3  on 7996  degrees of freedom
```

## Example: Customer Churn

- The p-value of 'Married' in model M2 is quite large (0.331), it indicates that 'Married' doesn't contribute significantly to the model when predicting the response.
- We consider to drop it and simplify the model to only two features, model M3.

```
> M3<- glm( Churned ~ Age + Churned_contacts,  
+           data = churn, family = binomial)
```

```
> summary(M3)
```

```
Call:
```

```
glm(formula = Churned ~ Age + Churned_contacts, family = binomial,  
     data = churn)
```

```
Deviance Residuals:
```

Min	1Q	Median	3Q	Max
-2.4599	-0.5214	-0.1960	-0.0736	3.3671

```
Coefficients:
```

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	3.502716	0.128430	27.27	<2e-16 ***
Age	-0.156551	0.004085	-38.32	<2e-16 ***
Churned_contacts	0.381857	0.027297	13.99	<2e-16 ***
---				

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(Dispersion parameter for binomial family taken to be 1)
```

```
Null deviance: 8387.3  on 7999  degrees of freedom  
Residual deviance: 5359.2  on 7997  degrees of freedom  
AIC: 5365.2
```

## Example: Customer Churn

- The fitted model M3 is then

$$\hat{p} = \frac{e^{3.5-0.157 A+0.382 C}}{1 + e^{3.5-0.157 A+0.382 C}}$$

where A stands for Age and C stands for Churned\_contacts.

- We then can predict for a customer who is 50 years old with 5 churned contacts, the estimate probability of churning is

$$\hat{p} = \frac{e^{3.5-0.157 \times 50 + 0.382 \times 5}}{1 + e^{3.5-0.157 \times 50 + 0.382 \times 5}} = 0.08.$$

- We predict the outcome  $Y$  be 0 or 1 based on a threshold, such as 0.5.
- ROC curve can be constructed for logistic model.

1 Introduction

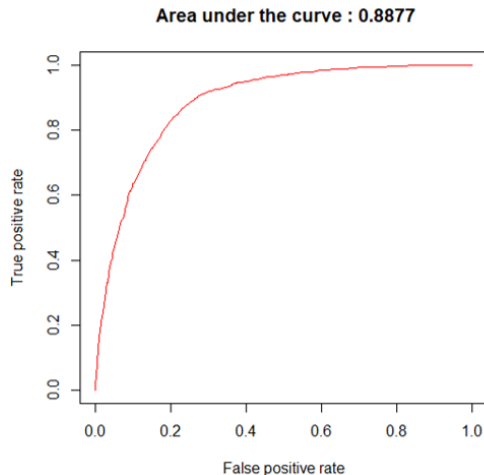
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```
> library(ROCR)
> prob = predict(M3, type = "response")
> # above is to predict probability  $\Pr(Y = 1)$ 
> #for each point in the training data set, using M3
>
> pred = prediction(prob , Churned )
> roc = performance(pred , "tpr", "fpr")
> auc = performance(pred , measure = "auc")
> auc@y.values[[1]] # gives value of AUC
[1] 0.8876509
> plot(roc , col = "red",
+      main = paste(" Area under the curve :",
+      round(auc@y.values[[1]] ,4)))
>
```

# ROC and AUC

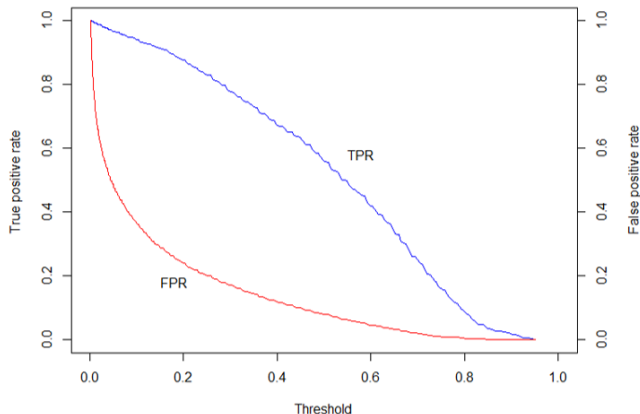


This ROC curve is created based on 328 different values of threshold.



# How TPR, FPR Change when Threshold Changes?

- We can plot to see how TPR and FPR change along with threshold.



## For you to try

- In the example above, we used the whole data set as training data. After that, we evaluated the model (M3) by comparing the real response versus the predict response using M3.
- Can you try to split the full data set into two sets: train set and test set; then build a logistic model on the train set; then check the goodness of the model (using ROC and AUC) using the test set?