Lab 09 - LogisticRegression

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Prompt

This dataset contain the data related to the passengers of the Titanic. The goal is to see if we are able to build a model that can predict whether or not a passenger would survive. The Survived column of this dataset reflects if they really survived or not (1: survived, and 0: didn't survive).

- Load the attached csv file into your Markdown accounting for the missing values.
- 2. Preprocess the data.
- Check for the missing values for each variable.
- Use missmap() to visualize the missing values.
- Reasonably ignore variables that probably do not affect the predictions.
- Impute the missing values in the numeric (continuous variables)
- Remove any of the rows with NA in their categorical variables (any categorical variable that is left in your data after leaving the unnecessary ones out).

1. Import/read the data.

Load the attached csv file into your Markdown accounting for the missing values.

Import the csv file

##

##

```
Let's import the training data here:
training.data.raw <- read.csv('train.csv', na.strings=c(""]</pre>
head(training.data.raw)
```

Braund

Name

Mr Owan Harris

##	1	1	0	3	
##	2	2	1	1	
##	3	3	1	3	
##	4	4	1	1	
##	5	5	0	3	
##	6	6	0	3	

PassengerId Survived Pclass

2. Preprocess the data. Missing values

C : L C --

Check for the missing values for each variable.

Now we need to check for missing values and look how many unique values there are for each variable using the sapply() function.

sapply(training.data.raw,function(x) sum(is.na(x))) ## PassengerId Survived Pclass Name ##

Ticket

D - -- - l-

SibSp Parch ## Fare ## 0 0 0

sapply(training.data.raw, function(x) length(unique(x)))

PassengerId Survived Pclass Name ## 891 891

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3. Split the data

Split the data up into the training and test set. (set.seed = 1).

Split the data

➤ Sample 100 rows from the data and use as your test set. Use the rest as your training set.

```
set.seed(1)
test.ID <- sample(dim(data)[1], 100)
train <- data[-test.ID,]
test <- data[test.ID,]</pre>
```

4. Fit a logistic regression model onto the training set Fit and summarize

Fit and summarize the model to predict Survived as a function of all other remaining variables.

```
model <- glm(Survived ~ ., family=binomial(link="logit"), (</pre>
summary(model)
```

```
##
## Call:
## glm(formula = Survived ~ ., family = binomial(link = "lo
```

data = train) ## ##

```
## Deviance Residuals:
                                Max
##
     Min
            10 Median
                          30
```

Estimate Std. Error z value Pr(>|z|)

-2.6224 -0.5968 -0.4434 0.6336 2.4269 ## ## Coefficients:

##

- 5. Evaluate your model on the test set. Misclassification error
 - Use your model on the test set to make predictions of Survived.

```
Calculate the misclassification error on your test set.
```

- pred.results <- predict(model, newdata=subset(test, select=</pre>
- pred.results <- ifelse(pred.results > .5, 1, 0)
- misClassificationError <- mean(pred.results!= test\$Survive print(paste("Accuracy = ", 1-misClassificationError))

- ## [1] "Accuracy = 0.83" Calculate the CV error
- the accuracy result your obtained depends on the test set so, now re-estimate the accuracy with a 10-fold CV

6. Fit a new model

Redo the last two steps with only Pclass and Sex.

Implement a new fit with Pclass and Sex

```
Do the fit again with only Pclass and Sex;
```

```
model2 <- glm(Survived ~ Pclass + Sex, family=binomial(lin)
summary(model2)
```

Call:

##

##

Deviance Residuals: ## Min

data = train) ##

glm(formula = Survived ~ Pclass + Sex, family = binomia

30

10 Median Max ## -2.1696 -0.7104 -0.4680 0.6804 2.1288

Coefficients: Eqtimate Ctd Error g value Dr(\|g|)