# STATS 769 Data Science Workflow

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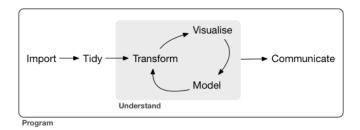
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#### Overview

- This section of the course provides an overview of the activities involved in Data Science.
- Our primary computing environment will be R.

#### Data Science Workflow



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#### Data Science Workflow

- This course will focus on computational concepts and tools that impact on each of the stages in the Data Science Workflow.
- Some topics (Code Efficiency and Parallel Code) have an impact at all stages (because we are always writing code).
- This course will not focus on Visualisation or Modelling or Communication because those topics are covered in other Statistics courses.
- This topic will provide some basics on Visualisation and Modelling and Communication so that everyone has at least one tool to perform those stages of the Data Science Workflow.

#### Data Import

- The important thing here is to be able to deal with a multitude of data formats.
- We will review some basics in the next topic (Data Tech Review).
- We will look at some more advanced import scenarios in later topics (Data Formats and Web Scraping).

# Tidying and Transforming Data

- The important thing here is to be comfortable with data structures in R and converting between different data structures.
- We will review some basics in the next topic (Data Tech Review).
- We will look at some tools outside of R in the Linux topics.
- We will consider some issues with Large Data in the second half of the course.

#### Visualisation

- Base graphics in R.
- plot(), barplot(), boxplot(), plot(density())
- lines(), abline()
- par(mfrow)

- We will provide a basic common framework in this topic.
- We may introduce some additional analysis techniques in later sections.
- We will consider issues with Large Data, Code Efficiency, and Parallel Computing in the second half of the course.

- We will consider a simplified modelling framework where we are only interested in prediction (no inference).
- We have an outcome variable, Y, and one or more predictor variables,  $X_1$ ,  $X_2$ , etc.
- We assume that  $Y = f(X) + \epsilon$ .
- Our problem is to find  $\hat{f}(X)$ , an estimate of f(X).
- epsilon is "irreducible" error (assumed to have mean zero).

- We will distinguish between **training** data, which we use to find  $\hat{f}$ , and **test** data, which we use to evaluate  $\hat{f}$ .
- We will distinguish between when Y is continuous (or quantitative) and when Y is categorical (or discrete or qualitative).
- Linear regression for continuous Y.
- (Multinomial) Logistic regression for categorical Y.

• Linear regression assumes ...

$$f(X) = \beta_0 + \beta_1 * X_1 + ... + \beta_p * X_p$$

- We can fit a linear regression model in R with lm().
- We can obtain predictions from the model with predict().
- We will evaluate models using Root Mean Square Error (RMSE) for linear regression.

$$RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{f}(x_i))^2}$$

• The "worst" RMSE is the (population) standard deviation.

$$RMSE_{min} = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (y_i - \mu_y)^2}$$

- The "best" RMSE is zero, but this will almost certainly require a more flexible model than linear regression and would almost certainly correspond to an overfitted model (which is why we evaluate on a test set).
- We can also use visualisations to explore where the model performs better or worse.

- Logistic regression does not model *Y* directly.
- Instead we model the probability that Y takes one of the possible values.
- Logistic regression assumes ...

$$P(X) = \frac{e^{\beta_0 + \beta_1 * X}}{1 + e^{\beta_0 + \beta_1 * X}}$$

$$\log\left(\frac{P(X)}{1-P(X)}\right) = \beta_0 + \beta_1 * X$$

 Multinomial logistic regression extends this to more than two possible outcomes for Y.

- We can fit a logistic model with glm(..., family="binomial").
- We can obtain predictions from the model with predict(..., type="response").
- These predictions are probabilities that must be converted to values of Y.

- We can fit a multinomial logistic model with nnet::multinom().
- We can obtain predictions from the model with predict(..., type="prob").
- These predictions are probabilities that must be converted to values of Y.
- We can obtain predictions from the model with predict().
- These are values of Y.

- We can evaluate models by how many predictions are "correct."
- Accuracy is the proportion of correct predictions. It can be misleading if the proportion of Y values in each category are not even.
- When there are only two possible Y values, sensitivity measures the proportion of category 1 that are correct.
- When there are only two possible Y values, specificity measures the proportion of category 2 that are correct.
- A confusion matrix is a table of counts for all combinations of actual Y values and predicted Y values.

- The "worst" accuracy is the overall (population) proportion, which leads to always predicting the most common category.
- The "best" accuracy is 1, but this will almost certainly require a more flexible model than logistic regression and would almost certainly correspond to an overfitted model (which is why we evaluate on a test set).
- We can also use visualisations to explore where the model performs better or worse.

#### Communication

- Literate documents.
- R Markdown documents.

# Template for 769 Lab submissions.

- Collect data (import and tidy often the largest part in this course).
- EDA (distributions, outliers, missing values, correlations), especially if first use of data set(s) or variable(s).
- Model fit.
- Model assessment.
- Reflect (briefly) on the course topic that the lab is built around; what have we used, why, and how successful were we?
- No more than 10 pages in total.

#### Reading

- Introduction to Statistical Learning (Chapter 2)
   http://www-bcf.usc.edu/~gareth/ISL/
- R Markdown Cheat Sheet https://www.rstudio.com/wp-content/uploads/2015/02/ rmarkdown-cheatsheet.pdf