# Data Science for Medicine and Public Health

National Preventive Medicine Residency Program

Resident-Led Session

29 March 2018

Guest Speaker: Asst. Prof. Mengling Feng

Facilitator: Dr Calvin Chiew

## Asst. Prof. Mengling 'Mornin' Feng



- Assistant Professor at NUS School of Public Health, School of Medicine and School of Computing
- Research Affiliate with Lab of Computational Physiology,
  Harvard-MIT Health Sciences and Technology Division
- Senior Assistant Director at NUH Informatics Office, championing big data analytics to extract actionable knowledge and improve quality of care
- Expert in machine learning, optimization, signal processing,
  statistical causal inference and big data management

## Today's Agenda

- Guest Lecture: Data Science and AI to Improve Our Understanding of Health and Disease
- Workshop I: Introduction to Python and Jupyter Notebook
  - Hands-on Exercise
- Break
- Workshop II: Applied Statistical Learning in Python
  - Sample Code Review
  - Hands-on Exercise







## Introduction to Python and Jupyter Notebook

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https://github.com/calvinjchiew/datasci

## Terminology

- A Jupyter notebook consists of cells, which can contain either Markdown or code.
  - Each cell can be executed independently, but once a block of code is executed, it lives in the memory of the kernel.
- A comma-separated values (CSV) file stores tabular data in plain text.
  - Each record consists of values (can be numeric or text) separated by commas.
- pandas (typically imported as pd) can load csv data into dataframes which optimize storage
  and manipulation of data. Dataframes have useful methods eg. head, shape, merge etc.
- The pyplot module (typically imported as plt) in matplotlib contains useful functions for generating simple plots eg. plot, scatter, hist, title, show etc.

## Basic Built-in Data Types

Integers	7
Floats	7.0
Booleans	True, False
Strings	'Hi', "7.0"
Lists	[], ['Hello', 70, 2.1, True]

- You can use either single or double quotation marks to enclose strings.
- Lists are collections of items, which can be of different types. They are indicated by square brackets, with items separated by commas.
- You do not need to declare the types of your variables. The type is inferred from the value assigned to the variable.

## Beginner's Cheat Sheet

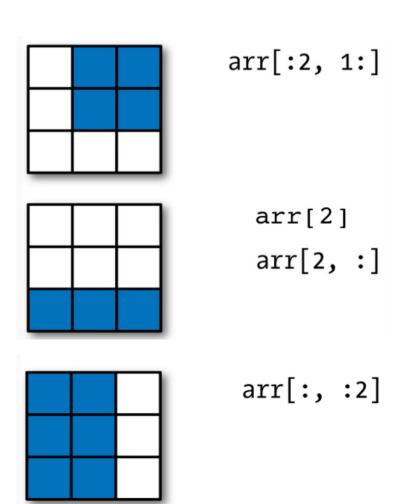
Arithmetic	+, -, *, /, %, **, //
Comparison	==, !=, >, <, >=, <=
Boolean logic	and, or, not
Indexing lists/strings	[n], [n:m], [n:], [:n]
Selection	if, elif, else
Iteration/loop	for, in, range
Create function	def, return
Call function	function(arg1, arg2,)
Call object's method or library's function	object.method(arg1, arg2,) library.function(arg1, arg2,)
Get length of list/string	len ()
Import library	import as
Print	print()

## **Indexing 2D Arrays**

Shape of 2D array is written as

(number of rows, number of columns)

- The element at the n-th row and the m-th column is indexed as [n, m]. Just like lists, you can also get multiple array values at a time.
- Remember that Python is zero-indexed, ie. counting starts from zero, not one!



#### **Best Practices**

- Code readability is key. Python syntax is close to plain English.
- Variables should be given descriptive names.
- Intersperse your code with comments, which are indicated by #.
- Proper indentation is non-negotiable in Python. Code blocks are not indicated by delimiters eg. { }, only by indentation.
- When in doubt, Google for help and read the documentation of libraries used.

## Hands-on Exercise

https://github.com/calvinjchiew/datasci >> 'python' folder >> Python.ipynb









## **Applied Statistical Learning** in Python

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https://github.com/calvinjchiew/datasci

## Important Concepts for Today

- 1. Model fit
- 2. Random forest
- 3. Support vector machine
- 4. Cross-validation

Summarized in handout

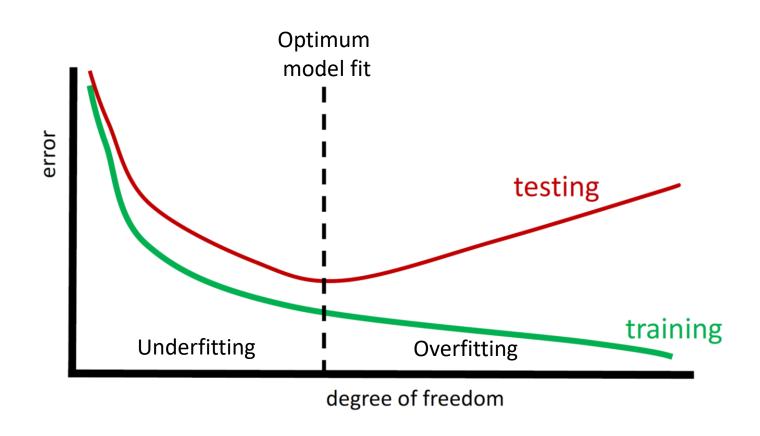
https://github.com/calvinjchiew/datasci >> 'lecture' folder >> handout.md

## Model Fitting

We want to estimate f where

$$Y = f(X_1, X_2, X_3 \dots) + \varepsilon$$

- X: feature, predictor, independent var
- Y: outcome, response, dependent var
- ε: error
- Data is split into distinct training and testing sets to prevent overfitting
- Loss (error) function depends on prediction task



## Popular ML Methods

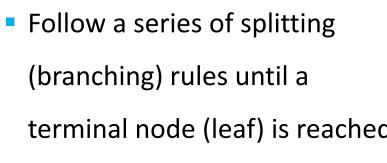
#### Supervised Learning

- K-nearest neighbours
- Regression (linear, logistic, polynomial, spline etc.) ± regularization
- Linear/quadratic discriminant analysis
- Tree-based approaches: decision tree, bagging, random forest, boosting
- Support vector machine
- Neural network

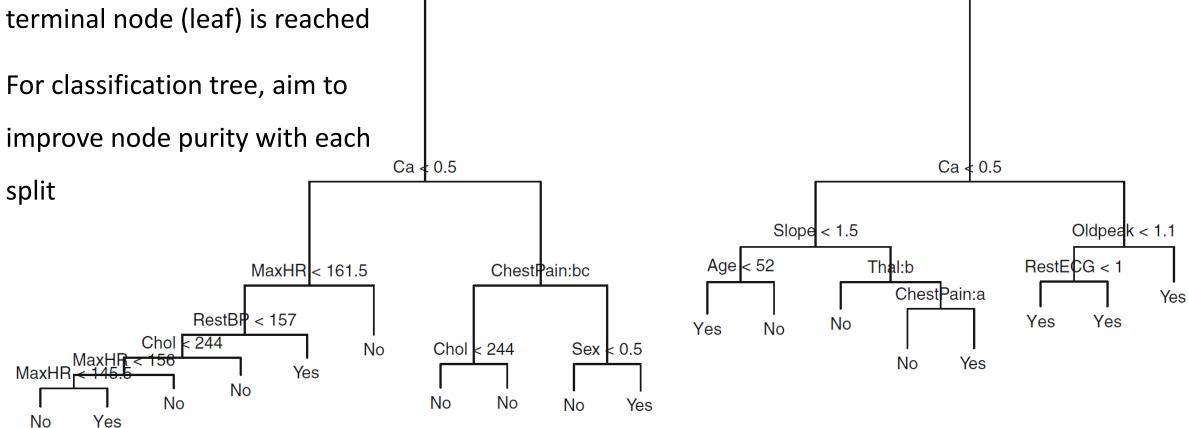
#### Unsupervised Learning

- Principal components analysis
- Clustering
- Neural network

#### **Decision Tree**



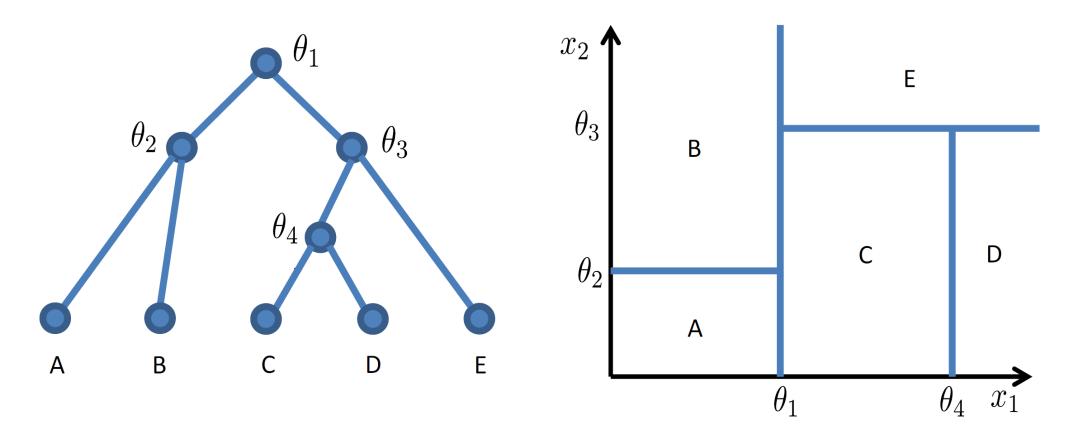
For classification tree, aim to



Thal:a

## **Decision Tree**

- The feature space is split into rectangular regions (boxes)
- We use the mean or majority class of observations in each region for prediction



### Random Forest

• Multiple trees which are combined to yield a single consensus prediction

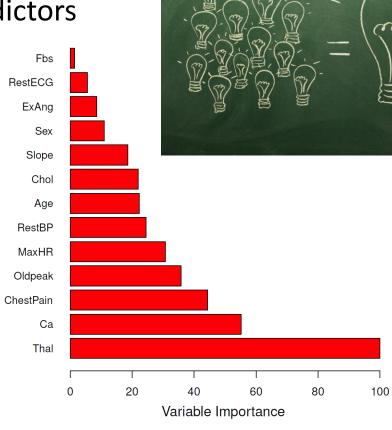
Averaging multiple onerous predictions produce less uncertain results

At each branch, only a random subset of all the predictors

are considered as potential split candidates

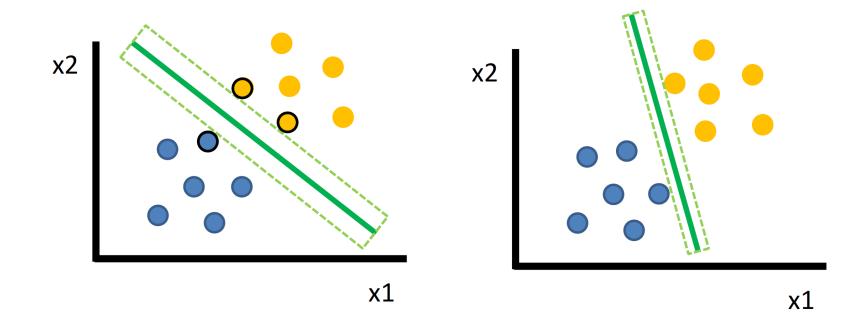
To obtain trees that are less similar to each other

 Feature importance can be visualized by total decrease in Gini index due to splits over the feature



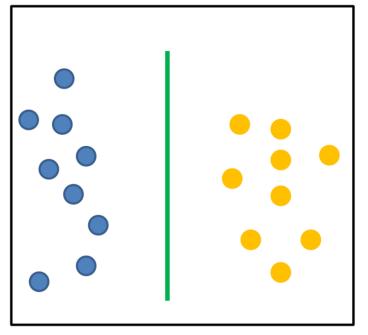
## Support Vector Machine

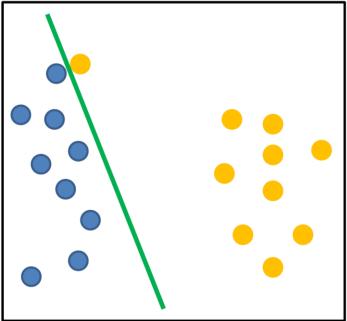
- The wider the margin, the more confident we are in the separating hyperplane
- Separating hyperplane depends only on the support vectors

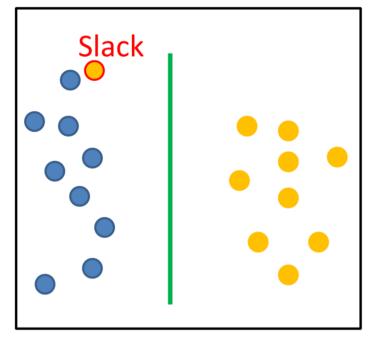


## **Support Vector Machine**

 We allow some slack for data points to be on the "wrong" side of the hyperplane in exchange for a more robust hyperplane (against outliers)

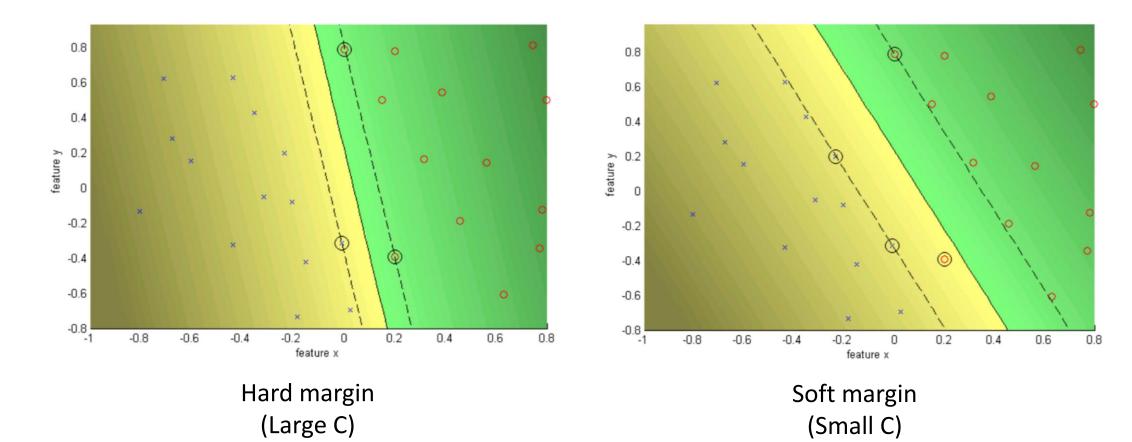






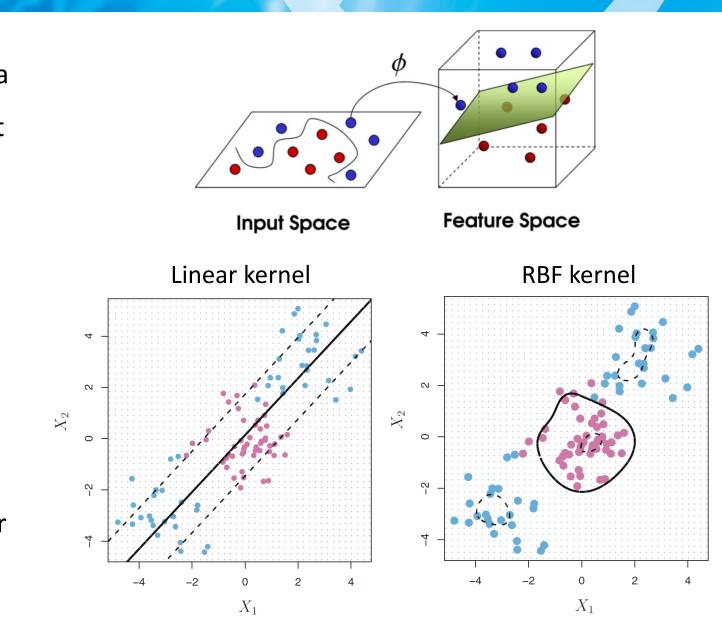
## **SVM** Regularization

■ When C is small, more slack is allowed, resulting in a softer (but wider) margin

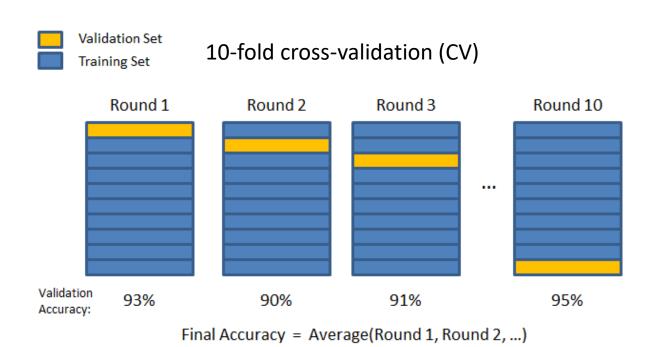


### **SVM Kernel**

- Non-linear kernels allow us to project data points not linearly separable (on the input space) onto a higher-dimensional (feature) space where a linear separating hyperplane can be drawn
- This projection is done by a kernel function eg. radial basis function (RBF)
- When projected back onto the input space, the decision boundary is non-linear



## k-fold Cross-validation

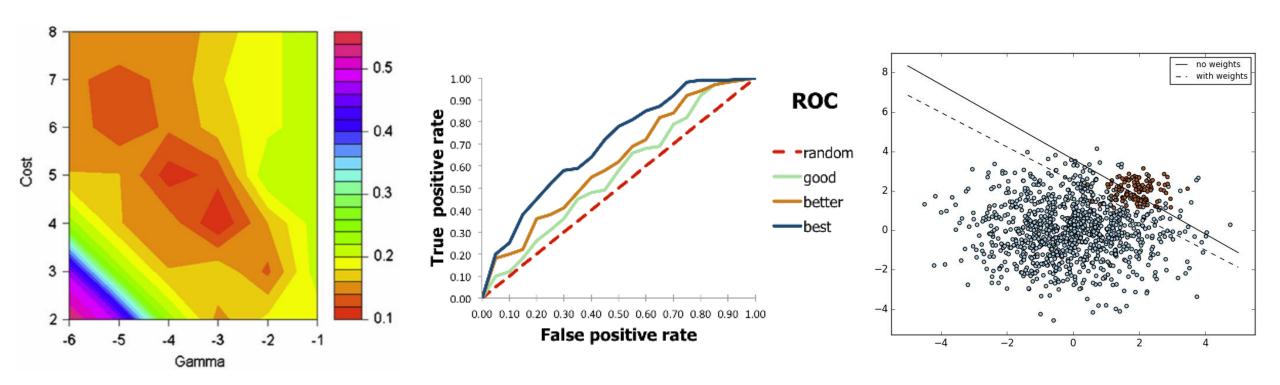


#### Examples of parameters to optimize:

- Random forest
  - Number of trees, B
  - Number of predictors considered at each split, m
  - Max tree depth / min node size
- Support vector machine
  - Penalty / amount of slack tolerated, C
  - Kernel coefficient, γ
- Boosting
  - Learning rate, λ
  - Number of splits in each tree, d

## Others

Grid Search CV, Receiver Operating Characteristic (ROC) curve, class weighting





https://github.com/calvinjchiew/datasci >> 'sample' folder >> Lung.ipynb



https://github.com/calvinjchiew/datasci >> 'exercise' folder >> Leukemia.ipynb