Introduction to Modeling in Python

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What is a model?

- A **model** is a mathematical function of form y = f(X)
 - y is some quantity of interest that we want to predict
 - X is a collection of individual observations and some information about them
 - f is our model
- Statistical models derive insight about a real-world event while quantifying uncertainty about predictions
- Machine learning models output the best prediction possible without regard for interpretability
- "All models are wrong, but some are useful"
 - A model is a simplified representation of a potentially complex real-world phenomenon

How does a model work?

- Models "learn" by associating a set of various observations of different feature inputs (e.g. weight, age) with a set of corresponding outputs (e.g. height) in a way such that an arbitrary cost function is minimized
 - This combination is known as the "training set"
- Modeling is typically one of the last steps in a data science pipeline
 - o Make sure data is cleaned and relevant features are selected or processed beforehand
 - Typical feature engineering tasks include removal of missing values,
 normalization/standardization, or one-hot encoding depending on model type
- The process of modeling includes many potentially subjective steps that need to be justified by the modeler, so be careful!

Model performance

- Models are ultimately judged on their ability to generalize to never-before seen data (frequently known as the "test set")
 - A train/test partition of 80%/20% is often used in practice
- A fundamental concept in modeling is the bias-variance tradeoff
 - o **Bias** error introduced in the specification of the form of the model
 - Variance error introduced by inherent variability in the data itself
- Too much bias indicates the model has underfit to the data, while too
 much variance indicates the model is overfit to the set of observations it
 has seen
- The best models balance both!

Train/test code

```
from sklearn.datasets import load_boston

data = load_boston()
df = pd.DataFrame(data.data, columns = data.feature_names)
df['target'] = data.target
df.head()
```

| \$ | CRIM ♦ | ZN 🕏 | INDUS \$ | CHAS \$ | NOX ≑ | RM 🕈 | AGE ♦ | DIS \$ | RAD \$ | TAX \$ | PTRATIO \$ | В \$ | LSTAT \$ | target 🕏 |
|-----------|---------|------|----------|---------|--------------|-------|--------------|--------|--------|--------|------------|--------|----------|----------|
| 0 | 0.00632 | 18.0 | 2.31 | 0.0 | 0.538 | 6.575 | 65.2 | 4.0900 | 1.0 | 296.0 | 15.3 | 396.90 | 4.98 | 24.0 |
| 1 | 0.02731 | 0.0 | 7.07 | 0.0 | 0.469 | 6.421 | 78.9 | 4.9671 | 2.0 | 242.0 | 17.8 | 396.90 | 9.14 | 21.6 |
| 2 | 0.02729 | 0.0 | 7.07 | 0.0 | 0.469 | 7.185 | 61.1 | 4.9671 | 2.0 | 242.0 | 17.8 | 392.83 | 4.03 | 34.7 |
| 3 | 0.03237 | 0.0 | 2.18 | 0.0 | 0.458 | 6.998 | 45.8 | 6.0622 | 3.0 | 222.0 | 18.7 | 394.63 | 2.94 | 33.4 |
| 4 | 0.06905 | 0.0 | 2.18 | 0.0 | 0.458 | 7.147 | 54.2 | 6.0622 | 3.0 | 222.0 | 18.7 | 396.90 | 5.33 | 36.2 |

Train/test code

(379, 13) (379,) (127, 13) (127,)

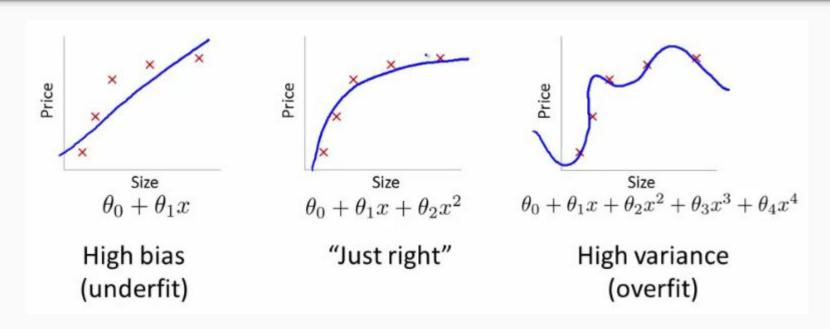
```
from sklearn.model_selection import train_test_split

X = data.data
y = data.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.25)

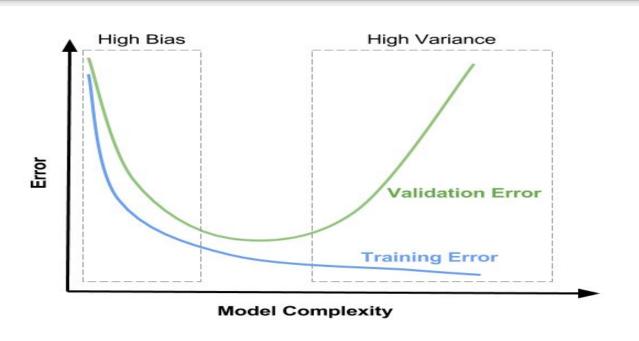
print(X_train.shape, y_train.shape, X_test.shape, y_test.shape)
```

Bias-variance tradeoff

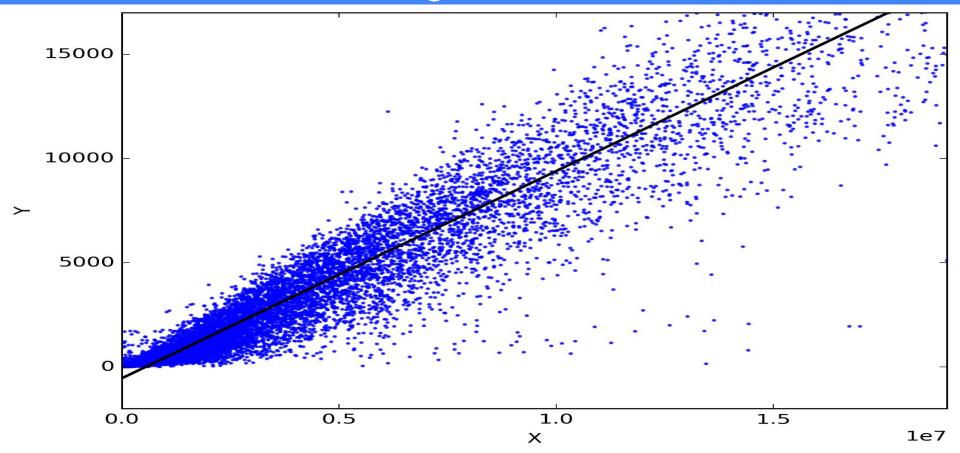


http://www.turingfinance.com/perils-optimization-in-investment-management/

Bias-variance tradeoff



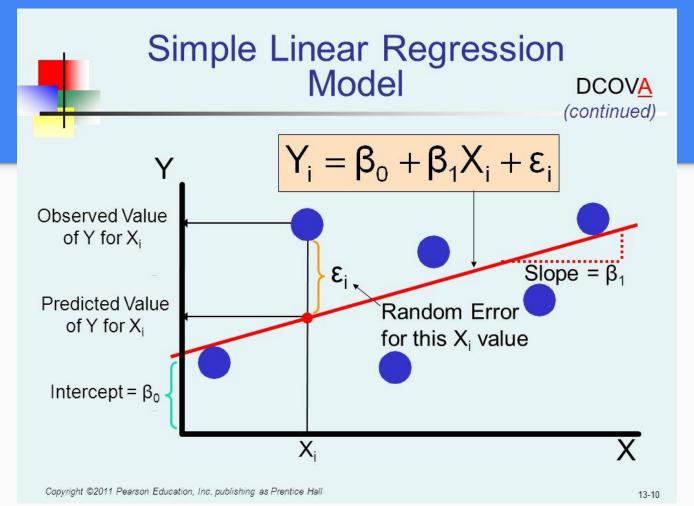
Regression



https://medium.com/@amarbudhiraja/ml-101-linear-regression-tutorial-1e40e29f1934

Regression background

- "Regression" is the act of predicting a continuous quantity (e.g. the price of something)
- An example is *linear regression*, which fits an (N-1)-dimensional hyperplane to an MxN numeric matrix of observations such that distance between the hyperplane and all observations is minimized
 - A typically used cost function is mean-squared error (MSE) e.g. the average of the sum of (y - y_pred)^2 across all M observations
- Linear regression assumes normally distributed error terms, and is therefore a *parametric* model
 - Other assumptions include independence and even variance of errors, a true underlying linear relationship, and uncorrelated input variables



Linear regression code - fitting the model

```
import numpy as np
 from sklearn.linear model import LinearRegression
 lin reg = LinearRegression()
 lin reg.fit(X train, y train)
- print(
      np.round(lin reg.coef ,1),
      '\n'.
      np.round(lin reg.intercept , 1)
\begin{bmatrix} -0.1 & 0. & -0.2 & 0. & -0. & 0.1 & -0.3 & -0.2 & -0. & 0.1 & -0.2 & -0.3 & -0. \end{bmatrix}
3.5
```

y = -0.1(CRIM) - 0.2(INDUS) + 0.1(RM) - 0.3(AGE) - 0.2(DIS) + 0.1(TAX) - 0.2(PTRATIO) - 0.3(B) + 3.5

Linear regression code - evaluation

```
from sklearn.metrics import mean_squared_error
y_pred = lin_reg.predict(X_test)

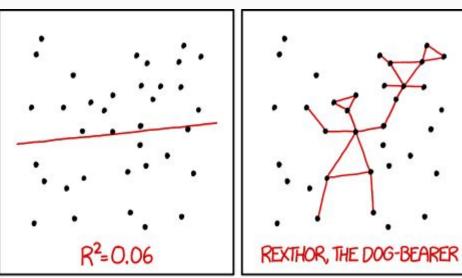
print('R^2 Score:', lin_reg.score(X_test, y_test))
print('Mean squared error: ', mean_squared_error(y_test, y_pred))
```

R^2 Score: 0.8850825151531964 Mean squared error: 0.06843974653098528

The **score()** method uses R², or the coefficient of determination, which measures how much of the total variation in the data is explained successfully by the model.

Values range from 0 to 1

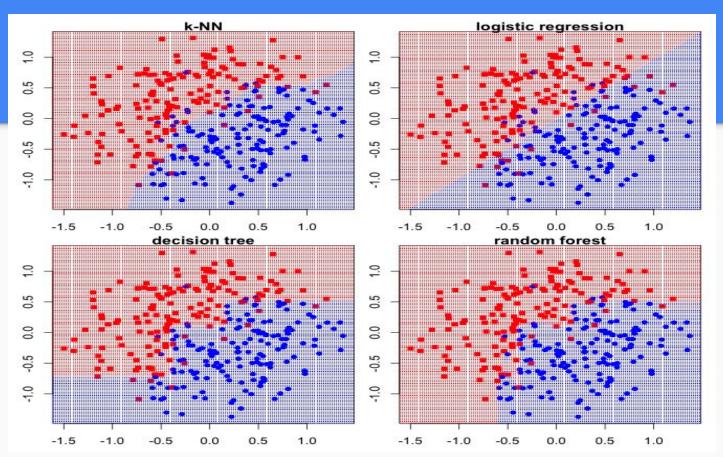
Relevant XKCD



I DON'T TRUST LINEAR REGRESSIONS WHEN IT'S HARDER TO GUESS THE DIRECTION OF THE CORRELATION FROM THE SCATTER PLOT THAN TO FIND NEW CONSTELLATIONS ON IT.

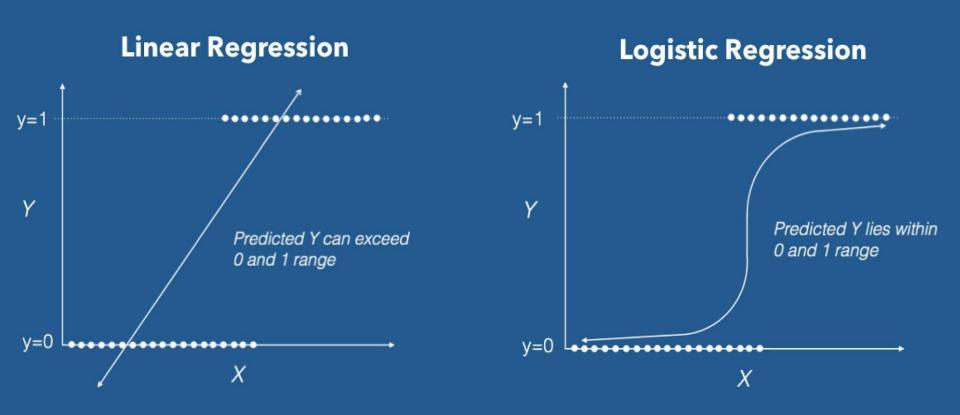
https://www.explainxk cd.com/wiki/index.php /1725:_Linear_Regress ion

Classification

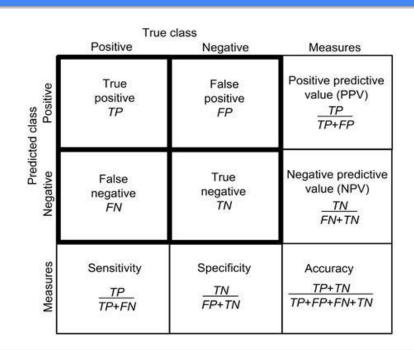


Classification background

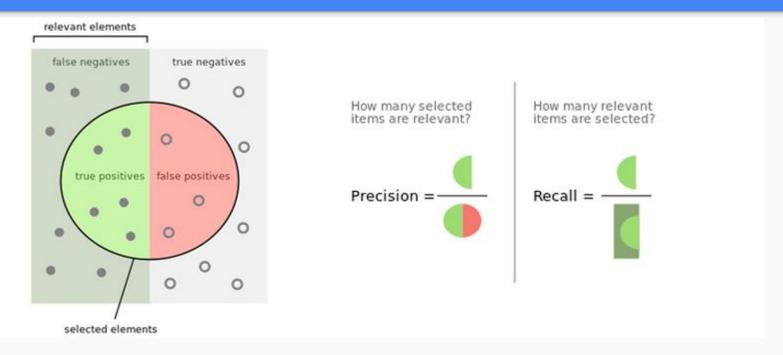
- In *classification*, our interest is a discrete class rather than a continuous quantity (e.g. a label indicating whether a sale was made or not), and our goal is to find a *decision boundary* that accurately separates classes
- A common classification model is *logistic regression*, which outputs a probability of each observation belonging to a class
 - Like linear regression, it consists of a linear set of coefficients which are fed into a link function ensuring they output a real number between 0 and 1
- Classifiers are generally evaluated through metrics like accuracy, precision, and recall
 - The right metric to optimize will depend on your modeling situation



Classification evaluation - confusion matrix



Classification evaluation - precision/recall



Classification Code

```
from sklearn.datasets import load_wine

wine = load_wine()
wine_df = pd.DataFrame(wine.data, columns = wine.feature_names)
wine_df['target'] = wine.target
wine_df.head()
```

| nesium 🗢 | total_phenols \$ | flavanoids \$ | nonflavanoid_phenols \$ | proanthocyanins \$ | color_intensity \$ | hue 🕏 | od280/od315_of_diluted_wines \$ | proline \$ | target \$ |
|----------|------------------|---------------|-------------------------|--------------------|--------------------|-------|---------------------------------|------------|-----------|
| 127.0 | 2.80 | 3.06 | 0.28 | 2.29 | 5.64 | 1.04 | 3.92 | 1065.0 | 0 |
| 100.0 | 2.65 | 2.76 | 0.26 | 1.28 | 4.38 | 1.05 | 3.40 | 1050.0 | 0 |
| 101.0 | 2.80 | 3.24 | 0.30 | 2.81 | 5.68 | 1.03 | 3.17 | 1185.0 | 0 |
| 113.0 | 3.85 | 3.49 | 0.24 | 2.18 | 7.80 | 0.86 | 3.45 | 1480.0 | 0 |
| 118.0 | 2.80 | 2.69 | 0.39 | 1.82 | 4.32 | 1.04 | 2.93 | 735.0 | 0 |

```
X = wine.data
y = wine.target

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.25, stratify = y)
print(X_train.shape, y_train.shape, X_test.shape, y_test.shape)

(133, 13) (133,) (45, 13) (45,)
```

Classification Code - fitting the model

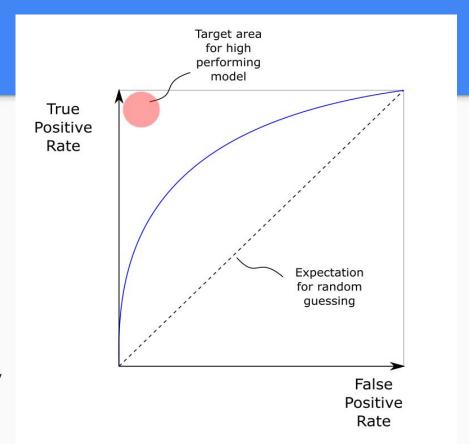
```
from sklearn.linear model import LogisticRegression
 log reg = LogisticRegression()
 log reg.fit(X train, y train)
▼ print(
     np.round(log reg.coef , 2),
      '\n'.
     np.round(log reg.intercept )
0.02]
 [ 0.68 -1.04 -0.7 \quad 0.26 \quad 0.01 \quad 0.32 \quad 0.52 \quad 0.21 \quad 0.45 \quad -1.87 \quad 0.73 \quad 0.15
  -0.011
 \begin{bmatrix} -0.08 & 0.72 & -0.08 & 0.06 & 0.01 & -0.61 & -1.49 & -0.01 & -0.64 & 1.33 & -0.41 & -1.26 \end{bmatrix}
 -0.11
 [-0. 0. -0.1]
```

Classification code - evaluation

```
print('Accuracy: ', log reg.score(X test, y test))
 y pred = log reg.predict(X test)
 conf matrix = confusion matrix(y test, y pred)
print('\nConfusion Matrix:\n',
       pd.DataFrame(
           confusion matrix(y test, y pred),
           columns = ['pred: 0', 'pred: 1', 'pred: 2'],
           index = ['true: 0', 'true: 1', 'true: 2']),
       '\n\nClassification Report:\n',
       classification report(y test, y pred))
Accuracy: 0.95555555555556
Confusion Matrix:
         pred: 0 pred: 1 pred: 2
true: 0
true: 1
                               12
true: 2
Classification Report:
             precision
                          recall f1-score
                                             support
                 1.00
                           1.00
                                     1.00
                                                 15
                 1.00
                           0.89
                                     0.94
                                                 18
                 0.86
                           1.00
                                     0.92
                                                 12
                 0.96
                           0.96
                                     0.96
                                                  45
avg / total
```

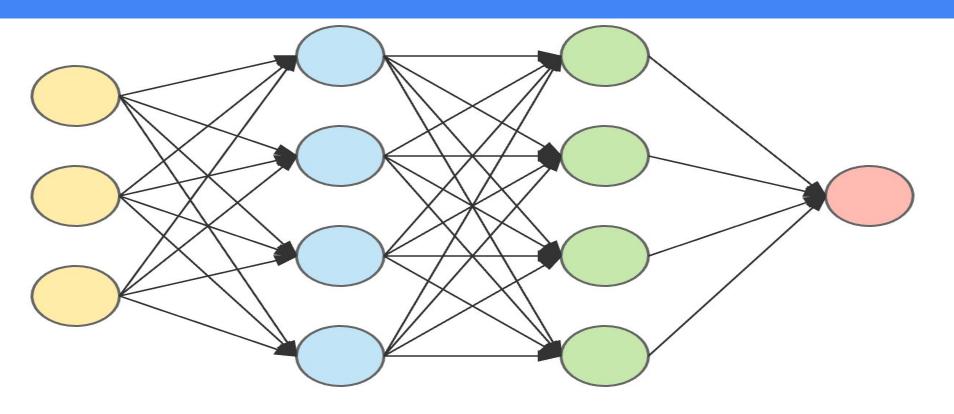
from sklearn.metrics import confusion matrix, classification report

ROC curve



https://deparkes.co.uk/2018/02/ 16/the-roc-curve/

Advanced Methods



hidden layer 2

output layer

https://towardsdatascience.com/applied-deep-learning-part-1-artificial-neural-networks-d7834f67a4f6

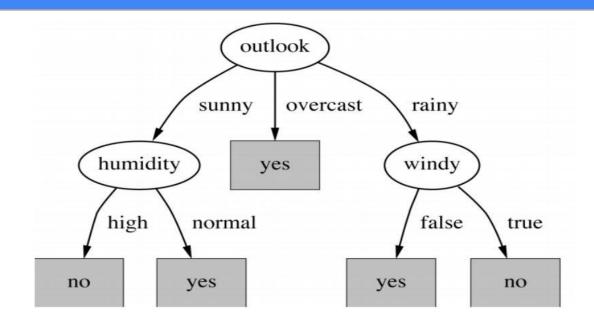
hidden layer 1

input layer

Decision tree background

- Some advanced methods can perform both regression and classification
- Decision trees attempt to split observations into distinct groupings based on feature values such that bin purity is maximized, ultimately creating a binary tree in the process
 - The predicted class will then be the maximally-represented class in the leaf nodes for classification, or an average of continuous values for a regression
- Decision trees are prone to overfitting, but the number of splits (e.g. tree depth) can be specified beforehand by the modeler
- Non-parametric, meaning it makes no assumption of the underlying form of the data

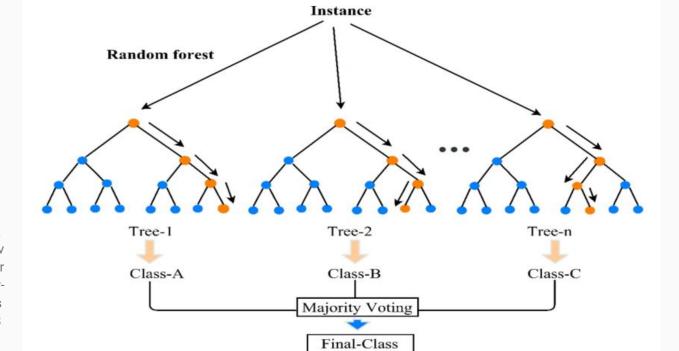
Decision tree (playing tennis)



Random forest background

- In order to reduce variance and chance of overfitting, a *random forest* is an average of the results of many decision trees splitting on random subsets of features
- Random forests are typically a *robust* model working well in many situations (including high-dimensional data), but they may lack interpretability as a result
- Other forms of tree-based learning include **boosting** and **bagging** (beyond the scope of this presentation, but worth looking into for those interested!)

Random forest example



https://www.resear chgate.net/figure/R andom-Forests-Naiv e-Bayes-NB-NB-appr oaches-are-a-familyof-simple-probabilis tic_fig2_326722598

Random forest code example: Boston

```
from sklearn.ensemble import RandomForestRegressor
 rf reg = RandomForestRegressor(max depth = 4)
 rf reg.fit(X train, y train)
 y pred rf = rf reg.predict(X test)
 print(rf reg.feature importances )
 print('\nR^2 Score:', rf reg.score(X test, y test))
 print('Mean squared error: ', mean squared error(y test, y pred rf))
[0.06543473 0.00104719 0.00340049 0. 0.00376146 0.
 0.38850392 0. 0.00343326 0.08661245 0. 0.22357216
 0.224234341
R^2 Score: 0.9059701492537314
Mean squared error: 0.056
```

Additional types of models

- We have only covered a small subset of modeling approaches that exist!
- Regression
 - Regularized linear regression (LASSO, Ridge, Elastic Net), polynomial regression/splines
- Classification
 - Naive Bayes, linear/quadratic discriminant analysis
- Regression/classification
 - Support vector machines, K-nearest neighbors, XGBoost
- Other approaches
 - Survival analysis, time series, structural equation modeling

Other types of modeling

- Unsupervised learning infer y from the data (no explicit optimization criteria)
 - E.g. k-means clustering (split data in k distinct groupings, useful for market segmentation)
 - Also encompasses dimensionality reduction techniques like principal component analysis
- Deep learning send input through a nested series of "hidden" activation functions (or layers) before reaching the output
 - E.g. neural networks (require lots of data, often used for language or vision related tasks)
- Reinforcement learning choose the best actions in a system where many are possible in order to optimize some reward function over time
 - E.g. q-learning (useful for robotics and training AI to play video games)

Additional resources

- Linear regression:
 https://scikit-learn.org/stable/modules/generated/sklearn.linear_model.LinearRegression.html
- Logistic regression: https://scikit-learn.org/stable/modules/generated/sklearn.linear_model.LogisticRegression.html
- More methods and explanations:https://scikit-learn.org/stable/tutorial/statistical_inference/s upervised_learning.html

Exercises

Link to respective Jupyter Notebook / RMD files here