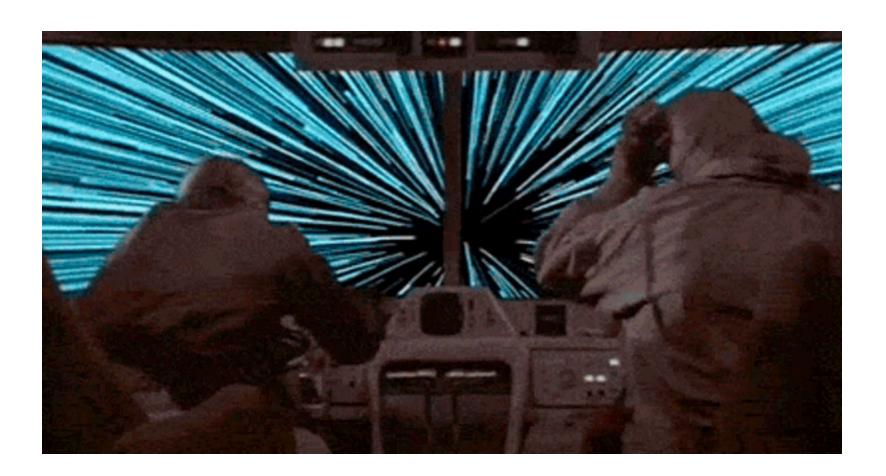
Boosted Trees and xgboost



Ensemble Flavors - Extended

Single Model Paradigm

Standard Model:

$$\hat{f}(x) = f^*(x) pprox f(x)$$

All training data is used to generate our single best estimate of the true functional form, f(x).

Bagging (Refresh)

$$\hat{f}_{bag}(x) = rac{1}{B} \sum_{b=1}^{B} f_b^*(x)$$

In bagging, each estimate utilizes a bootstrap (random) sample of the training data

The bagged estimate is then based on the weighted average of all of the models

Boosting

If we boost an algorithm using M stages, then we need to define $f_m(x)$ at each stage

$$\hat{f}_0(x) = 0$$

At each subsequent stage, we solve for

$$\hat{f}_m(x) = \hat{f}_{m-1}(x) - f_m^*(x)$$

So that each stage adds more information to our model.

Q: Why do we subtract??

Boosting vs Bagging

Bagging:

 An averaged model utilizing bootstrapped samples of the complete dataset

Boosting:

An "additive" model, where the predictions are incrementally improved

Boosting vs Bagging

Bagging:

- Much easier to implement
- Less Overfitting

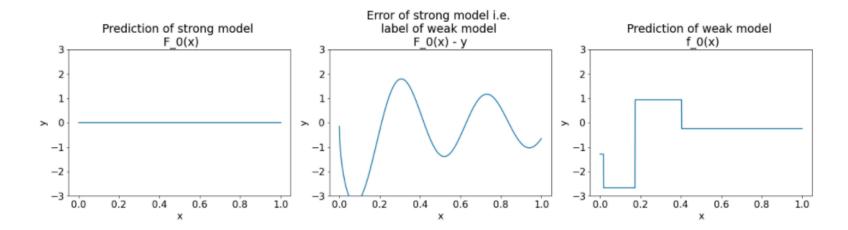
Boosting:

- Better Performance (generally)
- More vulnerable to overfitting

A visual example

Start with

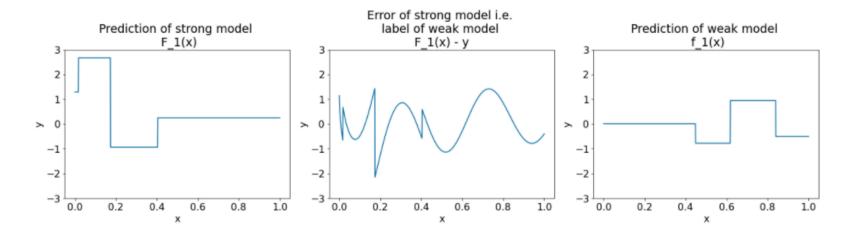
- A baseline model
- Error
- Tree fitted to the error of the baseline



A visual example

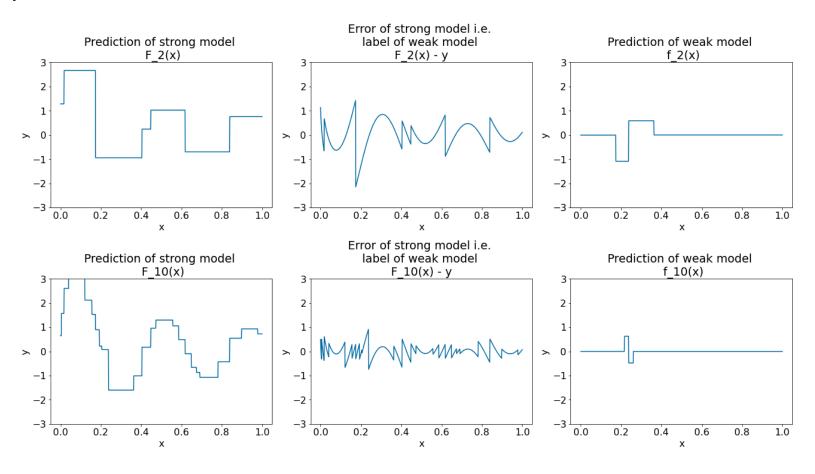
Our new model is the old baseline - the new tree, then we repeat:

- Baseline
- Error
- Tree predicting error



A visual example

We keep doing it! Over time, our model converges toward the patterns observed in the data:



Why xgboost? It enables parallel computation of the model and distributed training!

This makes it a useful production model. $\ensuremath{\mbox{\ensuremath{$\oplus}}}$

First things first, let's import all of our other libraries and read in our MNIST data:

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score

data = pd.read_csv("https://github.com/dustywhite7/
Econ8310/raw/master/DataSets/mnistTrain.csv")
```

Create those train and test splits!

```
# Upper case before split, lower case after
Y = data['Label']
# make sure you drop a column with the axis=1 argument
X = data.drop('Label', axis=1)

x, xt, y, yt = train_test_split(X, Y, test_size=0.1, random_state=42)
```

Implement the model:

```
from xgboost import XGBClassifier

xgb = XGBClassifier(n_estimators=50, max_depth=3,
   learning_rate=0.5, objective='multi:softmax')
```

Fit the model and test its performance:

```
xgb.fit(x, y)
pred = xgb.predict(xt)
print(accuracy_score(yt, pred)*100)
```

```
93.4
```

Nice! Low effort, high accuracy!

A Note on Cross-Validation

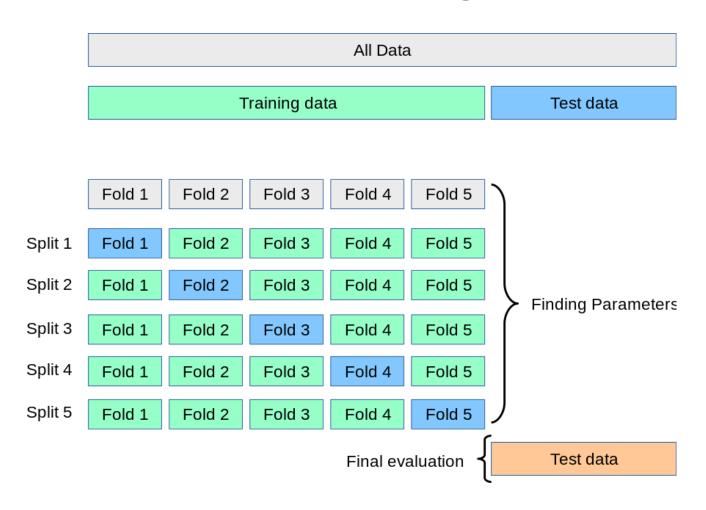
Cross-Validation

In order to maximize the information that we have about our model's performance, we will often test the model on many draws of our existing data.

This is called cross-validation. We resample our training and testing data k times, and compare the performance of the models.

If the models perform more or less equally well, then we can treat our model as well-specified. If not, then we know performance was sample-dependent.

Cross-Validation Diagram



credit to sklearn for the image

Cross-Validation Code

```
from sklearn.model selection import KFold
# If we have imported data and created x, y already:
kf = KFold(n_splits=10) # 10 "Folds"
models = [] # We will store our models here
for train, test in kf.split(x): # Iterate over folds
  model = model.fit(x[train], y[train]) # Fit model
  accuracy = accuracy_score(y[test],  # Store accuracy
    model.predict(x[test]))
  print("Accuracy: ", accuracy_score(y[test],
  model.predict(x[test])))  # Print results
models.append([model, accuracy])  # Store it all
print("Mean Model Accuracy: ",  # Print aggregate
  np.mean([model[1] for model in models]))
```

One More Note

After we complete our cross-validation, we do not use the cross-validation models. When we are satisfied with the results of the cross-validation, we

- Recombine all training data
- Train the model on all training data
- Test the model on withheld testing data (should not have been used at all in cross-validation)
- If results are still positive, implement model!

Lab Time!