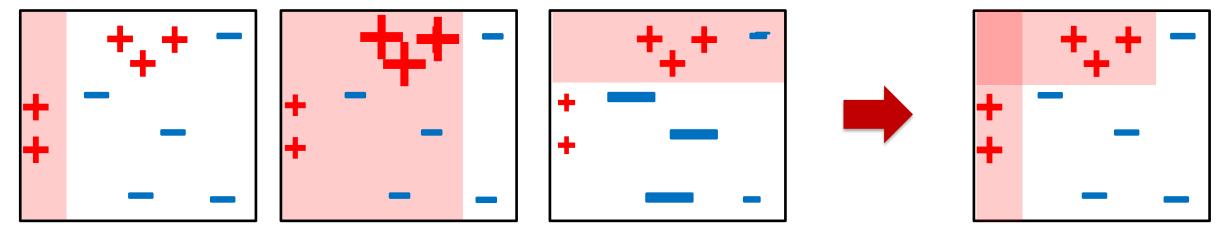
Machine Learning for Biomedical/Healthcare Applications

Combining weak classifiers....



....to make strong ones

Ensemble Learning: Random Forests and Boosting

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Learning Outcomes from the tutorial

- 1. Understanding of how to implement Bagged ensembles of trees through
 - 1. Creating bootstrapped samples of a data set
 - 2. Training large numbers of trees
 - 3. Aggregating test predictions
- 2. Compare the performance of ensembles relative to decision Trees
- 3. Be able to deploy bagging, forests and boosting in Scikit-Learn
- 4. Be able to perform parameter optimization for ensemble methods

Solutions for decision trees

1.1 estimate gini index for a given split

- 1. Estimate the total number of samples by summing over the list branch
- 2. Proportion in each branch equals (total number of samples in branch)/(total number of samples)
 - Total number of samples in branch is given by variable class_total (each item in list branch)
 - Total overall is given by output of step 1 variable split_size
- 3. Gini= $1 \sum_{y_k \in Y} p(y_k)^2$

```
def gini coefficient(branch):
           Estimates Gini Coefficient for a given class split
            input:
                split: list of length k (where k= number of classes).
                       The values at each index reflect the toal number of instances
                      of each class, for this proposed branch split
 9
10
            output:
11
                gini: gini coefficient for this split
12
13
       # 1.1.1 estimating total number of samples in branch split (by summing contents of split list)
14
       split size=np.sum(branch)
       gini=1
15
16
       # iterating over all items in the array
17
       for class total in branch:
18
            # 1.1.2. estimating p*p for this class label; subtracting from current gini total
19
           proportion class k=(class total/split size)
           # 1.1.3. subract the squaree of the propoertion per class from current estimate of the gininindex
20
21
            gini--proportion class k*proportion class k
22
23
       return gini
```

1.2 Propose splits

- 1. check all the values of the features at that indexed position
 - Given by row[index] for each row (here index is the feature/attribute index)
- 2. split the data into a left branch (if that data example's feature is below the threshold) and into a right branch (if that data example's feature is below the threshold).

```
def test split(index, value, dataset):
           Split a dataset based on an attribute and an attribute value
           input:
               index = feature/attribute index (i.e. data column index) on which to split on
               value = threshold value (everything below this goes to left split,
                        everything above goes to right)
                dataset = array (n samples, n features+1)
                        rows are examples
10
                        last column indicates class membership
11
                        remaining columns reflect features/attributes of data
12
13
           output:
14
               left, right: data arrays reflecting data split into left and right branches, respectively
        0.00
15
16
17
       # create empty list that you will populate with rows of dataset
       left=[]
19
       right = []
       # the loop below will slice rows from data set
21
       for row in dataset:
           # if the value of this feature for this row is less than
23
           # the (threshold) value split into left branch, else split into right
24
           if row[index] < value:</pre>
25
               left.append(row)
26
           else:
27
                right.append(row)
28
29
       return np.asarray(left), np.asarray(right)
```

1.2 Propose splits

- Then test for outcome of splitting on first feature (column index=6)
- With threshold equal to 7th value (or row)

```
index=0
   rowindex=6
 3 threshold=X[rowindex,index]
    print('the value of the feature {} at row {} of the data set is {}'.format(index,rowindex,threshold))
the value of the feature 0 at row 6 of the data set is 9.00220326
    branches=test split(index, X[rowindex,index], X)
   print('Our left branch is \n {}'.format(branches[0]))
   print('Our right branch is \n {}'.format(branches[1]))
Our left branch is
[[2.77124472 1.78478393 0.
 [1.72857131 1.16976141 0.
[3.67831985 2.81281357 0.
[3.96104336 2.61995032 0.
[2.99920892 2.20901421 0.
[7.49754587 3.16295355 1.
[7.44454233 0.47668338 1.
 [6.64228735 3.31998376 1.
Our right branch is
[[ 9.00220326 3.33904719 1.
[10.12493903 3.23455098 1.
                                     11
```

1.3: Estimate total cost of split

- 1. Iterate over class IDs (line 26)
- 2. Slices rows corresponding to that class (line 34)
- 3. Count number of rows for that slice (line 36)
- 4. Add these to list: class_counts_for_branch
 - input to gini_coefficient function
- 5. Estimate gini_coefficient for that branch
- 6. Weight by proportion of samples in branch
 - branch.shape[0]/total samples

```
Estimates the cost for a proposed split
                splits: tuple or form (L,R) where L reflects the data for the left split and
                        R reflects data for left split
                classes: list of class values i.e. [0,1]
10
11
                cost: sum of gini coefficient for left and right sides of the split
12
13
        cost=0
14
        total samples=0
15
16
        # estimate the relative size of each branch
17
        for branch in split:
18
            total samples += branch. shape[0]
19
20
        # for each (left/right) split on the proposed tree
21
        for br index, branch in enumerate(split):
            # initialise list of class counts for this branch
23
            class counts for branch=[]
24
            # for each class value, count total of data examples (rows)
25
            # that have for this class, in this branch
26
            for class val in classes:
27
28
                if branch.shape[0] == 0: # don't continue if size of split is 0
29
                    continue
30
31
                # 1.3.1 slice data to return only rows from branch which have this specific class value
32
                # here branch[:,-1] returns the column containing the labels and we want to slice all rows
33
                # for class=class val
                branch per class=branch[branch[:,-1]==class_val]
35
                # 1.3.2 count the number of rows with this class in this branch and append
36
                total rows=branch per class.shape[0]
37
                # this is generating list of class counts per branch which get fed to
38
                # the gini coefficient function
39
                class counts for branch.append(total rows)
40
41
            # 1.3.3. estimate the gini coefficient for this split (or branch)
42
            gini split=gini coefficient(class counts for branch)
43
            # 1.3.4. estimated the weighted contribution for this split
44
            weighted by sample size=gini split*(branch.shape[0]/total samples)
45
            # total cost is a weighted average of gini coefficients for both splits
46
            cost+=weighted by sample size
47
48
49
        return cost
          class values=[0,1]
          splitcost=split_cost(branches,class_values)
          print('The cost of the proposed split is: ', splitcost)
```

The cost of the proposed split is: 0.375

def split cost(split, classes):

1.4 Choose optimal feature/threshold split

- 1. Line 28 Iterate over all but last column:
 np.arange(dataset.shape[1]-1)
- 2. Iterate over all rows to propose threshold value (line 31)
- 3. Line 33 Create split accordingly:
 - Function test split
 - Arguments: index (feature)
 row[index] (threshold) dataset
- 4. Estimate cost using split_cost (line 35)
- 5. If improved (cost < best_cost) as Gini must be minimized (l. 36)
- 6. If cost<best_cost then save current set of parameters as 'best'

```
def get best split(dataset):
            Search through all attributes and all possible thresholds to find the best split for the data
            input:
                dataset = array (n samples, n features+1)
                        rows are examples
                        last column indicates class membership
                        remaining columns reflect features/attributes of data
10
11
            output:
12
                dict containing: 1) 'index' : index of feature used for splittling on
13
                                 2) 'value': value of threshold split on
14
                                 3) 'branches': tuple of data arrays reflecting the optimal split into left and right
15
16
17
18
        # estimating the total number of classes by looking for the total number of different unique values
        # in the final column of the data set (which represents class labels)
19
20
        class values=np.unique(dataset[:,-1])
21
22
        # initalising optimal values prior to refinment
23
        best cost=sys.float info.max # initialise to max float
24
       best value=sys.float info.max # initialise to max float
25
        best index=dataset.shape[1]+1 # initialise as greater than total number of features
                               # the best split variable should contain the output of test split that corresponds to
26
27
        #1.4.1 iterating over all features/attributes (columns of dataset)
28
        for index in np.arange(dataset.shape[1]-1):
29
30
            #Trialling splits defined by each row value for this attribute
31
            for r index,row in enumerate(dataset):
32
               # 1.4.2. return branches corresponding to thresholding on feaure (index) and threshold value (for row.
33
                branches=test split(index, row[index], dataset)
34
35
                cost=split cost(branches, class values) # 1.4.3 estimate cost for this split
36
                if cost < best_cost: # 1.4.4. if this cost is an improvement on previous costs then save the
37
                    best cost=cost # cost
38
                    best split=branches #branches
39
                    best index=index # feature index
40
                    best value=row[index] # threshold value
41
                    print('Best cost={}; Best feature={}; Best row={}'.format(best cost,index,r index)
42
43
        return {'index':best index, 'value':best value, 'branches':best split}
44
```

1.4 Choose optimal feature/threshold split

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37
                    best cost=cost # cost
38
                    best split=branches #branches
39
                    best index=index # feature index
40
                    best value=row[index] # threshold value
41
                    print('Best cost={}; Best feature={}; Best row={}'.format(best cost,index,r index)
42
43
        return {'index':best index, 'value':best value, 'branches':best split}
44
```

- 3 Running Decision Trees with Scikit-Learn
- 1. instantiate a decision tree classifier model (line 29)
- 2. fit the model to training data (line 30)
- 3. Predict test labels (line 31)
- 4. Return performance score on test examples (line 32)

```
from sklearn.tree import DecisionTreeClassifier # import the scikit-Learn Decision Tree module
   from sklearn.model selection import train test split
   from sklearn.preprocessing import StandardScaler
   from sklearn.datasets import make moons
    # CREATE a Random Data set using the sklearn Make Moons dataset
   DATA, LABELS =make moons(noise=0.3, random state=0)
10
   # Plot the data
11
   figure = plt.figure(figsize=(5, 5))
   cm = plt.cm.RdBu
   cm bright = ListedColormap(['#FF0000', '#0000FF'])
   ax = plt.subplot(1,1, 1)
16
   ax.set_title("Input data")
   ax.scatter(DATA[:, 0], DATA[:, 1], c=LABELS, cmap=cm bright,
19
              edgecolors='k')
20
21 # randomly split the data
   X train, X test, y train, y test = train test split(DATA, LABELS, test size=.4, random state=42)
23
   # **to DO** implement scikit learn decision tree classifier on this data
   #**** complete the above steps for the scikit learn classifier *****
   model = DecisionTreeClassifier(random state=0)
   model.fit(X train, y train)
   pred=model.predict(X test)
   score = model.score(X test, y test)
30
31
   print("Scikit-learn's decision tree Score", score)
32
33 # OPTIONALLY plot your results
34 # suggest plotting with with different colours for each class
   #and different markers for test and train data in order to aid visualisation
36
   # just plot the dataset first
38 f, (ax1, ax2) = plt.subplots(2, 1, sharey=True, figsize=(5,10))
39 cm = plt.cm.RdBu
40 cm bright = ListedColormap(['#FF0000', '#0000FF'])
42 ax1.set title("True labels")
   ax1.scatter(X test[:, 0], X test[:, 1], c=y test, cmap=cm bright,
44
              edgecolors='k')
45
   ax2.set title("Predicted labels")
   ax2.scatter(X_test[:, 0], X_test[:, 1], c=pred, cmap=cm_bright,
              edgecolors='k')
```

Scikit-learn's decision tree Score 0.95

• 3.2 – optional compare own tree against Scikit-Learn

```
# first combine X_train and y_train togeher (and X_test, y_test) to put data into form expected by our tree
dataset=np.concatenate((X_train,y_train.reshape((y_train.shape[0],1))),axis=1)

test_dataset=np.concatenate((X_test,y_test.reshape((y_test.shape[0],1))),axis=1)

# #3.2.1 train your tree - set max depth to 5 and min size to 1

tree = build_tree(dataset, 5,1)

# #3.2.2 Get a prediction from your test data
prediction_DT1=predict(tree, test_dataset)

# #3.2.3 Score the accuracy of your decision tree classifier
score_DT1=tree_score(y_test,prediction_DT1)
print('Our Decision Tree Score', score_DT1)
```

Our Decision Tree Score 0.875

Scikit learn applied pruning (see notebook)

Ensembles exercises

Exercises

- 1. Exercise 2 Building a Bagging Classifier (30 mins)
 - 1. Complete create_bagged_ensemble to bootstrap samples from data and train ensemble of decision trees
 - 2. Using decision tree code programmed last week (imported through DecisionTree.py)
 - 3. Complete bagging_predict to aggregate test predictions through majority voting
- 2. Exercise 3 Comparing our Bagged Predictor against our Decision Tree (just run code 5 mins)
- Exercise 4 Comparing against Scikit learn (10-15 mins)
 - 1. Build a decision tree and bagging classifier with scikit learn; fit model
 - 2. Return test prediction and accuracy (score) -> compare
- 4. Exercise 5: Training Random Forests to Predict Gestational Age from Regional Brain Volumes (20 mins)
 - implement the training and testing of the DecisionTreeRegressor()
 - implement the training and testing of the RandomForestRegressor()
 - (optional) Return and plot feature importances see the <u>scikit-learn tutorial</u> for guidance

Exercises

- 5. Exercise 6 (optional): Perform Parameter Optimisation for Random Forests using GridSearchCV (10-15 mins)
- 6. Exercise 7 (optional/homework) Building a Random Forest Classifier from scratch (20-30 mins)
- 7. Exercise 8: Training Adaboost to Predict Gestational Age from Regional Brain Volumes (20-30 mins)¶
 - Implement adaboost regression using scikit learn
 - Apply GridCV optimisation of the ensemble and base learner paramters