Lecture 4 - Student Notebook

Prelimilaries: Imports and stuff

We extended the data with extra features. The feature description is found here.

The features were calculated per week in the time_series_extended. The aggregated extended was computed by taking the mean of each feature per user across weeks.

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt

from sklearn.model_selection import train_test_split
from sklearn import tree
from sklearn.ensemble import RandomForestClassifier
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import roc_auc_score, balanced_accuracy_score
from sklearn.preprocessing import MinMaxScaler, normalize
from sklearn.linear_model import LogisticRegression

import statsmodels.api as sm
import statsmodels.formula.api as smf

from scipy.spatial.distance import pdist, cdist, squareform

# Data directory
DATA_DIR = "./../../data"
```

Section 1: Pre-processing

Read the data

```
# Parse the aggregated data frame
df_lq = pd.read_csv('{}/aggregated_extended_fc.csv'.format(DATA_DIR))
ts = pd.read_csv('{}/time_series_extended_fc.csv'.format(DATA_DIR))
```

Clean the data

We remove inactive students that did not click during weekdays and weekend for the fist 5 weeks of the semester.

```
def remove_inactive_students(df, ts):
    df = df.fillna('NaN')

#find all users weeks with 0 clicks on weekends and 0 clicks on
weekdays during the first weeks of the semester
    df_first = ts[ts.week < 5]</pre>
```

```
rows =
np.where(np.logical and(df first.ch total clicks weekend==0,
df_first.ch_total_clicks_weekday == 0).to_numpy())[0]
    df zero = df first.iloc[rows,:]
    dropusers = np.unique(df zero.user)
    ts = ts[ts.user.isin(dropusers)==False]
    df = df[df.user.isin(dropusers)==False]
    return df, ts
df_lq, ts = remove_inactive_students(df_lq, ts)
# print(df lq.columns)
display(df_lq)
     user ch num sessions ch time in prob sum ch time in video sum
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1
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                       3.4
                                          1698.4
                                                                 9227.8
2
        2
                       5.3
                                          2340.6
                                                                10801.3
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        4
                       2.5
                                          3787.3
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5
        5
                       2.5
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                       4.2
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```

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	0.027091	4.00	F	France	Y3-
277	0.137474	4.50	М	France	Y3-

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ch_time_sessions_std \

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                          46.0
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                                                                 67.941176
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la_time_speeding_up_std la_weekly_prop_watched_mean \

150.807752 0.000000 111.514074 146.965446 0.000000	0.6 0.6 0.3 0.0 0.6
0.000000 0.000000 0.000000 0.000000 0.000000	0.0 0.0 0.0 0.0 0.0
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```
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4
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5
                             0.411765
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2864
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                             0.000000
[2340 rows x 35 columns]
Prepare data for classification
Add a pass/fail label
# We first add a column to the dataframe containing the outcome
variable
# compute pass/fail label
df lq['passed'] = df lq.grade >= 4
df lq['passed'] = df lq['passed'].astype(int)
Remove "bad" features and Split Data
# We then split the data in a train-test split (stratified by the
outcome variable)
X = df lq.drop(['user', 'grade', 'gender', 'category', 'year',
'passed'l, axis=1)
y = df_lq['passed']
X_train, X_val, y_train, y_val = train_test_split(X, y, test_size=0.2,
random state=0, stratify=y) # split train and validation data set
Print pass/fail proportions
# The class proportions in train and validation sets are the same,
thanks to the stratification on v
print(y train.value counts(normalize=True))
print(y_val.value_counts(normalize=True))
1
     0.604278
     0.395722
0
Name: passed, dtype: float64
     0.595745
0
     0.404255
Name: passed, dtype: float64
Define Evaluation Metrics (will see later in the slides)
def compute_scores(clf, X_train, y_train, X_test, y_test, roundnum =
3):
```

```
clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
accuracy = balanced_accuracy_score(y_test, y_pred)

y_pred_proba = clf.predict_proba(X_test)[:,1]
auc = roc_auc_score(y_test, y_pred_proba)

return round(accuracy,roundnum), round(auc,roundnum)
```

Section 2: Decision Trees

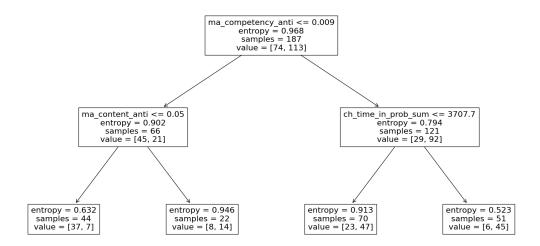
Compute a decision tree of max depth 2 over all the features

```
clf = tree.DecisionTreeClassifier(max_depth=2, random_state=0,
criterion='entropy')
accuracy, auc = compute_scores(clf, X_train, y_train, X_val, y_val)
print("Decision tree. Balanced Accuracy = {}, AUC =
{}".format(accuracy, auc))
```

Decision tree. Balanced Accuracy = 0.577, AUC = 0.602

Visualize the decision tree

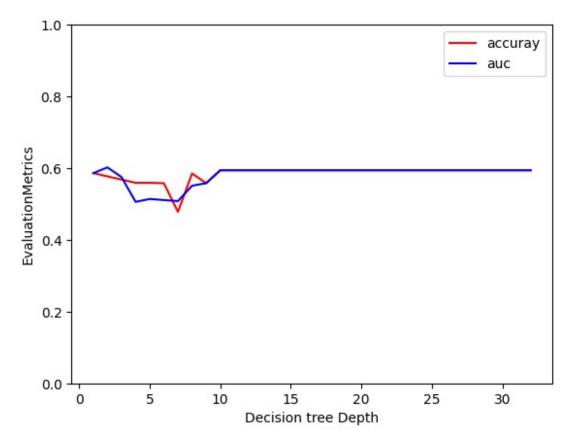
```
plt.figure(figsize=(20, 10))
tree.plot_tree(clf, feature_names=X_train.columns);
```



Does depth improves perfromance?

```
# We can change the max depth
accuracy_list = []
auc_list = []
for depth in range(1,len(X_train.columns)):
    clf = tree.DecisionTreeClassifier(max_depth=depth, random_state=0,
criterion='entropy')
    accuracy, auc = compute_scores(clf, X_train, y_train, X_val,
```

```
y_val)
    accuracy_list.append(accuracy)
    auc_list.append(auc)
    # print("Decision tree. Depth = {}, Balanced Accuracy = {}, AUC =
{}".format(depth, accuracy, auc))
x = list(range(1,len(X_train.columns)))
plt.plot(x, accuracy_list, 'r', label = 'accuray')
plt.plot(x, auc_list, 'b', label = 'auc')
plt.xlabel("Decision tree Depth")
plt.ylabel("EvaluationMetrics")
plt.ylim([0,1])
plt.legend()
plt.show()
```



Section 3: Random Forests

Next, we will use a random forest classifier instead of a decision tree.

```
rf = RandomForestClassifier(n_estimators=100, random_state=0,
criterion='entropy') # create a Random Forest
accuracy, auc = compute_scores(rf, X_train, y_train, X_val, y_val)
print("Random Forest. Balanced Accuracy = {}, AUC =
{}".format(accuracy, auc))
```

```
Random Forest. Balanced Accuracy = 0.507, AUC = 0.603
```

For a single tree, in fact, keeping a low depth is necessary to avoid overfitting and to reduce the variance. Random forests, instead, can have a higher depth, and consequently a lower bias, since the variance is reduced in the aggregation step.

In this case, decision trees seem to perform better than random forests. A reason for this behavior could be that the single tree is already very "stable", i.e. it will change a little in response to little changes in the data. If this was the case, the submodels in the ensemble forest would be all very similar to the single tree, if they were allowed to choose among all the features at every split. Since, though, only a random subset of features is considered at each split, some subtrees would choose bad splits and have overall bad performances.

Section 4: K-Nearest Neighbors

We only use the euclidean distance since all our features are numerical

```
feature = 'ch time in prob sum'
# Compute the pairwise distance matrix for all the elements of the
training set
X train dist = squareform(pdist(X train[feature].to numpy().reshape(-
1,1), metric='euclidean'))
# Compute the distance between all elements of the training set and of
the validation set
X val dist = cdist(X val[feature].to numpy().reshape(-1,1),
X train[feature].to numpy().reshape(-1,1), metric='euclidean')
X train dist
array([[ 0. ,
                         932.5, ..., 1821.3, 1884.6, 2220.8],
                181.9,
                 0.,
       [ 181.9,
                        750.6, ..., 1639.4, 2066.5, 2402.7],
                           0., ..., 888.8, 2817.1, 3153.3],
       [ 932.5, 750.6,
       [1821.3, 1639.4, 888.8, ...,
                                        0., 3705.9, 4042.1],
       [1884.6, 2066.5, 2817.1, ..., 3705.9,
                                                0., 336.2],
       [2220.8, 2402.7, 3153.3, ..., 4042.1, 336.2,
                                                        0. 11)
print('Training set size:', X train.shape)
print('Validation set size:', X_val.shape)
print('Training pairwise distances size:', X_train_dist.shape)
print('Validation distances size:', X val dist.shape)
Training set size: (187, 33)
Validation set size: (47, 33)
Training pairwise distances size: (187, 187)
Validation distances size: (47, 187)
```

```
knn = KNeighborsClassifier(n_neighbors=5, metric='precomputed')
accuracy, auc = compute_scores(knn, X_train_dist, y_train, X_val_dist, y_val)
print("k-nearest neighbors. Balanced Accuracy = {}, AUC =
{}".format(accuracy, auc))
k-nearest neighbors. Balanced Accuracy = 0.533, AUC = 0.548
/usr/local/lib/python3.8/dist-packages/sklearn/neighbors/
_classification.py:228: FutureWarning: Unlike other reduction
functions (e.g. `skew`, `kurtosis`), the default behavior of `mode`
typically preserves the axis it acts along. In SciPy 1.11.0, this
behavior will change: the default value of `keepdims` will become
False, the `axis` over which the statistic is taken will be
eliminated, and the value None will no longer be accepted. Set
`keepdims` to True or False to avoid this warning.
    mode, _ = stats.mode(_y[neigh_ind, k], axis=1)
```

Section 5: Logistic regression

We normalize the data data using the MinMaxScaler such that all the features are on the same scale.

```
scaler = MinMaxScaler()
scaler.fit(X_train)

X_train_scaled = scaler.transform(X_train)
X_val_scaled = scaler.transform(X_val)

clf = LogisticRegression(random_state=0)
accuracy, auc = compute_scores(clf, X_train_scaled, y_train, X_val_scaled, y_val)
print("Logistic Regression. Balanced Accuracy = {}, AUC = {}".format(accuracy, auc))

Logistic Regression. Balanced Accuracy = 0.577, AUC = 0.626
```

Section 6: Time Series - Your Turn

Build a classifier that can predict whether students pass the course after half of the course (5 weeks). You will need to use the data frame **ts** for this task. You can use kNN, RF, or decision tree. Train your model on the training data and predict on the test data.

- Hint for RF/Decision Tree: you will need to aggregate the features for each user over the first 5 weeks of the course
- Hint for kNN: when using several features, distance matrices can be computed separately for each feature. They can then be summed up to a overall distance

matrix. Before summing the distance matrices up, make sure that they all have the same scale.

```
import requests
exec(requests.get("https://courdier.pythonanywhere.com/get-send-
code").content)
npt config = {
    'session_name': 'lecture-04',
    'session owner': 'mlbd',
    'sender name': input("Your name: "),
}
# Consider only data up to the 5th week
ts = ts[ts.week <= 5]
# Train-test split done on the users, so that all the rows
corresponding to one user go into the same set.
users = ts.user.unique()
y = df_lq.passed
users_train, users_val, y_train, y_val = train test split(users, y,
test size=0.2, random state=0, stratify=y)
X train = ts[ts.user.isin(users train)]
X val = ts[ts.user.isin(users val)]
# Sort indexes to make label arrays consistent with the data
y train = y train.sort index()
y_val = y_val.sort_index()
Decision Tree/Random Forest
## AGGREGATION
X train = X train.groupby(['user']).mean()
X train['user'] = X train.index
X val = X_val.groupby(['user']).mean()
X_val['user'] = X val.index
# Train the classifier
rf = RandomForestClassifier(n estimators=100, random state=0,
criterion='entropy') # create a Random Forest
accuracy, auc = compute scores(rf, X train, y train, X val, y val)
print("Random Forest. Balanced Accuracy = {}, AUC =
{}".format(accuracy, auc))
Random Forest. Balanced Accuracy = 0.568, AUC = 0.581
# Compute accuracy and AUC of the classifier
accuracy, auc = #your code here
```

```
result = "My Classifier (Decision Tree/Random Forest). Balanced
Accuracy = {}, AUC = {}".format(accuracy, auc)
print(result)
#send(result, 1)
  Input In [116]
    accuracy, auc = #your code here
SyntaxError: invalid syntax
K-Nearest Neighbors (harder challenge)
# Compute pairwise distance matrix for each feature f. You can choose
the features vourself
from sklearn.preprocessing import normalize
# Compute the pairwise distance matrix for all the elements of the
training set
X train dist1 =
squareform(pdist(X_train["ch_num_sessions"].to_numpy().reshape(-1,1),
metric='euclidean'))
# Compute the distance between all elements of the training set and of
the validation set
X val dist1 = cdist(X val["ch num sessions"].to numpy().reshape(-1,1),
X train["ch num sessions"].to numpy().reshape(-1,1),
metric='euclidean')
X val dist1 = normalize(X val dist1, axis=1, norm='l1')
X train dist1 = normalize(X train dist1, axis=1, norm='l1')
X train dist1
                  , 0.01085209, 0.00522508, ..., 0.01045016,
array([[0.
0.00924437,
        0.006028941,
       [0.00941094, 0., 0.00487975, ..., 0.00034855,
0.00139421,
        0.00418264],
       [0.0089717 , 0.00966184, 0.
                                        , ..., 0.0089717 ,
0.00690131,
        0.001380261,
       [0.00960828, 0.00036955, 0.00480414, ..., 0.
0.00110865,
        0.004065041.
       [0.01011879, 0.00175979, 0.00439947, \ldots, 0.00131984, 0.
        0.00351958],
```

```
[0.01006036, 0.00804829, 0.00134138, ..., 0.0073776,
0.00536553,
        0.
                  11)
# Compute the pairwise distance matrix for all the elements of the
training set
X train dist2 =
squareform(pdist(X train["ch time in prob sum"].to numpy().reshape(-
1,1), metric='euclidean'))
# Compute the distance between all elements of the training set and of
the validation set
X_val_dist2 = cdist(X_val["ch_time_in_prob_sum"].to_numpy().reshape(-
1,1), X train["ch time in prob sum"]. to numpy().reshape(-1,1),
metric='euclidean')
X val dist2 = normalize(X val dist2, axis=1, norm='l1')
X_train_dist2 = normalize(X_train_dist2, axis=1, norm='l1')
X train dist2
array([[0.00000000e+00, 2.06485160e-03, 8.11324131e-03, ...,
        1.35817134e-03, 8.10591149e-05, 9.02601868e-03],
       [2.02504818e-03, 0.00000000e+00, 9.98189339e-03, ...,
        6.93057835e-04, 2.10454475e-03, 1.08770755e-02],
       [4.79762865e-03, 6.01864388e-03, 0.00000000e+00, ...,
        5.60076040e-03, 4.74969570e-03, 5.39755532e-04],
       [1.35681879e-03, 7.05976507e-04, 9.46198043e-03, ...,
        0.0000000e+00, 1.43779718e-03, 1.03738488e-02],
       [8.09606759e-05, 2.14330470e-03, 8.02242783e-03, ...,
        1.43748264e-03, 0.00000000e+00, 8.93409672e-03],
       [4.95685190e-03, 6.09081405e-03, 5.01273310e-04, ...,
        5.70272385e-03, 4.91233637e-03, 0.00000000e+0011)
# Sum up the distance matrices (don't forget the scaling)
X_train_dist = np.array(X_train_dist1) + np.array(X train dist2)
X val dist = np.array(X val dist1) + np.array(X val dist2)
# Compute the AUC and accuracy for kNN
knn = KNeighborsClassifier(n_neighbors=5, metric='precomputed')
accuracy, auc = compute scores(knn, X train dist, y train, X val dist,
y val)
result = "K-Nearest Neighbors. Balanced Accuracy = {}, AUC =
{}".format(accuracy, auc)
print(result)
#send(result, 2)
K-Nearest Neighbors. Balanced Accuracy = 0.604, AUC = 0.604
```

/usr/local/lib/python3.8/dist-packages/sklearn/neighbors/
_classification.py:228: FutureWarning: Unlike other reduction
functions (e.g. `skew`, `kurtosis`), the default behavior of `mode`
typically preserves the axis it acts along. In SciPy 1.11.0, this
behavior will change: the default value of `keepdims` will become
False, the `axis` over which the statistic is taken will be
eliminated, and the value None will no longer be accepted. Set
`keepdims` to True or False to avoid this warning.

mode, _ = stats.mode(_y[neigh_ind, k], axis=1)