

# Project 2: Fire in the nature park

Intelligent Data Analysis & Machine Learning I

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# **Data Preperation**

```
import numpy as np
import pandas as pd

original_df = pd.read_csv('fires.csv',sep=',',header=0)
header = original_df.columns
print(header)
original_df.head()
```

Index(['X', 'Y',	'month', '	day', 'F	FMC', 'DI	чс', 'DC'	', 'ISI',	'temp',	'RH',
'wind', 'r	ain', 'are	a'],					
dtype='obje	ct')						

	Х	Υ	month	day	FFMC	DMC	DC	ISI	temp	RH	wind	rain	area
0	7	5	mar	fri	86.2	26.2	94.3	5.1	8.2	51	6.7	0.0	0.0
1	7	4	oct	tue	90.6	35.4	669.1	6.7	18.0	33	0.9	0.0	0.0
2	7	4	oct	sat	90.6	43.7	686.9	6.7	14.6	33	1.3	0.0	0.0
3	8	6	mar	fri	91.7	33.3	77.5	9.0	8.3	97	4.0	0.2	0.0
4	8	6	mar	sun	89.3	51.3	102.2	9.6	11.4	99	1.8	0.0	0.0

• Fire data: 13 Features

• Two categorical features: 'month' and 'day'

• Objective: prediction of the brunt forest 'area'

# **Data Preprocessing**

```
from sklearn.preprocessing import LabelEncoder
# transform the 'area' column to log(area+1)
df = original_df.copy()
original_area = df['area']
df['area'] = np.log(df['area']+1)
print(df)
num df = df.copy()
label_encoders = {} # Dictionary initialization
for column in ['month', 'day']:
    label encoders[column] = LabelEncoder()
    num df[column] = label encoders[column].fit transform(df[column])
    # Convert each string to an integer. Days of the week and months are
    automatically converted to integers.
```

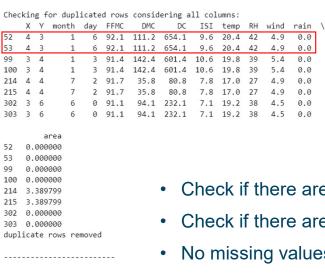
```
DC ISI temp RH
                       35.4 669.1
                                  6.7 18.0 33
                       43.7 686.9
                                6.7 14.6 33
                       33.3 77.5 9.0
                                      8.3 97 4.0
                       51.3 102.2 9.6 11.4 99
                      56.7 665.6 1.9 27.8 32 2.7 0.0 2.006871
             sun 81.6
         aug sun 81.6
                      56.7 665.6 1.9 21.9 71 5.8 0.0 4.012592
                      56.7 665.6 1.9 21.2 70
         aug sat 94.4 146.0 614.7 11.3 25.6 42 4.0 0.0 0.000000
516 6 3 nov tue 79.5
                       3.0 106.7 1.1 11.8 31 4.5 0.0 0.000000
```

[517 rows x 13 columns]

- Log-transformation of feature 'area'
- Two categorical features are transformed to integer for further process
- `LabelEncoder()` maps an unique integer value to categorical value

# **Data Preprocessing**

```
print("Checking for duplicated rows considering all columns:")
duplicates = num_df[num_df.duplicated(keep=False)]
if not duplicates.empty:
    print(duplicates)
else:
   print("No duplicated data found when considering all columns.")
num df = num df.drop duplicates()
print("duplicate rows removed")
print("\n----\n")
# to check if there is any missing value in the data
print(df.isnull().sum())
# no missing data
```



wind

dtype: int64

- Check if there are any missing values
- Check if there are any duplicated values
- No missing values
- Duplicated values are removed

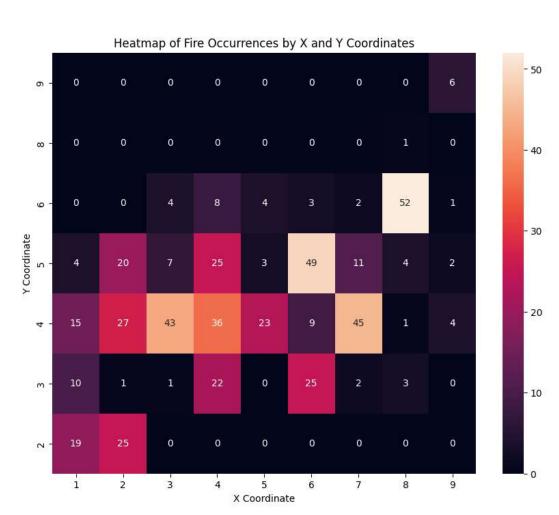
```
import matplotlib.pyplot as plt
                                                                                 Histogram representation using pyplot
num_features = len(header)
                                                                                  library
fig, axes = plt.subplots(nrows=(num_features + 2) // 3, ncols=3, figsize=
(15, 5 * ((num features + 2) // 3)))
axes = axes.flatten() # Convert the axes in the form of a two-dimensional
array to a one-dimensional array.
                       # This makes it convenient to use a for loop.
for i, col in enumerate(df.columns):
   axes[i].hist(df[col], bins=30, color='blue', alpha=0.7)
   axes[i].set_title(f'Histogram of {col}')
    axes[i].set_xlabel(col)
   axes[i].set_ylabel('Frequency')
for i in range(len(df.columns), len(axes)):
   fig.delaxes(axes[i])
# Adjusting the subplot layout
plt.tight layout()
plt.show()
```

```
import matplotlib.pyplot as plt
                                                                                  Histogram representation using pyplot
num_features = len(header)
                                                                                   library
fig, axes = plt.subplots(nrows=(num_features + 2) // 3, ncols=3, figsize=
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    axes[i].set_xlabel(col)
    axes[i].set_ylabel('Frequency')
for i in range(len(df.columns), len(axes)):
    fig.delaxes(axes[i])
# Adjusting the subplot layout
plt.tight layout()
plt.show()
                                                                                                                   Histogram of area
```

```
# Create a heatmap using X, Y coordinates
heatmap_data = df.pivot_table(index='Y', columns='X', values='area',
aggfunc='count', fill_value=0)
# The pivot_table function reorganizes a data frame to create a new table.

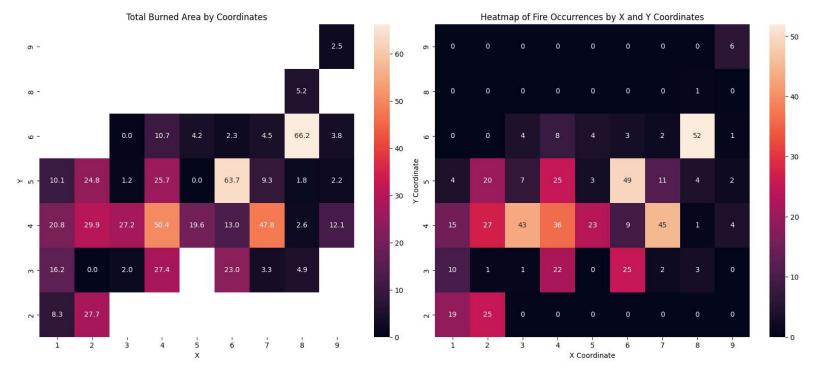
plt.figure(figsize=(10, 8))
sns.heatmap(heatmap_data, annot=True, cbar=True) # Create a heatmap using the
pivot table we created earlier.
plt.gca().invert_yaxis()
plt.title('Heatmap of Fire Occurrences by X and Y Coordinates')
plt.xlabel('X Coordinate')
plt.ylabel('Y Coordinate')
plt.show()
```

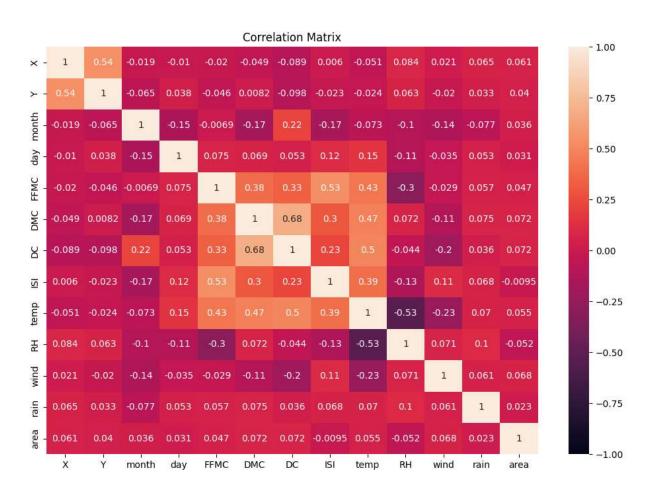
- Find relationship between coordinate and fire occurrences: location specificity
- The most frequent fires occur in (8,6), (6,5), (7,4), and (3,4) (n>40).
- Problem: most of value of 'area' are very low
- There might not be a relationship between location and 'area'



```
area_sum_by_coordinates = df.groupby(['X', 'Y'])['area'].sum().reset_index()
pivot_table = area_sum_by_coordinates.pivot(index='Y', columns='X', values='area')
plt.figure(figsize=(10, 8))
sns.heatmap(pivot_table, annot=True, fmt=".1f")
plt.gca().invert_yaxis()
plt.title('Total Burned Area by Coordinates')
plt.show()
```

 There seems to be a correlation between occurrence and sum of 'area', but low

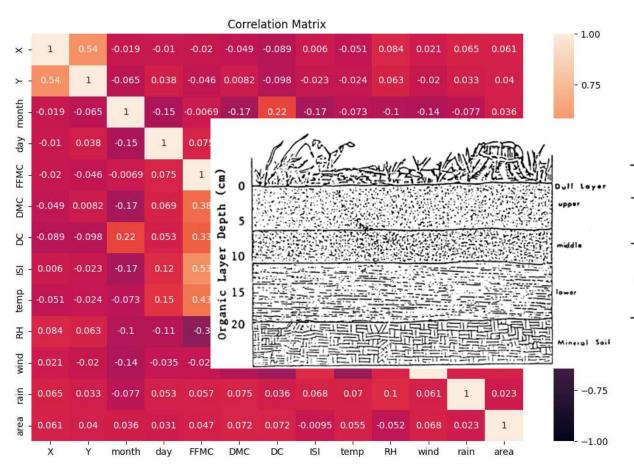




```
# Calculate correlation matrix
corr_matrix = num_df.corr()

plt.figure(figsize=(12, 8))
sns.heatmap(corr_matrix, annot=True, vmin=-1, vmax=1)
plt.title('Correlation Matrix')
plt.show()
```

- DMC and DC show the highest correlations
- These two show a relatively low correlation with FFMC
- Temperature has a strong correlation with these three features, and RH



<pre># Calculate correlation matrix corr_matrix = num_df.corr()</pre>
<pre>plt.figure(figsize=(12, 8)) sns.heatmap(corr_matrix, annot=True, vmin=-1, vmax=1) slt_title('Correlation Matrix')</pre>

Weight	Fuel Moisture Code					
5 t/ha	FFMC	the highest				
50 t/ha	DMC	relatively low				
440 t/ha	DC	strong correlation with s, and RH				

### **Model Selection**

### Random Forest

- Linear Regression, Decision Tree, SVM, etc.. Many options
- Our data have only 13 features -> no reason to use SVM
- Our data set is small: only ca. 500 size
- Decision Tree has a higher risk of overfitting than Random Forest
- Stojanova et al. (2012) compared and evaluated models such as KNN, DT, LR and SVM for prediction of forest fires
- RF showed the highest performance

### Random Forest

```
from sklearn.model_selection import train_test_split
  from sklearn.metrics import mean_squared_error
  from sklearn.ensemble import RandomForestRegressor
  X = num_df.drop('area', axis=1)
  y = num_df['area']
  # split the data into training and test sets
  X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
  random_state=42)
  # Random Forest
  RFmodel = RandomForestRegressor(random_state=42)
  RFmodel.fit(X_train, y_train)
  # prediction
  RF_pred_test = RFmodel.predict(X_test)
  RF_pred_train = RFmodel.predict(X_train)
  # evaluation
  RF_mse = mean_squared_error(y_test, RF_pred_test)
  RF mse train = mean squared error(y train, RF pred train)
  results = {
      "Metric": ["Mean Squared Error"],
      "Train": [RF_mse_train],
      "Test": [RF_mse]
  results_df = pd.DataFrame(results)
  print(results_df)
√ 0.1s
```

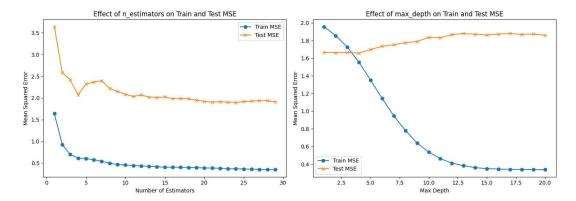
Metric Train Test
0 Mean Squared Error 0.337283 1.869937

- A large difference in MSE between the training set and the test set
- It might be caused by overfitting!

### Parameter modification

```
# Evaluate model performance by changing n estimators. Use n estimators values
between 1 and 30.
n_estimators_range = range(1, 30)
train mse values = []
test mse values = []
for n_estimators in n_estimators_range:
   model = RandomForestRegressor(n estimators=n estimators, random state=42)
    model.fit(X train, y train)
    train_pred = model.predict(X_train)
    test_pred = model.predict(X_test)
    train mse = mean squared error(y train, train pred)
    test mse = mean squared error(y test, test pred)
    train mse values.append(train mse)
    test mse values.append(test mse)
# Evaluate model performance according to changes in max depth. Use max depth
values between 1 and 20.
max_depth_range = range(1, 21)
train accuracies depth = []
test_accuracies_depth = []
for max_depth in max_depth_range:
   model = RandomForestRegressor(max_depth=max_depth, random_state=42)
    model.fit(X train, y train)
   train pred = model.predict(X train)
    test_pred = model.predict(X_test)
    train mse = mean squared error(y train, train pred)
    test_mse = mean_squared_error(y_test, test_pred)
    train accuracies depth.append(train mse)
    test accuracies depth.append(test mse)
```

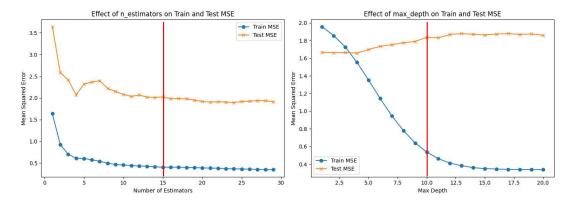
```
# Visualization
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(14, 5))
# First graph: MSE as a function of n estimators
axes[0].plot(n estimators range, train mse values, label='Train MSE', marker='o')
axes[0].plot(n_estimators_range, test_mse_values, label='Test_MSE', marker='x')
axes[0].set_xlabel('Number of Estimators')
axes[0].set_ylabel('Mean Squared Error')
axes[0].set_title('Effect of n_estimators on Train and Test MSE')
axes[0].legend()
# Second graph: MSE according to max depth
axes[1].plot(max_depth_range, train_accuracies_depth, label='Train MSE',
marker='o')
axes[1].plot(max depth range, test accuracies depth, label='Test MSE',
marker='x')
axes[1].set_xlabel('Max Depth')
axes[1].set ylabel('Mean Squared Error')
axes[1].set_title('Effect of max_depth on Train and Test MSE')
axes[1].legend()
plt.tight_layout()
plt.show()
```



### Parameter modification

```
# Evaluate model performance by changing n estimators. Use n estimators values
between 1 and 30.
n_estimators_range = range(1, 30)
train mse values = []
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for n_estimators in n_estimators_range:
   model = RandomForestRegressor(n estimators=n estimators, random state=42)
    model.fit(X train, y train)
    train_pred = model.predict(X_train)
    test_pred = model.predict(X_test)
    train mse = mean squared error(y train, train pred)
    test mse = mean squared error(y test, test pred)
    train mse values.append(train mse)
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for max_depth in max_depth_range:
   model = RandomForestRegressor(max_depth=max_depth, random_state=42)
    model.fit(X train, y train)
   train pred = model.predict(X train)
    test_pred = model.predict(X_test)
    train mse = mean squared error(y train, train pred)
    test_mse = mean_squared_error(y_test, test_pred)
    train accuracies depth.append(train mse)
    test accuracies depth.append(test mse)
```

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# First graph: MSE as a function of n estimators
axes[0].plot(n estimators range, train mse values, label='Train MSE', marker='o')
axes[0].plot(n_estimators_range, test_mse_values, label='Test_MSE', marker='x')
axes[0].set_xlabel('Number of Estimators')
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axes[0].set_title('Effect of n_estimators on Train and Test MSE')
axes[0].legend()
# Second graph: MSE according to max depth
axes[1].plot(max_depth_range, train_accuracies_depth, label='Train MSE',
marker='o')
axes[1].plot(max depth range, test accuracies depth, label='Test MSE',
marker='x')
axes[1].set_xlabel('Max Depth')
axes[1].set ylabel('Mean Squared Error')
axes[1].set_title('Effect of max_depth on Train and Test MSE')
axes[1].legend()
plt.tight_layout()
plt.show()
```



### **Evaluation**

```
from sklearn.model_selection import cross_val_score, KFold
from sklearn.metrics import make_scorer, r2_score
from sklearn.dummy import DummyRegressor
# K-fold settings
kf = KFold(n splits=5, shuffle=True, random state=42)
# Model setup
RFmodel = RandomForestRegressor(n_estimators= 15, max_depth=10, random_state=42)
baseline_mean = DummyRegressor(strategy='mean')
baseline_median = DummyRegressor(strategy='median')
# Setting up MSE scorer for performance evaluation
mse_scorer = make_scorer(mean_squared_error)
r2 scorer = make scorer(r2 score)
# Perform Random Forest cross validation
mse_scores_rf = cross_val_score(RFmodel, X, y, cv=kf, scoring=mse_scorer)
r2_scores_rf = cross_val_score(RFmodel, X, y, cv=kf, scoring=r2_scorer)
# Perform Dummy Regressor cross validation
mse_scores_dummy_mean = cross_val_score(baseline_mean, X, y, cv=kf,
scoring=mse_scorer)
r2 scores dummy mean = cross val score(baseline mean, X, y, cv=kf,
scoring=r2 scorer)
mse_scores_dummy_median = cross_val_score(baseline_median, X, y, cv=kf,
scoring=mse scorer)
r2_scores_dummy_median = cross_val_score(baseline_median, X, y, cv=kf,
scoring=r2 scorer)
# Output
comparision = {
   "" : ["MSE", "r2_score"],
    "Random_Forest": [np.mean(mse_scores_rf), np.mean(r2_scores_rf)],
   "Baseline_Mean": [np.mean(mse_scores_dummy_mean), np.mean
   (r2_scores_dummy_mean)],
   "Baseline_Median": [np.mean(mse_scores_dummy_median), np.mean
   (r2 scores dummy median)]
print(round(pd.DataFrame(comparision),2))
```

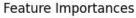
```
Random_Forest Baseline_Mean Baseline_Median
0 MSE 2.29 1.96 2.45
1 r2_score -0.19 -0.01 -0.25
```

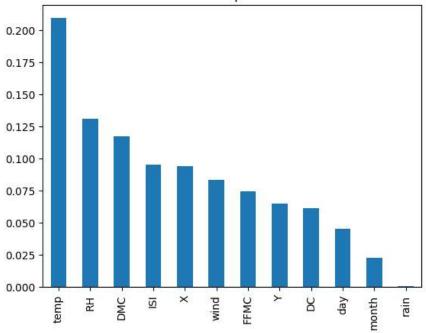
- Baseline Model: Dummy Regressor for prediction of mean and median
- Evaluation using MSE and r2 score
- MSE(RF) < MSE(dummy\_mean) & negative r2 score
- Our model doesn't showing good performance

# Feature Importances

```
RFmodel.fit(X_train, y_train)

feature_importances = RFmodel.feature_importances_
RF_importances = pd.Series(feature_importances, index=X.columns)
RF_importances = RF_importances.sort_values(ascending=False)
RF_importances.plot(kind='bar')
plt.title('Feature Importances')
plt.show()
```





- The 3 most important role in prediction were `temp`, `RH`, and `DMC`
- Remarkable contribution of `temp`
- It is assumed to be due to high correlation with other important features.
- Very low contribution of month and rain
- Most of the data is distributed in a narrow range
- Does not play a significant role in prediction

## Additional Analysis

MSE

1 r2 score

2.29

-0.19

```
from sklearn.preprocessing import StandardScaler
from sklearn.linear_model import LinearRegression
# normalization for linear regression
scaler = StandardScaler()
numeric_features = ['X', 'Y', 'FFMC', 'DMC', 'DC', 'ISI', 'temp', 'RH', 'wind',
'rain', 'area']
num df[numeric features] = scaler.fit transform(num df[numeric features])
LRmodel = LinearRegression()
mse scores lr = cross val score(RFmodel, X, y, cv=kf, scoring=mse scorer)
r2 scores lr = cross val score(RFmodel, X, y, cv=kf, scoring=r2 scorer)
additional comparision = {
    "" : ["MSE", "r2_score"],
    "Random Forest": [np.mean(mse scores rf), np.mean(r2 scores rf)],
    "Baseline Mean": [np.mean(mse scores dummy mean), np.mean
    (r2 scores dummy mean)],
    "Baseline Median": [np.mean(mse scores dummy median), np.mean
    (r2_scores_dummy_median)],
    "Linear_Regression": [np.mean(mse_scores_lr), np.mean(r2_scores_lr)]
print(round(pd.DataFrame(additional_comparision),2))
         Random_Forest Baseline_Mean Baseline_Median Linear_Regression
```

1.96

-0.01

2.45

-0.25

2.29

-0.19

- In case our RF model does not fit the data well, Linear regression is also tested
- For Linear Regression, normalization is needed
- · Here, z-score normalization is used
- It shows exactly same bad performance

