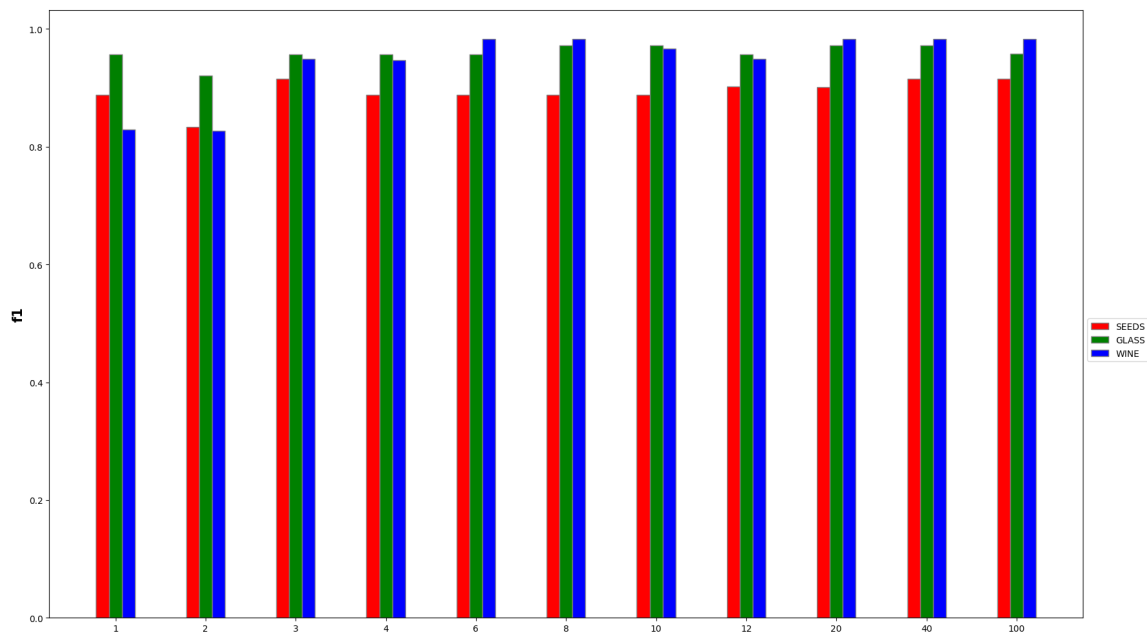


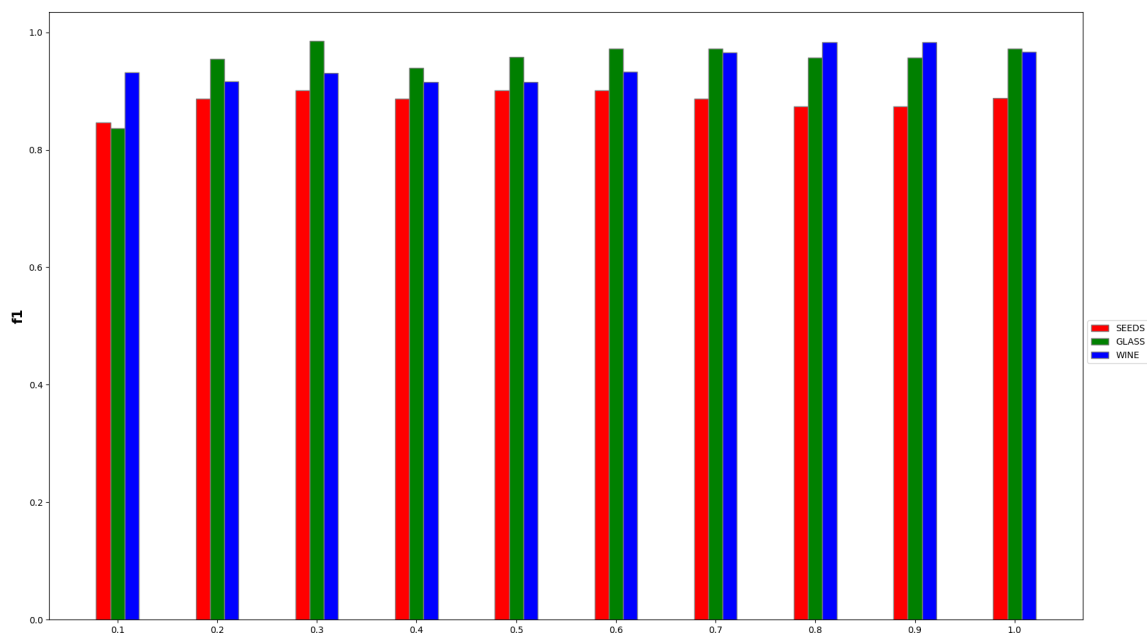
Wnioski

Bagging

Badając wpływ liczby klasyfikatorów. Ze wzrostem tego parametru wzrasta f1-score, jednak optymalna wartość jest w środku zakresu badanych wartości. Zbiór GLASS się nie zmienia, bo ten zbiór jest specyficzny.



Max samples też ogólnie polepsza wyniki, ale dla co datasetu możemy wyznaczyć optymalną wartość, jednak możliwie to spowodowane wadanie wartości podanej jako %.

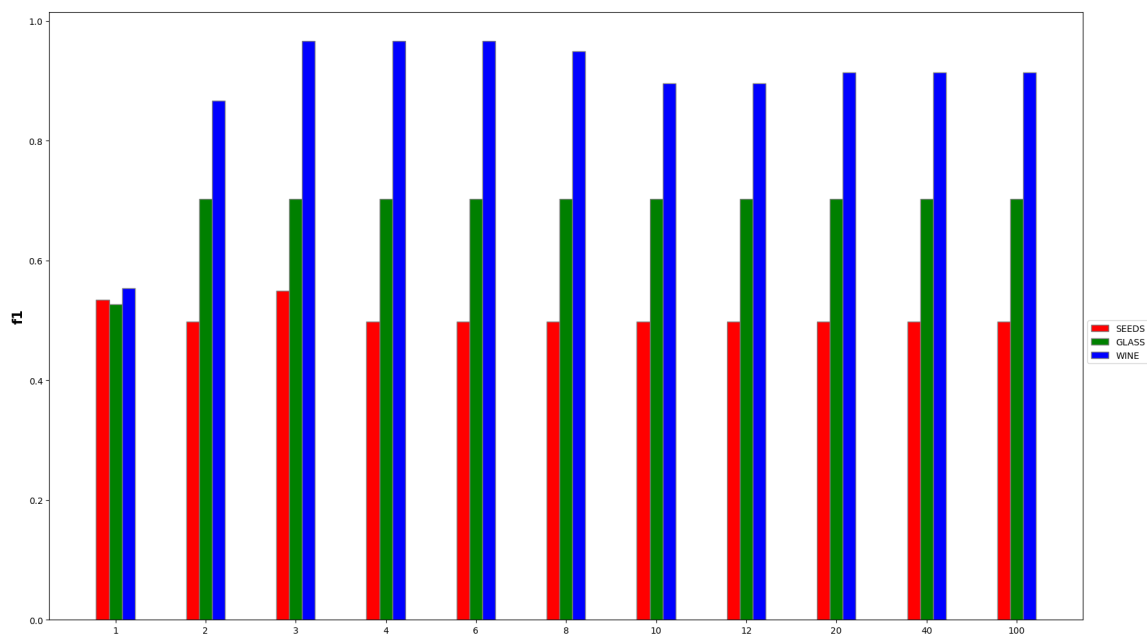


Liczba atrybutów daje gorsze wyniki dla małych wartości, ale optymalne wartości są w środku zakresu.

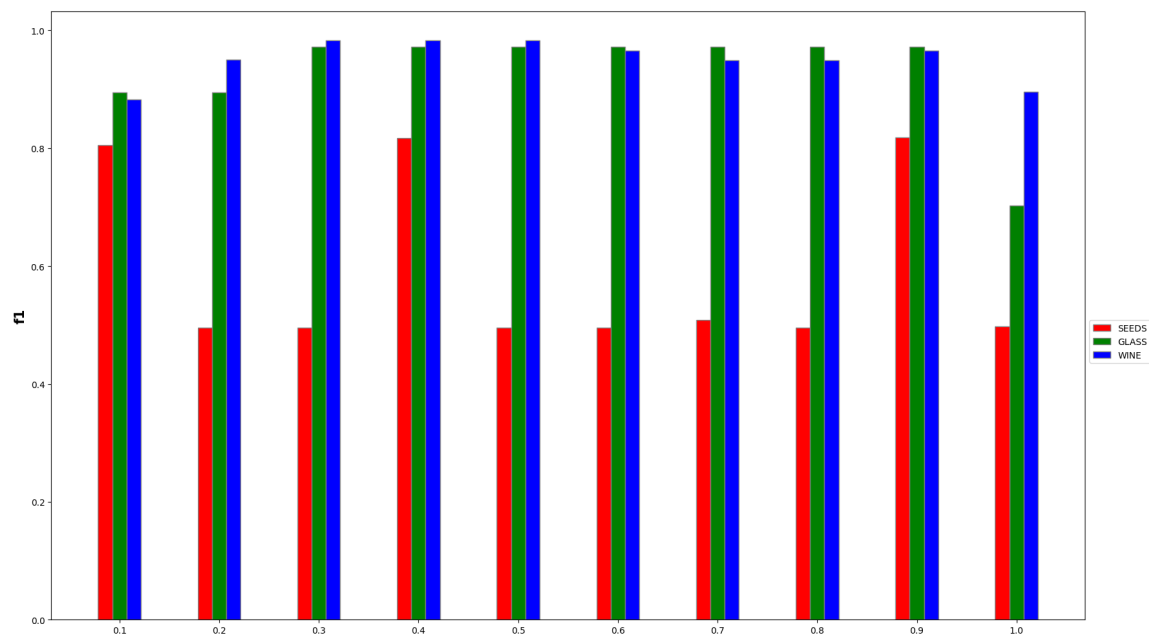
Zmiany spowodowane różnymi wartościami bootstrap nie są znaczące.

AdaBoost

W przypadku liczby klasyfikatorów tutaj też możemy zobaczyć, że wartości w środku, 2,4,6, dają same optymalne wyniki dla WINE. GLASS robi swoje rzeczy, a SEEDS się nie udało polepszyć.

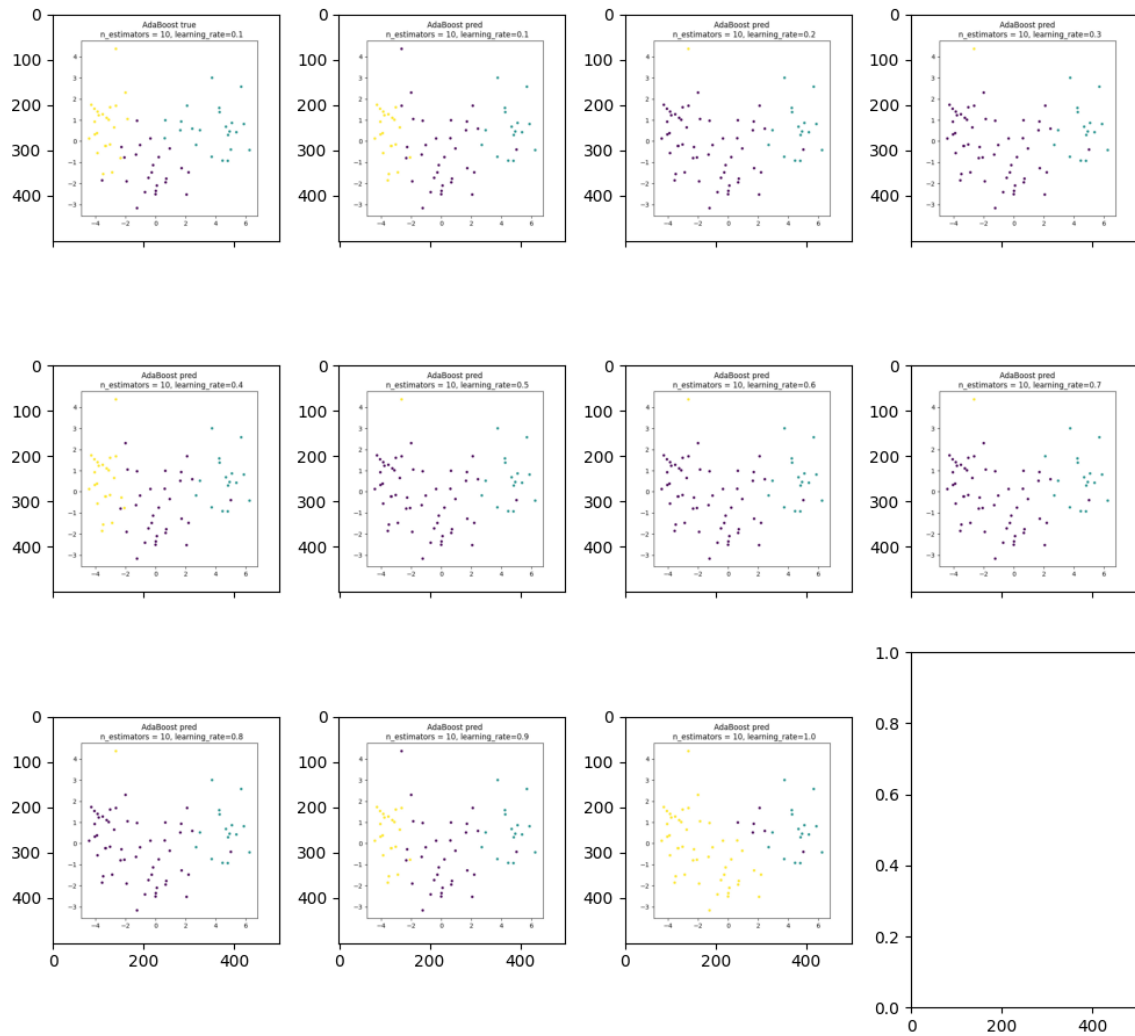


Badając współczynnik uczeni dla zbiorów GLASS i WINE nic dziwnego się nie zdażyło, niektóre wartości lepsze, niektóre gorsze, jednak SEEDS zachowywał się dziwno



Aby zrozumieć, co się dzieje, popatrzyłem na wizualizację

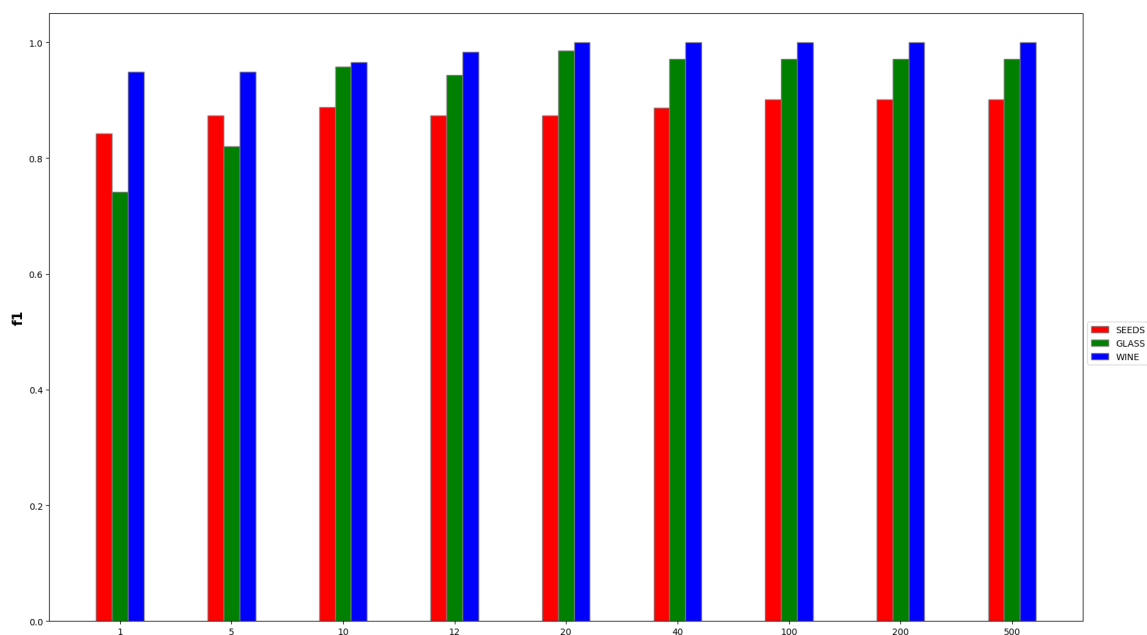
SEEDS



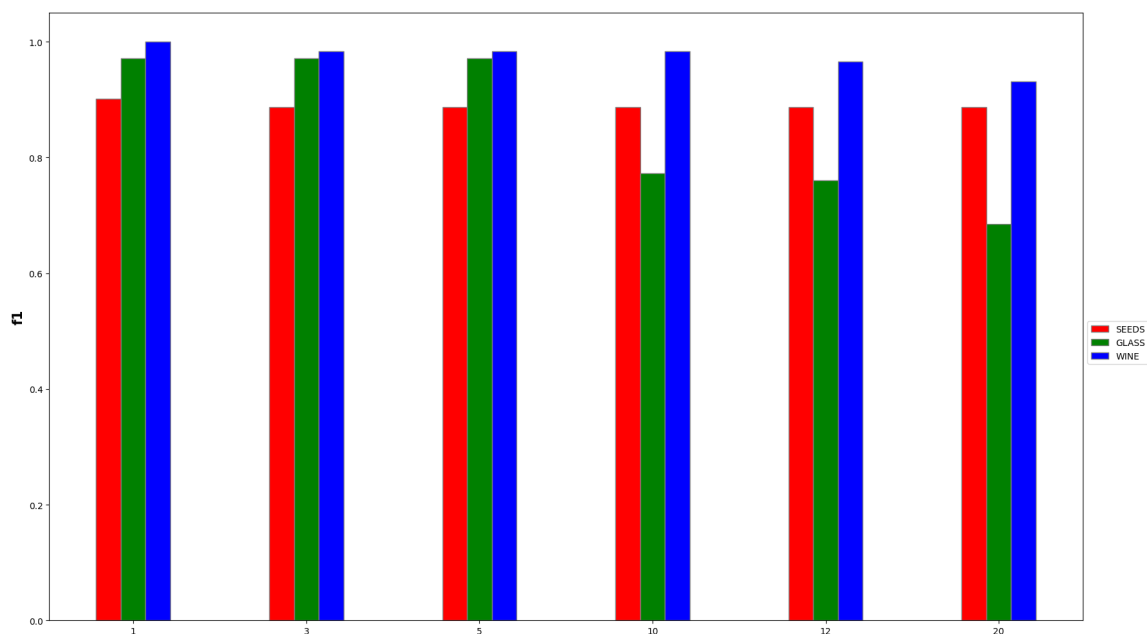
Widać, że dla wartości oprócz 0.1, 0.4 i 0.9 adaboost pracuje bardzo średnio i nie ma nawet wszystkich trzech klas.

Random forest

Nic ciekawego z liczbą drzew, można znaleźć optymalną wartość dla co zbioru



Duża wartość liczby samplów pogorszyła rezultat dla GLASS, ale pozostałe zbiory mało odczuły zmiany.



Grając z liczbą cech można polepszyć wynik, ale rezultat jest +- taki sam dla wszystkich zbiorów

Głębokość też nie dała duże zmiany

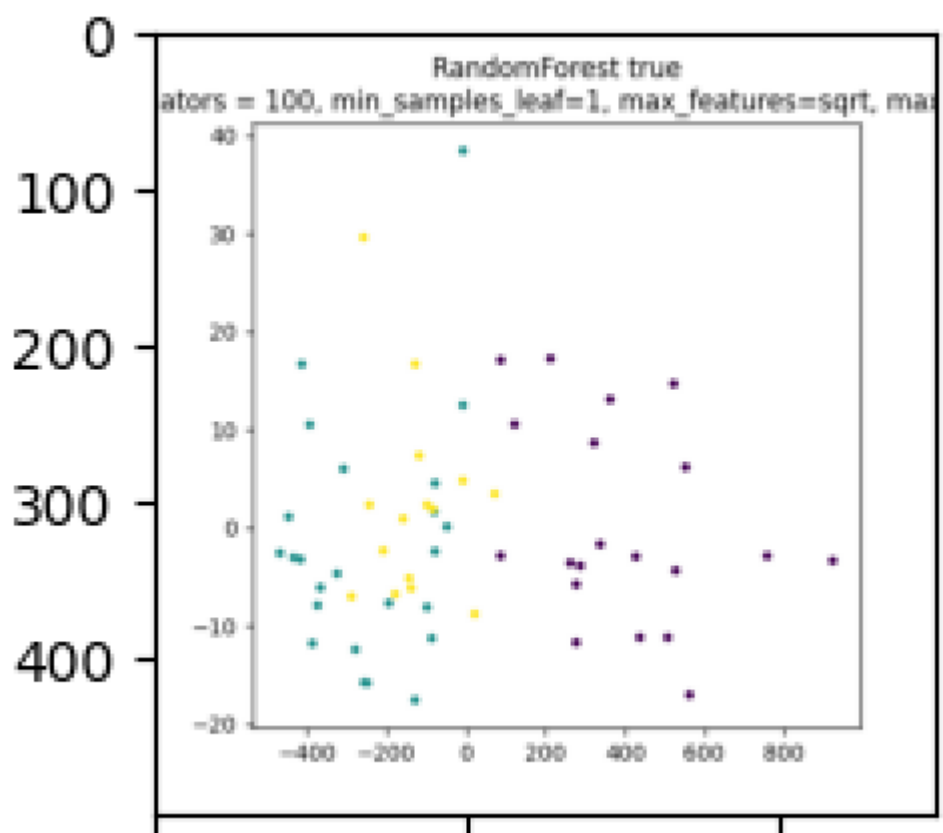
Porównanie z klasyfikatorem bazowym

Sprawdzimy wartości f1 dla parametrów domyślnych tech klasyfikatorów

	f1 KNN	f1 bagging	f1 boost	f1 forest
IRIS	0.980000	1.000000	0.878487	0.980000
SEEDS	0.885714	0.887773	0.498168	0.901039
GLASS	0.957746	0.971702	0.702701	0.971702
WINE	0.677966	0.966270	0.895931	1.000000

IRIS, SEEDS i GLASS powodują sobie podobnie: wszystkie metody ensemble oprócz boosting dają lepsze wyniki.

Ale najciekawsze są wyniki dla WINE, gdzie ensemble dają znacznie lepsze wyniki. Zobaczmy jak wygląda ten dataset.



I od razu widać dlaczego, bo dane tutaj są zmieszane (dla 2 klas). A metody ensemble, ogólna idea których to wykorzystanie kilku klasyfikatorów aby jedne klasyfikatory negocjowały błędy innych klasyfikatorów, jest w stanie złapać takie struktury. w odróżnieniu od klasyfikatorów bazowych

Część techniczna

```
In [ ]: import matplotlib.pyplot as plt
import matplotlib as matplotlib

# unused but required import for doing 3d projections with matplotlib < 3.2
import mpl_toolkits.mplot3d # noqa: F401
import numpy
import pandas as pd
from sklearn import datasets
from sklearn.decomposition import PCA
from sklearn.model_selection import train_test_split
from scipy.io import arff
```

```
In [ ]: iris = datasets.load_iris()
df_iris = pd.DataFrame(iris.data, columns=['sepal length', 'sepal width', 'petal le
df_iris = pd.concat([df_iris, pd.DataFrame(iris.target, columns=['name'])], axis=1
```

```
In [ ]: import requests

data_seeds_raw = requests.get('https://archive.ics.uci.edu/ml/machine-learning-data
data_seeds = ''
for data in (data_seeds_raw.iter_content()):
    data_seeds = data_seeds + data.decode("utf-8")

data_seeds_split = data_seeds.split('\n')
data_seeds = []

for x in data_seeds_split:
    if x != '':
        data_seeds.append([float(xx) for xx in x.split('\t') if xx != ''])

data_seeds = numpy.array(data_seeds)

df_seeds = pd.DataFrame(data_seeds)

df_seeds = df_seeds.astype({7: int})
df_seeds = df_seeds.rename(columns = {7: 'name'})
df_seeds.columns = df_seeds.columns.astype(str)
```

```
In [ ]: import requests

data_glass_raw = requests.get('https://archive.ics.uci.edu/ml/machine-learning-data

data_glass = ''
for data in (data_glass_raw.iter_content()):
    data_glass = data_glass + data.decode("utf-8")
data_glass_split = data_glass.split('\n')
data_glass = []
for x in data_glass_split:
    if x != '':
        data_glass.append([float(xx) for xx in x.split(',')])

data_glass = numpy.array(data_glass)

df_glass = pd.DataFrame(data_glass)

df_glass = df_glass.astype({10: int})
df_glass = df_glass.rename(columns = {10: 'name'})
df_glass.columns = df_glass.columns.astype(str)
```

```
In [ ]: dict_attr_wine = {
    0 : 'Alcohol',
    1 : 'Malic acid',
    2 : 'Ash',
    3 : 'Alcalinity of ash' ,
    4 : 'Magnesium',
    5 : 'Total phenols',
    6 : 'Flavanoids',
    7 : 'Nonflavanoid phenols',
    8 : 'Proanthocyanins',
    9 : 'Color intensity',
    10: 'Hue',
    11: 'OD280/OD315 of diluted wines',
    12: 'Proline'
}

wine = datasets.load_wine()
df_wine = pd.DataFrame(wine.data, columns=dict_attr_wine.values())
df_wine = pd.concat([df_wine, pd.DataFrame(wine.target, columns=['name'])], axis=1)
```



```

In [ ]: import math
import numpy as np
def plotStatistics(df, metrics, col_names, dataset_names, colours):
    size_height = math.ceil(len(metrics)/2)
    size_width = math.ceil((len(metrics))/size_height)
    fig, axes = plt.subplots(size_height, size_width, sharex=True, sharey=False, fi

    for i in range(len(metrics)):
        col_id = i % size_width
        row_id = i // size_width

        y = [[exp_data[metrics[i]] for exp_data in ds_data ] for ds_data in df]

        barWidth = 1 / (len(y) + 4)

        brs = []
        brs.append(np.arange(len(y[0])))

        for j in range(1, len(y)):
            brs.append([x + barWidth for x in brs[j-1]])

        for j in range(len(y)):
            if len(metrics) == 1:
                axes.bar(brs[j], y[j], color = colours[j], width = barWidth,
                        edgecolor = 'grey', label =dataset_names[j])
            elif size_height > 1:
                axes[row_id][col_id].bar(brs[j], y[j], color = colours[j], width =
                        edgecolor = 'grey', label =dataset_names[j])
            else:
                axes[col_id].bar(brs[j], y[j], color = colours[j], width = barWidth
                        edgecolor = 'grey', label =dataset_names[j])

        # axes[row_id][col_id].set_xlabel('Value', fontweight = 'bold', fontsize = 1
        if len(metrics) == 1:
            axes.set_ylabel(metrics[i], fontweight = 'bold', fontsize = 15)
            axes.set_xticks([r + barWidth for r in range(len(y[0]))],
                            col_names)

            axes.legend(bbox_to_anchor=(1, 0.5), fancybox=True)
        elif size_height > 1:
            axes[row_id][col_id].set_ylabel(metrics[i], fontweight = 'bold', fontsiz
            axes[row_id][col_id].set_xticks([r + barWidth for r in range(len(y[0]))]
                            col_names)

            axes[row_id][col_id].legend(bbox_to_anchor=(1, 0.5), fancybox=True)
        else:
            axes[col_id].set_ylabel(metrics[i], fontweight = 'bold', fontsize = 15)
            axes[col_id].set_xticks([r + barWidth for r in range(len(y[0]))],
                            col_names)

            axes[col_id].legend(bbox_to_anchor=(1, 0.5), fancybox=True)

plt.tight_layout()

```

```
# plotStatistics(results_list_criterion, ['silhouette_score', 'davies_bouldin_score
```

```
In [ ]: from io import BytesIO

def combineFigsInOnePlot(dict_list, plot_name = ''):
    size_width = min(4, len(dict_list) + 1)
    size_height = math.ceil((len(dict_list) + 1)/size_width)
    fig, axes = plt.subplots( size_height, size_width, sharex=True, sharey=False, f

    buffer_tru = BytesIO()
    dict_list[0]['true_fig'].savefig(buffer_tru, format='png')
    buffer_tru.seek(0)
    image_true_data = plt.imread(buffer_tru)

    if size_height > 1:
        axes[0][0].imshow(image_true_data)
    else:
        axes[0].imshow(image_true_data)

    for i in range(len(dict_list)):
        col_id = (i + 1) % size_width
        row_id = (i + 1) // size_width

        buffer = BytesIO()
        dict_list[i]['pred_fig'].savefig(buffer, format='png')
        buffer.seek(0)
        image_data = plt.imread(buffer)

        if size_height > 1:
            axes[row_id][col_id].imshow(image_data)
        else:
            axes[col_id].imshow(image_data)

    fig.suptitle(plot_name, fontsize=16)
    plt.tight_layout()
```

Klasyfikator bazowy

```

In [ ]: from sklearn.model_selection import StratifiedKFold
        from sklearn.model_selection import KFold
        from sklearn.neighbors import KNeighborsClassifier
        from sklearn.model_selection import train_test_split
        from sklearn.model_selection import cross_val_score
        import sklearn.metrics as skmet

def RunKNN (df, n_neighbors = 5, weights = 'uniform', metric = 'minkowski', cv = 3,
            neigh = KNeighborsClassifier(n_neighbors = n_neighbors, metric = metric, weight

df_train, df_test = train_test_split( df, test_size=0.33, random_state=42)

neigh.fit(df_train.iloc[:, :-1], df_train.iloc[:, -1])

y_true = df_test.iloc[:, -1].tolist()
y_pred = neigh.predict(df_test.iloc[:, :-1])

strat_k_fold = None
if isStratified:
    strat_k_fold = StratifiedKFold(n_splits=cv, shuffle=True, random_state=42)
else:
    strat_k_fold = KFold(n_splits=cv, shuffle=True, random_state=42)

cross_val = cross_val_score(neigh, df.iloc[:, :-1], df.iloc[:, -1], cv=strat_k_
cross_val = cross_val.mean()

if print_metrics:
    print(skmet.classification_report(y_true, y_pred))
    print('Cross val: ', cross_val)

    return {'f1':skmet.f1_score(y_true, y_pred, average='micro'), 'cross-val': cros

RunKNN(df_iris, cv = 3)

```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	19
1	0.94	1.00	0.97	15
2	1.00	0.94	0.97	16
accuracy			0.98	50
macro avg	0.98	0.98	0.98	50
weighted avg	0.98	0.98	0.98	50

Cross val: 0.9666666666666667

```
Out[ ]: {'f1': 0.98, 'cross-val': 0.9666666666666667}
```

```
In [ ]: score_metric = 'f1'
results_df = pd.DataFrame([
    RunKNN (df_iris, print_metrics = False)[score_metric], RunKNN (df_iris, print_metr
    RunKNN (df_seeds, print_metrics = False)[score_metric], RunKNN (df_seeds, print_me
    RunKNN (df_glass, print_metrics = False)[score_metric], RunKNN (df_glass, print_me
    RunKNN (df_wine, print_metrics = False)[score_metric], RunKNN (df_wine, print_metr
], index=['IRIS', 'SEEDS', 'GLASS', 'WINE'], columns = ['f1', 'cross_val'])

results_df
```

```
Out[ ]:
```

	f1	cross_val
IRIS	0.980000	0.966667
SEEDS	0.885714	0.904762
GLASS	0.957746	0.976721
WINE	0.677966	0.640207

Bagging

```
In [ ]: from sklearn.metrics import f1_score
from sklearn.cluster import KMeans
from sklearn.pipeline import Pipeline
from sklearn.impute import SimpleImputer
from sklearn.preprocessing import StandardScaler
import numpy as np
from sklearn.ensemble import BaggingClassifier

def run_bagging(df, n_estimators = 10, max_samples = 1.0, max_features = 1.0, boots
df_train, df_test = train_test_split( df, test_size=0.33, random_state=42)
clf = BaggingClassifier(n_estimators = n_estimators, max_samples = max_samples,
clf = Pipeline(['impute', SimpleImputer( strategy='mean')),
                ('standardization', StandardScaler()),
                ('clf', clf),])

clf.fit(df_train.iloc[:, :-1], df_train.iloc[:, -1])

y_true = df_test.iloc[:, -1].tolist()
y_pred = clf.predict(df_test.iloc[:, :-1])

X_reduced = PCA(n_components=2).fit_transform(df_test)

pred_fig = plt.figure(1, figsize=(5, 5))
ax1 = pred_fig.add_subplot()
ax1.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_pred)
ax1.set_title(f"Bagging pred\n n_estimators = {n_estimators}, max_samples={max
plt.close()

true_fig = plt.figure(2, figsize=(5, 5))
ax2 = true_fig.add_subplot()
ax2.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_true)
ax2.set_title(f"Bagging true\n n_estimators = {n_estimators}, max_samples={max
plt.close()

f1 = f1_score(y_true, y_pred, average='weighted')

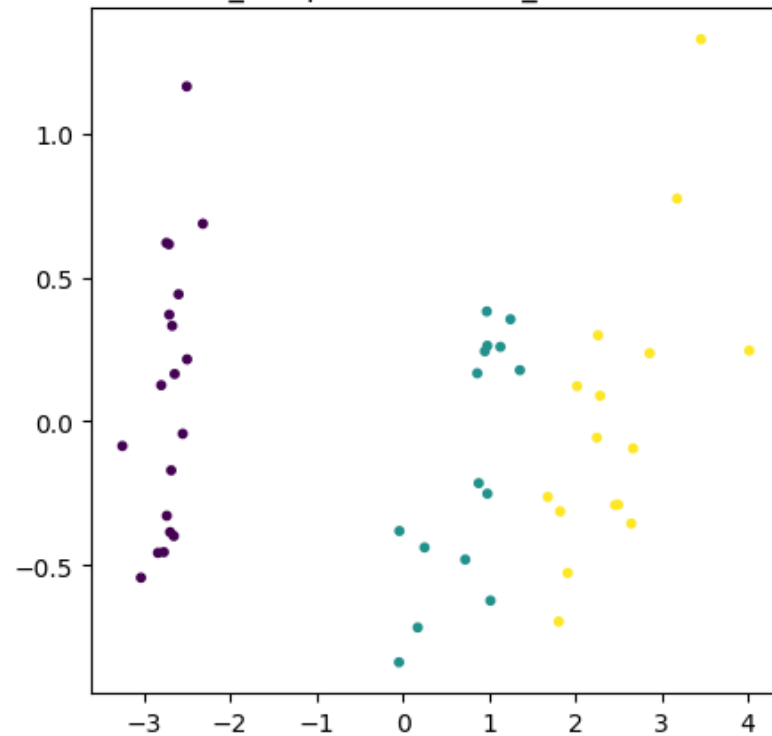
return {'f1':f1,
        'pred_fig': pred_fig, 'true_fig': true_fig,}

test_run = run_bagging(df_iris)
print(test_run['f1'])
test_run['pred_fig']
```

1.0

Out[]:

Bagging pred
n_estimators = 10, max_samples=1.0, max_features = 1.0, bootstrap = True

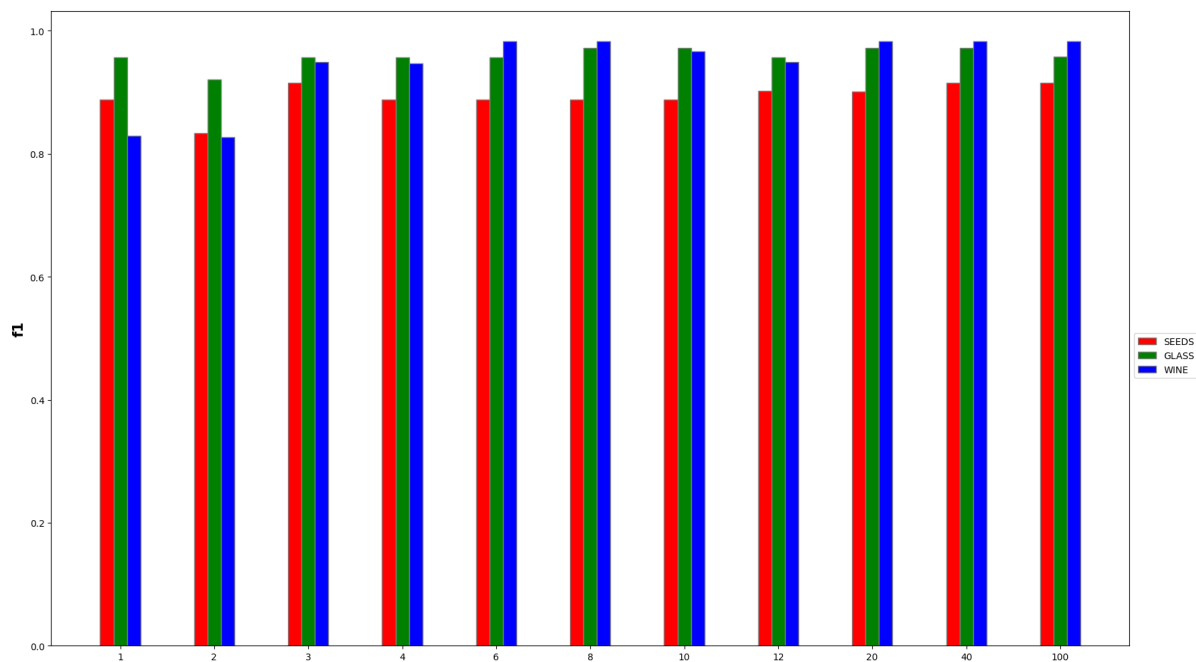


```
In [ ]: metrics_list = ['f1']  
dataset_list = ['SEEDS', 'GLASS', 'WINE']  
colour_list = ['r', 'g', 'b']
```

Bagging - liczba classifikatorów

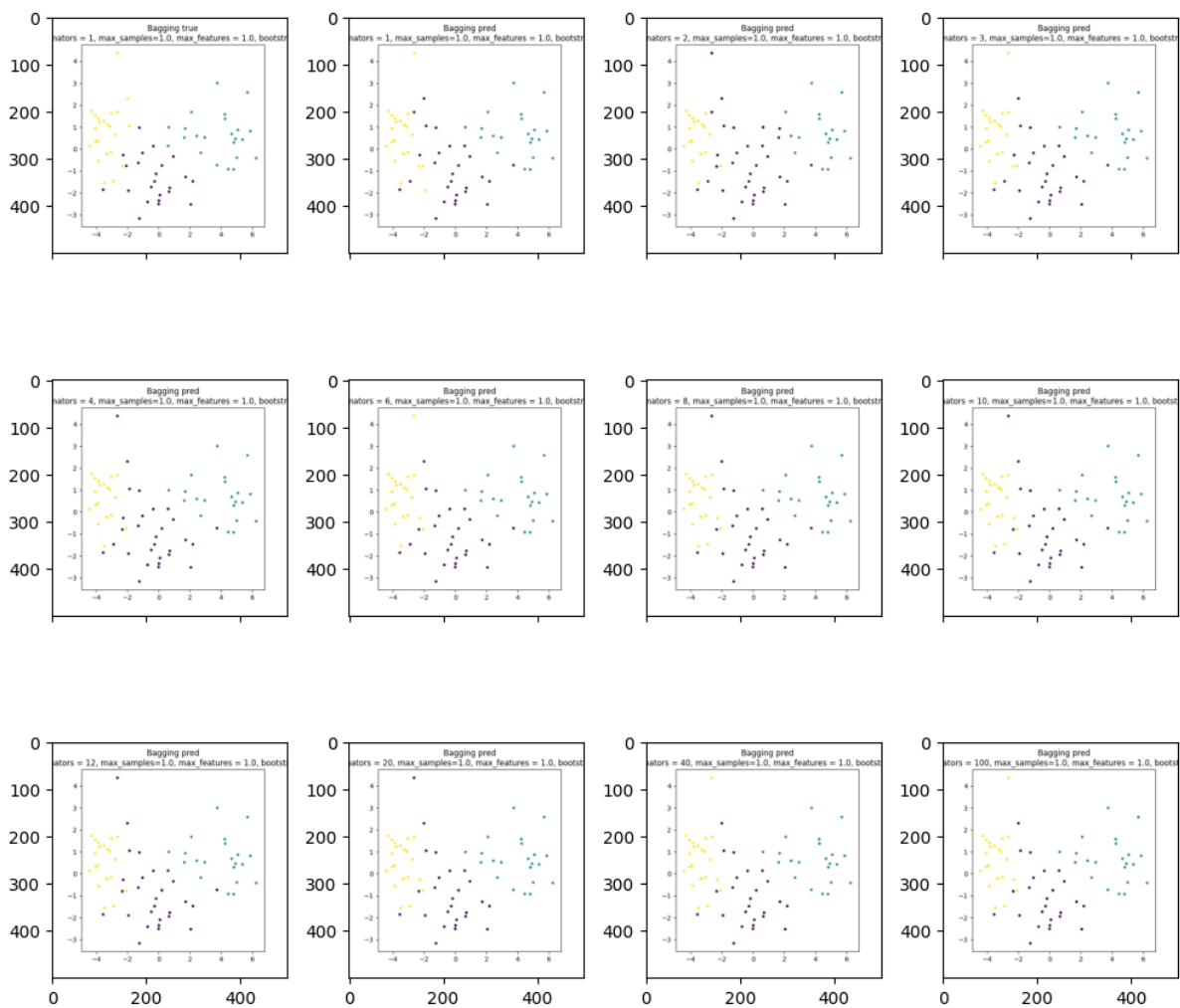
```
In [ ]: study_vals = [1, 2, 3, 4, 6, 8, 10, 12, 20, 40, 100]  
results_list = [  
    [run_bagging(df_seeds, n_estimators = c) for c in study_vals],  
    [run_bagging(df_glass, n_estimators = c) for c in study_vals],  
    [run_bagging(df_wine, n_estimators = c) for c in study_vals],  
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



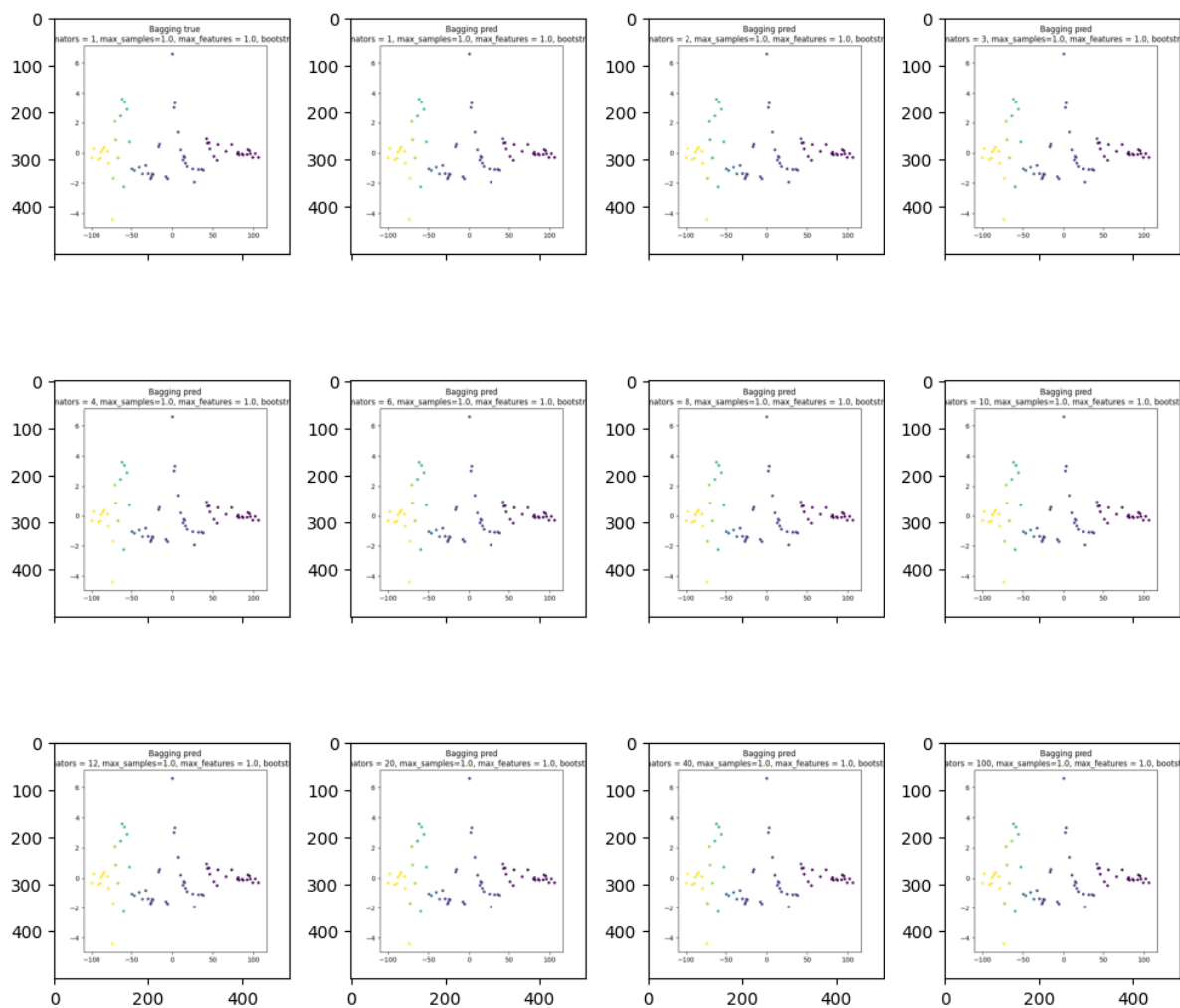
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



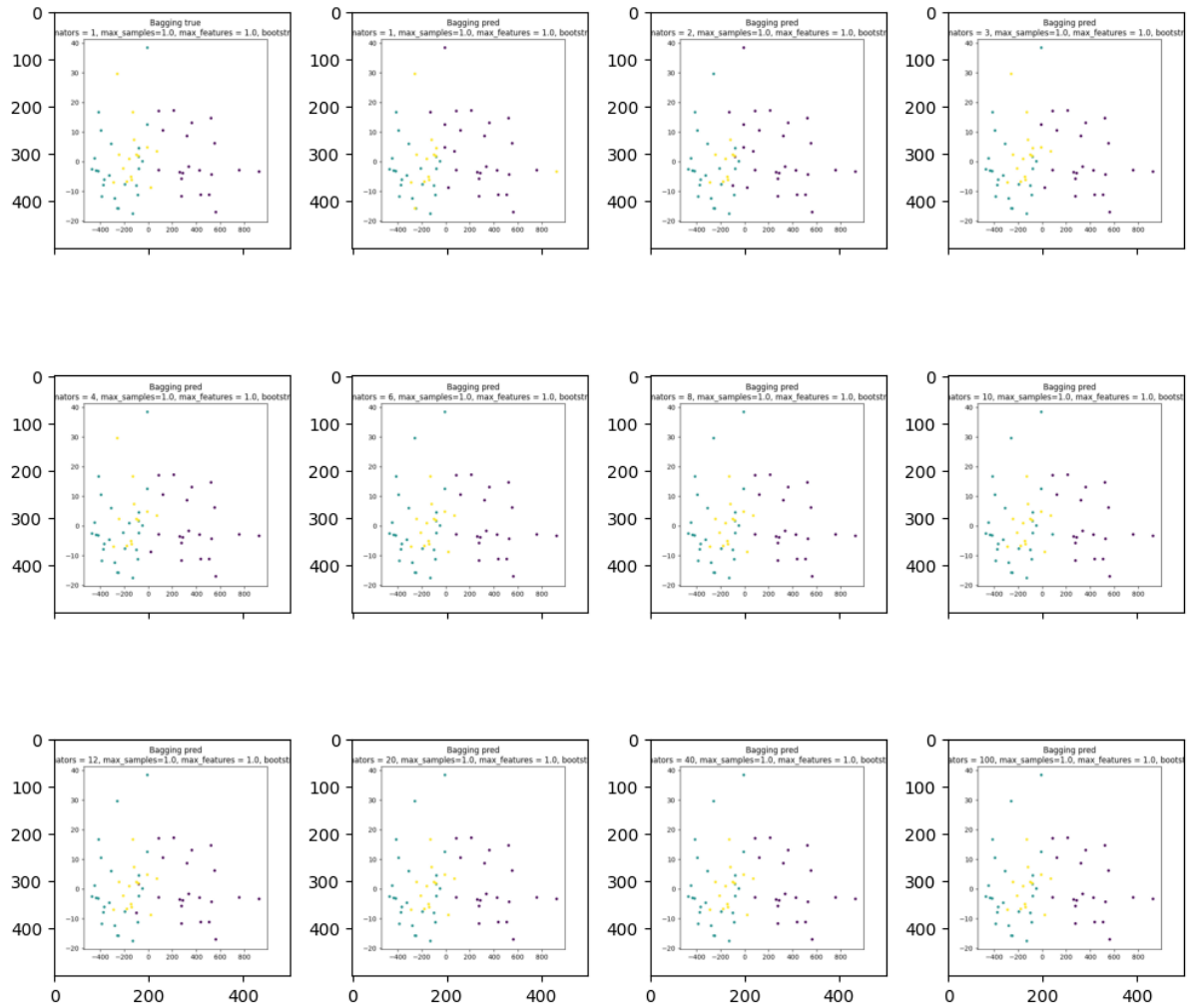
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

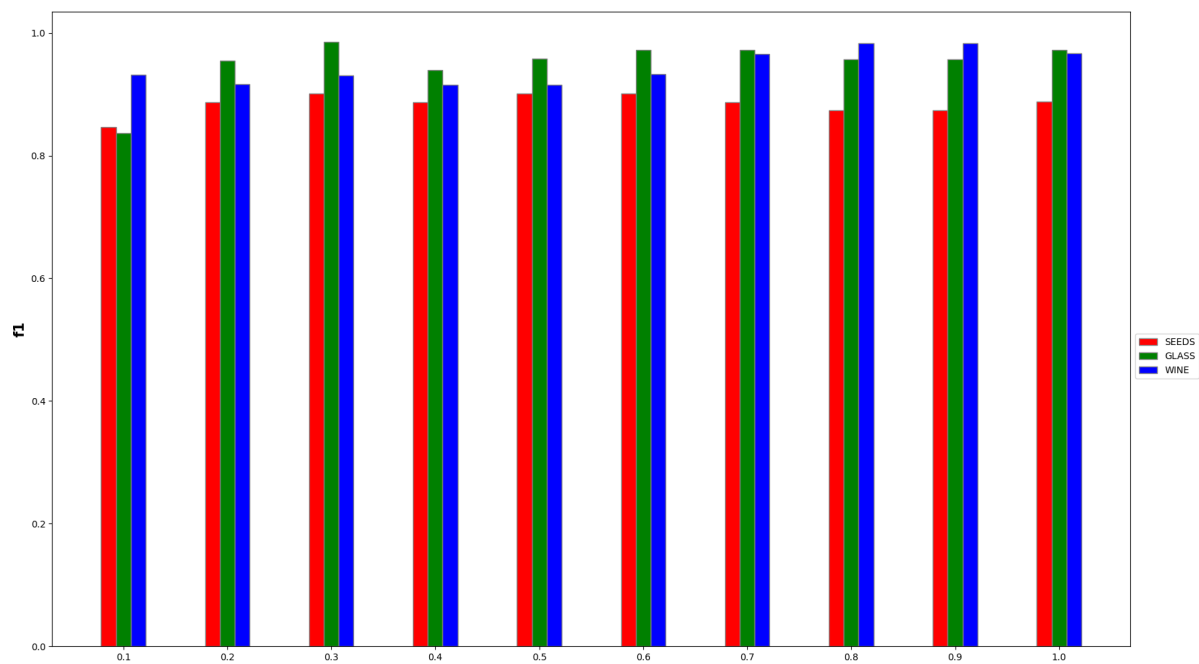

WINE



Bagging - liczba probek

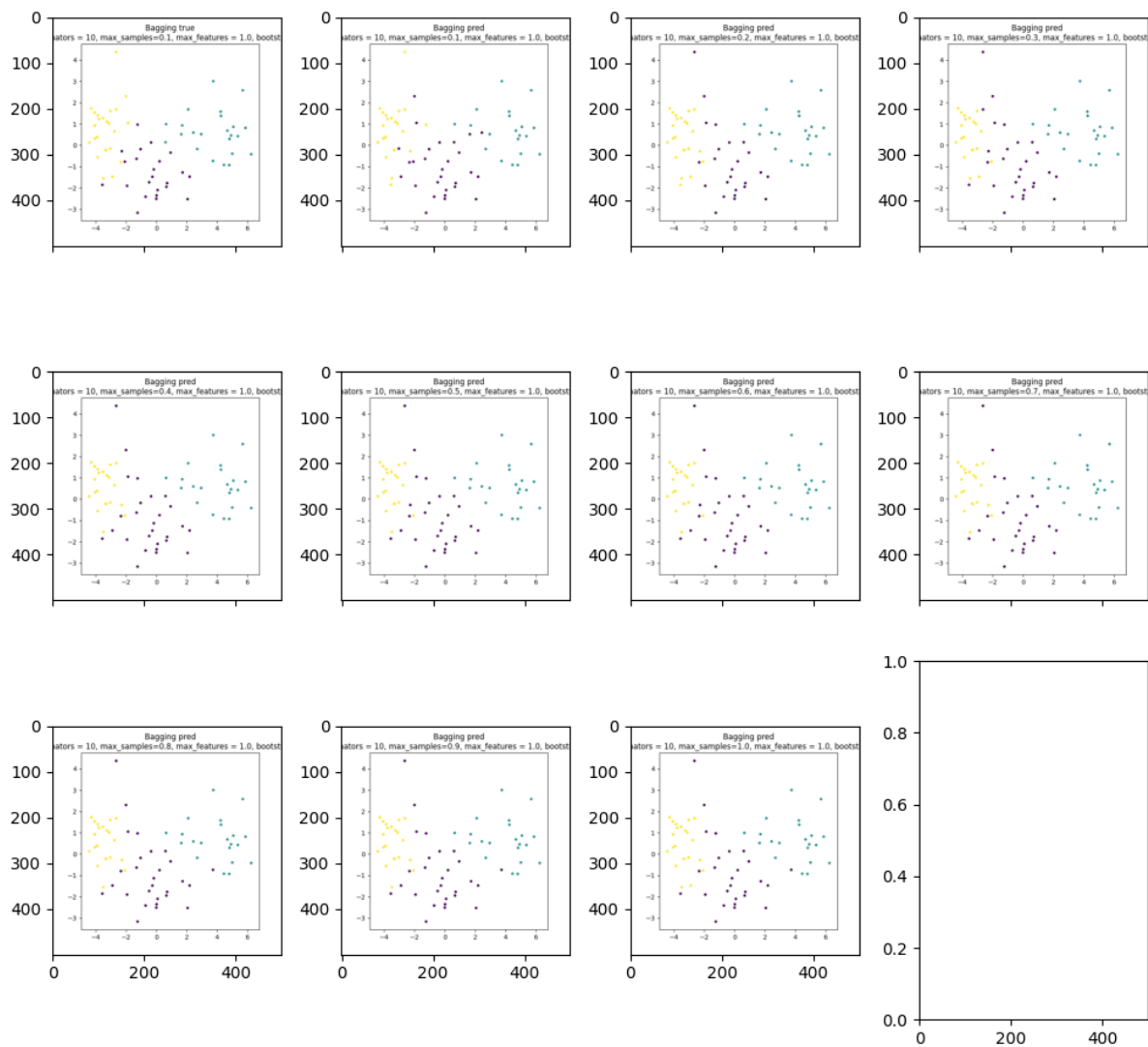
```
In [ ]: study_vals = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.]
results_list = [
    [run_bagging(df_seeds, max_samples = c) for c in study_vals],
    [run_bagging(df_glass, max_samples = c) for c in study_vals],
    [run_bagging(df_wine, max_samples = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



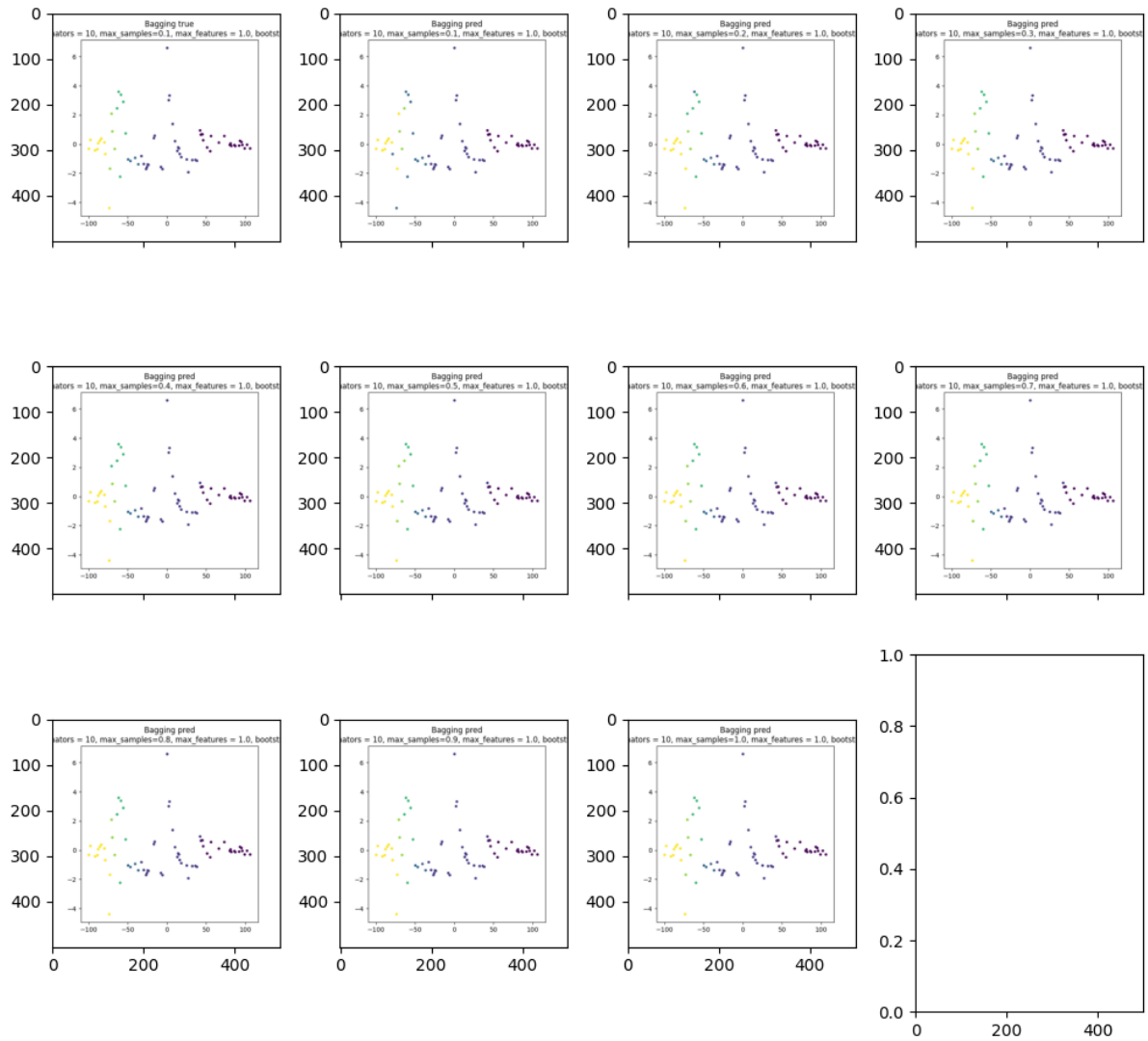
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



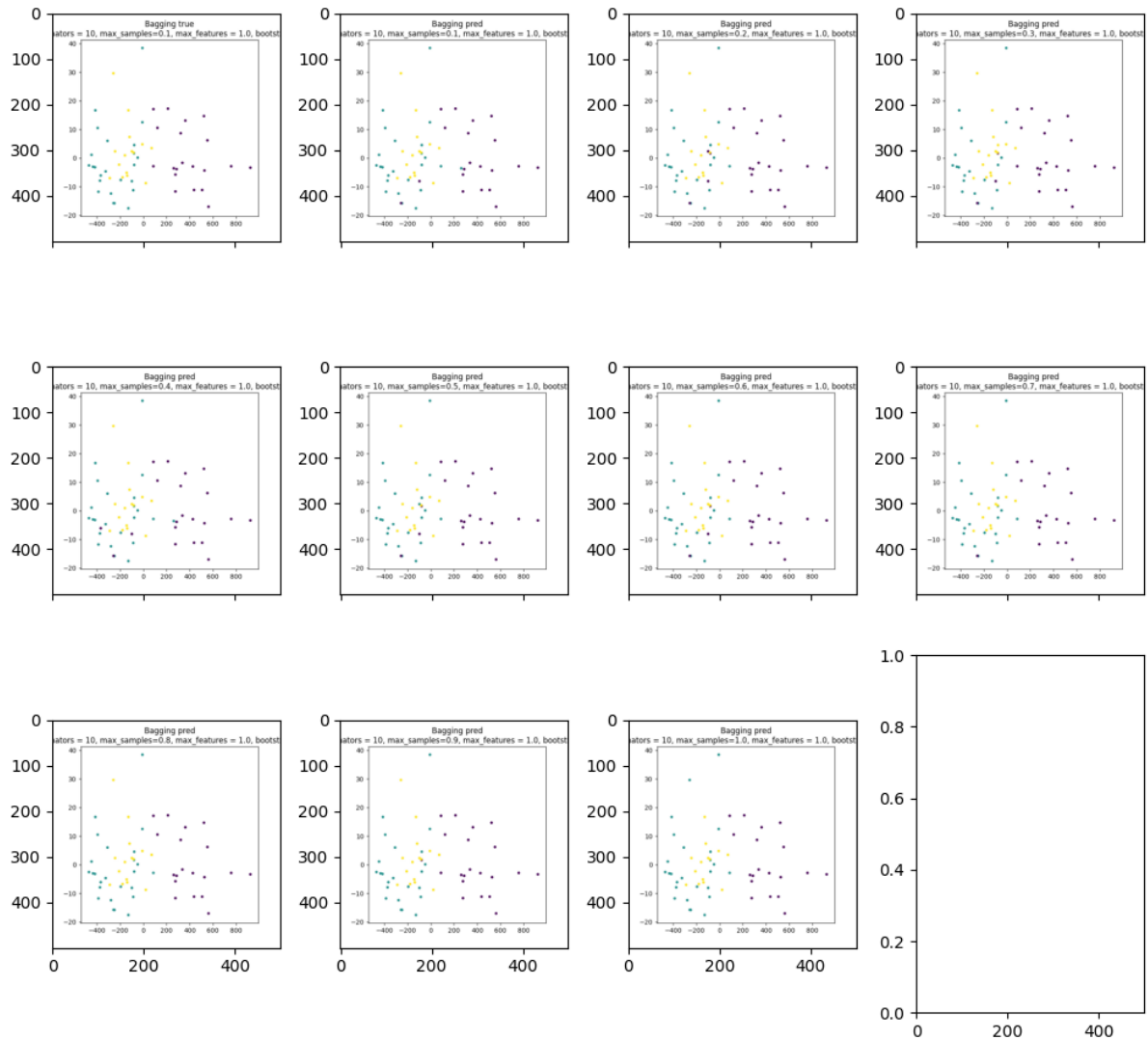
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

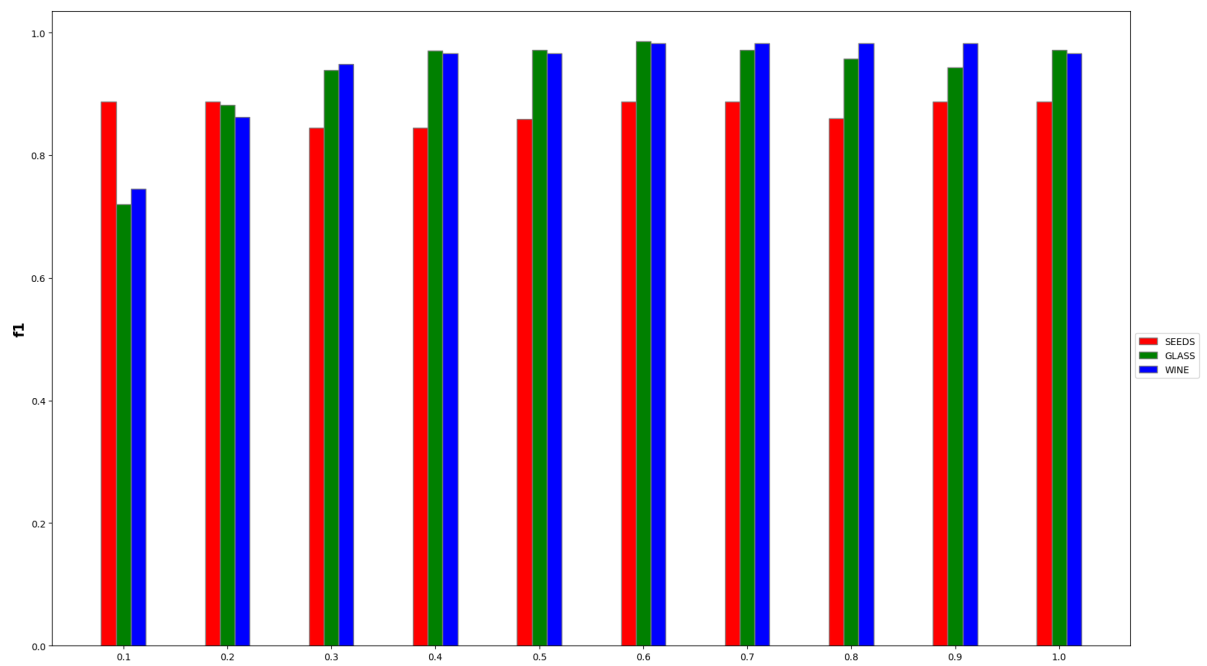
WINE



Bagging - liczba atrybutow

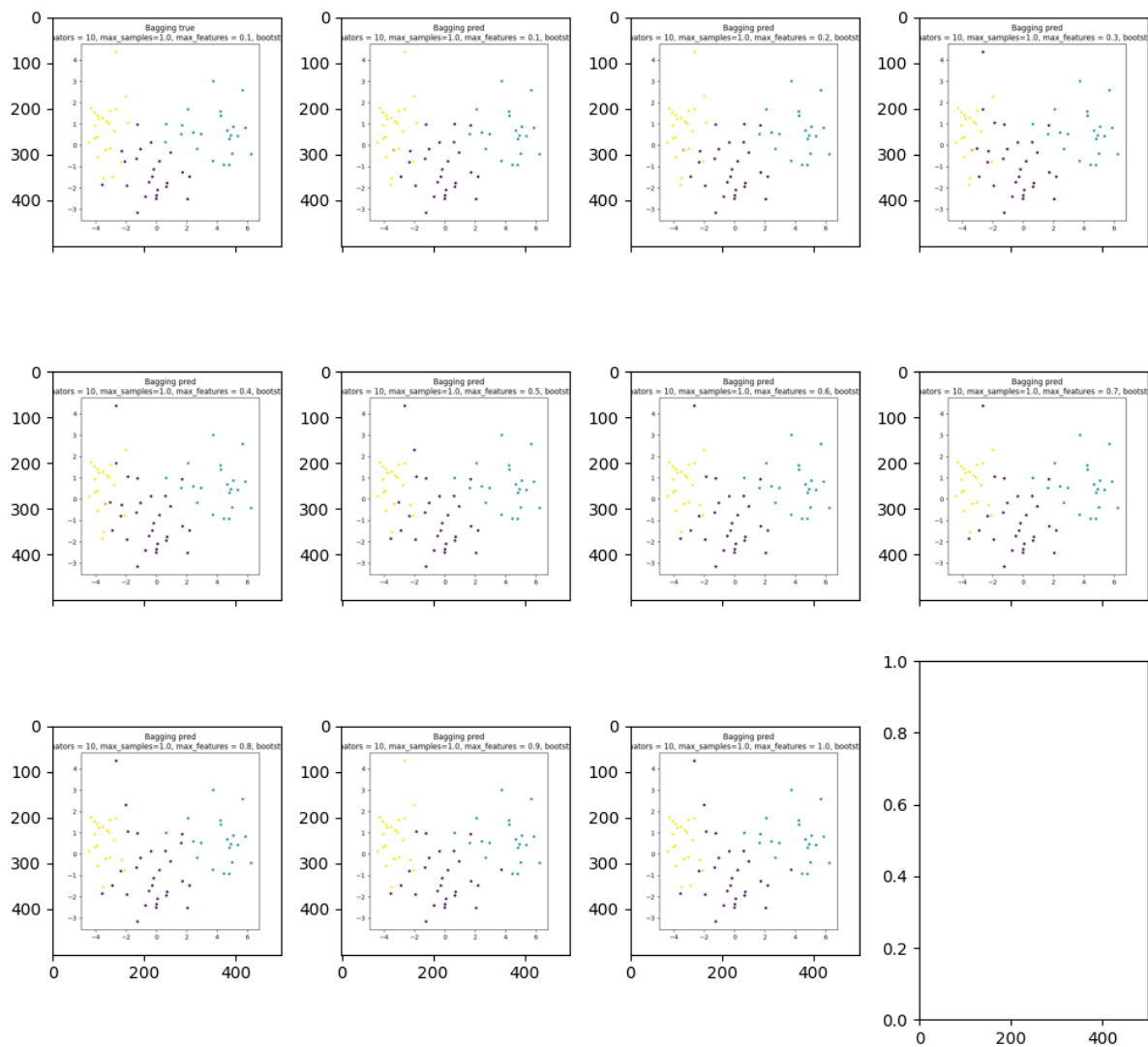
```
In [ ]: study_vals = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.]
results_list = [
    [run_bagging(df_seeds, max_features = c) for c in study_vals],
    [run_bagging(df_glass, max_features = c) for c in study_vals],
    [run_bagging(df_wine, max_features = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



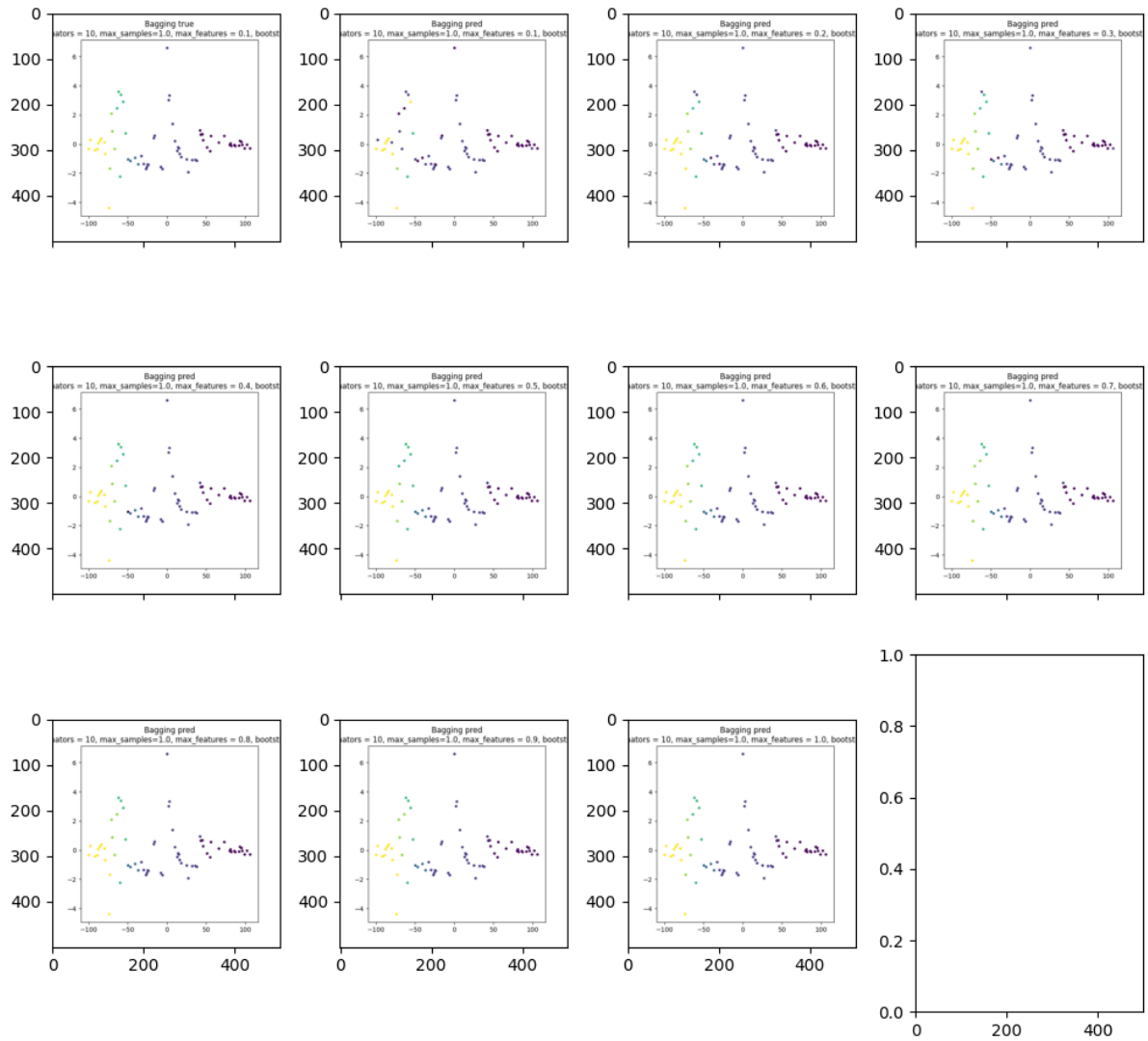
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



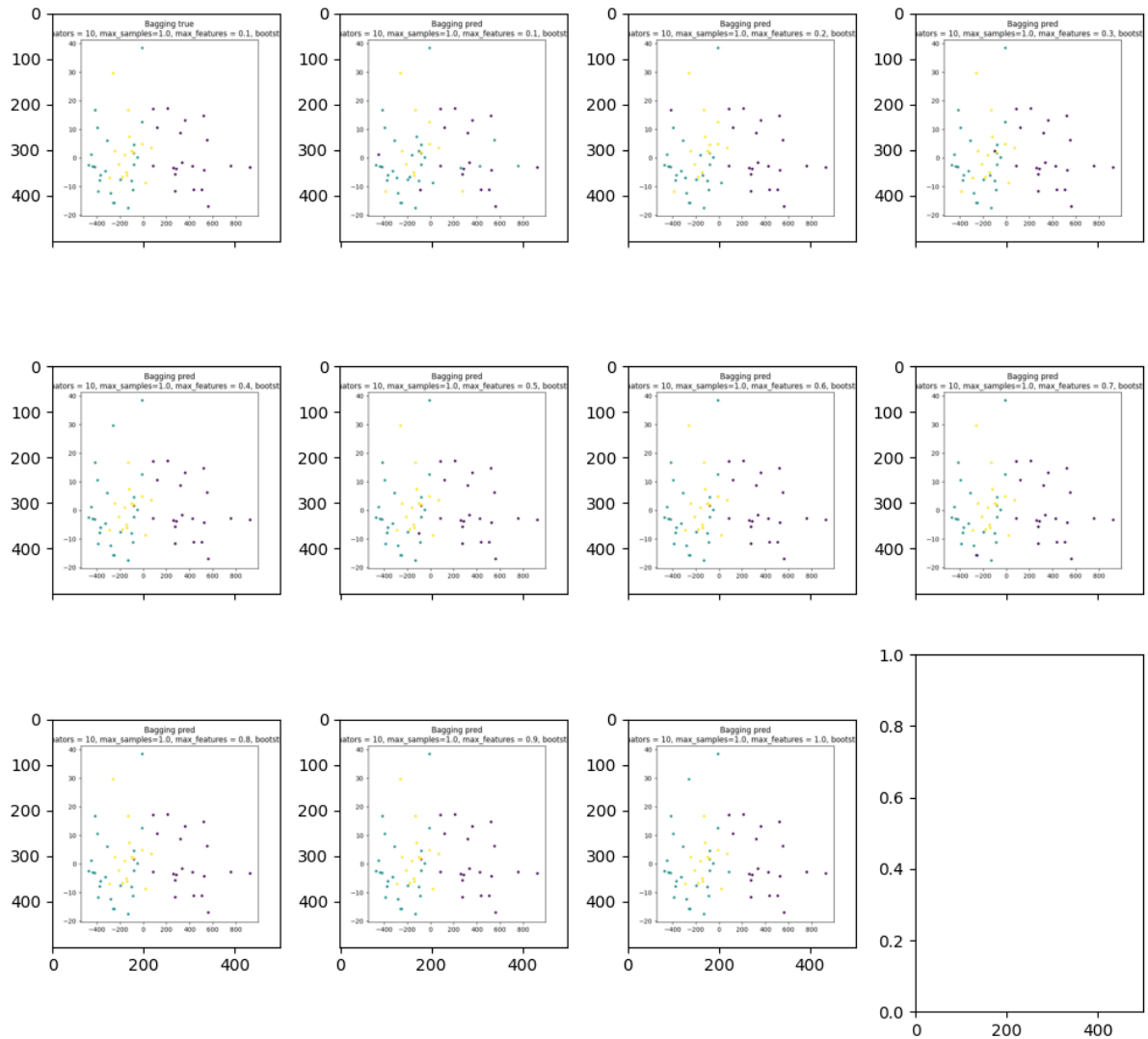
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

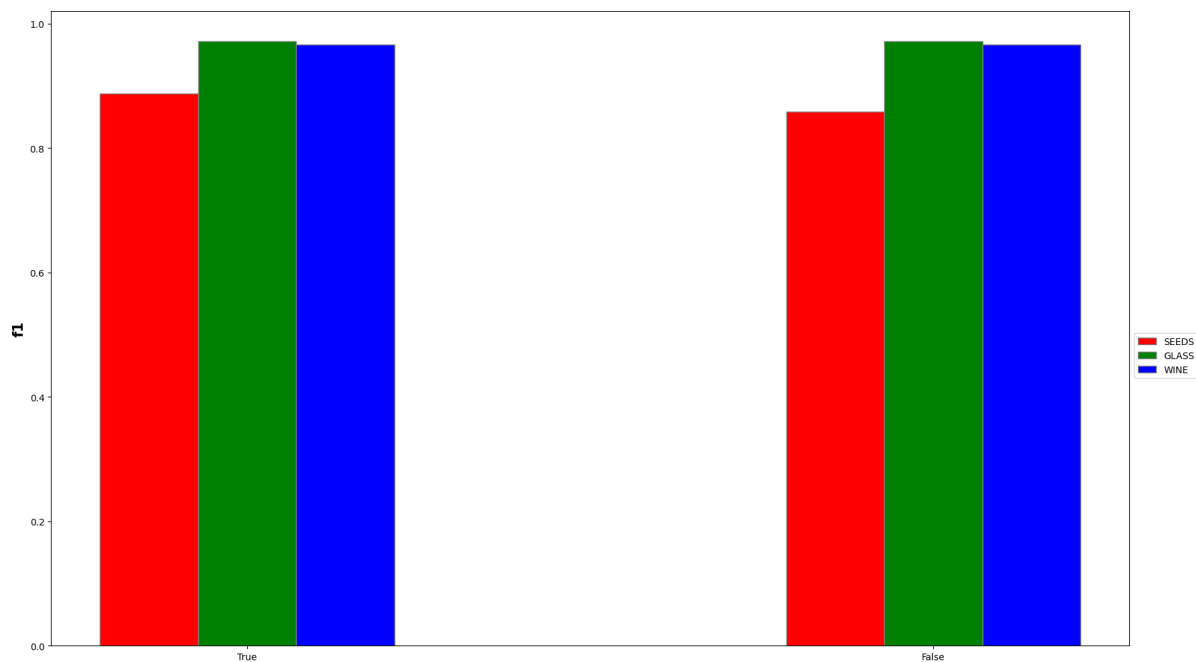

WINE



Bagging - bootstrap

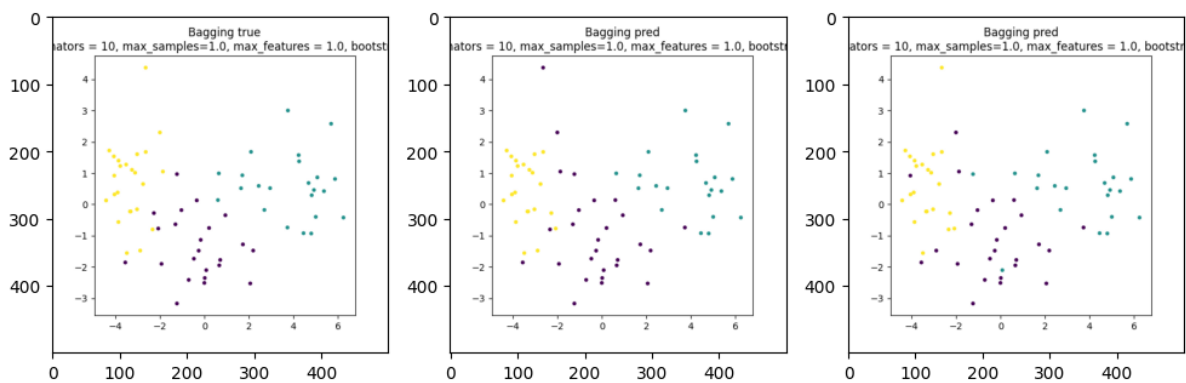
```
In [ ]: study_vals = [True, False]
results_list = [
    [run_bagging(df_seeds, bootstrap = c) for c in study_vals],
    [run_bagging(df_glass, bootstrap = c) for c in study_vals],
    [run_bagging(df_wine, bootstrap = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



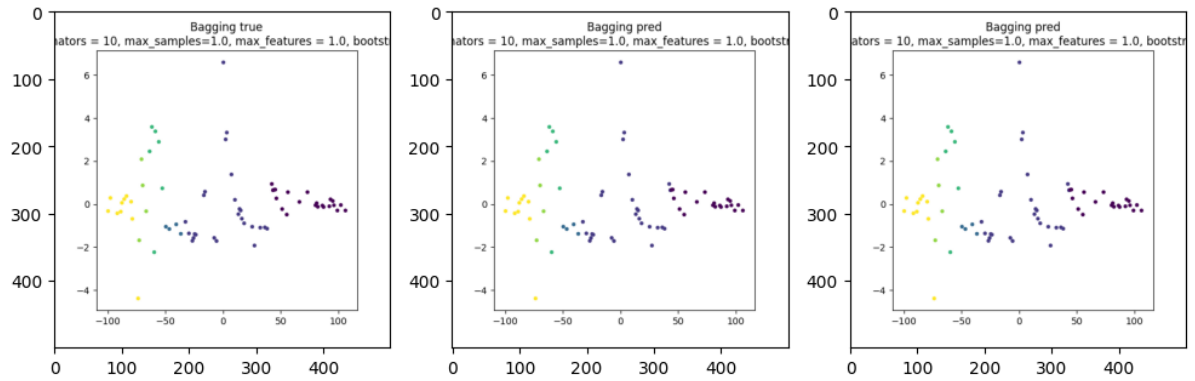
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



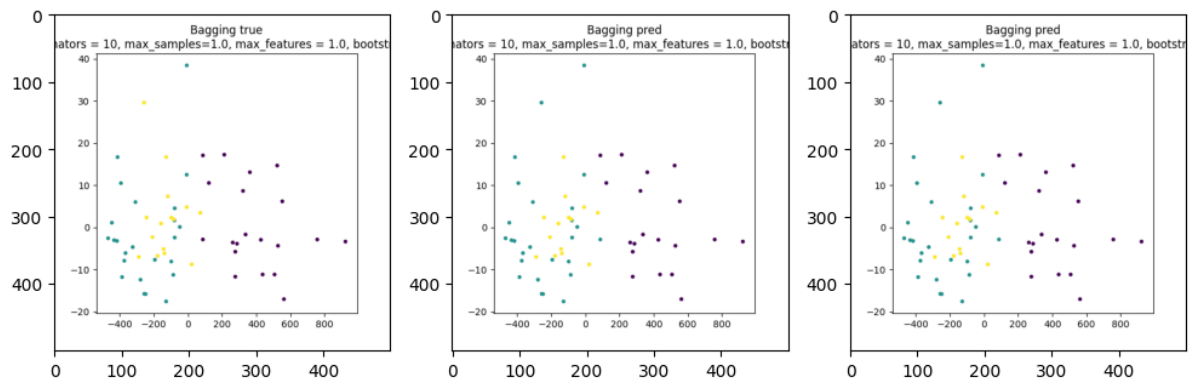
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

WINE



Boosting

```
In [ ]: from sklearn.metrics import f1_score
from sklearn.cluster import KMeans
from sklearn.pipeline import Pipeline
from sklearn.impute import SimpleImputer
from sklearn.preprocessing import StandardScaler
import numpy as np
from sklearn.ensemble import AdaBoostClassifier

def run_adaboost(df, n_estimators = 10, learning_rate = 1.0):
    df_train, df_test = train_test_split(df, test_size=0.33, random_state=42)
    clf = AdaBoostClassifier(n_estimators = n_estimators, learning_rate = learning_rate)
    clf = Pipeline([('impute', SimpleImputer(strategy='mean')),
                    ('standardization', StandardScaler()),
                    ('clf', clf),])

    clf.fit(df_train.iloc[:, :-1], df_train.iloc[:, -1])

    y_true = df_test.iloc[:, -1].tolist()
    y_pred = clf.predict(df_test.iloc[:, :-1])

    X_reduced = PCA(n_components=2).fit_transform(df_test)

    pred_fig = plt.figure(1, figsize=(5, 5))
    ax1 = pred_fig.add_subplot()
    ax1.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_pred)
    ax1.set_title(f"AdaBoost pred\n n_estimators = {n_estimators}, learning_rate={learning_rate}")
    plt.close()

    true_fig = plt.figure(2, figsize=(5, 5))
    ax2 = true_fig.add_subplot()
    ax2.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_true)
    ax2.set_title(f"AdaBoost true\n n_estimators = {n_estimators}, learning_rate={learning_rate}")
    plt.close()

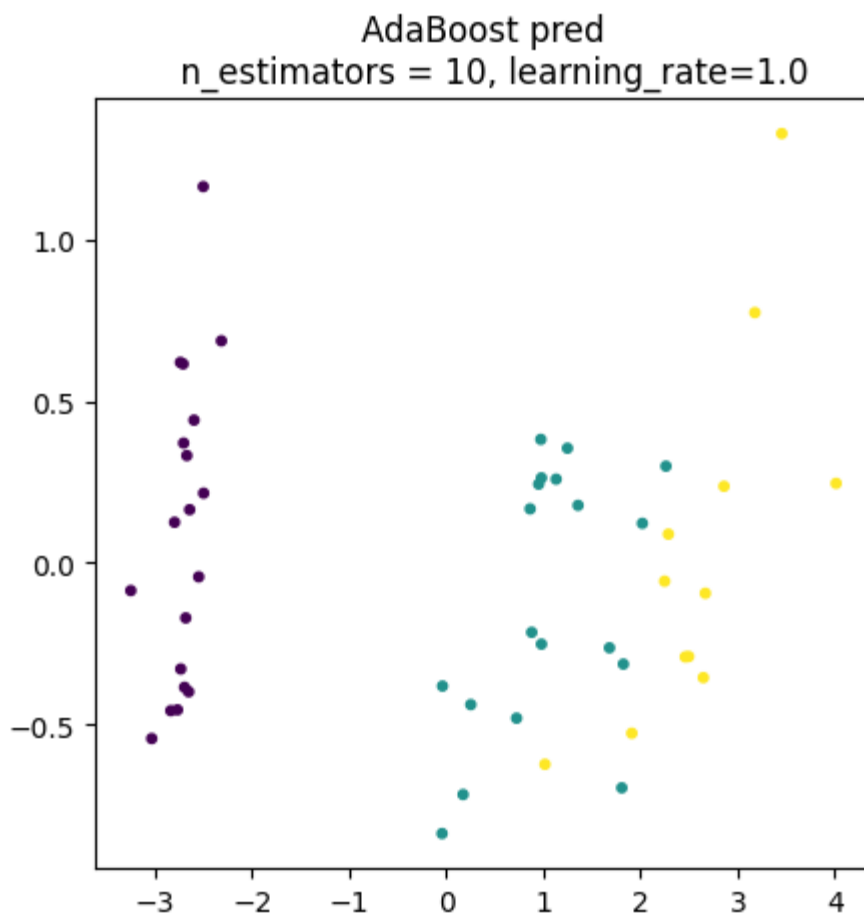
    f1 = f1_score(y_true, y_pred, average='weighted')

    return {'f1': f1,
            'pred_fig': pred_fig, 'true_fig': true_fig,}

test_run = run_adaboost(df_iris)
print(test_run['f1'])
test_run['pred_fig']
```

0.8784873949579831

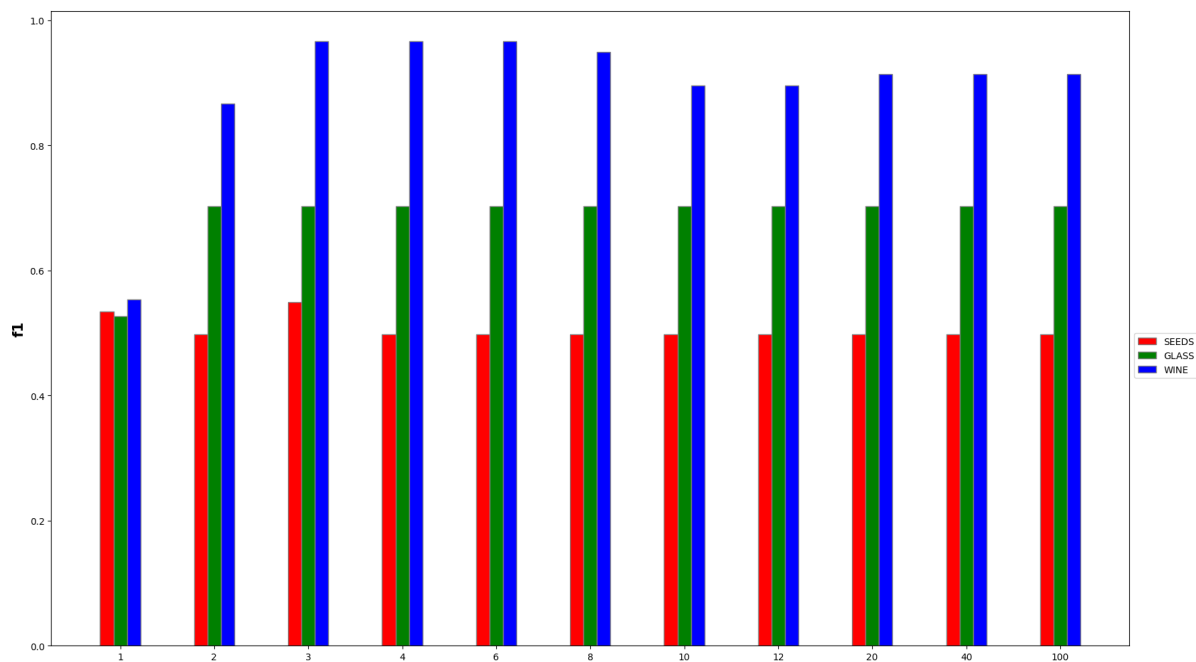
Out[]:



AdaBoost - liczba klasyfikatorow

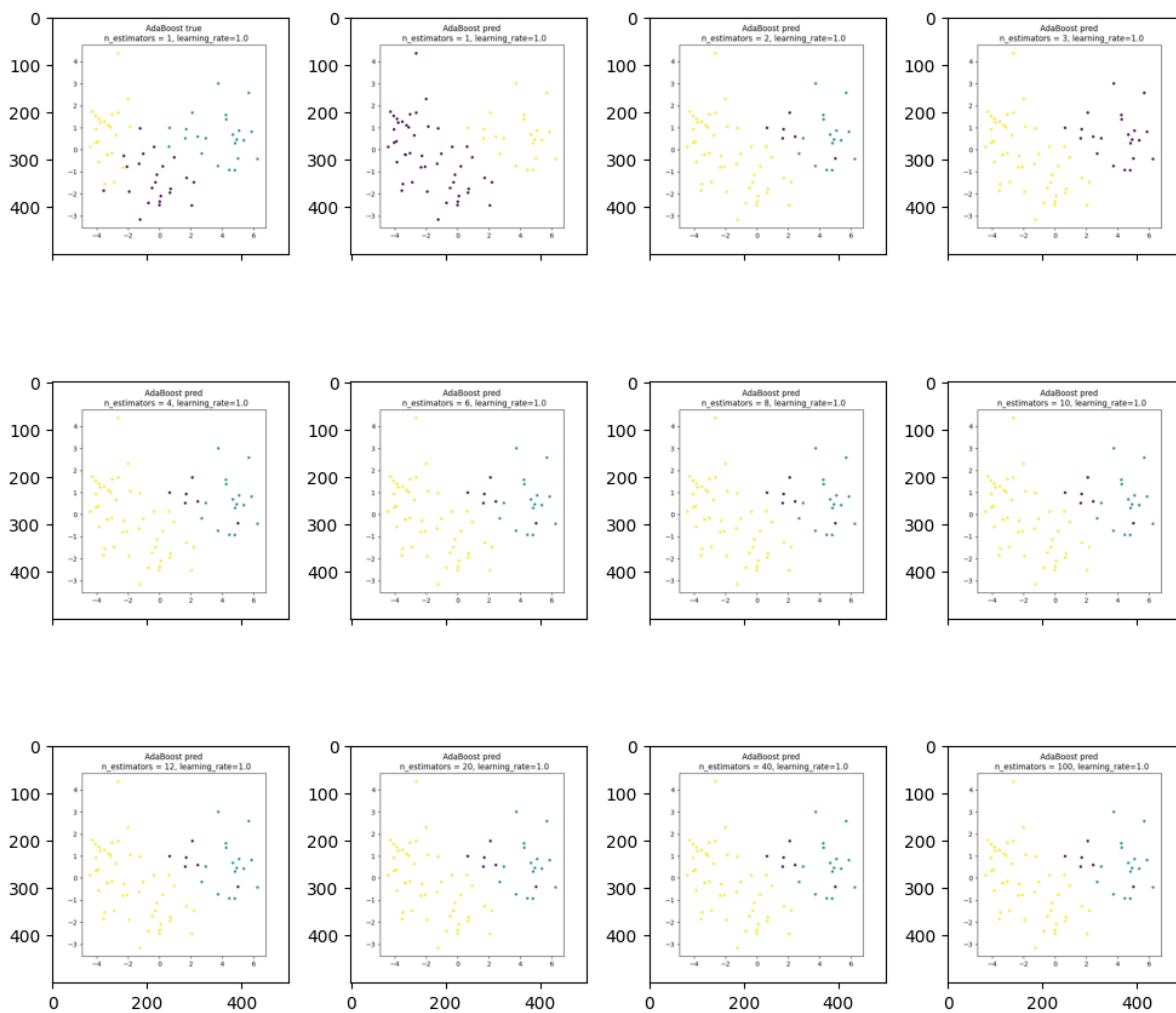
```
In [ ]: study_vals = [1, 2, 3, 4, 6, 8, 10, 12, 20, 40, 100]
results_list = [
    [run_adaboost (df_seeds, n_estimators = c) for c in study_vals],
    [run_adaboost (df_glass, n_estimators = c) for c in study_vals],
    [run_adaboost (df_wine, n_estimators = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



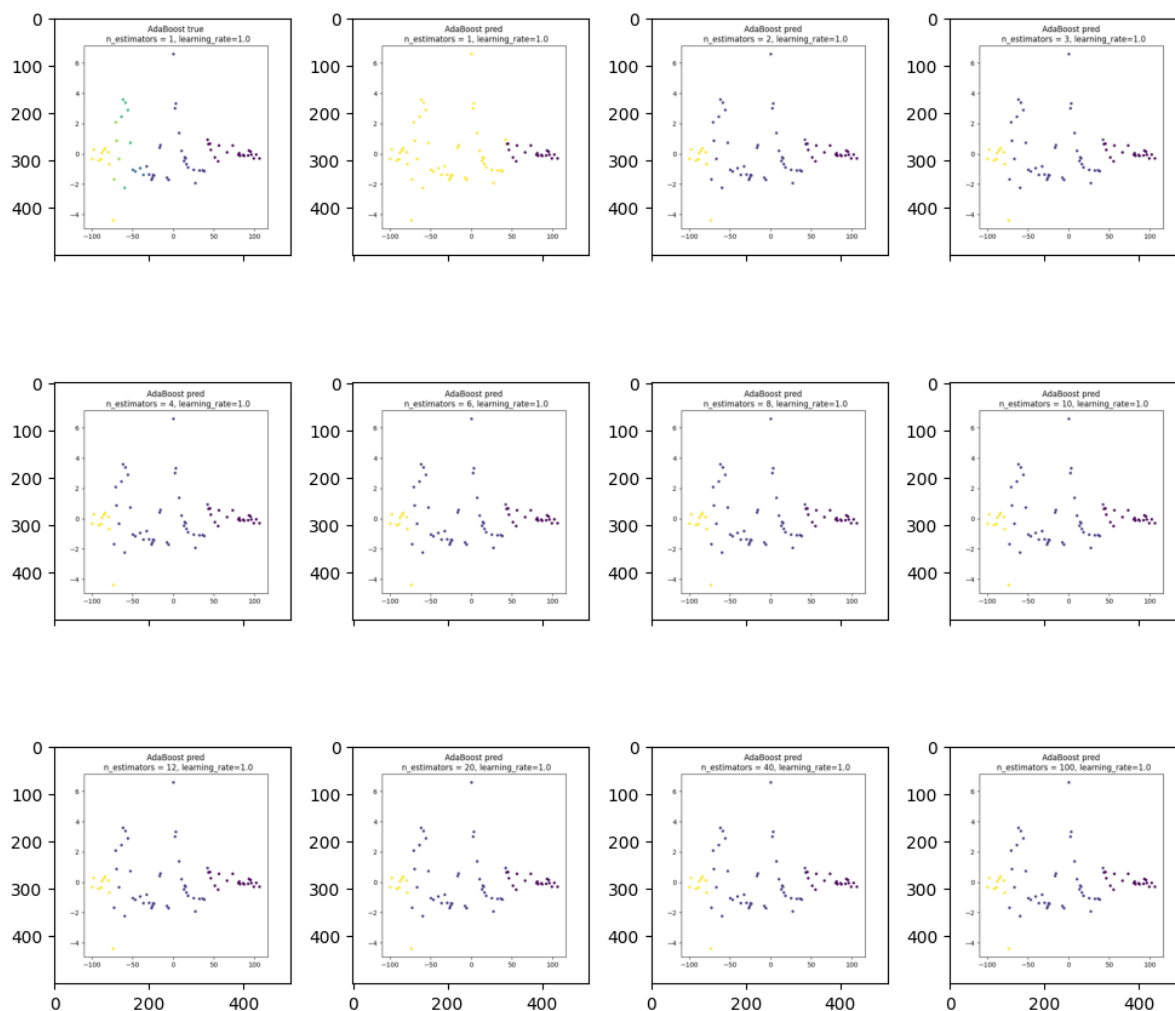
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



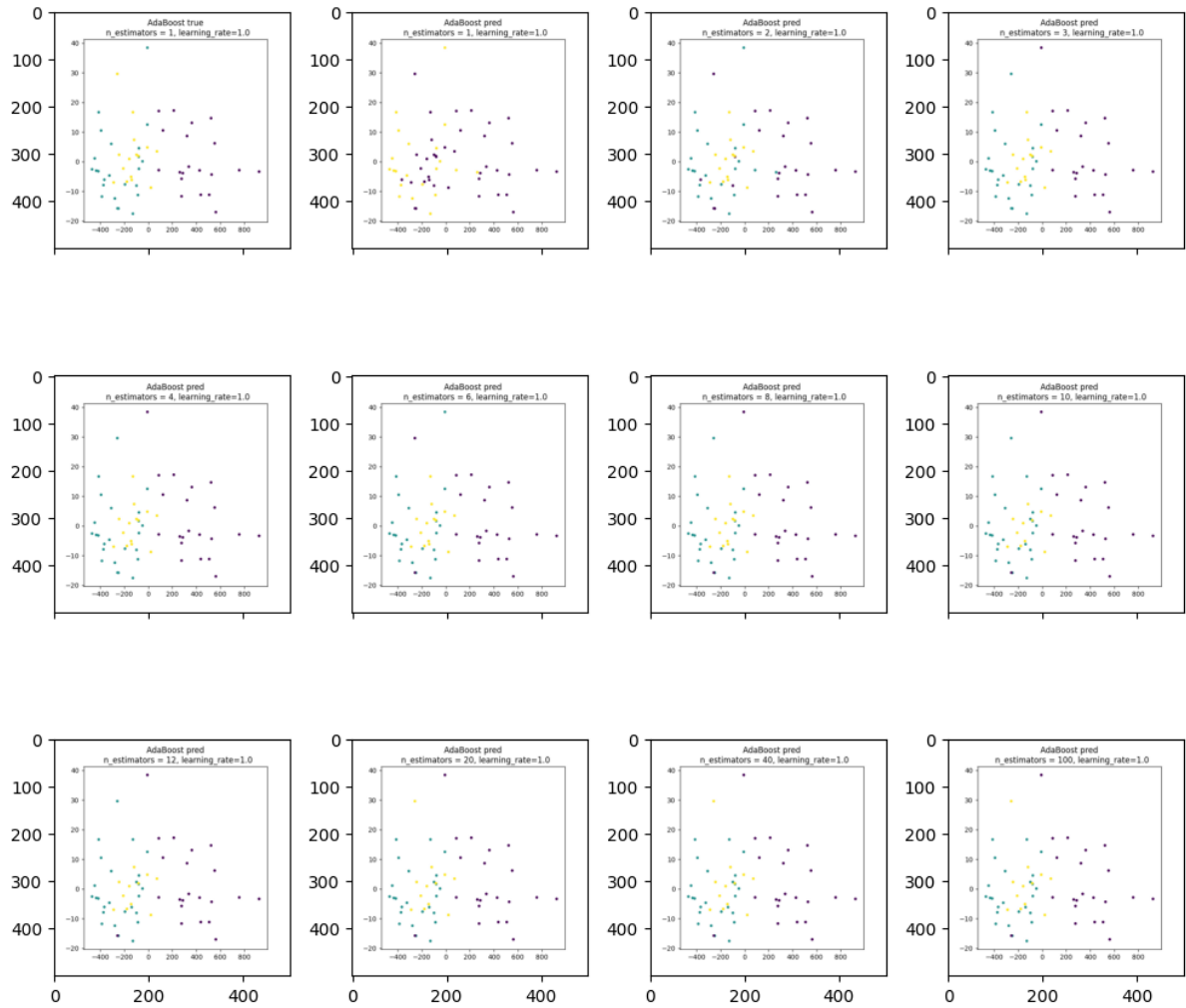
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

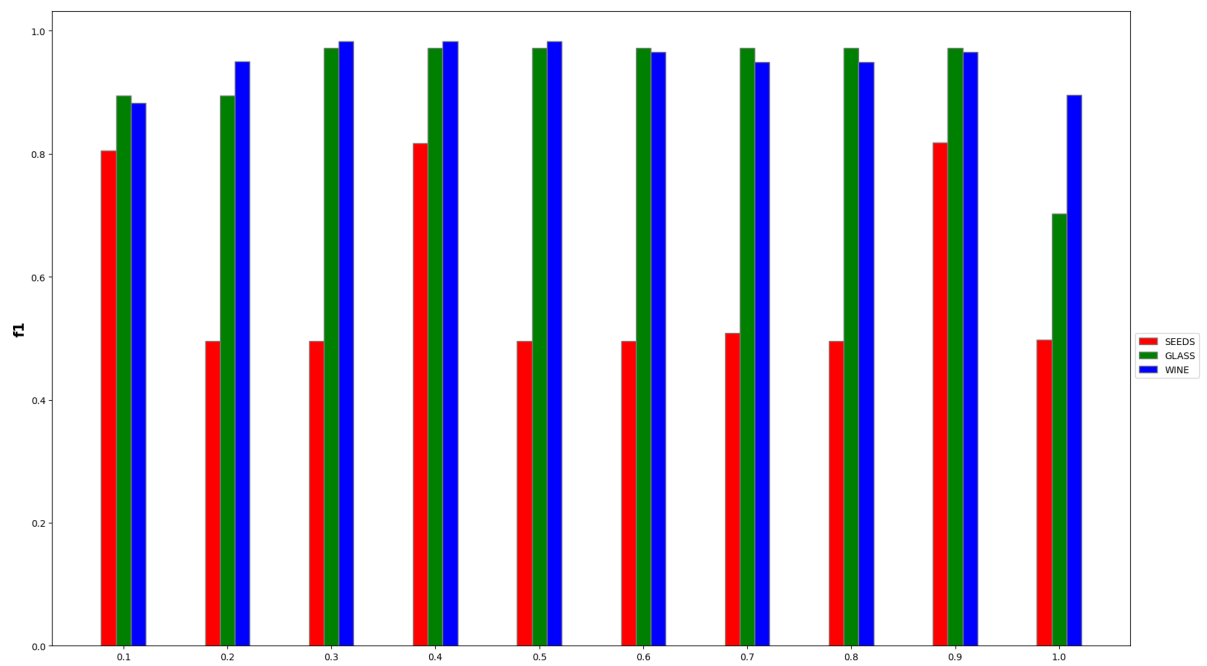
WINE



AdaBoost - wspl uczenia

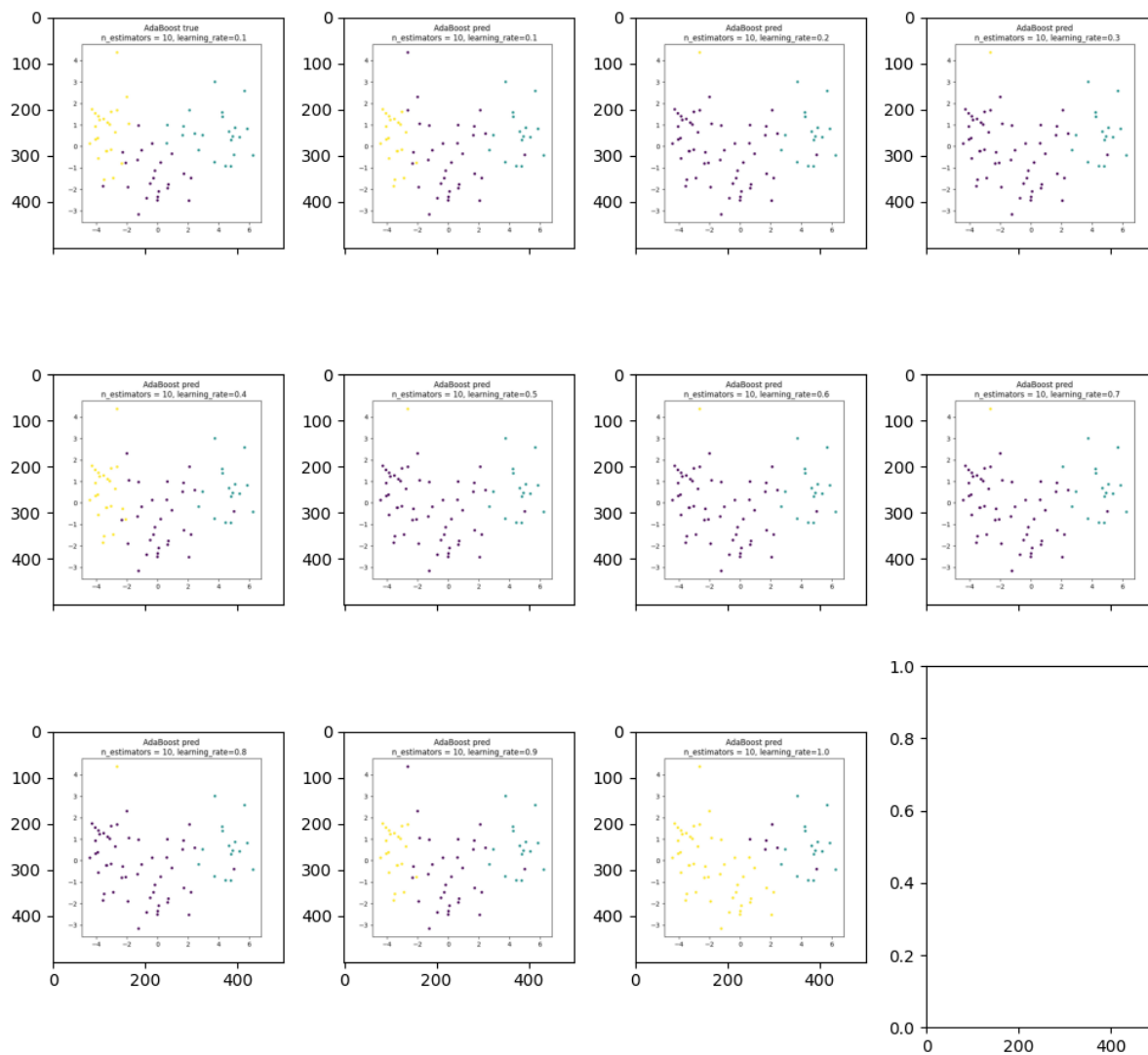
```
In [ ]: study_vals = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.]
results_list = [
    [run_adaboost (df_seeds, learning_rate = c) for c in study_vals],
    [run_adaboost (df_glass, learning_rate = c) for c in study_vals],
    [run_adaboost (df_wine, learning_rate = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```

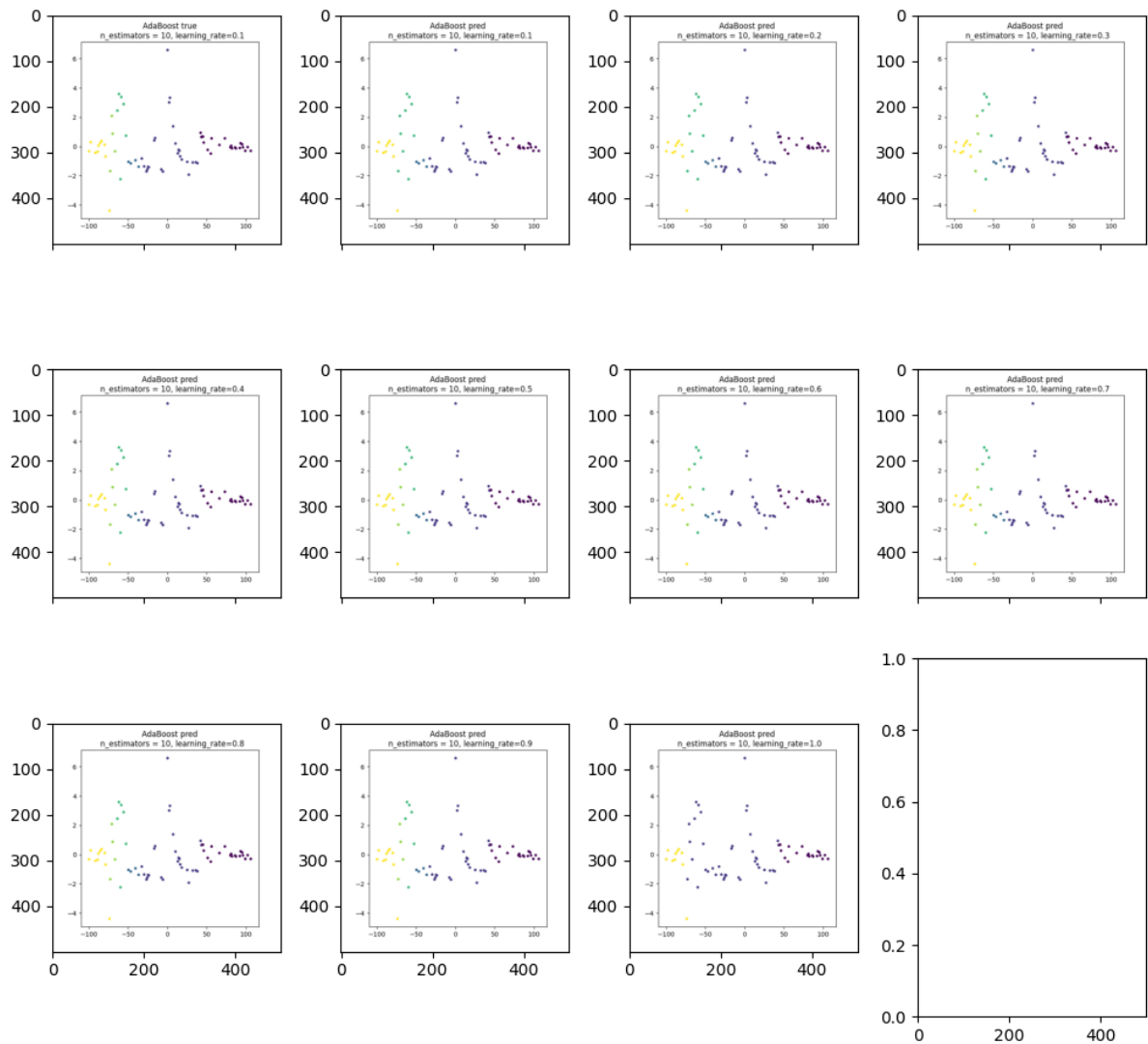
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



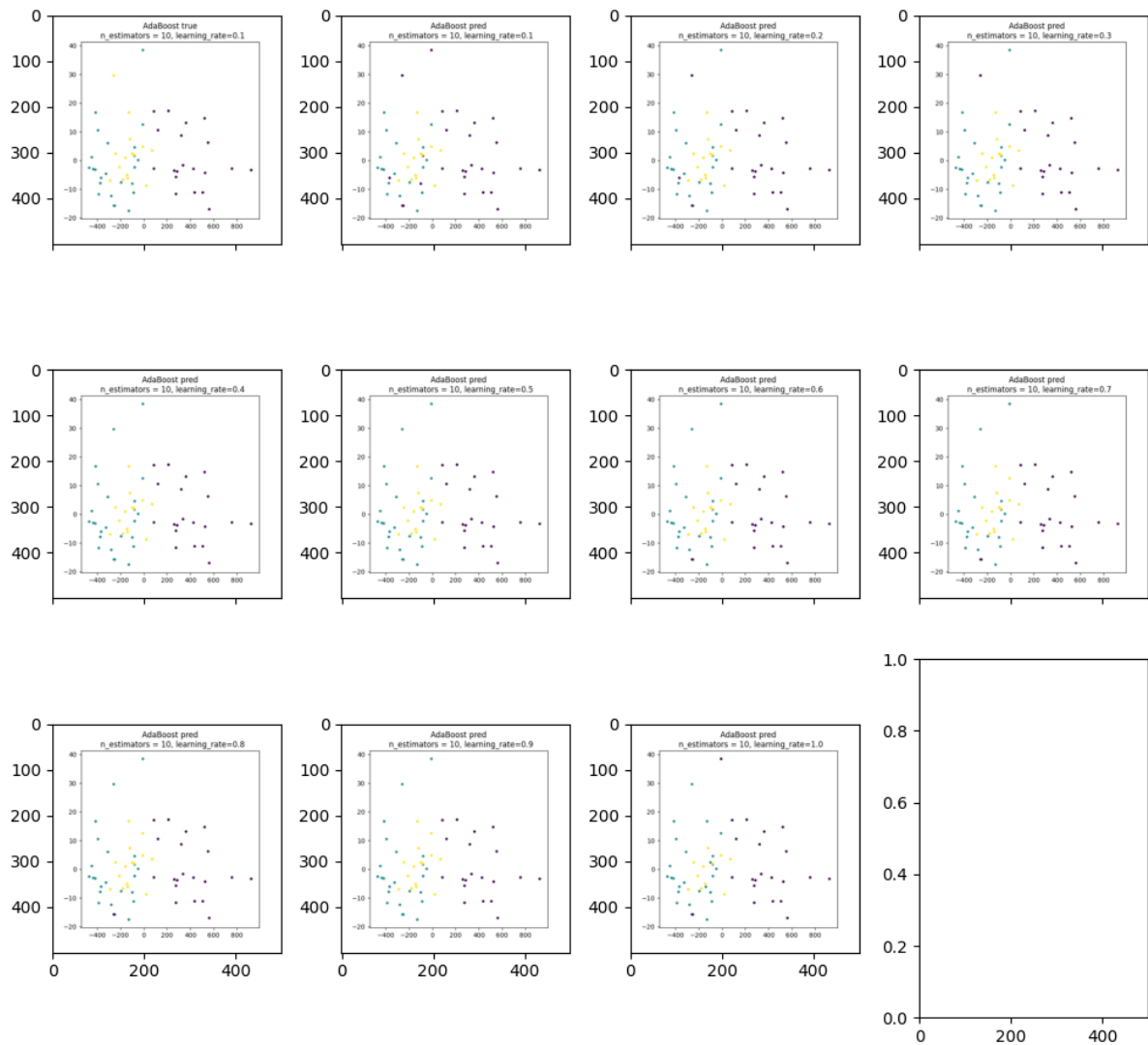
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

WINE



Random forest

```
In [ ]: from sklearn.metrics import f1_score
from sklearn.cluster import KMeans
from sklearn.pipeline import Pipeline
from sklearn.impute import SimpleImputer
from sklearn.preprocessing import StandardScaler
import numpy as np
from sklearn.ensemble import RandomForestClassifier

def run_random_forest(df, n_estimators = 100, min_samples_leaf = 1, max_features = '
df_train, df_test = train_test_split( df, test_size=0.33, random_state=42)
clf = RandomForestClassifier(n_estimators = n_estimators, min_samples_leaf = mi
max_features = max_features, max_depth = max_depth, r
clf = Pipeline([('impute', SimpleImputer( strategy='mean')),
                ('standardization', StandardScaler()),
                ('clf', clf),])

clf.fit(df_train.iloc[:, :-1], df_train.iloc[:, -1])

y_true = df_test.iloc[:, -1].tolist()
y_pred = clf.predict(df_test.iloc[:, :-1])

X_reduced = PCA(n_components=2).fit_transform(df_test)

pred_fig = plt.figure(1, figsize=(5, 5))
ax1 = pred_fig.add_subplot()
ax1.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_pred)
ax1.set_title(f"RandomForest pred\ n_estimators = {n_estimators}, min_samples
plt.close()

true_fig = plt.figure(2, figsize=(5, 5))
ax2 = true_fig.add_subplot()
ax2.scatter(X_reduced[:, 0], X_reduced[:, 1], s=10, c=y_true)
ax2.set_title(f"RandomForest true\ n_estimators = {n_estimators}, min_samples
plt.close()

f1 = f1_score(y_true, y_pred, average='weighted')

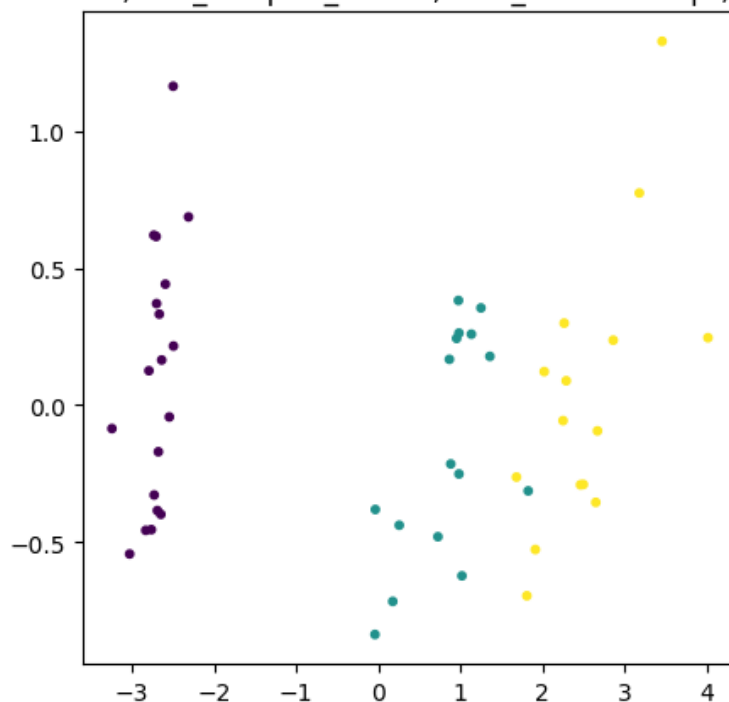
return {'f1':f1,
        'pred_fig':pred_fig, 'true_fig': true_fig,}

test_run = run_random_forest(df_iris)
print(test_run['f1'])
test_run['pred_fig']
```

0.98

Out[]:

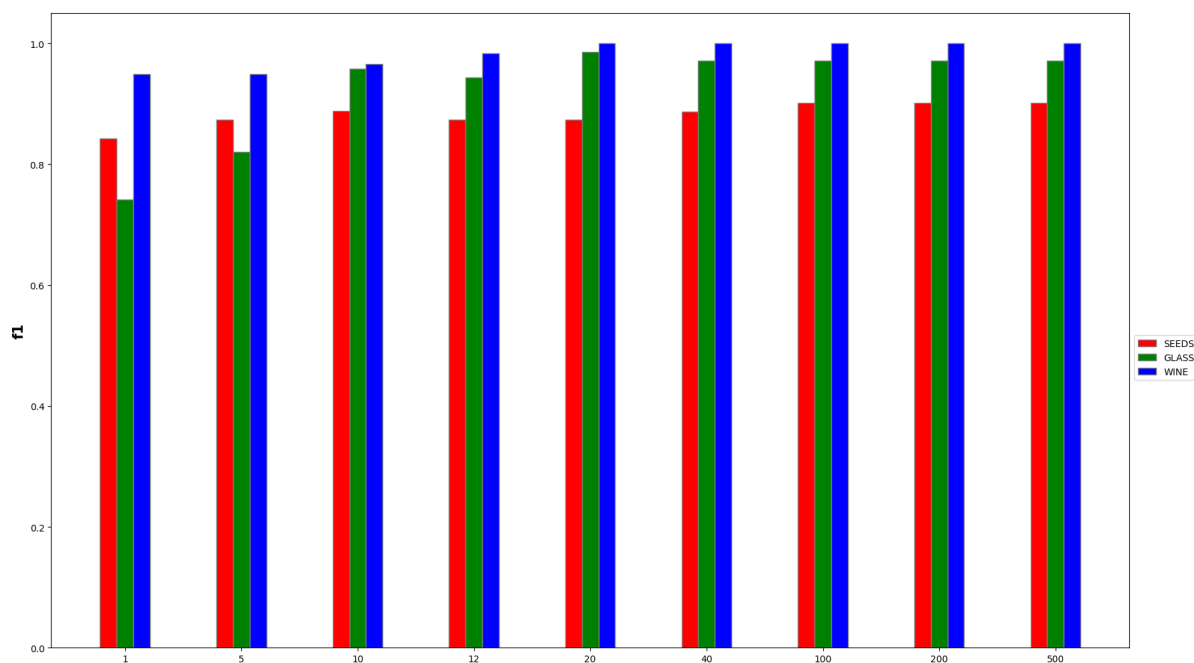
RandomForest pred
n_estimators = 100, min_samples_leaf=1, max_features=sqrt, max_depth=None



Random forest - liczba drzew

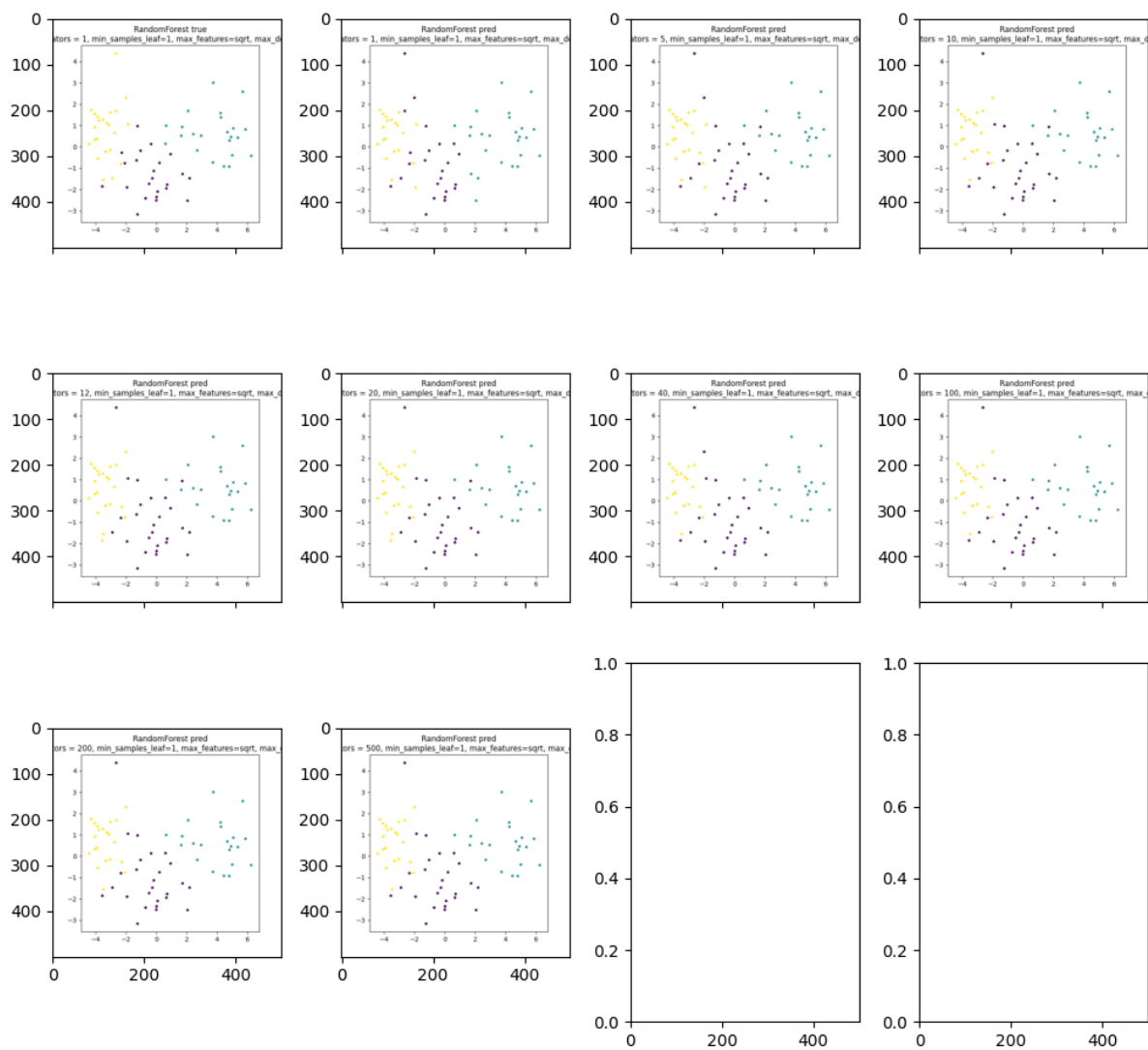
```
In [ ]: study_vals = [1, 5, 10, 12, 20, 40, 100, 200, 500]
results_list = [
    [run_random_forest (df_seeds, n_estimators = c) for c in study_vals],
    [run_random_forest (df_glass, n_estimators = c) for c in study_vals],
    [run_random_forest (df_wine, n_estimators = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



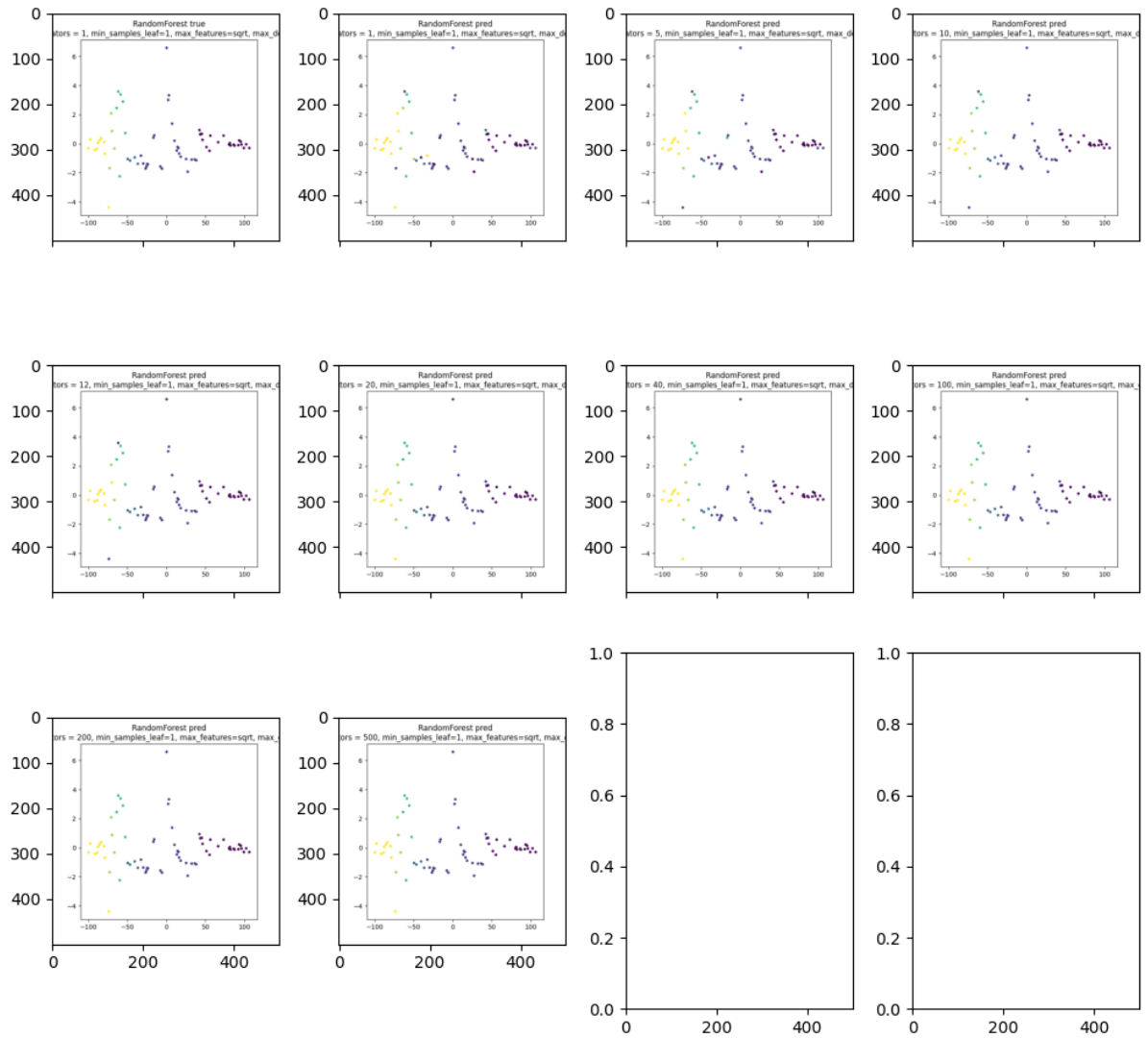
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



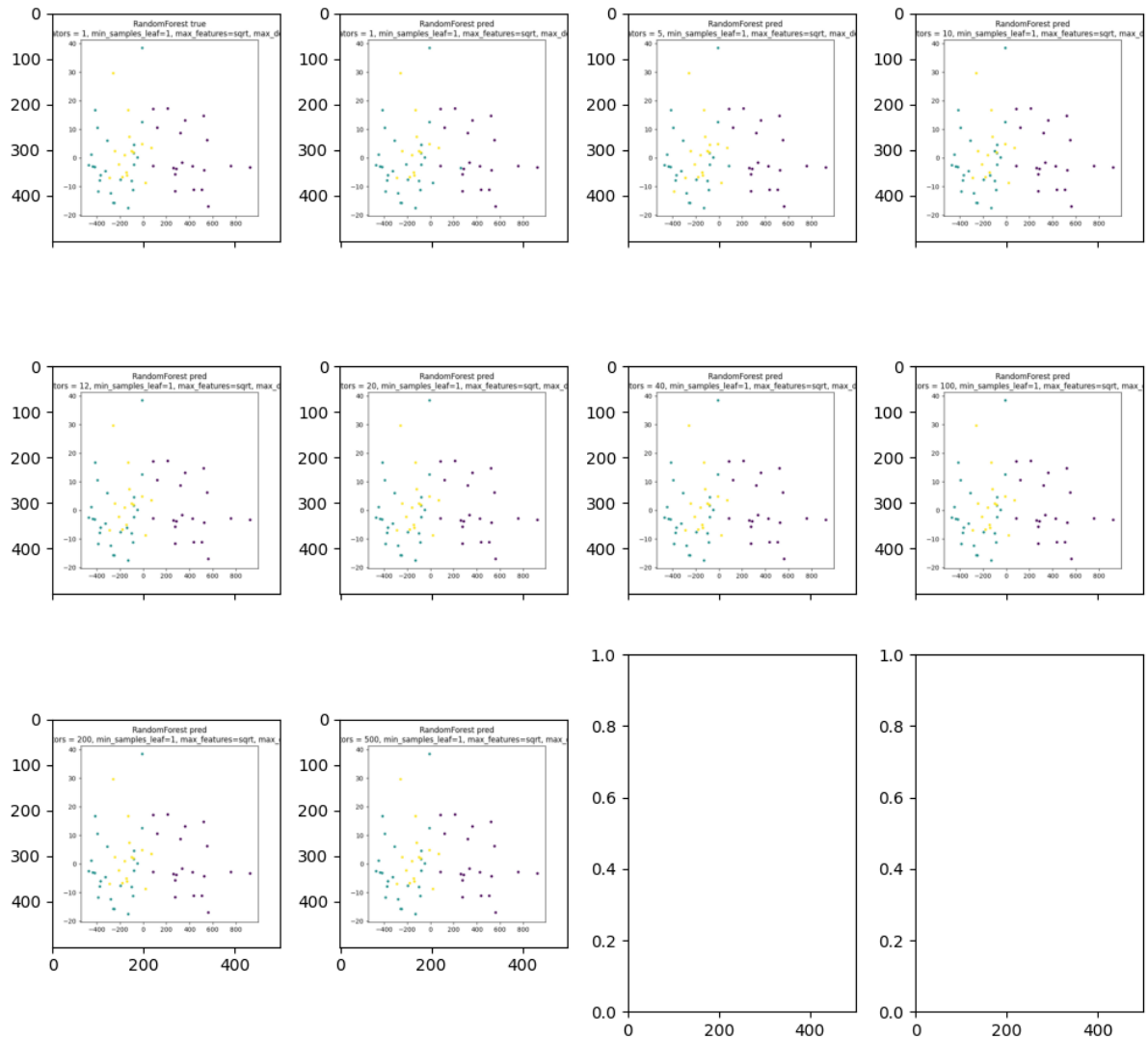
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

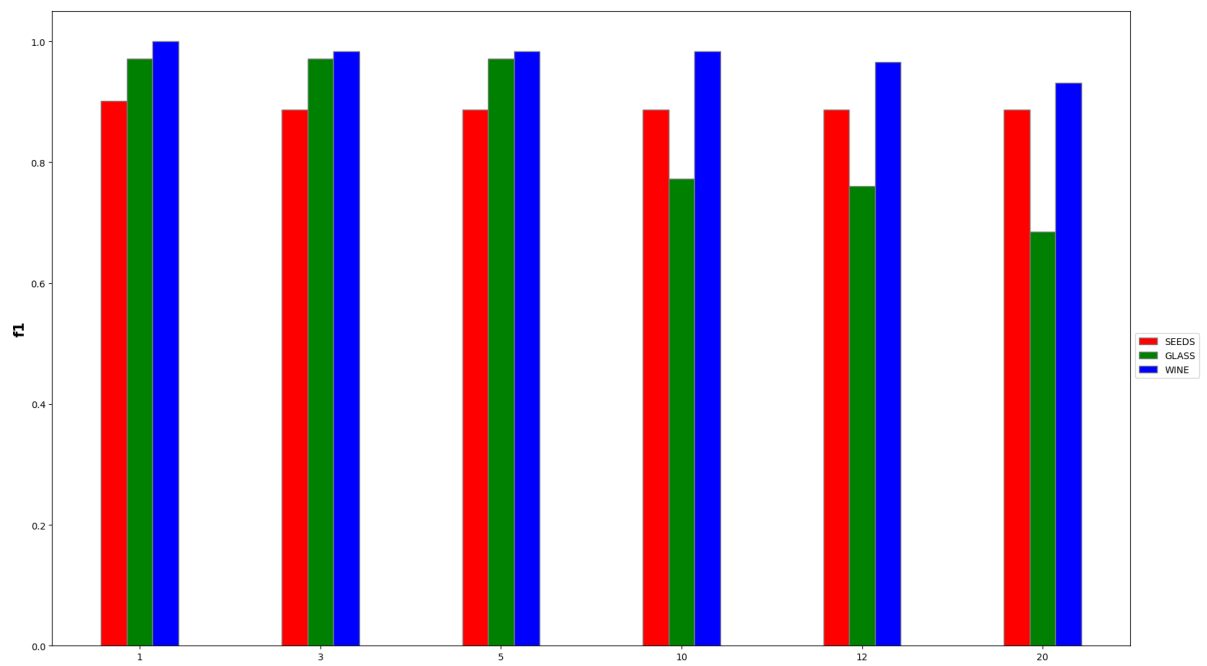

WINE



Random forest - liczba samplow

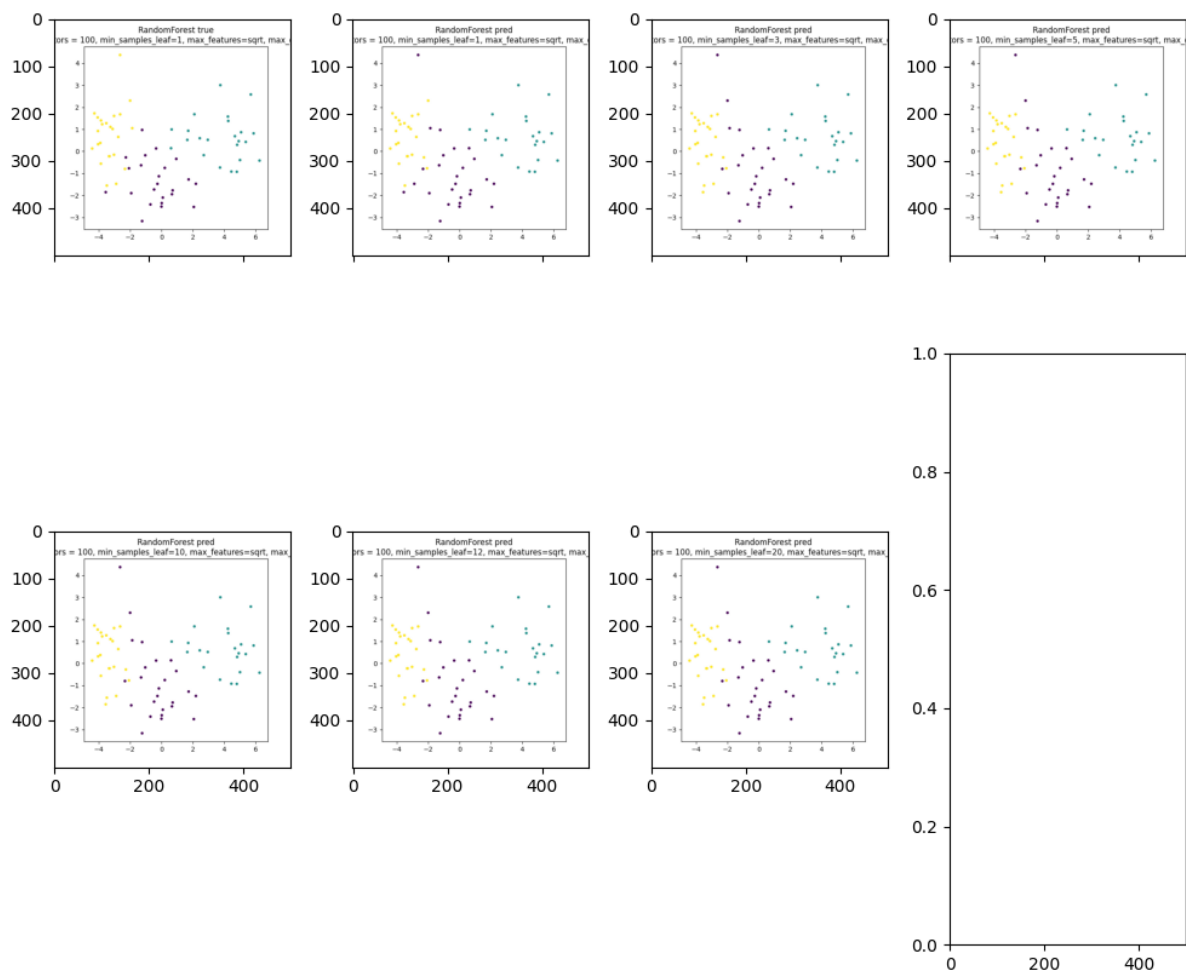
```
In [ ]: study_vals = [1, 3, 5, 10, 12, 20]
results_list = [
    [run_random_forest(df_seeds, min_samples_leaf = c) for c in study_vals],
    [run_random_forest(df_glass, min_samples_leaf = c) for c in study_vals],
    [run_random_forest(df_wine, min_samples_leaf = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



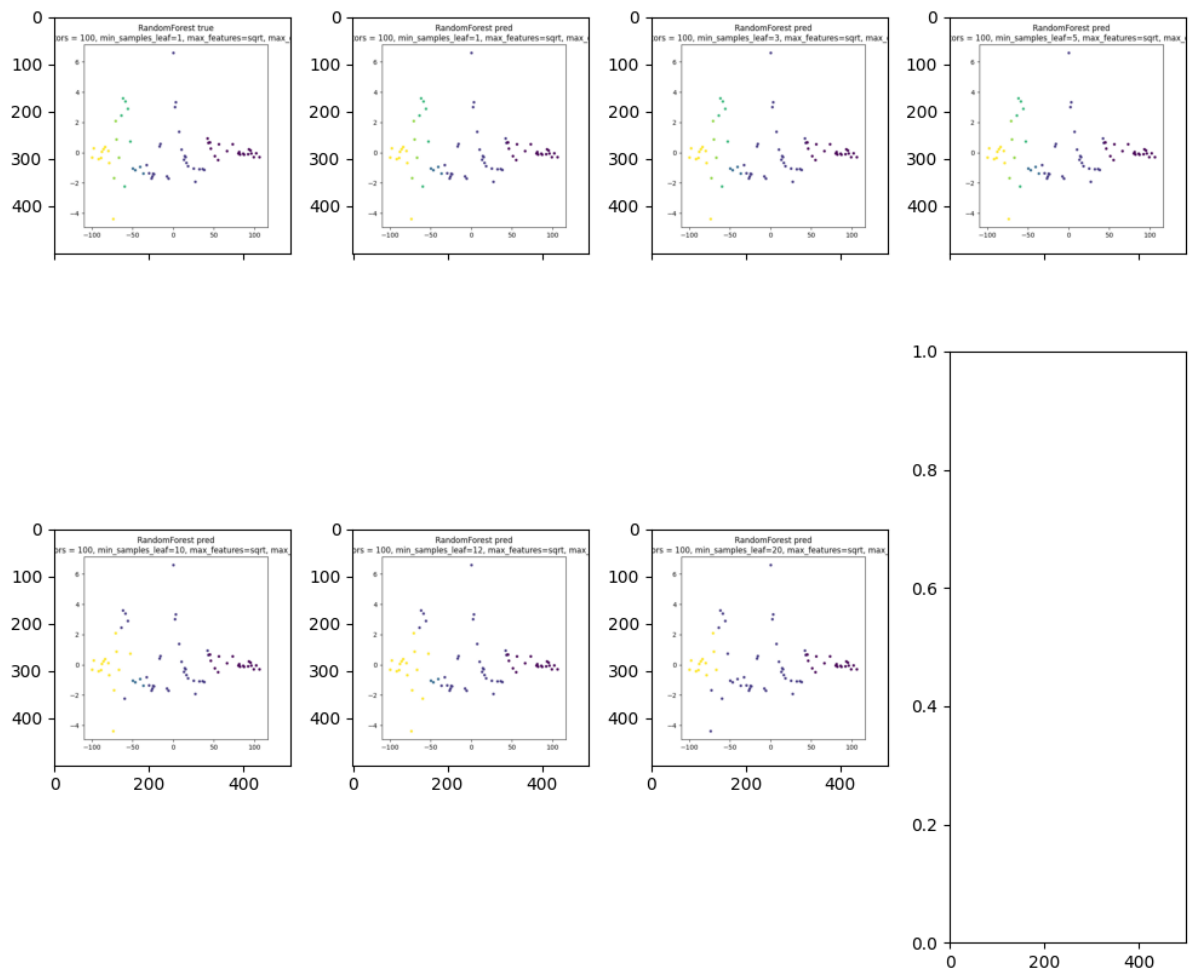
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



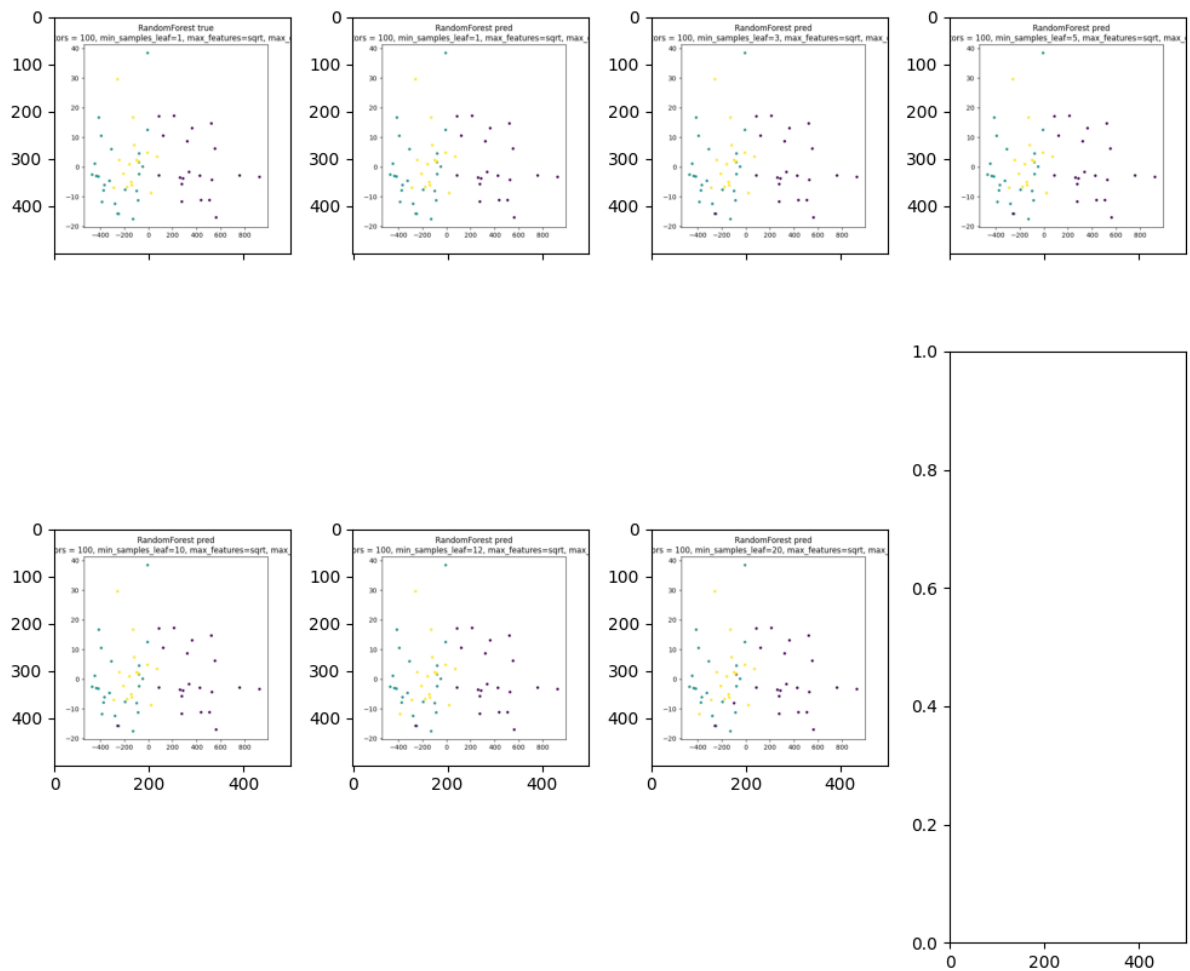
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

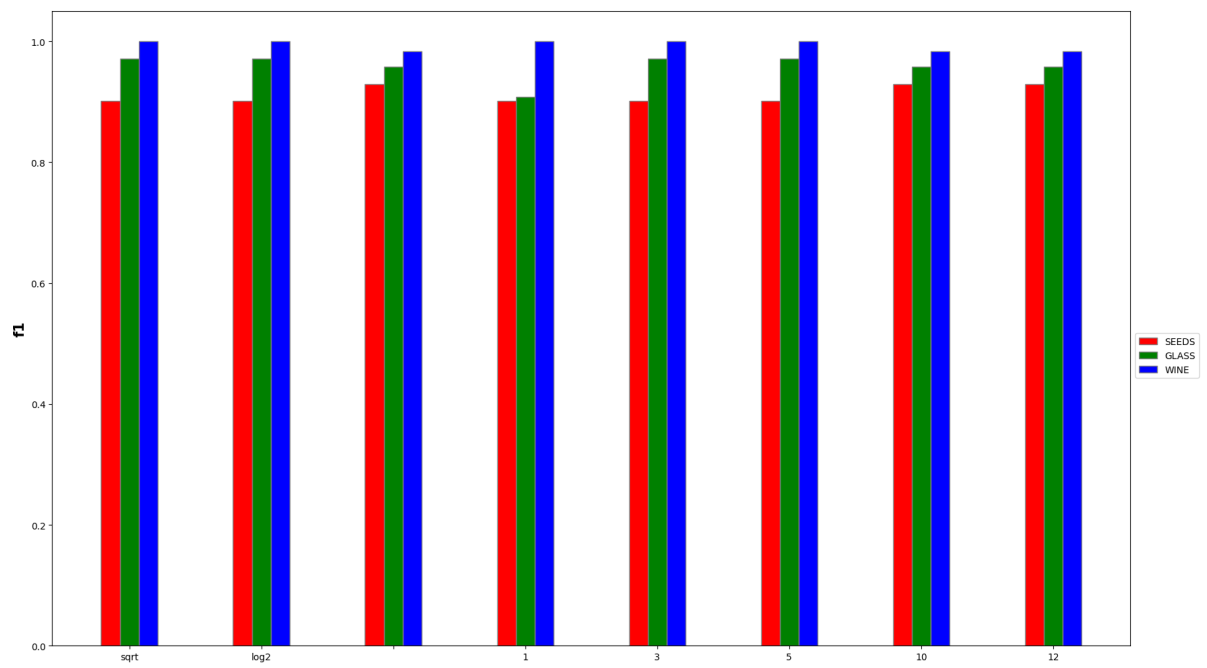
WINE



Random forest - liczba cech

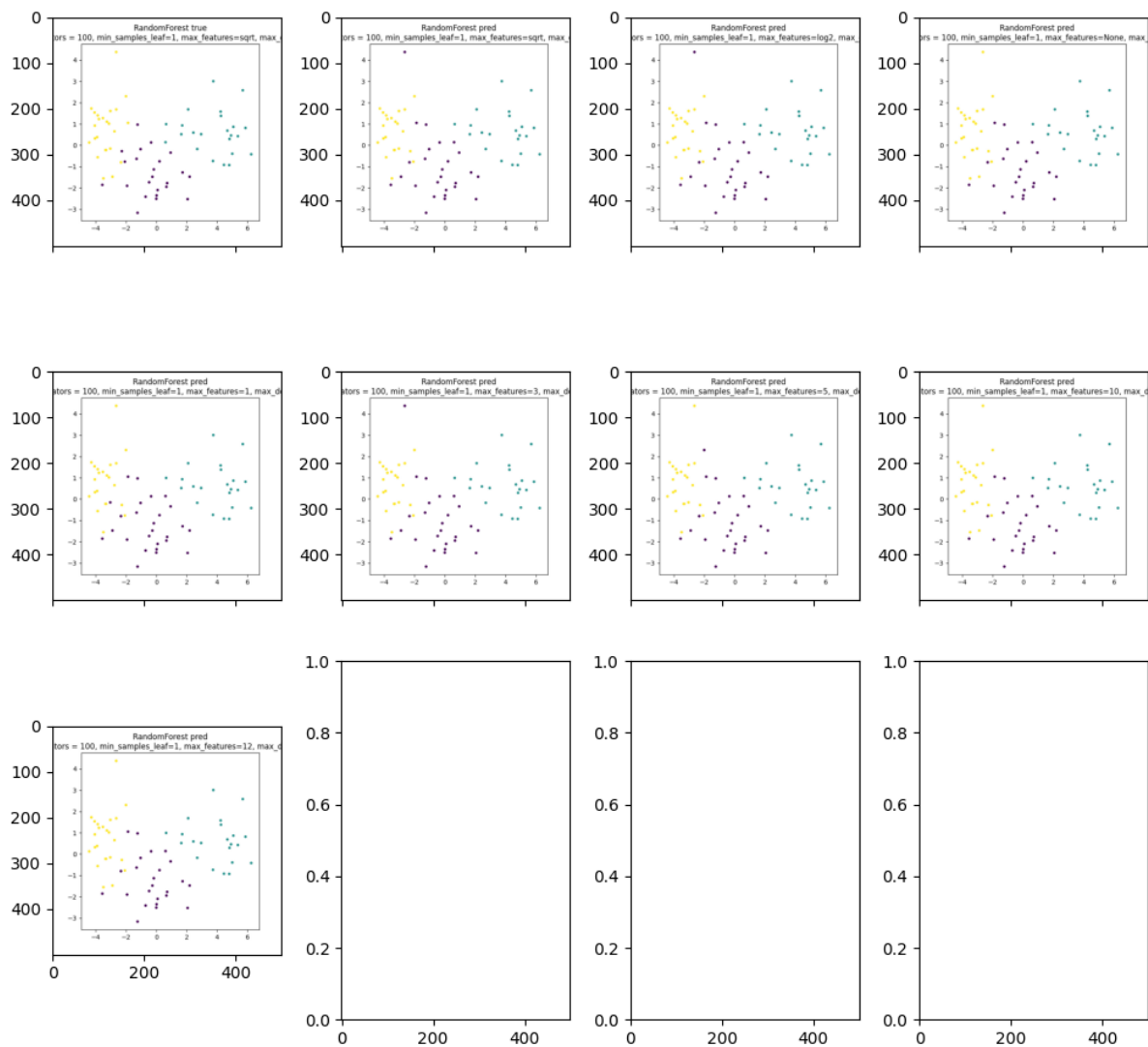
```
In [ ]: study_vals = ['sqrt', 'log2', None, 1, 3, 5, 10, 12]
results_list = [
    [run_random_forest (df_seeds, max_features = c) for c in study_vals],
    [run_random_forest (df_glass, max_features = c) for c in study_vals],
    [run_random_forest (df_wine, max_features = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



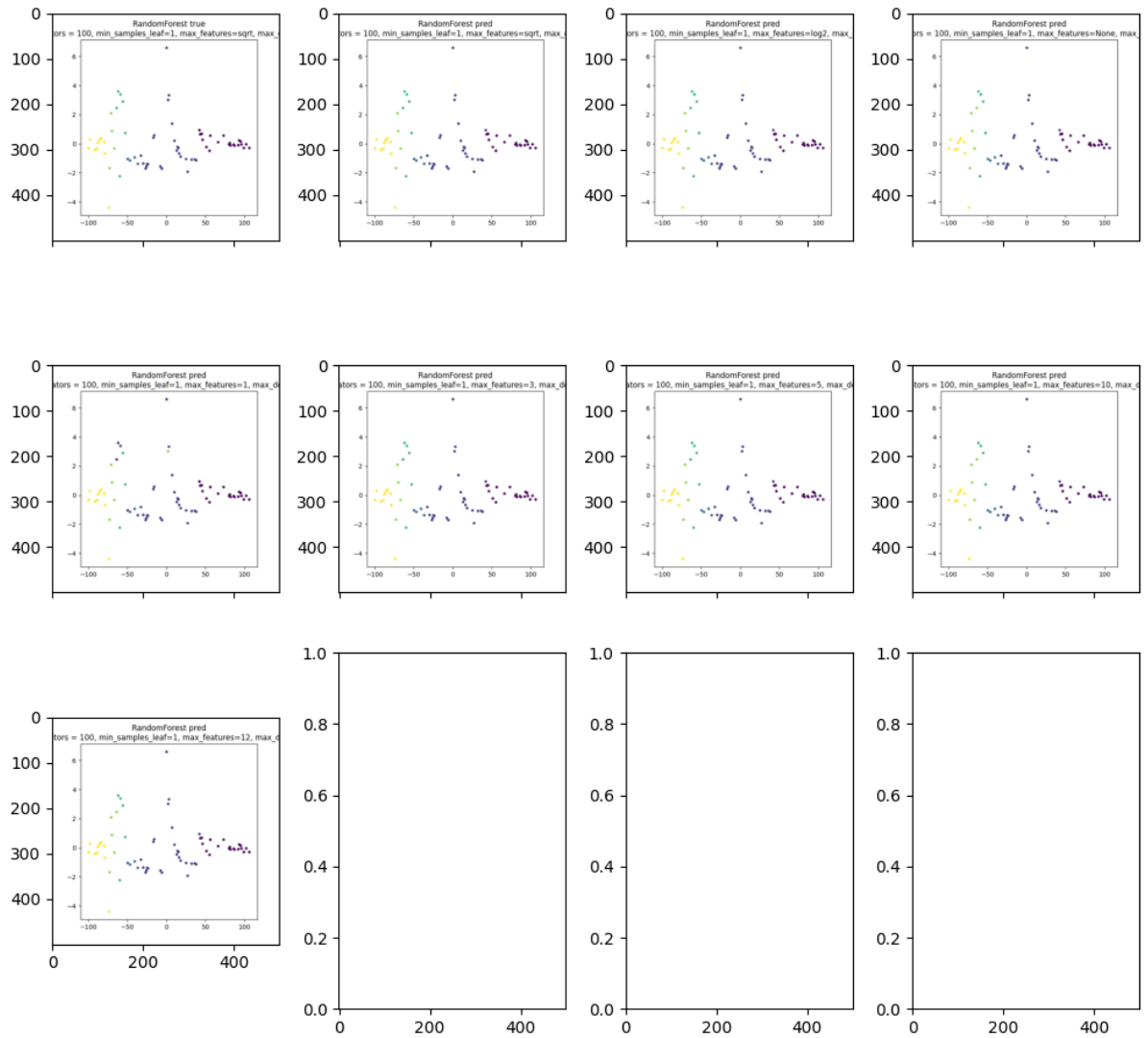
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



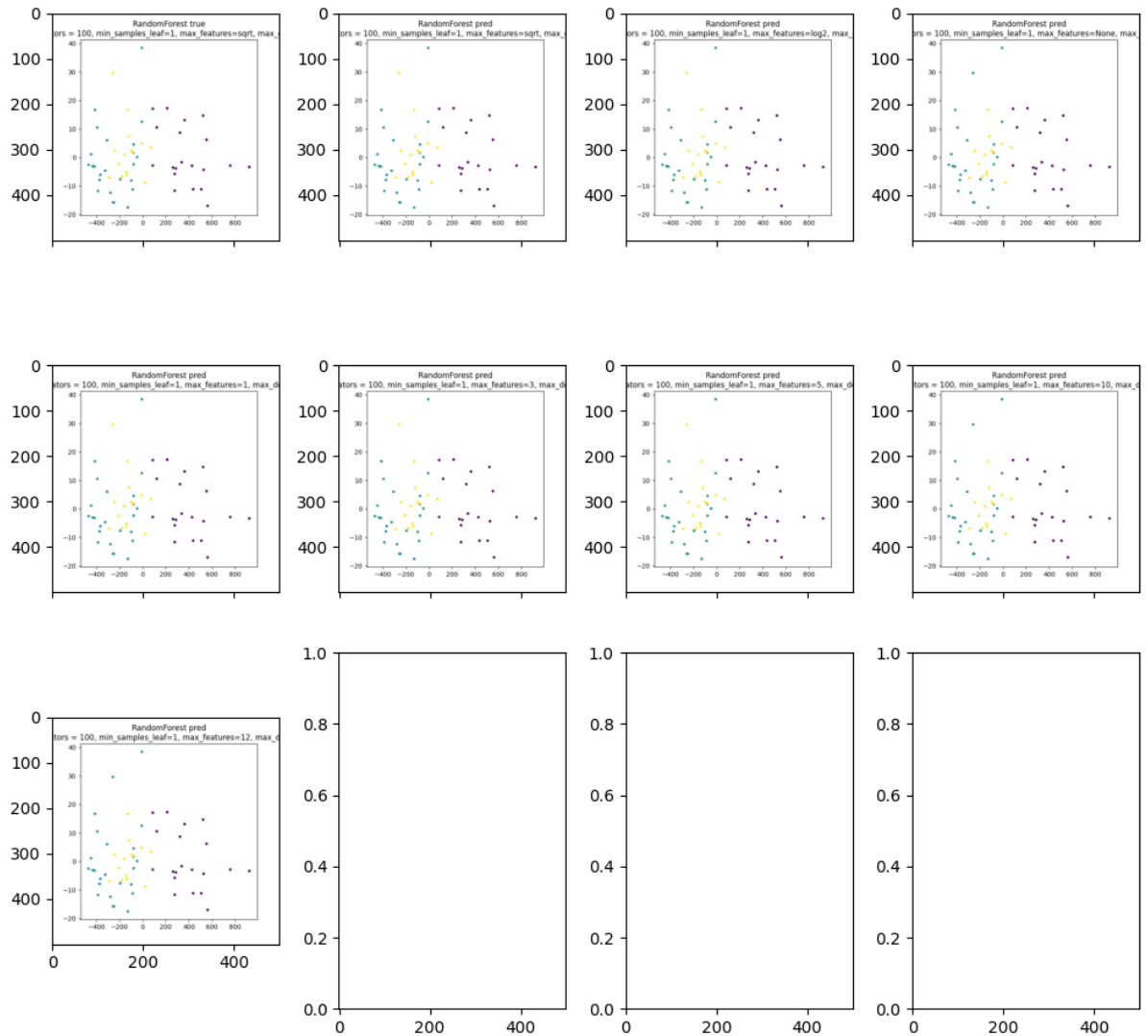
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

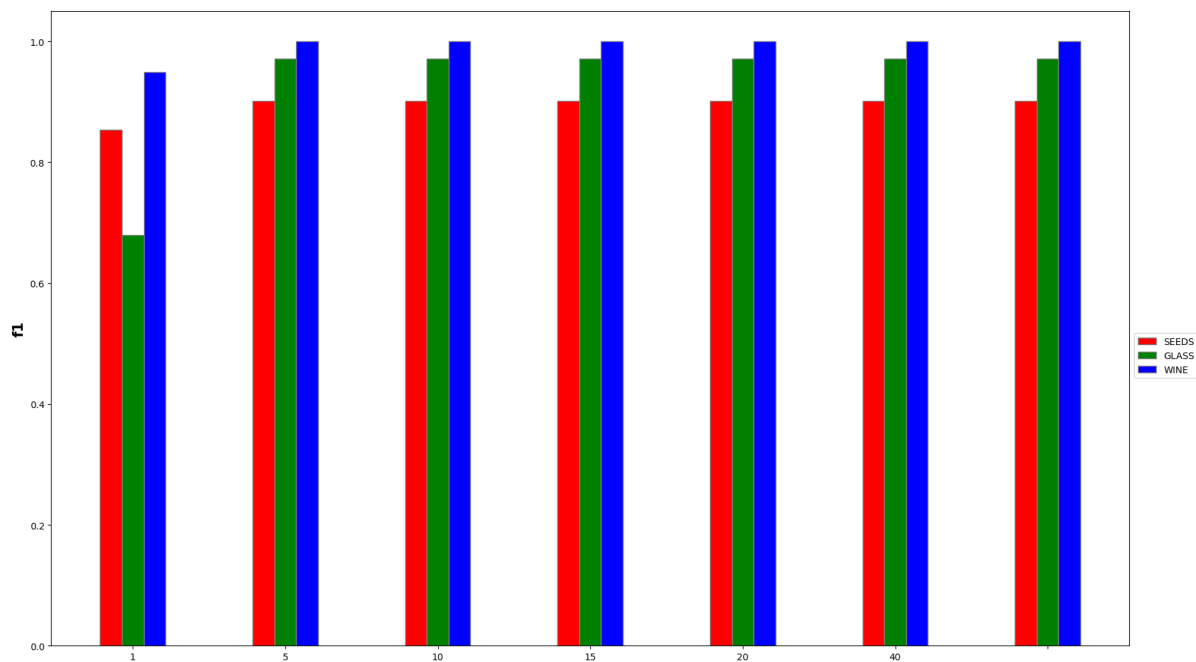

WINE



Random forest - glebokosc

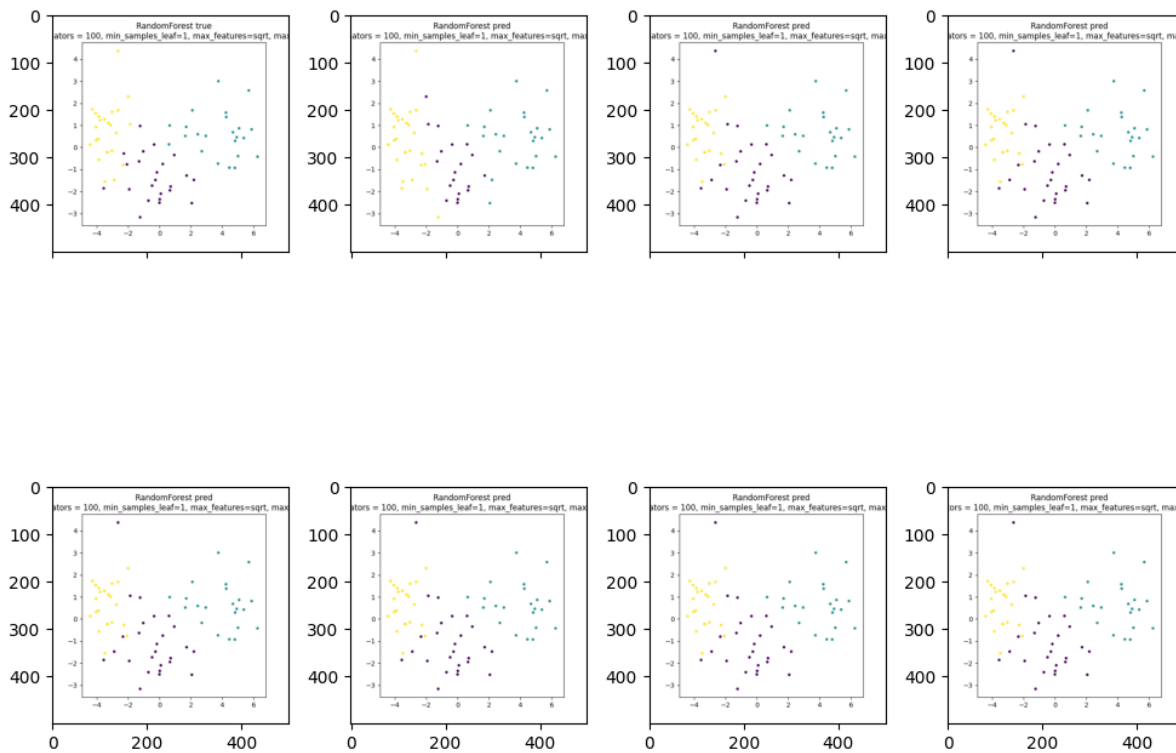
```
In [ ]: study_vals = [ 1, 5, 10, 15, 20, 40, None]
results_list = [
    [run_random_forest (df_seeds, max_depth = c) for c in study_vals],
    [run_random_forest (df_glass, max_depth = c) for c in study_vals],
    [run_random_forest (df_wine, max_depth = c) for c in study_vals],
]
```

```
In [ ]: plotStatistics(results_list, metrics_list, study_vals, dataset_list, colour_list)
```



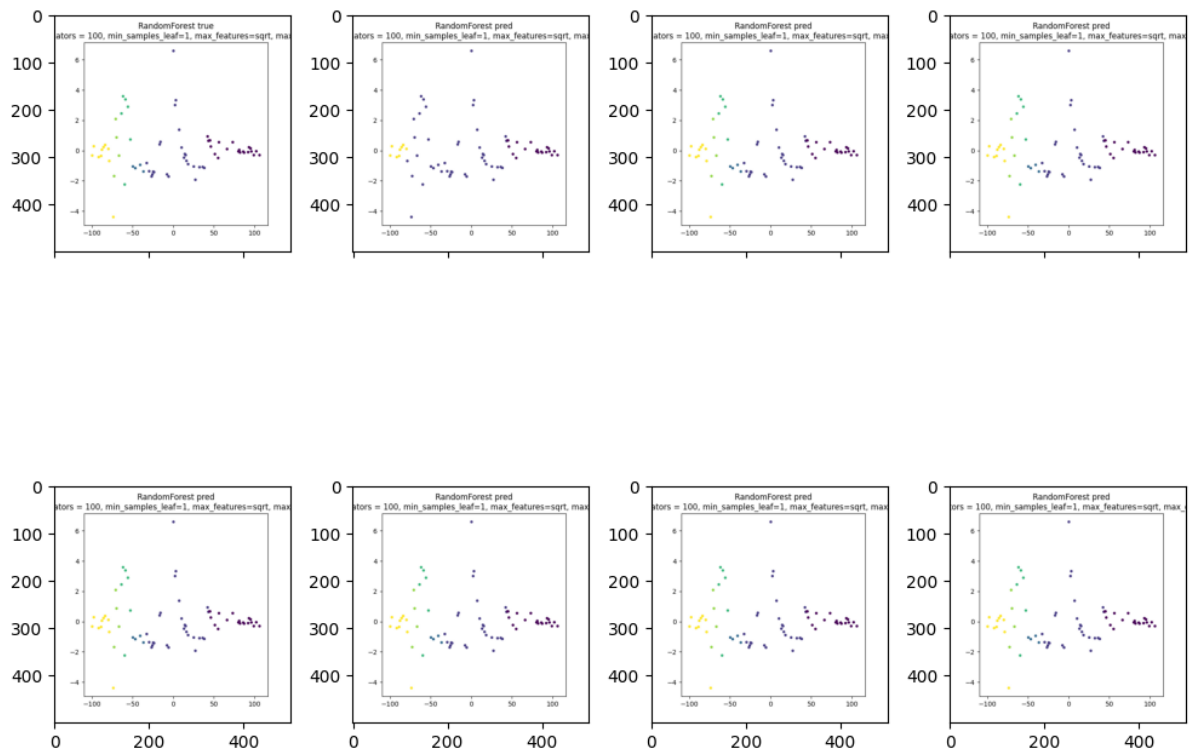
```
In [ ]: combineFigsInOnePlot(results_list[0], dataset_list[0])
```

SEEDS



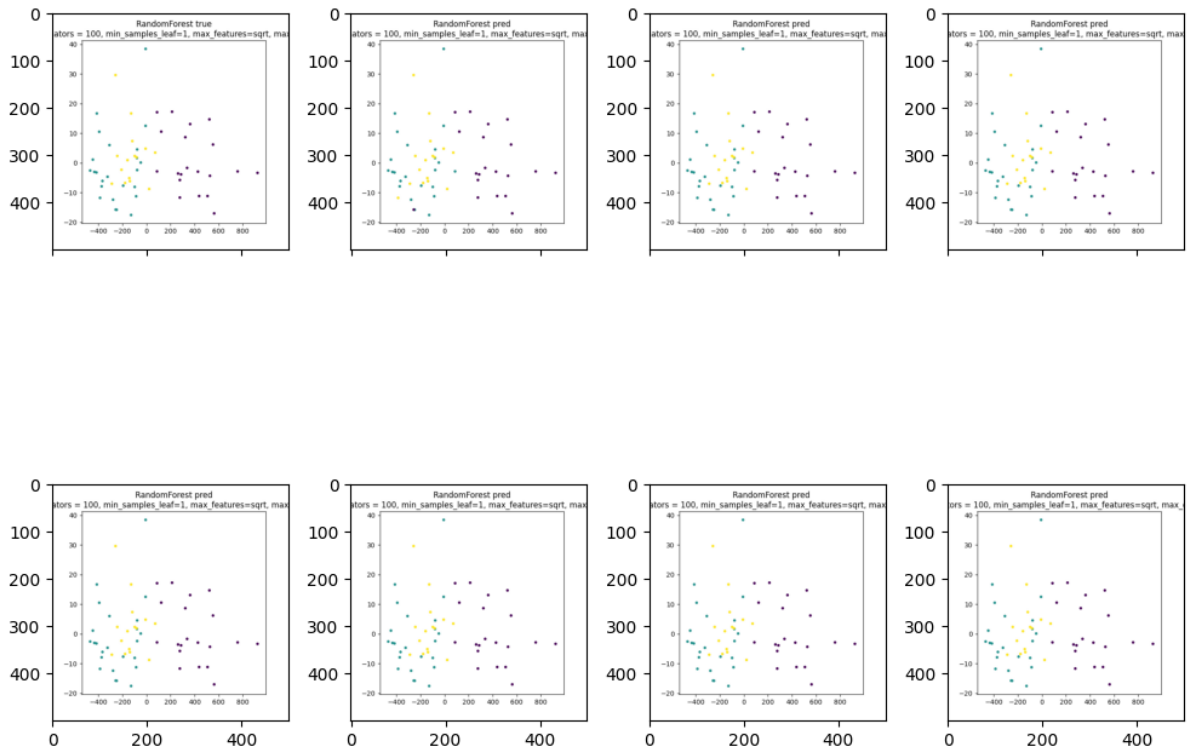
```
In [ ]: combineFigsInOnePlot(results_list[1], dataset_list[1])
```

GLASS



```
In [ ]: combineFigsInOnePlot(results_list[2], dataset_list[2])
```

WINE



```
In [ ]: score_metric = 'f1'
results_df = pd.DataFrame([
    [RunKNN(df_iris)[score_metric], run_bagging(df_iris)[score_metric], run_adaboost
    [RunKNN(df_seeds)[score_metric], run_bagging(df_seeds)[score_metric], run_adaboos
    [RunKNN(df_glass)[score_metric], run_bagging(df_glass)[score_metric], run_adaboos
    [RunKNN(df_wine)[score_metric], run_bagging(df_wine)[score_metric], run_adaboost
    ], index=['IRIS', 'SEEDS', 'GLASS', 'WINE'], columns = ['f1 KNN', 'f1 bagging', 'f1
results_df
```

	precision	recall	f1-score	support
0	1.00	1.00	1.00	19
1	0.94	1.00	0.97	15
2	1.00	0.94	0.97	16

accuracy			0.98	50
macro avg	0.98	0.98	0.98	50
weighted avg	0.98	0.98	0.98	50

Cross val: 0.9666666666666667

	precision	recall	f1-score	support
1	0.80	0.87	0.83	23
2	1.00	0.91	0.95	23
3	0.88	0.88	0.88	24

accuracy			0.89	70
macro avg	0.89	0.89	0.89	70
weighted avg	0.89	0.89	0.89	70

Cross val: 0.9047619047619048

	precision	recall	f1-score	support
1	1.00	0.91	0.95	22
2	0.93	1.00	0.96	25
3	1.00	1.00	1.00	4
5	1.00	0.83	0.91	6
6	0.80	1.00	0.89	4
7	1.00	1.00	1.00	10

accuracy			0.96	71
macro avg	0.95	0.96	0.95	71
weighted avg	0.96	0.96	0.96	71

Cross val: 0.9767214397496088

	precision	recall	f1-score	support
0	0.85	0.85	0.85	20
1	0.67	0.67	0.67	24
2	0.47	0.47	0.47	15

accuracy			0.68	59
macro avg	0.66	0.66	0.66	59
weighted avg	0.68	0.68	0.68	59

Cross val: 0.6402071563088512

Out[]:	f1 KNN	f1 bagging	f1 boost	f1 forest
IRIS	0.980000	1.000000	0.878487	0.980000
SEEDS	0.885714	0.887773	0.498168	0.901039
GLASS	0.957746	0.971702	0.702701	0.971702
WINE	0.677966	0.966270	0.895931	1.000000

In []: