Лабораторна робота #1

Базові алгоритми класифікації з використанням бібліотеки Scikit-learnt

Завдання роботи

- 1. [+] Завантажити дані, вивести назви колонок і розмір датасета
- 2. [+] Опрацювати пропуски (по можливості заповнити їх або видалити)
- 3. [+] Візуалізувати дані: побудувати графік (heatmap), що відображає кореляції ознак між собою і з цільовою змінною (розміткою); побудувати гістограми розподілу ознак і boxplot-и ознак відносно цільової змінної (якщо ознак занадто багато обмежитися декількома)
- 4. [+] Нормалізувати дані
- 5. Провести навчання наступних класифікаторів:
- [+] kNN
- [+] дерево ухвалення рішень
- [+] SVM
- [+] Random Forest
- [+] AdaBoost

Підібрати оптимальні параметри • для kNN • для SVM за допомогою GridSearch підібрати оптимальні «С» і «gamma» Серед обраних оптимальних моделей кожного класу вибрати найкращу.

[+] Відобразити sklearn.metrics.classification_report i sklearn.metrics.confusion_matrix

Хід роботи

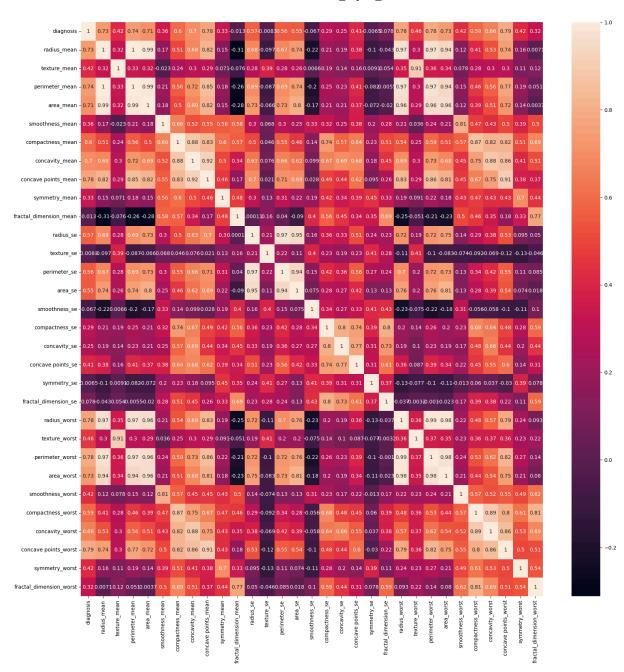
Для роботи я вирішив обрати наступний датасет https://www.kaggle.com/datasets/erdemtaha/cancer-data, оскільки вже використовував його у попередніх курсах із машинного навчання.

```
In [133... import pandas as pd

data = pd.read_csv('data/Cancer_Data.csv', delimiter=',')

In [134... print("Column Names:")
    print(data.columns.tolist())
    print("\nDataset Size:")
    print(data.shape)
```

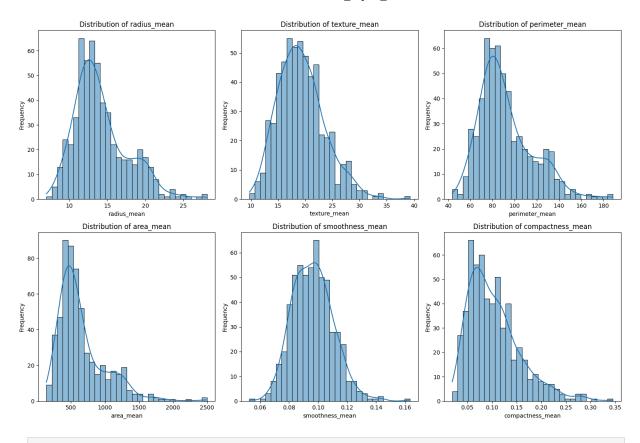
```
Column Names:
         ['id', 'diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean', 'area_mean', 's
         moothness_mean', 'compactness_mean', 'concavity_mean', 'concave points_mean', 'symme
         try_mean', 'fractal_dimension_mean', 'radius_se', 'texture_se', 'perimeter_se', 'are
         a_se', 'smoothness_se', 'compactness_se', 'concavity_se', 'concave points_se', 'symm
         etry_se', 'fractal_dimension_se', 'radius_worst', 'texture_worst', 'perimeter_wors
         t', 'area_worst', 'smoothness_worst', 'compactness_worst', 'concavity_worst', 'conca
         ve points_worst', 'symmetry_worst', 'fractal_dimension_worst', 'Unnamed: 32']
         Dataset Size:
         (569, 33)
In [135...
          data["diagnosis"] = data["diagnosis"].replace({"B": 0, "M": 1})
          print("\nSurv_status: \n0 = Benign cancer \n1 = Malignant cancer\n")
          y_column = 'diagnosis'
          data = data.dropna(subset=[y_column])
          data = data.drop(columns=['id', "Unnamed: 32"], errors='ignore')
          X_data = data.drop(y_column, axis=1)
          X_column = X_data.columns
          print("\nProcessed Data Size:")
          print(data.shape)
         Surv_status:
         0 = Benign cancer
         1 = Malignant cancer
         Processed Data Size:
         (569, 31)
         C:\Users\user\AppData\Local\Temp\ipykernel_5344\319634148.py:1: FutureWarning: Downc
         asting behavior in `replace` is deprecated and will be removed in a future version.
         To retain the old behavior, explicitly call `result.infer_objects(copy=False)`. To o
         pt-in to the future behavior, set `pd.set_option('future.no_silent_downcasting', Tru
         e)`
          data["diagnosis"] = data["diagnosis"].replace({"B": 0, "M": 1})
In [136...
          import seaborn as sns
          import matplotlib.pyplot as plt
          plt.figure(figsize=(20, 20))
          sns.heatmap(data.corr(), annot=True)
          plt.show()
```



```
In [137... features_to_plot = data.select_dtypes(include=['float64', 'int64']).columns[1:7] #

plt.figure(figsize=(15, 10))
for feature in features_to_plot:
    plt.subplot(2, 3, list(features_to_plot).index(feature) + 1)
    sns.histplot(data[feature], bins=30, kde=True)
    plt.title(f'Distribution of {feature}')
    plt.xlabel(feature)
    plt.ylabel('Frequency')

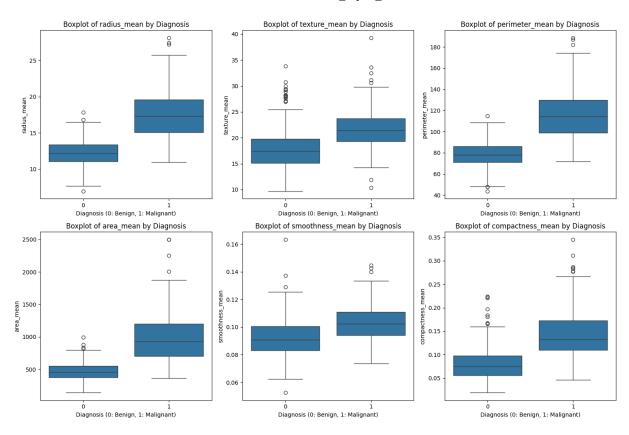
plt.tight_layout()
plt.show()
```



```
In [138... features_to_boxplot = data.select_dtypes(include=['float64', 'int64']).columns[1:7]

# Plot boxplots
plt.figure(figsize=(15, 10))
for feature in features_to_boxplot:
    plt.subplot(2, 3, list(features_to_boxplot).index(feature) + 1)
    sns.boxplot(x='diagnosis', y=feature, data=data)
    plt.title(f'Boxplot of {feature} by Diagnosis')
    plt.xlabel('Diagnosis (0: Benign, 1: Malignant)')
    plt.ylabel(feature)

plt.tight_layout()
plt.show()
```



```
In [139... from sklearn.preprocessing import StandardScaler

# Normalize the data
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X_data)

# Update the dataset with scaled features
data[X_column] = X_scaled

# Optionally, if you want to view the updated DataFrame
print("\nUpdated Data Sample:")
print(data.head())
```

```
Updated Data Sample:
   diagnosis radius_mean texture_mean perimeter_mean area_mean \
           1
                 1.097064
                                                          0.984375
0
                              -2.073335
                                               1.269934
                                                          1.908708
1
           1
                1.829821
                              -0.353632
                                               1.685955
2
           1
                1.579888
                               0.456187
                                               1.566503
                                                          1.558884
3
           1
                -0.768909
                               0.253732
                                              -0.592687 -0.764464
4
                1.750297
                              -1.151816
                                               1.776573
                                                          1.826229
   smoothness mean compactness mean concavity mean concave points mean \
0
          1.568466
                            3.283515
                                            2.652874
                                                                 2.532475
         -0.826962
                           -0.487072
                                           -0.023846
                                                                 0.548144
1
2
          0.942210
                            1.052926
                                            1.363478
                                                                 2.037231
          3.283553
                            3.402909
                                            1.915897
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          0.280372
                            0.539340
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   symmetry_mean ... radius_worst texture_worst perimeter_worst \
0
        2.217515 ...
                           1.886690
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                                                           2.303601
1
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   area_worst smoothness_worst compactness_worst concavity_worst \
                       1.307686
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                                          3.893397
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    1.220724
                       0.220556
                                         -0.313395
                                                           0.613179
   concave points_worst symmetry_worst fractal_dimension_worst
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               2.296076
                               2.750622
                                                        1.937015
                              -0.243890
1
               1.087084
                                                        0.281190
2
              1.955000
                               1.152255
                                                        0.201391
3
               2.175786
                               6.046041
                                                        4.935010
              0.729259
                              -0.868353
                                                       -0.397100
```

[5 rows x 31 columns]

Шаблон виводу метрик

Написав функцію для виводів, щоб не повторювати код.

```
In [140... from sklearn.metrics import accuracy_score, classification_report, confusion_matrix

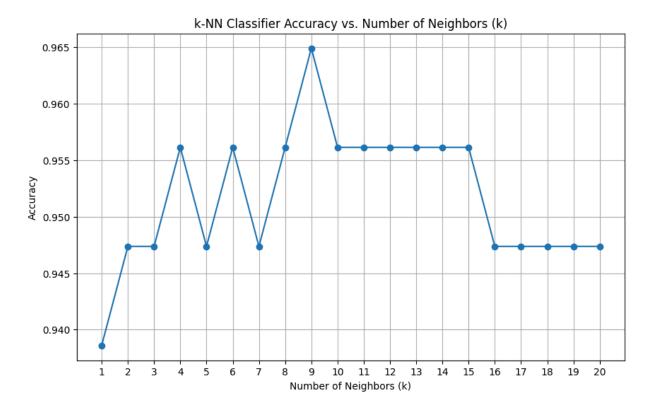
def print_evaluation_metrics(y_true, y_pred):
    accuracy = accuracy_score(y_true, y_pred)
    print(f"Accuracy: {accuracy:.2f}")

    print("Classification Report:")
    print(classification_report(y_true, y_pred))

    print("Confusion Matrix:")
    print(confusion_matrix(y_true, y_pred))
```

kNN

```
import numpy as np
In [141...
          import pandas as pd
          import matplotlib.pyplot as plt
          from sklearn.model_selection import train_test_split
          from sklearn.neighbors import KNeighborsClassifier
          from sklearn.metrics import accuracy score
          # Prepare the feature matrix (X) and target variable (y)
          X_data = data.drop('diagnosis', axis=1)
          y_data = data['diagnosis']
          X_train, X_test, y_train, y_test = train_test_split(X_data, y_data, test_size=0.2,
In [142...
          k_values = range(1, 21) # Trying k from 1 to 20
          accuracies = []
          # Evaluate k-NN for different values of k
          for k in k_values:
              knn = KNeighborsClassifier(n_neighbors=k)
              knn.fit(X_train, y_train)
              y pred = knn.predict(X test)
              accuracy = accuracy_score(y_test, y_pred)
              accuracies.append(accuracy)
              # print(f"Results for k = {k}:")
              # print_evaluation_metrics(y_test, y_pred)
              # print("\n" + "="*50 + "\n")
          # Plotting the accuracy vs. k
In [143...
          plt.figure(figsize=(10, 6))
          plt.plot(k_values, accuracies, marker='o')
          plt.title('k-NN Classifier Accuracy vs. Number of Neighbors (k)')
          plt.xlabel('Number of Neighbors (k)')
          plt.ylabel('Accuracy')
          plt.xticks(k_values)
          plt.grid()
          plt.show()
```



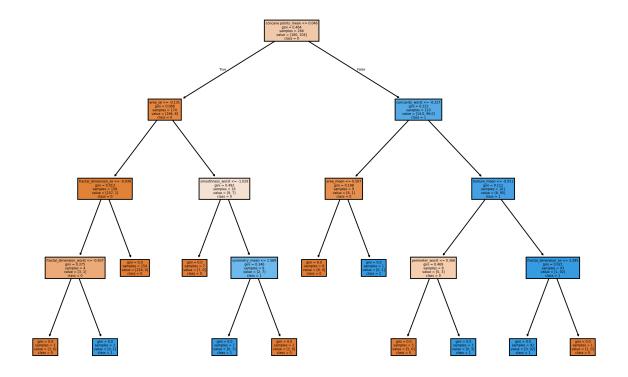
GridSearch для підбору оптимальних параметрів

```
In [144...
          param grid = {
              'n_neighbors': range(1, 21), # Testing neighbors from 1 to 25
              'weights': ['uniform', 'distance'],
              'algorithm': ['auto', 'ball_tree', 'kd_tree', 'brute'],
              'metric': ['euclidean', 'manhattan'], # You can add more metrics if desired
          knn = KNeighborsClassifier()
In [145...
          from sklearn.model_selection import train_test_split, GridSearchCV
          # Perform Grid Search with cross-validation
          grid_search = GridSearchCV(knn, param_grid, cv=3, scoring='accuracy', verbose=1, n_
          grid_search.fit(X_train, y_train)
          # Output the best parameters and the best score
          print("Best Parameters:", grid_search.best_params_)
          print("Best Cross-Validation Score:", grid_search.best_score_)
         Fitting 3 folds for each of 320 candidates, totalling 960 fits
         Best Parameters: {'algorithm': 'auto', 'metric': 'euclidean', 'n_neighbors': 7, 'wei
         ghts': 'uniform'}
         Best Cross-Validation Score: 0.9647670500755199
In [146...
          # Make predictions with the best parameters on the test set
          best_knn = grid_search.best_estimator_
          y_pred = best_knn.predict(X_test)
          print_evaluation_metrics(y_test, y_pred)
```

```
Accuracy: 0.95
Classification Report:
             precision
                         recall f1-score
                                            support
          0
                  0.96
                            0.96
                                      0.96
                                                 71
                  0.93
                                      0.93
                            0.93
                                                 43
                                      0.95
                                                 114
   accuracy
                  0.94
                            0.94
                                      0.94
  macro avg
                                                 114
weighted avg
                  0.95
                            0.95
                                      0.95
                                                 114
Confusion Matrix:
[[68 3]
[ 3 40]]
```

Дерево ухвалення рішень (Decision Tree) та ансамбль дерев ухвалення рішень (Random Forest)

Decision Tree



Accuracy: 0.93

Classification Report:

	precision	recall	f1-score	support
0	0.94	0.94	0.94	177
1	0.90	0.91	0.90	108
accuracy			0.93	285
macro avg	0.92	0.92	0.92	285
weighted avg	0.93	0.93	0.93	285

Confusion Matrix:

[[166 11] [10 98]]

Тестовий код для того, щоб побачити аутпут predict_proba

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In [148... predict_proba = clf.predict_proba(X_test)
    print(predict_proba)
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```

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Random Forest

```
In [149... from sklearn.ensemble import RandomForestClassifier

X_train, X_test, y_train, y_test = train_test_split(X_data, data[y_column], test_si

rf_classifier = RandomForestClassifier(n_estimators=10, random_state=31337)

rf_classifier.fit(X_train, y_train)

predictions = rf_classifier.predict(X_test)

print_evaluation_metrics(y_test, predictions)
```

Accuracy: 0.92

Classification Report:

	precision	recall	†1-score	support
0	0.90	0.98	0.94	186
1	0.96	0.80	0.87	99
accuracy			0.92	285
macro avg	0.93	0.89	0.91	285
weighted avg	0.92	0.92	0.92	285

Confusion Matrix:

[[183 3]

[20 79]]

AdaBoost

```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.ensemble import AdaBoostClassifier

X_data = data.drop('diagnosis', axis=1)
y_data = data['diagnosis']

X_train, X_test, y_train, y_test = train_test_split(X_data, y_data, test_size=0.2,
```

```
In [151... ada_boost_model = AdaBoostClassifier(n_estimators=50, random_state=42)
    ada_boost_model.fit(X_train, y_train)
```

C:\Users\user\AppData\Local\Packages\PythonSoftwareFoundation.Python.3.12_qbz5n2kfra 8p0\LocalCache\local-packages\Python312\site-packages\sklearn\ensemble_weight_boost ing.py:527: FutureWarning: The SAMME.R algorithm (the default) is deprecated and wil 1 be removed in 1.6. Use the SAMME algorithm to circumvent this warning. warnings.warn(

```
Out[151...
                  AdaBoostClassifier
          AdaBoostClassifier(random_state=42)
```

```
In [152... y_pred = ada_boost_model.predict(X_test)
          print_evaluation_metrics(y_test, y_pred)
```

Accuracy: 0.97

Classification Report:

	precision	recall	f1-score	support
0	0.97	0.99	0.98	71
1	0.98	0.95	0.96	43
accuracy			0.97	114
macro avg	0.97	0.97	0.97	114
weighted avg	0.97	0.97	0.97	114

Confusion Matrix:

[[70 1] [2 41]]

SVM

```
In [153...
         X_data = data.drop('diagnosis', axis=1)
          y_data = data['diagnosis']
          X_train, X_test, y_train, y_test = train_test_split(X_data, y_data, test_size=0.2,
In [154...
         from sklearn.svm import SVC
          # Set up the parameter grid for GridSearchCV
          param_grid = {
              'C': [0.01, 0.1, 1, 10, 100], # Regularization parameter
              'gamma': [0.001, 0.01, 0.1, 1], # Kernel coefficient
          }
          svm = SVC(kernel='rbf', random_state=31337)
In [155...
         grid_search = GridSearchCV(svm, param_grid, cv=5, scoring='accuracy', verbose=1, n_
          grid_search.fit(X_train, y_train)
          print("Best Parameters:", grid_search.best_params_)
          print("Best Cross-Validation Score:", grid_search.best_score_)
         Fitting 5 folds for each of 20 candidates, totalling 100 fits
         Best Parameters: {'C': 10, 'gamma': 0.01}
         Best Cross-Validation Score: 0.9736263736263737
In [156... best_svm = grid_search.best_estimator_
          y_pred = best_svm.predict(X_test)
```

print_evaluation_metrics(y_test, y_pred)

Accuracy: 0.98

Classification Report:

support	f1-score	recall	precision	
71	0.99	1.00	0.97	0
	0.98	0.95	1.00	1
3 114	0.98			2661192614
	0.98	0.98	0.99	accuracy macro avg
3 114	0.98	0.98	0.98	weighted avg

Confusion Matrix:

[[71 0] [2 41]]