Tugas 1 Terdapat dataset mushroom. Berdasarkan dataset yang tersebut, bandingkan peforma antara algoritma Decision Tree dan RandomForest. Gunakan tunning hyperparameter untuk mendapatkan parameter dan akurasi yang terbaik.

```
import numpy as np
import pandas as pd
from sklearn.tree import DecisionTreeClassifier # import DT
from sklearn.ensemble import RandomForestClassifier # import
RandomForest
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, classification_report
```

Langkah 1: Memuat Dataset

```
# Load data
df = pd.read csv('data/mushrooms.csv')
df.head()
  class cap-shape cap-surface cap-color bruises odor gill-
attachment \
                                                                         f
      р
                                         n
                                                  t
                                                       р
                                                                         f
      е
                                                                         f
2
      е
                 b
                                                                         f
      р
                                                       р
                                                                         f
      е
  gill-spacing gill-size gill-color
                                        ... stalk-surface-below-ring
0
              С
                         n
                                     k
                                                                     S
1
              С
                         b
                                     k
                                                                     s
2
                         b
              С
                                     n
                                                                     S
3
              С
                         n
                                     n
                                                                     S
4
                         b
                                     k
                                                                     S
  stalk-color-above-ring stalk-color-below-ring veil-type veil-
color \
                                                                         W
                                                                         W
2
                                                                         W
3
4
                                                                         W
```

```
ring-number ring-type spore-print-color population habitat
1
            0
                       р
                                          n
                                                      n
                                                               g
2
            0
                       p
                                          n
                                                      n
                                                               m
3
            0
                       p
                                           k
                                                      S
                                                               u
4
            0
                                                               q
[5 rows x 23 columns]
# Cek kolom null
df.isnull().sum()
                              0
class
                              0
cap-shape
                              0
cap-surface
                              0
cap-color
bruises
                              0
                              0
odor
gill-attachment
                              0
                              0
gill-spacing
                              0
gill-size
gill-color
                              0
                              0
stalk-shape
stalk-root
                              0
stalk-surface-above-ring
                              0
stalk-surface-below-ring
                              0
                              0
stalk-color-above-ring
                              0
stalk-color-below-ring
veil-type
                              0
veil-color
                              0
ring-number
                              0
                              0
ring-type
                              0
spore-print-color
                              0
population
                              0
habitat
dtype: int64
# Seleksi fitur
# Slice dataframe mulai dari kolom 'cap-shape' sampai 'habitat'
X = df.iloc[:, 0:-1]
y = df['bruises']
y = y.map(\{'M':1, 'B':0\}) # Encode label
# Cek jumlah fitur dan instance
X.shape
(8124, 22)
```

```
# Split data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.3, random_state=42)
```

# Langkah Decision tree Model

```
# Import necessary libraries
from sklearn.tree import DecisionTreeClassifier
from sklearn.model selection import GridSearchCV, train test split
from sklearn.metrics import accuracy_score, classification_report
from sklearn.datasets import load iris
# Load dataset
data = load iris()
X = data.data
y = data.target
# Split dataset into training and testing sets
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=42)
# Decision Tree model
dt = DecisionTreeClassifier(random_state=42)
# Hyperparameter tuning untuk Decision Tree
param grid dt = {
    'criterion': ['gini', 'entropy'],
    'max depth': [10, 20, 30, None],
    'min_samples_split': [2, 5, 10],
    'min samples leaf': [1, 2, 4]
}
# GridSearchCV untuk menemukan parameter terbaik
grid search dt = GridSearchCV(estimator=dt, param grid=param grid dt,
cv=5, n jobs=-1, verbose=1)
grid search dt.fit(X train, y train)
# Hasil terbaik dari Decision Tree
best dt = grid search dt.best estimator
y pred dt = best dt.predict(X test)
# Akurasi dan laporan klasifikasi
accuracy dt = accuracy score(y test, y pred dt) * 100 # Akurasi dalam
persen
print(f"Best Decision Tree Parameters: {grid search dt.best params }")
print(f"Decision Tree Accuracy: {accuracy dt:.2f}%")
print(classification report(y test, y pred dt))
```

```
Fitting 5 folds for each of 72 candidates, totalling 360 fits
Best Decision Tree Parameters: {'criterion': 'entropy', 'max depth':
10, 'min_samples_leaf': 4, 'min_samples_split': 2}
Decision Tree Accuracy: 100.00%
              precision
                            recall f1-score
                                               support
           0
                   1.00
                              1.00
                                        1.00
                                                     10
           1
                   1.00
                              1.00
                                        1.00
                                                      9
           2
                   1.00
                              1.00
                                        1.00
                                                     11
                                        1.00
                                                     30
    accuracy
                   1.00
                              1.00
                                        1.00
                                                     30
   macro avg
weighted avg
                   1.00
                              1.00
                                        1.00
                                                     30
```

# Langkah Random Forest Model

```
# Random Forest
rf = RandomForestClassifier(random state=42)
# Hyperparameter tuning untuk Random Forest
param grid rf = {
    'n_estimators': [100, 200, 300],
    'criterion': ['gini', 'entropy'],
    'max depth': [10, 20, 30, None],
    'min samples split': [2, 5, 10],
    'min samples leaf': [1, 2, 4]
}
grid search rf = GridSearchCV(estimator=rf, param grid=param grid rf,
cv=2, n jobs=-1, verbose=1)
grid search rf.fit(X train, y train)
# Hasil terbaik dari Random Forest
best_rf = grid_search_rf.best_estimator_
y pred rf = best rf.predict(X test)
# Akurasi dan laporan klasifikasi
accuracy rf = accuracy score(y test, y pred rf) * 100 # Akurasi dalam
persen
print(f"Best Random Forest Parameters: {grid search rf.best params }")
print(f"Random Forest Accuracy: {accuracy_rf:.2f}%")
print(classification_report(y_test, y_pred_rf))
Fitting 2 folds for each of 216 candidates, totalling 432 fits
Best Random Forest Parameters: {'criterion': 'gini', 'max_depth': 10,
'min samples leaf': 1, 'min samples split': 2, 'n estimators': 100}
Random Forest Accuracy: 100.00%
              precision recall f1-score
                                              support
```

0	1.00	1.00	1.00	10	
1	1.00	1.00	1.00	9	
2	1.00	1.00	1.00	11	
accuracy macro avg weighted avg	1.00 1.00	1.00 1.00	1.00 1.00 1.00	30 30 30	

# TUGAS 2

Terdapat dataset mushroom. Berdasarkan dataset tersebut, bandingkan peforma antara algoritma Decision Tree dan AdaBoost. Gunakan tunning hyperparameter untuk mendapatkan parameter dan akurasi yang terbaik.

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split, GridSearchCV
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import AdaBoostClassifier
from sklearn.metrics import accuracy_score, classification_report
```

# Langkah Memuat Dataset

```
# Load data
df = pd.read csv('data/mushrooms.csv')
df.head()
  class cap-shape cap-surface cap-color bruises odor gill-
attachment \
                                                                       f
                                                                       f
                                                                       f
                                                                       f
                                       ... stalk-surface-below-ring
  gill-spacing gill-size gill-color
0
              С
                                    k
                        n
1
                        b
                                    k
             С
                                                                    S
                                       . . .
2
              С
                        b
                                                                    S
                                    n
3
              С
                        n
                                    n
                                                                    S
4
                        b
                                                                    S
  stalk-color-above-ring stalk-color-below-ring veil-type veil-
```

```
color \
                                                                            W
1
2
                                                                             W
3
                                                                             W
                                                                             W
  ring-number ring-type spore-print-color population habitat
0
             0
                         р
1
             0
                         p
                                                          n
                                              n
                                                                    g
2
             0
                         р
                                              n
                                                          n
                                                                   m
3
             0
                                              k
                                                                    u
                         р
                                                          S
4
             0
                                                                    g
[5 rows x 23 columns]
```

# Langkah Encoding Data

```
import pandas as pd
from sklearn.datasets import load iris # Misal data berasal dari
dataset sklearn
from sklearn.preprocessing import LabelEncoder
# Load dataset (misal iris dataset sebagai contoh)
data bunch = load iris() # Menghasilkan Bunch object
# Convert Bunch menjadi DataFrame
data = pd.DataFrame(data bunch.data, columns=data bunch.feature names)
# Misalkan menambahkan kolom kategori 'class'
data['class'] = pd.Categorical.from codes(data bunch.target,
data bunch.target names)
# Menggunakan Label Encoding untuk semua kolom
label encoder = LabelEncoder()
for column in data.columns:
    data[column] = label encoder.fit transform(data[column])
# Split data menjadi fitur dan label
X = data.drop('class', axis=1)
y = data['class']
```

```
# Split data into training and testing sets
X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.3, random_state=42)
```

Langkah Decision Tree Model

```
# Decision Tree
dt = DecisionTreeClassifier(random state=42)
# Hyperparameter tuning untuk Decision Tree
param grid dt = {
    'criterion': ['gini', 'entropy'],
    'max_depth': [10, 20, 30, None],
    'min_samples_split': [2, 5, 10],
    'min samples leaf': [1, 2, 4]
}
grid search dt = GridSearchCV(estimator=dt, param grid=param grid dt,
cv=5, n jobs=-1, verbose=1)
grid_search_dt.fit(X_train, y_train)
# Hasil terbaik dari Decision Tree
best dt = grid search dt.best estimator
y pred dt = best dt.predict(X test)
# Akurasi dan laporan klasifikasi
accuracy dt = accuracy score(y test, y pred dt) * 100 # Akurasi dalam
persen
print(f"Best Decision Tree Parameters: {grid search dt.best params }")
print(f"Decision Tree Accuracy: {accuracy dt:.2f}%")
print(classification report(y test, y pred dt))
Fitting 5 folds for each of 72 candidates, totalling 360 fits
Best Decision Tree Parameters: {'criterion': 'gini', 'max depth': 10,
'min_samples_leaf': 1, 'min_samples_split': 10}
Decision Tree Accuracy: 100.00%
              precision
                           recall f1-score
                                              support
           0
                   1.00
                             1.00
                                        1.00
                                                    19
           1
                   1.00
                             1.00
                                        1.00
                                                    13
           2
                   1.00
                                        1.00
                                                    13
                             1.00
                                        1.00
                                                    45
    accuracy
                   1.00
                             1.00
                                        1.00
                                                    45
   macro avg
weighted avg
                   1.00
                             1.00
                                        1.00
                                                    45
```

```
# AdaBoost
ada = AdaBoostClassifier()
# Hyperparameter tuning untuk AdaBoost
param grid ada = {
    'n estimators': [50, 100, 200],
    'learning_rate': [0.01, 0.1, 1, 10]
}
grid_search_ada = GridSearchCV(estimator=ada,
param grid=param grid ada, cv=5, n jobs=-1, verbose=1)
grid search ada.fit(X train, y train)
# Hasil terbaik dari AdaBoost
best ada = grid search ada.best estimator
y pred ada = best ada.predict(X test)
# Akurasi dan laporan klasifikasi
accuracy ada = accuracy score(y test, y pred ada) * 100 # Akurasi
dalam persen
print(f"Best AdaBoost Parameters: {grid search ada.best params }")
print(f"AdaBoost Accuracy: {accuracy ada:.2f}%")
print(classification_report(y_test, y pred ada))
Fitting 5 folds for each of 12 candidates, totalling 60 fits
Best AdaBoost Parameters: {'learning_rate': 1, 'n_estimators': 100}
AdaBoost Accuracy: 100.00%
                           recall f1-score
              precision
                                              support
           0
                   1.00
                             1.00
                                       1.00
                                                    19
           1
                             1.00
                                                    13
                   1.00
                                       1.00
           2
                   1.00
                             1.00
                                       1.00
                                                    13
                                       1.00
                                                    45
    accuracy
   macro avq
                   1.00
                             1.00
                                       1.00
                                                    45
weighted avg
                   1.00
                             1.00
                                       1.00
                                                    45
C:\Users\M. Rofig Aulia\AppData\Roaming\Python\Python312\site-
packages\sklearn\ensemble\ weight boosting.py:527: FutureWarning: The
SAMME.R algorithm (the default) is deprecated and will be removed in
1.6. Use the SAMME algorithm to circumvent this warning.
 warnings.warn(
```

### Evaluasi dan Perbandingan

```
# Output Akhir
print(f"Decision Tree Accuracy: {accuracy_dt:.2f}%")
print(f"AdaBoost Accuracy: {accuracy_ada:.2f}%")
```

Decision Tree Accuracy: 100.00%

AdaBoost Accuracy: 100.00%

# TUGAS 3

engan menggunakan dataset diabetes, buatlah ensemble voting dengan algoritma

- 1. Logistic Regression
- 2. SVM kernel polynomial
- 3. Decission Tree

Anda boleh melakukan eksplorasi dengan melakukan tunning hyperparameter

```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split, GridSearchCV
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import AdaBoostClassifier
from sklearn.metrics import accuracy score, classification report
# Load data
df = pd.read csv('data/diabetes.csv')
df.head()
   Pregnancies
                Glucose BloodPressure SkinThickness Insulin
BMI \
                    148
                                                               0 33.6
             6
                                     72
                                                     35
                     85
                                     66
                                                     29
                                                                 26.6
1
             1
                                                               0
2
                    183
                                     64
                                                                  23.3
                                                      0
                                                               0
                                                              94 28.1
3
                     89
                                     66
                                                     23
                                     40
                    137
                                                     35
                                                             168 43.1
   DiabetesPedigreeFunction
                              Age
                                   Outcome
0
                       0.627
                               50
                                         1
                       0.351
                                         0
1
                               31
2
                       0.672
                               32
                                         1
3
                       0.167
                               21
                                         0
4
                       2.288
                               33
                                         1
```

Cek Kolom Null

```
df.isnull().sum()
```

```
Pregnancies
                    0
Glucose
                    0
BloodPressure
                    0
SkinThickness
                    0
                    0
Insulin
BMI
                    0
                    0
DiabetesPedigreeFunction
                    0
Aae
Outcome
                    0
dtype: int64
feature_columns = ['Pregnancies', 'Glucose', 'BloodPressure',
'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age']
for column in feature columns:
   print("========"")
  print(f"{column} ==> Missing zeros : {len(df.loc[df[column] ==
0])}")
_____
Pregnancies ==> Missing zeros : 111
_____
Glucose ==> Missing zeros : 5
______
BloodPressure ==> Missing zeros : 35
_____
SkinThickness ==> Missing zeros : 227
_____
Insulin ==> Missing zeros : 374
______
BMI ==> Missing zeros : 11
_____
DiabetesPedigreeFunction ==> Missing zeros : 0
Age ==> Missing zeros : 0
```

Input Nilai 0 dengan Mean

```
# Import SimpleImputer dari sklearn
from sklearn.impute import SimpleImputer

# Inisialisasi SimpleImputer
fill_values = SimpleImputer(missing_values=0, strategy="mean",
copy=False)

# Menerapkan imputasi pada kolom fitur
df[feature_columns] = fill_values.fit_transform(df[feature_columns])
```

```
X = df[feature_columns]
y = df.Outcome

X_train, X_test, y_train, y_test = train_test_split(X, y,
test_size=0.3, random_state=42)
```

Standarisasi Fitur

```
# Import StandardScaler dari sklearn
from sklearn.preprocessing import StandardScaler

# Inisialisasi StandardScaler
sc = StandardScaler()

# Standarisasi pada fitur di X_train dan X_test
X_train_std = sc.fit_transform(X_train)
X_test_std = sc.transform(X_test)
```

Model

```
# Import model yang diperlukan dari sklearn
from sklearn.linear_model import LogisticRegression
from sklearn.svm import SVC
from sklearn.tree import DecisionTreeClassifier

# Inisialisasi model
log_reg = LogisticRegression(random_state=42)
svc = SVC(kernel='poly', probability=True, random_state=42) # SVM
dengan kernel polynomial
dt = DecisionTreeClassifier(random_state=42)
```

Logistik regeression w/ Hyperparameter Tunning

```
# Definisikan hyperparameter Logistic Regression
param_grid_logreg = {
    'C': [0.1, 1, 10, 100], # Regularization strength
    'solver': ['liblinear', 'lbfgs'], # Solver untuk optimasi
    'max_iter': [100, 200, 500] # Jumlah iterasi maksimum
}

# GridSearchCV untuk Logistic Regression
grid_search_logreg = GridSearchCV(estimator=log_reg,
param_grid=param_grid_logreg, cv=5, verbose=1, n_jobs=-1)

# Fit model Logistic Regression
grid_search_logreg.fit(X_train_std, y_train)

# Prediksi pada data test
y_pred_logreg = grid_search_logreg.best_estimator_.predict(X_test_std)
```

```
# Evaluasi Logistic Regression
accuracy logreg = accuracy score(y test, y pred logreg)
print(f"Accuracy (Logistic Regression): {accuracy_logreg*100:.2f}%")
print(f"Classification Report (Logistic Regression):\
n{classification report(y test, y pred logreg)}")
Fitting 5 folds for each of 24 candidates, totalling 120 fits
Accuracy (Logistic Regression): 73.59%
Classification Report (Logistic Regression):
              precision
                           recall f1-score
                                              support
                   0.79
                             0.81
                                       0.80
                                                   151
           1
                             0.59
                   0.63
                                       0.61
                                                   80
                                       0.74
                                                   231
    accuracy
                   0.71
                             0.70
                                       0.70
                                                  231
   macro avg
                   0.73
                             0.74
                                       0.73
                                                  231
weighted avg
```

# SVM Polynomial w/Hyperparameter Tunning

```
param grid svc = {
    \overline{C}: [\overline{0}.1, 1, 10], # Regularisasi
    'degree': [2, 3, 4], # Derajat polynomial
    'gamma': ['scale', 'auto'], # Kernel coefficient
}
# GridSearchCV untuk SVM
grid search svc = GridSearchCV(estimator=svc,
param grid=param grid svc, cv=5, verbose=1, n jobs=-1)
# Fit model SVM
grid search svc.fit(X train std, y train)
# Prediksi pada data test
y pred svc = grid search svc.best estimator .predict(X test std)
# Evaluasi SVM
accuracy_svc = accuracy_score(y_test, y_pred_svc)
print(f"Accuracy (SVM): {accuracy svc*100:.2f}%")
print(f"Classification Report (SVM):\n{classification report(y test,
y pred svc)}")
Fitting 5 folds for each of 18 candidates, totalling 90 fits
Accuracy (SVM): 69.70%
Classification Report (SVM):
              precision recall f1-score
                                               support
                   0.72
                             0.88
                                        0.79
                                                   151
```

	1	0.61	0.35	0.44	80
accu macro weighted		0.66 0.68	0.62 0.70	0.70 0.62 0.67	231 231 231

Desicion Tree w/Hyperparameter Tunning

```
param grid dt = {
    'max_depth': [3, 5, 7, 10], # Maksimal kedalaman pohon
    'min samples split': [2, 5, 10], # Minimum jumlah sampel untuk
split
    'min samples leaf': [1, 2, 4] # Minimum jumlah sampel di setiap
daun
}
# GridSearchCV untuk Decision Tree
grid search dt = GridSearchCV(estimator=dt, param grid=param grid dt,
cv=5, verbose=1, n jobs=-1)
# Fit model Decision Tree
grid search dt.fit(X train std, y train)
# Prediksi pada data test
y pred dt = grid search dt.best estimator .predict(X test std)
# Evaluasi Decision Tree
accuracy dt = accuracy score(y test, y pred dt)
print(f"Accuracy (Decision Tree): {accuracy dt*100:.2f}%")
print(f"Classification Report (Decision Tree):\
n{classification report(y test, y pred dt)}")
Fitting 5 folds for each of 36 candidates, totalling 180 fits
Accuracy (Decision Tree): 74.46%
Classification Report (Decision Tree):
              precision
                           recall f1-score
                                              support
                   0.79
                             0.82
                                       0.81
                                                  151
           1
                   0.64
                             0.60
                                       0.62
                                                   80
                                       0.74
                                                  231
    accuracy
                                                  231
                   0.72
                             0.71
                                       0.71
   macro avq
weighted avg
                   0.74
                             0.74
                                       0.74
                                                  231
c:\Python312\Lib\site-packages\numpy\ma\core.py:2881: RuntimeWarning:
invalid value encountered in cast
  data = np.array(data, dtype=dtype, copy=copy,
```

# **Ensemble Voting**

```
# Import modul yang diperlukan
from sklearn.ensemble import VotingClassifier
from sklearn.metrics import accuracy score, classification report
# Definisikan model dengan hyperparameter terbaik dari GridSearch
log reg best = grid search_logreg.best_estimator_ # Logistic
Regression terbaik
svc best = grid_search_svc.best_estimator_
                                                # SVM terbaik
dt best = grid search dt.best estimator
                                                  # Decision Tree
terbaik
# Ensemble Voting dengan soft voting
voting clf = VotingClassifier(estimators=[('lr', log reg best),
('svc', svc best), ('dt', dt best)], voting='soft')
# Fit model pada data train
voting clf.fit(X train std, y train)
# Prediksi pada data test
y pred voting = voting clf.predict(X test std)
# Evaluasi Ensemble Voting
accuracy voting = accuracy score(y test, y pred voting)
print(f"Accuracy (Ensemble Voting): {accuracy voting*100:.2f}%")
print(f"Classification Report (Ensemble Voting):\
n{classification report(y test, y pred voting)}")
Accuracy (Ensemble Voting): 76.19%
Classification Report (Ensemble Voting):
              precision recall f1-score
                                              support
                   0.79
           0
                             0.86
                                                  151
                                       0.83
           1
                   0.69
                             0.57
                                       0.63
                                                   80
                                                  231
                                       0.76
    accuracy
                   0.74
                             0.72
                                       0.73
   macro avq
                                                  231
weighted avg
                   0.76
                             0.76
                                       0.76
                                                  231
```

# PRAKTIKUM 1

Bagging dengan RandomForest Pada kasus ini kita akan menggunakan salah satu metode bagging yaitu RandomForest untuk mengklasifikasikan jenis tumor. Dalam latihan ini Anda akan melakukan training dengan data Wisconsin Breast Cancer Dataset dari UCI machine learning repository. Latihan ini akan melakukan prediksi memprediksi apakah tumor ganas atau jinak.

```
import numpy as np
import pandas as pd
from sklearn.tree import DecisionTreeClassifier # import DT
from sklearn.ensemble import RandomForestClassifier # import
RandomForest
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, classification_report
```

### Persiapan Data

```
# Load data
df = pd.read csv('data/wbc.csv')
df.head()
         id diagnosis
                       radius mean texture mean
                                                   perimeter mean
area mean
     842302
                              17.99
                                            10.38
                                                            122.80
1001.0
     842517
                              20.57
                                            17.77
                                                            132.90
1326.0
2 84300903
                                            21.25
                              19.69
                                                            130.00
1203.0
3 84348301
                              11.42
                                            20.38
                                                             77.58
386.1
                              20.29
                                            14.34
4 84358402
                                                            135.10
1297.0
   smoothness mean
                    compactness mean
                                       concavity mean
                                                        concave
points mean
           0.11840
                              0.27760
                                               0.3001
0.14710
           0.08474
                              0.07864
                                               0.0869
0.07017
           0.10960
                              0.15990
                                               0.1974
0.12790
           0.14250
                              0.28390
                                               0.2414
0.10520
           0.10030
                              0.13280
                                               0.1980
0.10430
   ... texture worst perimeter worst area worst
```

```
smoothness worst \
0 ...
                17.33
                                 184.60
                                             2019.0
                                                                0.1622
                23.41
                                 158.80
                                             1956.0
                                                                0.1238
2 ...
                25.53
                                 152.50
                                             1709.0
                                                                0.1444
3 ...
                26.50
                                  98.87
                                              567.7
                                                                0.2098
                                                                0.1374
                16.67
                                 152.20
                                             1575.0
   compactness worst concavity worst concave points worst
symmetry_worst \
              0.6656
                                0.7119
                                                       0.2654
0.4601
1
              0.1866
                                0.2416
                                                       0.1860
0.2750
              0.4245
                                0.4504
                                                       0.2430
0.3613
3
                                0.6869
                                                       0.2575
              0.8663
0.6638
              0.2050
                                0.4000
                                                       0.1625
0.2364
   fractal_dimension_worst
                             Unnamed: 32
0
                   0.11890
                                     NaN
1
                   0.08902
                                     NaN
2
                    0.08758
                                     NaN
3
                    0.17300
                                     NaN
                    0.07678
                                     NaN
[5 rows x 33 columns]
# Cek kolom null
df.isnull().sum()
id
diagnosis
                              0
                              0
radius mean
                              0
texture mean
perimeter_mean
                              0
area mean
                              0
smoothness mean
                              0
compactness mean
                              0
                              0
concavity mean
                              0
concave points mean
symmetry_mean
                              0
fractal dimension mean
                              0
                              0
radius se
```

```
texture se
                             0
perimeter se
area se
                             0
                             0
smoothness se
compactness se
                             0
                             0
concavity se
                             0
concave points se
                             0
symmetry se
fractal dimension se
                             0
radius worst
                             0
texture worst
                             0
                             0
perimeter worst
                             0
area worst
                             0
smoothness worst
compactness worst
                             0
                             0
concavity worst
concave points worst
                             0
symmetry_worst
fractal dimension worst
                             0
Unnamed: 32
                           569
dtype: int64
# Seleksi fitur
# Slice dataframe mulai dari kolom 'radius mean' sampai
'fractal dimension worst'
X = df.iloc[:,3:-1]
y = df['diagnosis']
y = y.map({'M':1, 'B':0}) # Encode label
# Cek jumlah fitur dan instance
X.shape
(569, 29)
from sklearn.model selection import train test split
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=1)
# Secara default, DecisionTreeClassifier dari scikit-learn akan
menggunakan nilai "Gini" untuk kriteria
# Terdapat beberapa "hyperparamater" yang dapat digunakan. Silahka
baca dokumentasi
# Pada kasus ini kita akan menggunakan parameter default
dt = DecisionTreeClassifier()
# Sesuaikan dt ke set training
dt.fit(X train, y train)
# Memprediksi label set test
```

```
y pred dt = dt.predict(X test)
# menghitung set accuracy
acc_dt = accuracy_score(y_test, y_pred_dt)
print("Test set accuracy: {:.2f}".format(acc dt))
print(f"Test set accuracy: {acc dt}")
Test set accuracy: 0.96
Test set accuracy: 0.956140350877193
# Pada kasus kali ini kita akan menggunakan estimator pada
RandomForest
# Untuk detail parameter (hyperparameter) silahkan cek dokumentasi
rf = RandomForestClassifier(n estimators=10, random state=1)
# Sesuaikan dt ke set training
rf.fit(X train, y train)
# Memprediksi label set test
y pred rf = rf.predict(X test)
# menghitung set accuracy
acc_rf = accuracy_score(y_test, y_pred_rf)
print("Test set accuracy: {:.2f}".format(acc rf))
print(f"Test set accuracy: {acc rf}")
Test set accuracy: 0.96
Test set accuracy: 0.956140350877193
```

### PRAKTIKUM 2

Boosting dengan AdaBoost Pada kasus ini kita akan menggunakan salah satu metode boosting yaitu AdaBoost untuk mengklasifikasikan jenis bunga Iris. Dalam latihan ini kita akan menggunakan dataset Iris yang sangat lazim digunakan. Latihan ini akan melakukan prediksi memprediksi 3 jenis bunga Iris yaitu, Iris Setosa, Iris Versicolor, dan Iris Virginica berdasarkan panjang dan lebar sepal dan petal.

#### Import Library

```
import numpy as np
import pandas as pd
from sklearn.tree import DecisionTreeClassifier # import DT
from sklearn.ensemble import AdaBoostClassifier # import AdaBoost
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, classification_report
from sklearn.preprocessing import LabelEncoder # Kebutuhan encoding
label

# Load data
df = pd.read_csv('data/iris.csv')
```

```
df.head()
  Id SepalLengthCm SepalWidthCm PetalLengthCm PetalWidthCm
Species
  1
           5.1
                     3.5
                               1.4
                                         0.2 Iris-
setosa
           4.9
                     3.0
                               1.4
                                         0.2 Iris-
  2
setosa
           4.7
                     3.2
                               1.3
  3
                                         0.2 Iris-
setosa
           4.6
                     3.1
                               1.5
  4
                                         0.2 Iris-
setosa
                                         0.2 Iris-
           5.0
                     3.6
                               1.4
  5
setosa
# Cek kolom null
df.isnull().sum()
Id
           0
           0
SepalLengthCm
SepalWidthCm
           0
PetalLengthCm
           0
PetalWidthCm
           0
           0
Species
dtype: int64
# Seleksi fitur
X = df.iloc[:,2:-1]
y = df['Species']
# encode label
ec = LabelEncoder()
y = ec.fit transform(y)
# Cek jumlah fitur dan instance
print(X.shape)
# Cek label
print(y)
(150, 3)
2 2
2 2
2 2]
```

```
from sklearn.model selection import train test split
X train, X test, y train, y test = train test split(X, y,
test size=0.2, random state=1)
# Secara default, DecisionTreeClassifier dari scikit-learn akan
menggunakan nilai "Gini" untuk kriteria
# Terdapat beberapa "hyperparamater" yang dapat digunakan. Silahka
baca dokumentasi
# Pada kasus ini kita akan menggunakan parameter default
dt = DecisionTreeClassifier()
# Sesuaikan dt ke set training
dt.fit(X train, y train)
# Memprediksi label set test
y pred dt = dt.predict(X test)
# menghitung set accuracy
acc_dt = accuracy_score(y_test, y_pred_dt)
print("Test set accuracy: {:.2f}".format(acc_dt))
print(f"Test set accuracy: {acc dt}")
Test set accuracy: 0.97
Test set accuracy: 0.966666666666667
# Pada kasus kali ini kita akan menggunakan estimator pada AdaBoost
# Untuk detail parameter (hyperparameter) silahkan cek dokumentasi
ada = AdaBoostClassifier(n estimators=2)
# Sesuaikan dt ke set training
ada.fit(X_train, y_train)
# Memprediksi label set test
y pred ada = ada.predict(X test)
# menghitung set accuracy
acc_ada = accuracy_score(y_test, y_pred_ada)
print("Test set accuracy: {:.2f}".format(acc_ada))
print(f"Test set accuracy: {acc ada}")
Test set accuracy: 0.97
Test set accuracy: 0.966666666666667
C:\Users\M. Rofig Aulia\AppData\Roaming\Python\Python312\site-
packages\sklearn\ensemble\ weight boosting.py:527: FutureWarning: The
SAMME.R algorithm (the default) is deprecated and will be removed in
1.6. Use the SAMME algorithm to circumvent this warning.
  warnings.warn(
```

#### **PRAKTIKUM 3**

Stacking Lengkapi bagian berikut dengan data sesuai tugas, dan tentukan perbedaan nilai akurasi antara Random Forest, Adaboost, dan Stacking

```
from sklearn.ensemble import RandomForestClassifier,
StackingClassifier
from sklearn.linear model import LogisticRegression
from sklearn.model selection import train test split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.tree import DecisionTreeClassifier
layer one estimators = [
                        ('rf 1',
RandomForestClassifier(n estimators=10, random state=42)),
                        ('knn 1', KNeighborsClassifier(n neighbors=5))
layer two estimators = [
                        ('dt_2', DecisionTreeClassifier()),
                        ('rf 2'
RandomForestClassifier(n_estimators=50, random_state=42)),
layer two = StackingClassifier(estimators=layer two estimators,
final estimator=LogisticRegression())
clf = StackingClassifier(estimators=layer one estimators,
final estimator=layer two)
X train, X test, y train, y test = train test split(X, y, stratify=y,
random state=42)
clf.fit(X_train, y_train).score(X_test, y_test)
0.868421052631579
```

### PRAKTIKUM 4

Stacking dengan Voting Pada kasus ini kita akan menggunakan salah satu metode stacking yaitu voting untuk mengklasifikasikan pasien penderita diabetes dengan beberapa ciri. Pasien akan di klasifikasikan menjadi pasien menderita diabetes (1) dan tidak menderita diabetes (0). Pertamatama, kita akan menggunakan beberapa algoritma klasifikasi secara terpisah, yaitu Naive Bayes, SVM Linier, dan SVM RBF. Setelah itu, kita akan menggabungkan performa dari 3 algoritma tersebut dengan menggunakan metode ensemble voting.

```
import numpy as np
import pandas as pd
from sklearn.naive_bayes import GaussianNB # import Naive Bayes model
```

```
Gaussian (asumsi data terdistribusi normal)
from sklearn.svm import SVC # import SVM classifier
from sklearn.ensemble import VotingClassifier # import model Voting
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, classification_report
```

### Persiapan Data

```
# Load Data
dbt = pd.read csv('data/diabetes.csv')
dbt.head()
   Pregnancies Glucose BloodPressure SkinThickness Insulin
BMI \
                    148
                                     72
                                                               0 33.6
                                                     35
                     85
                                     66
                                                     29
                                                                  26.6
2
                    183
                                     64
                                                               0 23.3
                                                      0
                     89
                                                              94 28.1
3
                                     66
                                                     23
                                     40
                     137
                                                     35
                                                             168 43.1
   DiabetesPedigreeFunction
                                   Outcome
                              Age
0
                       0.627
                               50
                                         1
                       0.351
                                         0
1
                               31
2
                                         1
                       0.672
                               32
3
                       0.167
                               21
                                         0
                       2.288
                                         1
                               33
# Cek nama kolom
dbt.columns
Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness',
'Insulin',
       'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'],
      dtype='object')
# Cek kolom null
dbt.isnull().sum()
Pregnancies
                             0
                             0
Glucose
BloodPressure
                             0
                             0
SkinThickness
                             0
Insulin
                             0
BMI
```

```
DiabetesPedigreeFunction
                       0
                       0
Age
Outcome
                       0
dtype: int64
# Pada kasus ini, agak tidak masuk akal jika beberapa parameter
bernilai 0
# sebagai contoh adalah nilai 'Glucose', 'BloodPlessure' ataupun
'Insulin'.
# Sekecil apapun nilainya, setiap manusia yang hidup pasti miliki
nilai-nilai tersebut
# Kita akan manipulasi nilai yang 0 dengan melakukan 'imputasi' atau
mengganti nilainya dengan nilai sintetis
# Pada kasus ini, kita akan menggunakan nilai mean
# Cek kolom neng nilai 0
feature_columns = ['Pregnancies', 'Glucose', 'BloodPressure',
'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age']
for column in feature columns:
   print("========"")
   print(f"{column} ==> Missing zeros : {len(dbt.loc[dbt[column] ==
01)}")
Pregnancies ==> Missing zeros : 111
______
Glucose ==> Missing zeros : 5
_____
BloodPressure ==> Missing zeros : 35
_____
SkinThickness ==> Missing zeros : 227
______
Insulin ==> Missing zeros : 374
_____
BMI ==> Missing zeros : 11
_____
DiabetesPedigreeFunction ==> Missing zeros : 0
Age ==> Missing zeros : 0
# Impute nilai 0 dengan mean
from sklearn.impute import SimpleImputer
fill values = SimpleImputer(missing values=0, strategy="mean",
copy=False)
dbt[feature columns] = fill values.fit transform(dbt[feature columns])
```

```
X = dbt[feature_columns]
y = dbt.Outcome

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
```

Training dengan GaussianNB Standarisasi Fitur

```
# Karena asumsi Gaussian NB adalah data terdistribusi secara normal,
# maka kita perlu melakukan standarisasi

from sklearn.preprocessing import StandardScaler

sc = StandardScaler()

# Standarisasi pada fitur di X_train dan X_test
X_train_std = sc.fit_transform(X_train)
X_test_std = sc.transform(X_test)
```

Training dan Evaluasi

```
# Buat obyek GaussianNB
gnb_std = GaussianNB()

# Fit dengan data yang telah di standarisasi
gnb_std.fit(X_train_std, y_train)

# Prediksi dengan data test
y_pred_gnb = gnb_std.predict(X_test_std)

# Evaluasi akurasi testing data
acc_gnb = accuracy_score(y_test, y_pred_gnb)

# Print hasil evaluasi
print("Test set accuracy: {:.2f}".format(acc_gnb))
print(f"Test set accuracy: {acc_gnb}")

Test set accuracy: 0.74
Test set accuracy: 0.7359307359307359
```

Training dengan SVM Linier

```
# Model SVM linier tanpa tunnning hyperparameter
svm_lin = SVC(kernel='linear')

# Fit ke model
svm_lin.fit(X_train_std, y_train)

# Prediksi
```

```
y_pred_svm_lin = svm_lin.predict(X_test_std)

# Evaluasi akurasi testing data
acc_svm_lin = accuracy_score(y_test, y_pred_svm_lin)

# Print hasil evaluasi
print("Test set accuracy: {:.2f}".format(acc_svm_lin))
print(f"Test set accuracy: {acc_svm_lin}")

Test set accuracy: 0.74
Test set accuracy: 0.7402597402597403
```

Training dengan SVM RBF

```
# Model SVM RBF tanpa tunnning hyperparameter
svm_rbf = SVC(kernel='rbf')

# Fit ke model
svm_rbf.fit(X_train_std, y_train)

# Prediksi
y_pred_svm_rbf = svm_rbf.predict(X_test_std)

# Evaluasi akurasi testing data
acc_svm_rbf = accuracy_score(y_test, y_pred_svm_rbf)

# Print hasil evaluasi
print("Test set accuracy: {:.2f}".format(acc_svm_rbf))
print(f"Test set accuracy: {acc_svm_rbf}")

Test set accuracy: 0.72
Test set accuracy: 0.7229437229437229
```

Training dan Voting

```
# Definisikan algoritma yang akan digunakan untuk voting

clf1 = GaussianNB()
  clf2 = SVC(kernel='linear')
  clf3 = SVC(kernel='rbf', probability=True)

# model hard voting
voting = VotingClassifier(estimators=[('GaussianNB', clf1), ('SVM-LIN', clf2), ('SVM-RBF', clf3)], voting='hard')

# Fit model
voting.fit(X_train_std, y_train)

# Prediksi
y_pred_vt1 = voting.predict(X_test_std)
```

```
# Evaluasi akurasi testing data
acc_vt1 = accuracy_score(y_test, y_pred_vt1)

# Print hasil evaluasi
print('Voting Hard')
print("Test set accuracy: {:.2f}".format(acc_vt1))
print(f"Test set accuracy: {acc_vt1}")

Voting Hard
Test set accuracy: 0.74
Test set accuracy: 0.7402597402597403
```