EXPERIMENT 4A : Support Vector Machines (SVM)

AIM:

To build an SVM model for a binary classification task, tune its hyperparameters, and evaluate it using accuracy, precision, recall, F1-score, confusion matrix, and ROC-AUC.

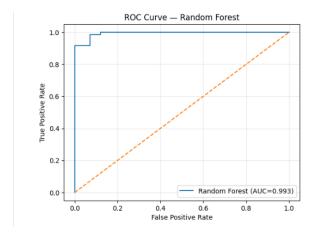
SOURCE CODE:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.datasets import load breast cancer
from sklearn.model selection import train test split, GridSearchCV
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVC
from sklearn.metrics import (
accuracy score, precision score, recall score, fl score,
confusion matrix, classification report, roc auc score, roc curve
)
data = load breast cancer()
X = pd.DataFrame(data.data, columns=data.feature names)
y = pd.Series(data.target, name="target") # 1 = malignant? (Check dataset doc: in this set,
0==malignant, 1==benign
X_train, X_test, y_train, y_test = train_test_split(
X, y, test size=0.20, random state=42, stratify=y
)
scaler = StandardScaler()
X_{train\_sc} = scaler.fit_transform(X_{train})
X \text{ test } sc = scaler.transform(X \text{ test})
svm = SVC(kernel='rbf', probability=True, random state=42)
param_grid = {
"C": [0.1, 1, 10, 100],
"gamma": ["scale", 0.01, 0.001, 0.0001]
}
grid = GridSearchCV(
```

```
estimator=svm,
param_grid=param_grid,
scoring='f1', # you can change to 'accuracy' or 'roc auc' as needed
cv=5,
n jobs=-1,
verbose=0
)
grid.fit(X_train_sc, y_train)
print("Best Params (CV):", grid.best params )
best_svm = grid.best_estimator_
best svm.fit(X train sc, y train)
y_pred = best_svm.predict(X_test_sc)
y prob = best svm.predict proba(X test sc)[:, 1]
acc = accuracy_score(y_test, y_pred)
prec = precision_score(y_test, y_pred, zero_division=0)
rec = recall_score(y_test, y_pred)
f1 = f1_score(y_test, y_pred)
auc = roc_auc_score(y_test, y_prob)
cm = confusion_matrix(y_test, y_pred)
print("\n=== SVM (RBF) — Test Metrics ===")
print(f"Accuracy : {acc:.4f}")
print(f"Precision: {prec:.4f}")
print(f"Recall : {rec:.4f}")
print(f"F1-Score : {f1:.4f}")
print(f"ROC-AUC : {auc:.4f}")
print("\nConfusion Matrix:\n", cm)
print("\nClassification Report:\n", classification report(y test, y pred, zero division=0))
fpr, tpr, thresholds = roc curve(y test, y prob)
plt.figure()
plt.plot(fpr, tpr, label=f"SVM (AUC={auc:.3f})")
plt.plot([0, 1], [0, 1], linestyle="--")
plt.xlabel("False Positive Rate")
```

```
plt.ylabel("True Positive Rate")
plt.title("ROC Curve — SVM (RBF)")
plt.legend()
plt.grid(True, alpha=0.3)
plt.show()
```

OUTPUT:



EXPERIMENT 4B: Ensemble Methods: Random Forest

AIM:

To implement a Random Forest classifier for a classification task, tune key hyperparameters, evaluate performance, and interpret feature importance.

SOURCE CODE:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn.datasets import load breast cancer
from sklearn.model selection import train test split, GridSearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import (
accuracy_score, precision_score, recall_score, f1_score,
confusion matrix, classification report, roc auc score, roc curve
)
data = load breast cancer()
X = pd.DataFrame(data.data, columns=data.feature names)
y = pd.Series(data.target, name="target")
X train, X test, y train, y test = train test split(
X, y, test size=0.20, random state=42, stratify=y
rf = RandomForestClassifier(random state=42, n jobs=-1)
param_grid = {
"n estimators": [100, 300, 500],
"max depth": [None, 5, 10, 20],
"min samples split": [2, 5, 10],
"min samples leaf": [1, 2, 4]
grid = GridSearchCV(
estimator=rf,
param grid=param grid,
```

```
scoring="f1",
cv=5,
n jobs=-1,
verbose=0
)
grid.fit(X train, y train)
print("Best Params (CV):", grid.best params )
best rf = grid.best estimator
best rf.fit(X train, y train)
y pred = best rf.predict(X test)
y prob = best rf.predict proba(X test)[:, 1]
acc = accuracy score(y test, y pred)
prec = precision score(y test, y pred, zero division=0)
rec = recall_score(y_test, y_pred)
f1 = f1_score(y_test, y_pred)
auc = roc auc score(y test, y prob)
cm = confusion matrix(y test, y pred)
print("\n=== Random Forest — Test Metrics ===")
print(f"Accuracy : {acc:.4f}")
print(f"Precision: {prec:.4f}")
print(f"Recall : {rec:.4f}")
print(f"F1-Score : {f1:.4f}")
print(f"ROC-AUC : {auc:.4f}")
print("\nConfusion Matrix:\n", cm)
print("\nClassification Report:\n", classification report(y test, y pred, zero division=0))
importances = pd.Series(best rf.feature importances , index=X.columns)
top10 = importances.sort_values(ascending=False).head(10)
plt.figure()
top10[::-1].plot(kind="barh")
plt.xlabel("Importance")
plt.title("Top 10 Feature Importances — Random Forest")
plt.grid(axis="x", alpha=0.3)
```

```
plt.show()

from sklearn.metrics import roc_curve

fpr, tpr, thresholds = roc_curve(y_test, y_prob)

plt.figure()

plt.plot(fpr, tpr, label=f"Random Forest (AUC={auc:.3f})")

plt.plot([0, 1], [0, 1], linestyle="--")

plt.xlabel("False Positive Rate")

plt.ylabel("True Positive Rate")

plt.title("ROC Curve — Random Forest")

plt.legend()

plt.grid(True, alpha=0.3)

plt.show()
```

OUTPUT:

