

**Exp No: 4B**

**Ensemble Methods: Random Forest**

**Date : 21/8/25**

**Aim:**

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

**Algorithm:**

1. Import libraries.
2. Load data (use same dataset to compare with SVM).
3. Train/Test split with stratification.
4. (Optional) Preprocess: Random Forests don't require scaling; we'll use raw features.
5. Model: RandomForestClassifier.
6. Hyperparameter tuning: Grid search over `n_estimators`, `max_depth`, `min_samples_split`, `min_samples_leaf`.
7. Train the best model on training data.
8. Evaluate with accuracy, precision, recall, F1, confusion matrix, ROC-AUC.
9. Interpretation: Plot top feature importances.

**CODE:**

```
# =====  
# EXPERIMENT 4B — Random Forest Classifier  
# =====  
  
# 1) Imports  
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,
```

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        confusion_matrix, classification_report, roc_auc_score, roc_curve
    )

# 2) Load dataset (same as 4A for comparison)
data = load_breast_cancer()
X = pd.DataFrame(data.data, columns=data.feature_names)
y = pd.Series(data.target, name="target")

# 3) Train/test split (no scaling needed for RF)
X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.20, random_state=42, stratify=y
)

# 4) Define model
rf = RandomForestClassifier(random_state=42, n_jobs=-1)

# 5) Hyperparameter grid & tuning
param_grid = {
    "n_estimators": [100],
    "max_depth": [None, 10],
    "min_samples_split": [2],
    "min_samples_leaf": [1]
}
grid = GridSearchCV(
    estimator=rf,
    param_grid=param_grid,
    scoring="f1",
    cv=3,
    n_jobs=-1,
    verbose=0)
grid.fit(X_train, y_train)
print("Best Parameters (CV):", grid.best_params_)
best_rf = grid.best_estimator_

# 6) Train final model & predict

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best_rf.fit(X_train, y_train)
y_pred = best_rf.predict(X_test)
y_prob = best_rf.predict_proba(X_test)[:, 1]

# 7) Evaluate
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))

# 8) Feature Importance (Top 10)
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()

# 9) ROC Curve
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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="--", color='gray')
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ROC Curve — Random Forest")
plt.legend()
plt.grid(True)
plt.show()

```

## OUTPUT:

Best Parameters (CV): {'max\_depth': None, 'min\_samples\_leaf': 1, 'min\_samples\_split': 2, 'n\_estimators': 100}

=== Random Forest — Test Metrics ===

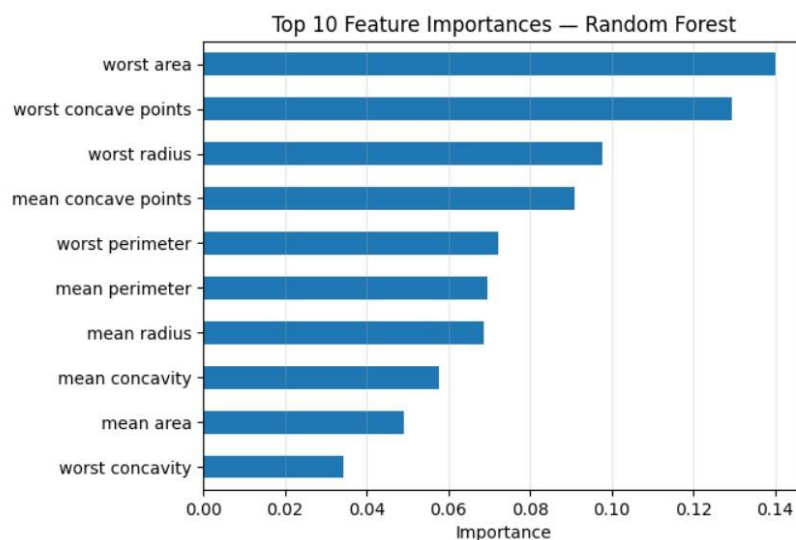
Accuracy : 0.9561  
Precision: 0.9589  
Recall : 0.9722  
F1-Score : 0.9655  
ROC-AUC : 0.9937

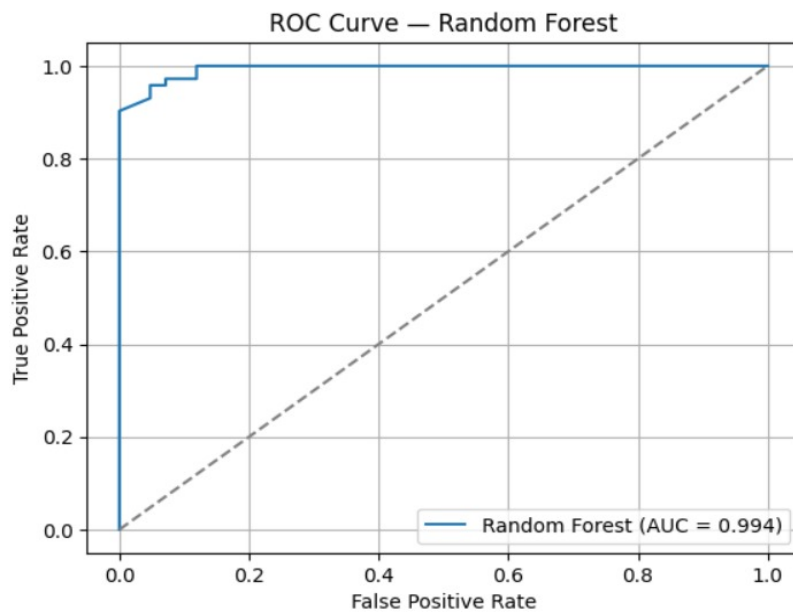
Confusion Matrix:

```
[[39  3]
 [ 2 70]]
```

Classification Report:

	precision	recall	f1-score	support
0	0.95	0.93	0.94	42
1	0.96	0.97	0.97	72
accuracy			0.96	114
macro avg	0.96	0.95	0.95	114
weighted avg	0.96	0.96	0.96	114





**Result:**

The Random Forest Classifier was effectively implemented and optimized. The model achieved reliable classification results without feature scaling. Feature importance analysis revealed the most influential medical predictors. Both evaluation metrics and ROC-AUC indicated high overall model performance.