Diabetes Prediction — 02 Modeling

Goal: Train and evaluate baseline and tuned models for the Pima Indians Diabetes dataset.

```
In [1]: # ==========
        # 0) Setup
        # ========
        import os
        import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        from sklearn.model_selection import train_test_split, cross_val_score, Stratifie
        from sklearn.preprocessing import StandardScaler
        from sklearn.pipeline import Pipeline
        from sklearn.compose import ColumnTransformer
        from sklearn.metrics import (
            accuracy_score, precision_score, recall_score, f1_score, roc_auc_score,
            confusion_matrix, RocCurveDisplay
        from sklearn.linear_model import LogisticRegression
        from sklearn.ensemble import RandomForestClassifier
        # Optional models
        try:
            from xgboost import XGBClassifier
            HAS_XGB = True
        except Exception:
            HAS XGB = False
        try:
            from lightgbm import LGBMClassifier
            HAS LGBM = True
        except Exception:
            HAS LGBM = False
        plt.rcParams['figure.figsize'] = (8, 5)
        plt.rcParams['axes.grid'] = True
        # Paths
        DATA_PROCESSED_PATH = os.path.join('C:/Users/nazar/OneDrive/Documentos/machine 1
        # If you saved elsewhere, override here:
        # DATA_PROCESSED_PATH = r'C:/Users/nazar/OneDrive/Documentos/machine Learning/di
        assert os.path.exists(DATA PROCESSED PATH), f"Processed CSV not found at {DATA P
In [2]: # ========
        # 1) Load & split
        # =========
        df = pd.read csv(DATA PROCESSED PATH)
        print('Shape:', df.shape)
        print('Columns:', df.columns.tolist())
```

```
target_col = 'Outcome'
 X = df.drop(columns=[target_col])
 y = df[target_col].astype(int)
 # numeric/ categorical split (dataset is numeric but keep generic)
 num_cols = X.select_dtypes(include=[np.number]).columns.tolist()
 cat_cols = [c for c in X.columns if c not in num_cols]
 # Preprocess: scale numeric features; passthrough categoricals (none expected he
 preprocess = ColumnTransformer(
     transformers=[
         ('num', StandardScaler(), num_cols),
         ('cat', 'passthrough', cat_cols) # keeps order; fine if empty
 # Train/test split (stratified)
 X_train, X_test, y_train, y_test = train_test_split(
     X, y, test_size=0.2, random_state=42, stratify=y
 print('Train size:', X_train.shape, ' Test size:', X_test.shape)
Shape: (768, 9)
Columns: ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin',
```

2) Baseline models

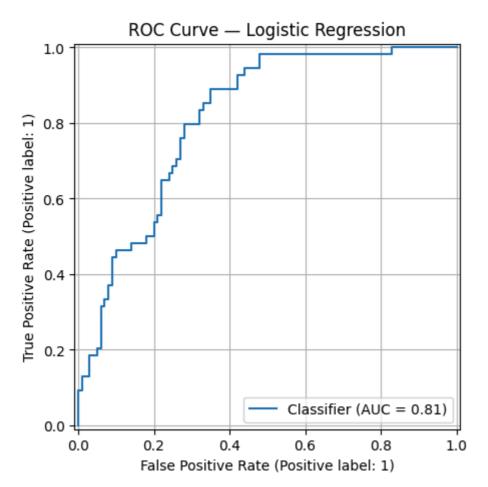
We train a few standard models with default settings and report metrics.

'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome']

Train size: (614, 8) Test size: (154, 8)

```
In [3]: def evaluate_model(name, model, X_tr, X_te, y_tr, y_te):
            model.fit(X_tr, y_tr)
            y pred = model.predict(X te)
            # Some models may not implement predict_proba; handle gracefully
            trv:
                y_proba = model.predict_proba(X_te)[:, 1]
                roc = roc_auc_score(y_te, y_proba)
            except Exception:
                # Fallback to decision_function if available
                    scores = model.decision_function(X_te)
                    from sklearn.preprocessing import MinMaxScaler
                    y proba = MinMaxScaler().fit transform(scores.reshape(-1,1)).ravel()
                    roc = roc_auc_score(y_te, y_proba)
                except Exception:
                    y_proba = None
                    roc = np.nan
            acc = accuracy_score(y_te, y_pred)
            prec = precision_score(y_te, y_pred, zero_division=0)
            rec = recall_score(y_te, y_pred, zero_division=0)
            f1 = f1_score(y_te, y_pred, zero_division=0)
            cm = confusion_matrix(y_te, y_pred)
            print(f"\n=== {name} ===")
            print(f"Accuracy: {acc:.3f} Precision: {prec:.3f} Recall: {rec:.3f} F1: {
            print("Confusion matrix:\n", cm)
```

```
if y_proba is not None:
         fig, ax = plt.subplots()
         RocCurveDisplay.from_predictions(y_te, y_proba, ax=ax)
         ax.set_title(f'ROC Curve - {name}')
         plt.show()
 # Pipelines
 logreg = Pipeline([('prep', preprocess), ('clf', LogisticRegression(max_iter=100)
 rf = Pipeline([('prep', preprocess), ('clf', RandomForestClassifier(random_state
 evaluate_model('Logistic Regression', logreg, X_train, X_test, y_train, y_test)
 evaluate_model('Random Forest', rf, X_train, X_test, y_train, y_test)
 if HAS XGB:
     xgb = Pipeline([('prep', preprocess), ('clf', XGBClassifier(
         n_estimators=300, learning_rate=0.05, max_depth=3, subsample=0.8, colsam
         random_state=42, eval_metric='logloss', n_jobs=-1
     ))])
     evaluate_model('XGBoost', xgb, X_train, X_test, y_train, y_test)
 else:
     print("XGBoost not available - install with: pip install xgboost")
 if HAS LGBM:
     lgbm = Pipeline([('prep', preprocess), ('clf', LGBMClassifier(
         n_estimators=300, learning_rate=0.05, num_leaves=31, subsample=0.8, cols
         random_state=42
     ))])
     evaluate_model('LightGBM', lgbm, X_train, X_test, y_train, y_test)
     print("LightGBM not available - install with: pip install lightgbm")
=== Logistic Regression ===
Accuracy: 0.708 Precision: 0.600 Recall: 0.500 F1: 0.545 ROC-AUC: 0.813
Confusion matrix:
[[82 18]
[27 27]]
```



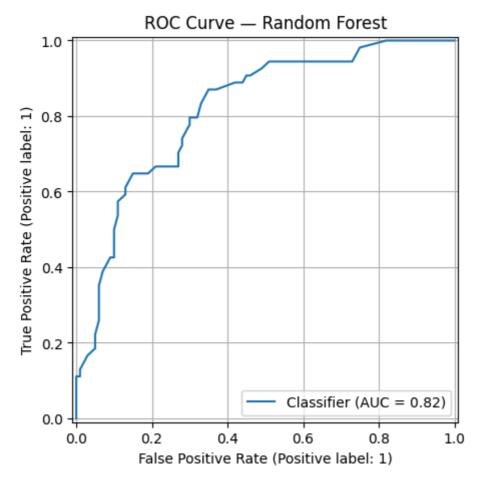
=== Random Forest ===

Accuracy: 0.779 Precision: 0.717 Recall: 0.611 F1: 0.660 ROC-AUC: 0.818

Confusion matrix:

[[87 13]

[21 33]]



XGBoost not available — install with: pip install xgboost LightGBM not available — install with: pip install lightgbm

3) Cross-validation (Logistic Regression example)

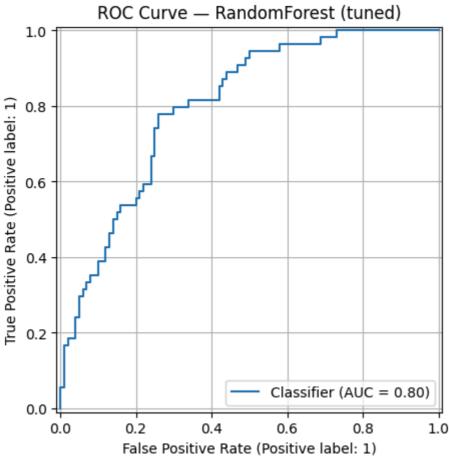
```
In [4]: cv = StratifiedKFold(n_splits=5, shuffle=True, random_state=42)
    cv_scores = cross_val_score(logreg, X, y, cv=cv, scoring='roc_auc', n_jobs=-1)
    print('LogReg ROC-AUC CV:', np.round(cv_scores, 3), '→ mean:', cv_scores.mean().
    LogReg ROC-AUC CV: [0.825 0.867 0.852 0.829 0.81] → mean: 0.837
```

4) Hyperparameter tuning (RandomizedSearchCV on RandomForest)

```
In [5]: param_distributions = {
    'clf__n_estimators': [100, 200, 400, 600],
    'clf__max_depth': [None, 3, 5, 7, 10],
    'clf__min_samples_split': [2, 4, 6, 10],
    'clf__min_samples_leaf': [1, 2, 4],
    'clf__max_features': ['sqrt', 'log2', None]
}

rf_pipe = Pipeline([('prep', preprocess), ('clf', RandomForestClassifier(random_search = RandomizedSearchCV(
    rf_pipe,
    param_distributions=param_distributions,
    n_iter=20,
    scoring='roc_auc',
```

```
cv=cv,
     random_state=42,
     n_{jobs=-1}
     verbose=1
 search.fit(X_train, y_train)
 print('Best ROC-AUC CV:', search.best_score_.round(3))
 print('Best params:', search.best_params_)
 best_rf = search.best_estimator_
 evaluate_model('RandomForest (tuned)', best_rf, X_train, X_test, y_train, y_test
Fitting 5 folds for each of 20 candidates, totalling 100 fits
Best ROC-AUC CV: 0.84
Best params: {'clf__n_estimators': 200, 'clf__min_samples_split': 4, 'clf__min_sa
mples_leaf': 2, 'clf__max_features': 'log2', 'clf__max_depth': 3}
=== RandomForest (tuned) ===
Accuracy: 0.721 Precision: 0.641 Recall: 0.463 F1: 0.538 ROC-AUC: 0.801
Confusion matrix:
 [[86 14]
 [29 25]]
```



5) Save the best model (optional)

```
In [7]: import joblib

best_model = best_rf # or choose your favorite model above
out_dir = os.path.join('...', 'models')
os.makedirs(out_dir, exist_ok=True)
```

```
model_path = os.path.join(out_dir, 'C:/Users/nazar/OneDrive/Documentos/machine l
joblib.dump(best_model, model_path)
print(f'Model saved to: {model_path}')
```

Model saved to: C:/Users/nazar/OneDrive/Documentos/machine learning\diabetes_best _model.joblib