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In [1]: import pandas as pd
        import numpy as np
        from sklearn.ensemble import RandomForestRegressor
        from sklearn.metrics import mean_squared_error, mean_absolute_error
        from sklearn.model selection import KFold
In [2]: # Load the dataset
        data = pd.read_csv('1M_ahead_dataset.csv')
In [3]: # Separate predictors and target; assuming 'Yt.1M' is the target.
        X = data.drop(['Yt.1M'], axis=1)
        y = data['Yt.1M']
In [4]: # Set up 5-Fold cross-validation
        kf = KFold(n_splits=5, shuffle=True, random_state=42)
        fold\ metrics = []
        fold counter = 1
In [5]: # Loop over each fold
        for train index, test index in kf.split(X):
            X train, X test = X.iloc[train index], X.iloc[test index]
            y train, y test = y.iloc[train index], y.iloc[test index]
            # Initialize the Random Forest Regressor (you can adjust parameters if needed)
            model = RandomForestRegressor(n_estimators=100, random_state=42)
            model.fit(X_train, y_train)
            # Predict on the test set
            y_pred = model.predict(X_test)
            # Compute evaluation metrics
            mse = mean_squared_error(y_test, y_pred)
            mae = mean absolute error(y test, y pred)
            rmse = np.sqrt(mse)
            fold_metrics.append({'Fold': fold_counter, 'MSE': mse, 'RMSE': rmse, 'MAE': mae})
            # Print feature importances and metrics for this fold
            print(f"\nFold {fold_counter} Feature Importances:")
            print(model.feature_importances_)
            print(f"Fold {fold counter} -- MSE: {mse:.4f}, RMSE: {rmse:.4f}, MAE: {mae:.4f}")
            fold_counter += 1
       Fold 1 Feature Importances:
       [0.13509883 0.04889066 0.0397044 0.11138954 0.07242384 0.04762999
        0.0807937 0.06448376 0.08452475 0.06029241 0.05509629 0.12047118
        0.07920066]
       Fold 1 -- MSE: 0.0159, RMSE: 0.1260, MAE: 0.0758
       Fold 2 Feature Importances:
       [0.12023042 0.04771412 0.03790176 0.11896093 0.09513735 0.06779251
        0.05484233 \ 0.05283297 \ 0.04447057 \ 0.06549216 \ 0.05721671 \ 0.12615097
        0.1112572 ]
       Fold 2 -- MSE: 0.0184, RMSE: 0.1357, MAE: 0.0755
       Fold 3 Feature Importances:
       [0.11773791 0.04895031 0.04391118 0.10410141 0.11568593 0.06091138
        0.06695169\ 0.05460215\ 0.06155982\ 0.04989205\ 0.05474247\ 0.08716689
        0.1337868 ]
       Fold 3 -- MSE: 0.0140, RMSE: 0.1182, MAE: 0.0752
       Fold 4 Feature Importances:
        [0.09528621 \ 0.04475375 \ 0.04524532 \ 0.10368317 \ 0.10129583 \ 0.05781189 
        0.11120091 \ 0.0642027 \quad 0.08263978 \ 0.04649983 \ 0.04249197 \ 0.10819634
        0.0966923 1
       Fold 4 -- MSE: 0.0133, RMSE: 0.1152, MAE: 0.0758
       Fold 5 Feature Importances:
        \hbox{\tt [0.12029715\ 0.04005138\ 0.04161956\ 0.13746674\ 0.12284139\ 0.05889433] }
        0.08350061\ 0.05623217\ 0.06060666\ 0.04487732\ 0.05308742\ 0.1160085
       Fold 5 -- MSE: 0.0190, RMSE: 0.1377, MAE: 0.0767
In [6]: # Summarize all results into a DataFrame
        results df = pd.DataFrame(fold metrics)
        print("\n0verall Cross-Validation Results:")
        print(results_df)
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0ve	rall	Cross-Validation Results:		
	Fold	MSE	RMSE	MAE
0	1	0.015876	0.126000	0.075766
1	2	0.018417	0.135709	0.075462
2	3	0.013976	0.118220	0.075187
3	4	0.013261	0.115158	0.075764
4	5	0.018955	0.137677	0.076690

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