## Final Code

## December 6, 2023

```
[]: # Import necessary libraries for data visualization
     import matplotlib.pyplot as plt # Matplotlib for plotting
     import seaborn as sns # Seaborn for statistical data visualization
     # Import machine learning libraries
     from sklearn.ensemble import GradientBoostingRegressor # Gradient Boosting
      \hookrightarrowRegressor
     import pandas as pd # Pandas for data manipulation
     import numpy as np # NumPy for numerical operations
[]: # Read the dataset into a Pandas DataFrame
     df = pd.read_csv('C:/Users/rashi/OneDrive/Documents/4th sem/Capstone Project/
      →Final Dataset.csv')
[]: # Display the first few rows of the DataFrame to inspect the data
     df.head()
[]: # Provide information about the DataFrame, including data types and missing,
     \hookrightarrow values
     df.info()
[]: # Remove non-numeric characters from the "Weighted salary" column and convert_{\sqcup}
      ⇔it to float
     df["Weighted salary"] = df["Weighted salary"].str.replace('[^\d.]', '', u
      →regex=True).astype(float)
[]: # Check the number of missing values in each column
     df.isnull().sum()
[]: # Define a list of columns to impute with the median
     columns_to_impute = ["GMAT/GRE Required"]
[]: # Impute missing values in selected columns with the median
     for column in columns_to_impute:
         median value = df[column].median()
         df[column].fillna(median_value, inplace=True)
```

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[]: # Define a threshold for the maximum number of missing values allowed
     threshold = 250  # Adjust this threshold based on your dataset
[]: # Drop columns with a high number of missing values
     columns_to_drop = df.columns[df.isnull().sum() > threshold]
     df.drop(columns=columns_to_drop, inplace=True)
[]: # Check the number of missing values in each column
     df.isnull().sum()
[]: # Generate descriptive statistics for the dataset
     df.describe()
[]: # Setting the style for the plots
     sns.set_style("whitegrid")
     # Create a new figure
     plt.figure(figsize=(14, 6))
     # Create a subplot of 1 row and 2 columns for box plot and histogram
     plt.subplot(1, 2, 1)
     sns.boxplot(y=df['Weighted salary'])
     plt.title('Box plot of Weighted salary')
     plt.subplot(1, 2, 2)
     sns.histplot(df['Weighted salary'], kde=True, bins=30)
     plt.title('Histogram of Weighted salary')
     # Setting the style for the plots
     sns.set_style("whitegrid")
     # Create a new figure
     plt.figure(figsize=(14, 6))
     # Create a subplot of 1 row and 2 columns for box plot and histogram
     plt.subplot(1, 2, 1)
     sns.boxplot(y=df['International Diversity'])
     plt.title('Box plot of International Diversity')
     plt.subplot(1, 2, 2)
     sns.histplot(df['International Diversity'], kde=True, bins=30)
     plt.title('Histogram of International Diversity')
     # Setting the style for the plots
     sns.set_style("whitegrid")
     # Create a new figure
```

```
plt.figure(figsize=(14, 6))
# Create a subplot of 1 row and 2 columns for box plot and histogram
plt.subplot(1, 2, 1)
sns.boxplot(y=df['Salary percentage increase'])
plt.title('Box plot of Salary percentage increase')
plt.subplot(1, 2, 2)
sns.histplot(df['Salary percentage increase'], kde=True, bins=30)
plt.title('Histogram of Salary percentage increase')
# Setting the style for the plots
sns.set_style("whitegrid")
# Create a new figure
plt.figure(figsize=(14, 6))
# Create a subplot of 1 row and 2 columns for box plot and histogram
plt.subplot(1, 2, 1)
sns.boxplot(y=df['Faculty with doctorates (%)'])
plt.title('Box plot of Faculty with doctorates (%)')
plt.subplot(1, 2, 2)
sns.histplot(df['Faculty with doctorates (%)'], kde=True, bins=30)
plt.title('Histogram of Faculty with doctorates (%)')
# Setting the style for the plots
sns.set_style("whitegrid")
# Create a new figure
plt.figure(figsize=(14, 6))
# Create a subplot of 1 row and 2 columns for box plot and histogram
plt.subplot(1, 2, 1)
sns.boxplot(y=df['Effective Emp rate'])
plt.title('Box plot of Effective Emp rate')
plt.subplot(1, 2, 2)
sns.histplot(df['Effective Emp rate'], kde=True, bins=30)
plt.title('Histogram of Effective Emp rate')
plt.tight_layout()
plt.show()
```

```
[]: # Calculate feature correlations
    correlation_matrix = df.corr()
    correlations_with_target = correlation_matrix["#"].sort_values(ascending=False)
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print(correlations_with_target)
[]: # Multivariate Analysis (Correlation Heatmap)
     corr_matrix = df.corr()
     plt.figure(figsize=(25, 15))
     sns.heatmap(corr_matrix, annot=True, cmap='coolwarm')
     plt.title('Correlation Heatmap')
     plt.show()
[]: import pandas as pd
     from sklearn.ensemble import RandomForestRegressor
     import xgboost as xgb
     import matplotlib.pyplot as plt
     # Replace the target variable and feature columns as needed
     target_column = "#"
     feature_columns = [ "Weighted salary"
     , "Salary percentage increase"
     ,"Value for money rank"
     ,"Career progress rank"
     ,"Aims achieved (%)"
     "Effective Emp rate"
     ,"International work mobility rank"
     ,"International course experience rank"
     ,"Faculty with doctorates (%)"
     "Internship"
     ,"Overall Satisfaction"
     ,"International Diversity"
     ,"GMAT/GRE Required"
     ,"Female Empowerment Score"
     ]
     X = df[feature_columns] # Features
     y = df[target_column]
                             # Target variable
     # Method 1: Random Forest
     rf model = RandomForestRegressor()
     rf_model.fit(X, y)
     feature_importance_rf = rf_model.feature_importances_
     # Method 2: XGBoost
     xgb_model = xgb.XGBRegressor()
     xgb_model.fit(X, y)
     feature_importance_xgb = xgb_model.feature_importances_
```

# Display feature importance results for each method

```
print("Feature Importance (Random Forest):", feature_importance rf)
print("Feature Importance (XGBoost):", feature_importance_xgb)
# Visualize feature importances using bar plots
plt.figure(figsize=(12, 6))
plt.bar(range(len(feature_importance_rf)), feature_importance_rf)
plt.xlabel('Features')
plt.ylabel('Feature Importance Score (Random Forest)')
plt.title('Feature Importance Scores (Random Forest)')
plt.xticks(range(len(feature importance rf)), feature columns, rotation=90)
plt.show()
plt.figure(figsize=(12, 6))
plt.bar(range(len(feature_importance_xgb)), feature_importance_xgb)
plt.xlabel('Features')
plt.ylabel('Feature Importance Score (XGBoost)')
plt.title('Feature Importance Scores (XGBoost)')
plt.xticks(range(len(feature_importance_xgb)), feature_columns, rotation=90)
plt.show()
,"Salary percentage increase"
```

```
[]: metrics = [ "Weighted salary"
     ,"Value for money rank"
     ,"Career progress rank"
     #, "Career service rank"
     "Aims achieved (%)"
     #, "Employed at three months (main)"
     #, "Employed at three months (additional)"
     ,"Effective Emp rate"
     #, "Female faculty (%)"
     ,"Female students (%)"
     #, "Women on board (%)"
     #, "International faculty (%)"
     #, "International students (%)"
     #, "International board (%)"
     ,"International work mobility rank"
     ,"International course experience rank"
     ,"Faculty with doctorates (%)"
     #, "Three year average"
     ,"Internship"
     #, "Overall Satisfaction"
     #, "No. of Students per Staff"
     "International Diversity,
     #,"GMAT/GRE Required"
     #, "Female Empowerment Score"
```

```
from sklearn.datasets import make_regression
    from sklearn.model_selection import train_test_split
    from sklearn.ensemble import RandomForestRegressor
    from sklearn.svm import SVR
    from sklearn.metrics import mean_squared_error, mean_absolute_error, r2_score
    import numpy as np
    X= df[metrics]
    y= df["#"]
    # 2. Split data into training and testing sets
    →random_state=42)
    # 3. Train Random Forest and SVM models
    rf = RandomForestRegressor(n_estimators=100, random_state=42)
    rf.fit(X_train, y_train)
    svm = SVR(kernel='linear')
    svm.fit(X train, y train)
    # 4. Predict on test data
    y_pred_rf = rf.predict(X_test)
    y_pred_svm = svm.predict(X_test)
    # 5. Calculate and print metrics
    metrics = {
        "MSE": mean_squared_error,
        "MAE": mean_absolute_error,
        "RMSE": lambda y_true, y_pred: np.sqrt(mean_squared_error(y_true, y_pred)),
        "R^2": r2 score
    }
    print("Random Forest Metrics:")
    for name, func in metrics.items():
        print(f"{name}: {func(y_test, y_pred_rf)}")
    print("\nSVM Metrics:")
    for name, func in metrics.items():
        print(f"{name}: {func(y_test, y_pred_svm)}")
[]: # Assuming you have already defined your X and y as features and target variable
    # If not, use X = df[metrics] and y = df["#"] as you did in your previous code
    metrics = [ "Weighted salary"
    , "Salary percentage increase"
```

```
,"Value for money rank"
,"Career progress rank"
,"Aims achieved (%)"
,"Effective Emp rate"
,"International work mobility rank"
,"International course experience rank"
,"Faculty with doctorates (%)"
,"Internship"
,"Overall Satisfaction"
"International Diversity,
,"GMAT/GRE Required"
, "Female Empowerment Score"
from sklearn.datasets import make_regression
from sklearn.model_selection import train_test_split
from sklearn.ensemble import RandomForestRegressor
from sklearn.svm import SVR
from sklearn.metrics import mean_squared_error, mean_absolute_error, r2_score
import numpy as np
X= df[metrics]
y= df["#"]
# 2. Split data into training and testing sets
⇒random state=42)
# 3. Create and train the Gradient Boosting Regressor model
gb_regressor = GradientBoostingRegressor(n_estimators=100, random_state=42)
gb_regressor.fit(X_train, y_train)
# 4. Predict on the test data
y_pred_gb = gb_regressor.predict(X_test)
# 5. Calculate and print regression metrics
metrics = {
   "MSE": mean_squared_error,
   "MAE": mean_absolute_error,
   "RMSE": lambda y_true, y_pred: np.sqrt(mean_squared_error(y_true, y_pred)),
   "R^2": r2_score
}
print("Gradient Boosting Metrics:")
for name, func in metrics.items():
   print(f"{name}: {func(y_test, y_pred_gb)}")
```

```
[]: from sklearn.model_selection import RandomizedSearchCV
     # Simplified hyperparameter grid
     simple_param_grid = {
         'learning_rate': [0.01, 0.1, 0.3],
         'n_estimators': [50, 100, 200],
         'max_depth': [3, 5, 7],
         'min_child_weight': [1, 3],
         'gamma': [0, 0.2],
         'subsample': [0.8, 1.0],
         'colsample_bytree': [0.8, 1.0]
     }
     # Randomized search with cross-validation (fewer iterations for simplicity)
     xgb_simple random_search = RandomizedSearchCV(xgb.XGBRegressor(objective = 'reg:

¬squarederror', random_state=42),
                                                  1.1
      →param_distributions=simple_param_grid,
                                                    n_iter=20,
                                                    scoring='neg_mean_squared_error',
                                                    cv=3.
                                                    verbose=1.
                                                    n_jobs=-1,
                                                    random_state=42)
     xgb_simple_random_search.fit(X_train, y_train)
     # Best hyperparameters from the simplified search
     simple_best_params = xgb_simple_random_search.best_params_
     simple_best_params
[]: import xgboost as xgb
     from sklearn.metrics import mean_squared_error, r2_score
     from sklearn.metrics import mean_absolute_error
     # Given training and testing sets: X_train, X_test, y_train, y_test
     # Train XGBoost with the optimal hyperparameters
     optimal_xgb = xgb.XGBRegressor(
         subsample=0.8,
         n_estimators=200,
         min_child_weight=3,
         max depth=3,
         learning_rate=0.1,
         gamma=0.2,
         colsample_bytree=0.8,
         objective ='reg:squarederror',
```

```
random_state=42
)

optimal_xgb.fit(X_train, y_train)

# Predict on the test set
y_pred_optimal = optimal_xgb.predict(X_test)

# Evaluate the model's performance
mse_optimal = mean_squared_error(y_test, y_pred_optimal)
r2_optimal = r2_score(y_test, y_pred_optimal)
mae_optimal = mean_absolute_error(y_test, y_pred_optimal)
rmse_optimal = np.sqrt(mse_optimal)

print(f"MSE: {mse_optimal}")
print(f"R^2: {r2_optimal}")
print(f"MAE: {mae_optimal}")
print(f"RMSE: {rmse_optimal}")
```

[]: