human-resources-analytics

November 20, 2024

Human Resources Analytics

Imported All the necessary libraries:

```
[]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split

from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.ensemble import GradientBoostingClassifier
from sklearn.metrics import accuracy_score, confusion_matrix,u
classification_report
from sklearn.model_selection import GridSearchCV
```

Read the Data set and store it in a pandas Data Frame:

```
[]: df = pd.read_csv("Human_Resuorces_Analytics.csv")
```

How many rows and columns in the data frame:

```
[]: df.shape
```

[]: (311, 36)

Explore the names of columns in the data frame:

```
[]: df.columns
```

```
[]: Index(['Employee_Name', 'EmpID', 'MarriedID', 'MaritalStatusID', 'GenderID', 'EmpStatusID', 'DeptID', 'PerfScoreID', 'FromDiversityJobFairID', 'Salary', 'Termd', 'PositionID', 'Position', 'State', 'Zip', 'DOB', 'Sex', 'MaritalDesc', 'CitizenDesc', 'HispanicLatino', 'RaceDesc', 'DateofHire', 'DateofTermination', 'TermReason', 'EmploymentStatus', 'Department', 'ManagerName', 'ManagerID', 'RecruitmentSource', 'PerformanceScore', 'EngagementSurvey', 'EmpSatisfaction',
```

```
'SpecialProjectsCount', 'LastPerformanceReview_Date', 'DaysLateLast30', 'Absences'], dtype='object')
```

General Information About the Columns of the Data Frame :

[]: df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 311 entries, 0 to 310
Data columns (total 36 columns):

#	Column	Non-Null Count	Dtype
0	Employee_Name	311 non-null	object
1	EmpID	311 non-null	int64
2	MarriedID	311 non-null	int64
3	MaritalStatusID	311 non-null	int64
4	GenderID	311 non-null	int64
5	EmpStatusID	311 non-null	int64
6	DeptID	311 non-null	int64
7	PerfScoreID	311 non-null	int64
8	FromDiversityJobFairID	311 non-null	int64
9	Salary	311 non-null	int64
10	Termd	311 non-null	int64
11	PositionID	311 non-null	int64
12	Position	311 non-null	object
13	State	311 non-null	object
14	Zip	311 non-null	int64
15	DOB	311 non-null	object
16	Sex	311 non-null	object
17	MaritalDesc	311 non-null	object
18	CitizenDesc	311 non-null	object
19	HispanicLatino	311 non-null	object
20	RaceDesc	311 non-null	object
21	DateofHire	311 non-null	object
22	DateofTermination	104 non-null	object
23	TermReason	311 non-null	object
24	EmploymentStatus	311 non-null	object
25	Department	311 non-null	object
26	ManagerName	311 non-null	object
27	ManagerID	303 non-null	float64
28	RecruitmentSource	311 non-null	object
29	PerformanceScore	311 non-null	object
30	EngagementSurvey	311 non-null	float64
31	EmpSatisfaction	311 non-null	int64
32	${\tt SpecialProjectsCount}$	311 non-null	int64
33	${\tt LastPerformanceReview_Date}$	311 non-null	object
34	DaysLateLast30	311 non-null	int64

35 Absences 311 non-null int64

dtypes: float64(2), int64(16), object(18)

memory usage: 87.6+ KB

Display the first 10 rows of data frame:

[]: df.head(10)

[]:	Employee_	Name	EmpID	Marri	edID l	MaritalStatus	ID Gen	derI	D	\
0	Adinolfi, Wilso		10026		0		0		1	
1	Ait Sidi, Karthikeya		10084		1		1		1	
2	Akinkuolie, S		10196		1		1		0	
3	Alagbe,T	rina	10088		1		1		0	
4	Anderson, Ca		10069		0		2		0	
5	Anderson, Lin	da	10002		0		0		0	
6	Andreola, C	olby	10194		0		0		0	
7	Athwal,	Sam	10062		0		4		1	
8	Bachiochi, L	inda	10114		0		0		0	
9	Bacong, Alejan	dro	10250		0		2		1	
	EmpStatusID DeptID	Perf9	ScoreID	From	Divers	ityJobFairID	Salary		\	
0	1 5		4			0	62506		`	
1	5 3		3			0	104437			
2	5 5		3			0	64955			
3	1 5		3			0	64991			
4	5 5		3			0	50825			
5	1 5		4			0	57568			
6	1 4		3			0	95660			
7	1 5		3			0	59365			
8	3 5		3			1	47837			
9	1 3		3			0	50178	•••		
	ManagerName Man	agerII) Reci	ruitme	ntSour	ce Performanc	eScore	\		
0	Michael Albert	22.0			Linked		xceeds			
1	Simon Roup	4.0)		Inde	ed Fully	Meets			
2	Kissy Sullivan	20.0)		Linked	-	Meets			
3	Elijiah Gray	16.0)		Inde	•	Meets			
4	Webster Butler	39.0)	Googl	e Sear	•	Meets			
5	Amy Dunn	11.0)		Linked	In E	xceeds			
6	Alex Sweetwater	10.0)		Linked	In Fully	Meets			
7	Ketsia Liebig	19.0) Emp]	Loyee	Referra	al Fully	Meets			
8	Brannon Miller	12.0) Dive	rsity	Job Fa:	ir Fully	Meets			
9	Peter Monroe	7.0)		Inde	ed Fully	Meets			
	EngagementSurvey Emp	Satisi	faction	Speci	alProi	ectsCount \				
0	4.60		5	•	3	0				
1	4.96		3			6				
2	3.02		3			0				

3	4.84	5	0
4	5.00	4	0
5	5.00	5	0
6	3.04	3	4
7	5.00	4	0
8	4.46	3	0
9	5.00	5	6

LastPerformanceReview_Date DaysLateLast30 Absences

0	1/17/2019	0	1
1	2/24/2016	0	17
2	5/15/2012	0	3
3	1/3/2019	0	15
4	2/1/2016	0	2
5	1/7/2019	0	15
6	1/2/2019	0	19
7	2/25/2019	0	19
8	1/25/2019	0	4
9	2/18/2019	0	16

[10 rows x 36 columns]

What are the data types of the columns in the data frame:

[]: df.dtypes

[]:	Employee_Name	object
	EmpID	int64
	MarriedID	int64
	MaritalStatusID	int64
	GenderID	int64
	EmpStatusID	int64
	DeptID	int64
	PerfScoreID	int64
	${\tt FromDiversityJobFairID}$	int64
	Salary	int64
	Termd	int64
	PositionID	int64
	Position	object
	State	object
	Zip	int64
	DOB	object
	Sex	object
	MaritalDesc	object
	CitizenDesc	object
	HispanicLatino	object
	RaceDesc	object

DateofHire object DateofTermination object TermReason object EmploymentStatus object Department object ManagerName object ManagerID float64 RecruitmentSource object PerformanceScore object EngagementSurvey float64 ${\tt EmpSatisfaction}$ int64 SpecialProjectsCount int64 LastPerformanceReview_Date object DaysLateLast30 int64Absences int64

dtype: object

Check the missing values in the columns of the data frame:

[]: df.isnull().any()

[]: Employee_Name False EmpID False MarriedID False MaritalStatusID False GenderID False EmpStatusID False DeptID False PerfScoreID False FromDiversityJobFairID False False Salary Termd False PositionID False Position False State False False Zip DOB False Sex False MaritalDesc False CitizenDesc False HispanicLatino False RaceDesc False False DateofHire DateofTermination True TermReason False EmploymentStatus False Department False

False ManagerName ManagerID True RecruitmentSource False PerformanceScore False EngagementSurvey False EmpSatisfaction False SpecialProjectsCount False LastPerformanceReview_Date False DaysLateLast30 False Absences False dtype: bool

Count the missing values in the columns of the data frame:

[]: df.isnull().sum()

[]:	Employee_Name	0
	EmpID	0
	MarriedID	0
	MaritalStatusID	0
	GenderID	0
	EmpStatusID	0
	DeptID	0
	PerfScoreID	0
	${\tt FromDiversityJobFairID}$	0
	Salary	0
	Termd	0
	PositionID	0
	Position	0
	State	0
	Zip	0
	DOB	0
	Sex	0
	MaritalDesc	0
	CitizenDesc	0
	HispanicLatino	0
	RaceDesc	0
	DateofHire	0
	DateofTermination	207
	TermReason	0
	EmploymentStatus	0
	Department	0
	ManagerName	0
	ManagerID	8
	RecruitmentSource	0
	PerformanceScore	0
	EngagementSurvey	0

```
EmpSatisfaction 0
SpecialProjectsCount 0
LastPerformanceReview_Date 0
DaysLateLast30 0
Absences 0
dtype: int64
```

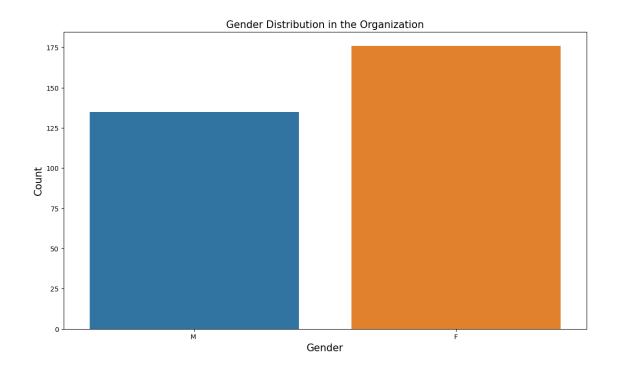
1 Section A (Data Exploration and Pre-processing):

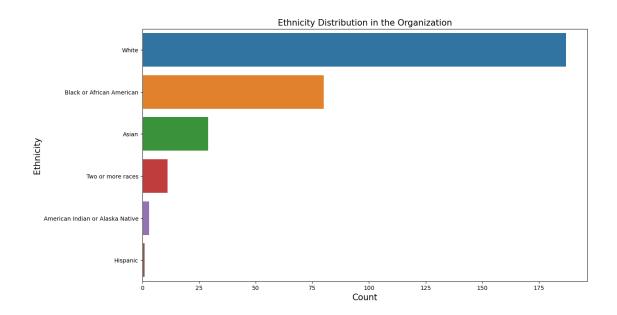
1.0.1 Explore the data using tables, visualizations, and other relevant methods

1. Overall Diversity Profile of the Organization:

To understand the diversity profile of the organization, we can analyze the distribution of employees based on their gender and ethnicity. We'll create a bar plot to visualize this

```
[]: # Count the number of employees in each gender category
     gender_counts = df['Sex'].value_counts()
     # Count the number of employees based on their ethnicity
     ethnicity_counts = df['RaceDesc'].value_counts()
     # Plot the gender distribution
     plt.figure(figsize=(14, 8))
     sns.countplot(data=df, x='Sex')
     plt.title('Gender Distribution in the Organization', fontsize = 15)
     plt.xlabel('Gender', fontsize = 15)
     plt.ylabel('Count', fontsize = 15)
     plt.show()
     # Plot the ethnicity distribution
     plt.figure(figsize=(14, 8))
     sns.countplot(data=df, y='RaceDesc', order=ethnicity_counts.index)
     plt.title('Ethnicity Distribution in the Organization', fontsize = 15)
     plt.xlabel('Count', fontsize = 15)
     plt.ylabel('Ethnicity', fontsize = 15)
     plt.show()
```



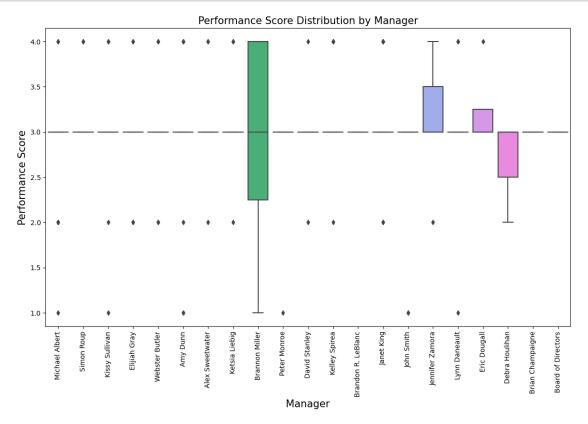


These plots will provide insights into the gender and ethnicity diversity within the organization.

2. Relationship between Manager and Performance Score:

We can investigate whether there is any relationship between the employee's manager and their performance score. To visualize this, we'll use a box plot to compare the performance scores across different managers.

```
[]: plt.figure(figsize=(14, 8))
    sns.boxplot(data=df, x='ManagerName', y='PerfScoreID')
    plt.xticks(rotation=90)
    plt.title('Performance Score Distribution by Manager', fontsize = 15)
    plt.xlabel('Manager', fontsize = 15)
    plt.ylabel('Performance Score', fontsize = 15)
    plt.show()
```

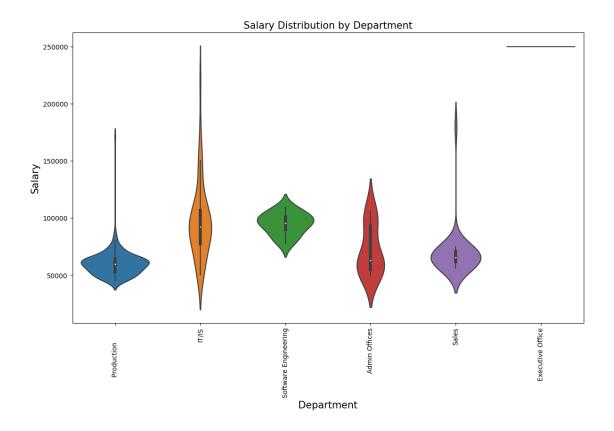


This box plot will help us identify any variations in performance scores based on the employee's manager.

3. Areas of Pay Inequity:

To identify areas of pay inequity within the company, we can compare the salaries across different departments. A violin plot can effectively showcase the distribution of salaries in each department.

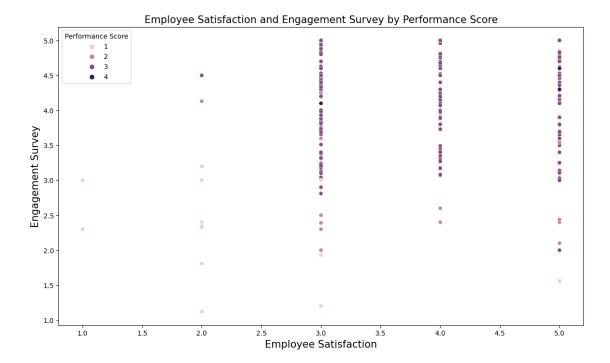
```
[]: plt.figure(figsize=(14, 8))
    sns.violinplot(data=df, x='Department', y='Salary')
    plt.xticks(rotation=90)
    plt.title('Salary Distribution by Department', fontsize = 15)
    plt.xlabel('Department', fontsize = 15)
    plt.ylabel('Salary', fontsize = 15)
    plt.show()
```



This visualization will highlight any variations in salary distribution across different departments, indicating potential areas of pay inequity.

4. Employee Satisfaction and Engagement Survey by Performance Score:

Let's create a scatter plot to visualize the relationship between employee satisfaction, engagement survey results, and performance scores. This will help us understand if there are any patterns or correlations among these variables.



This scatter plot will show how employee satisfaction and engagement survey results vary across different performance scores.

5. Salary Distribution by Department and Performance Score:

Let's visualize the distribution of salaries across different departments, considering the performance scores of employees as well. We'll use a box plot with color encoding to represent the performance scores.



In this plot, each box represents the salary distribution for a specific department, and the color of the boxes represents different performance scores. This allows us to simultaneously explore the salary distributions and performance scores across departments.

- 1) Overall Diversity Profile of the Organization:
- The gender distribution graph provides insights into the gender diversity within the organization.
- The ethnicity distribution graph offers an understanding of the ethnic diversity within the organization.
- These graphs can help us assess the overall diversity profile of the organization.
- 2) Relationship between Who a Person Works for and Their Performance Score:
- The box plot graph comparing performance scores across different managers provides insights into the potential relationship between an employee's manager and their performance score.
- This graph allows us to explore whether there is any correlation between the employee's manager and their performance score.
- 3) Areas of the Company Where Pay Is Not Equitable:
- The violin plot graph displaying the salary distribution by department can help identify potential areas where pay may not be equitable. This graph enables us to visually assess variations in salary distribution across different departments, indicating areas where pay inequity might exist.

1.0.2 Apply different methods of pre-processing

Handling the missing values in ManagerID column:

Convert date columns to datetime data type:

Replace 'no' and 'yes' with 'No' and 'Yes' in HispanicLatino column:

```
[]: df['HispanicLatino'] = df['HispanicLatino'].str.replace('no', 'No').

←replace('yes','Yes')
```

Create some new columns: AgeAtHire, EmploymentDuration , TimeSinceLastReview and Time-UntilTermination from date columns in the data frame:

Drop some unnecessary columns from data frame:

```
[]: df.drop(['Employee_Name','Position','State','TermReason','EmpStatusID',

→'Department','MaritalDesc','EmpID', 'Zip', 'ManagerName'], axis = 1,

→inplace = True)
```

Convert the values of categorical columns to numeric columns:

```
[]: df['Sex'] = df['Sex'].astype('category').cat.codes
    df['CitizenDesc'] = df['CitizenDesc'].astype('category').cat.codes
    df['HispanicLatino'] = df['HispanicLatino'].astype('category').cat.codes
    df['RaceDesc'] = df['RaceDesc'].astype('category').cat.codes
    df['RecruitmentSource'] = df['RecruitmentSource'].astype('category').cat.codes
    df['PerformanceScore'] = df['PerformanceScore'].astype('category').cat.codes
    #df['EmploymentStatus'] = df['EmploymentStatus'].astype('category').cat.codes
```

Display the updated dataset with the new features

```
[]: df.head(10)
```

[]:		MarriedID	MaritalStatu	sID	GenderID	\mathtt{DeptID}	PerfScoreI	D \
	0	0		0	1	5		4
	1	1		1	1	3		3
	2	1		1	0	5		3
	3	1		1	0	5		3
	4	0		2	0	5		3
	5	0		0	0	5		4
	6	0		0	0	4		3
	7	0		4	1	5		3
	8	0		0	0	5		3
	9	0		2	1	3		3
		EmamDirramai	+IohEoimID	Colo	rr Tormd	Dogiti	on TD Corr	\

	${\tt FromDiversityJobFairID}$	Salary	Termd	${\tt PositionID}$	Sex	 \
0	0	62506	0	19	1	
1	0	104437	1	27	1	
2	0	64955	1	20	0	
3	0	64991	0	19	0	
4	0	50825	1	19	0	
5	0	57568	0	19	0	
6	0	95660	0	24	0	
7	0	59365	0	19	1	
8	1	47837	0	19	0	
9	0	50178	0	14	1	

	PerformanceScore	EngagementSurvey	${\tt EmpSatisfaction}$	${\tt Special Projects Count}$	\
0	0	4.60	5	0	
1	1	4.96	3	6	
2	1	3.02	3	0	
3	1	4.84	5	0	
4	1	5.00	4	0	

5		0	5.00	5	
6		1	3.04	3	
7		1	5.00	4	
8		1	4.46	3	
9		1	5.00	5	
	DaysLateLast30	Absences	AgeAtHire	EmploymentDuration	\
0	0	1	28	1583	
1	0	17	39	444	
2	0	3	22	447	
3	0	15	19	2858	
4	0	2	21	1884	
5	0	15	34	1395	
6	0	19	35	359	
7	0	19	30	765	
8	0	4	-61	2312	
9	0	16	27	303	
	TimeSinceLastRe	view Time	UntilTermin	ation	
0		1620		1170	
1		2678		113	
2		4058		132	
3		1634		1156	
4		2701		218	
5		1630		1160	
6		1635		1155	
7		1581		1209	
8		1612		1178	

[10 rows x 26 columns]

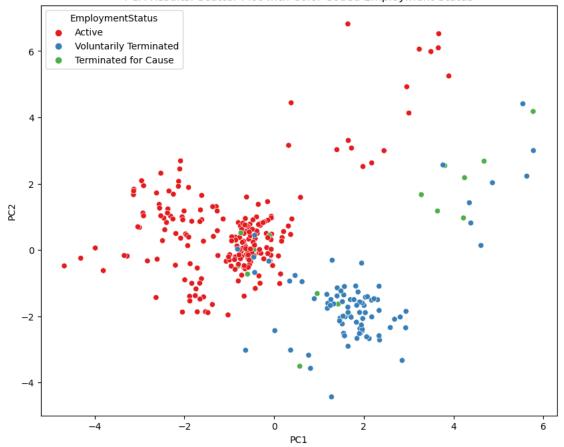
2 Section B (Dimensionality Reduction):

1. Apply PCA algorithm on the data.:

2. Create a scatter plot with color-coded observations based on employment status:

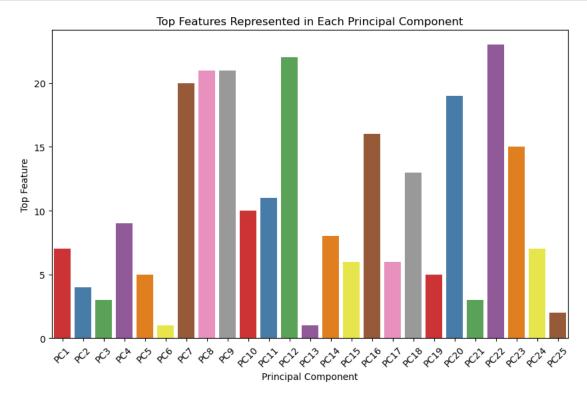
['Active', 'Voluntarily Terminated', 'Terminated for Cause'] = [0, 1, 2]

PCA Results: Scatter Plot with Color-Coded Employment Status



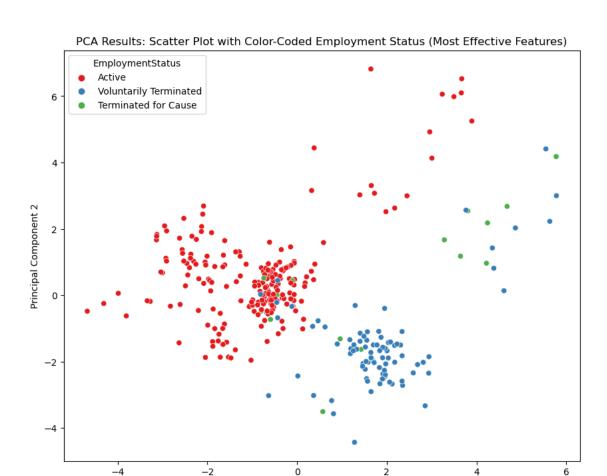
3. Identify the features most strongly represented in each component:

```
[]: plt.figure(figsize=(10, 6))
    sns.barplot(data=top_features.reset_index(), x='index', y=0, palette='Set1')
    plt.title('Top Features Represented in Each Principal Component')
    plt.xlabel('Principal Component')
    plt.ylabel('Top Feature')
    plt.xticks(rotation=45)
    plt.show()
```



The most effective features for separating employees by their employment status:

```
[]: # Visualize resulting clusters using top-ranked principal components
plt.figure(figsize=(10, 8))
sns.scatterplot(data=pca_df, x='PC1', y='PC2', hue=df['EmploymentStatus'],
palette='Set1')
plt.title('PCA Results: Scatter Plot with Color-Coded Employment Status (Most
Effective Features)')
plt.xlabel('Principal Component 1')
plt.ylabel('Principal Component 2')
plt.show()
```



This code will plot a scatter plot where each point represents an employee, and the color of the point will indicate their employment status. The principal components PC1 and PC2, which are among the most effective features in separating the employees, will be used as the x and y axes, respectively.

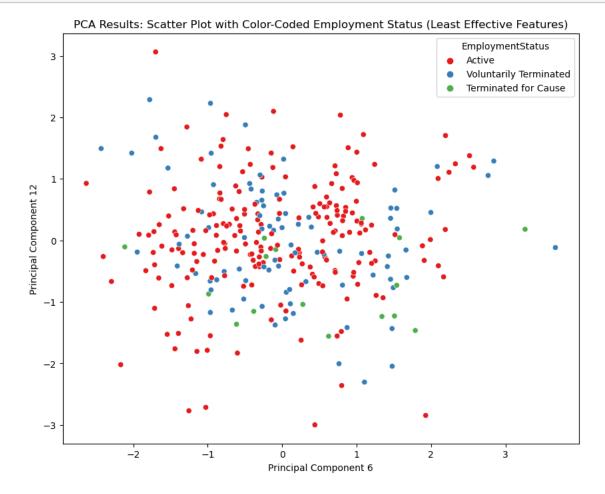
Principal Component 1

By visualizing the resulting clusters, you can observe how the most effective features contribute to the differentiation of employees based on their employment status. These features have a strong influence on the separation, leading to more distinct clusters for different employment statuses.

The least effective features for separating employees by their employment status:

```
[]: # Visualize resulting clusters using lower-ranked principal components
plt.figure(figsize=(10, 8))
sns.scatterplot(data=pca_df, x='PC6', y='PC12', hue=df['EmploymentStatus'],
□ □ palette='Set1')
plt.title('PCA Results: Scatter Plot with Color-Coded Employment Status (Least
□ □ Effective Features)')
plt.xlabel('Principal Component 6')
plt.ylabel('Principal Component 12')
```





This code will plot a scatter plot where each point represents an employee, and the color of the point will indicate their employment status. The principal components PC6 and PC12, which are among the least effective features in separating the employees, will be used as the x and y axes, respectively.

By visualizing the resulting clusters, you can observe how the least effective features contribute to the differentiation of employees based on employment status. However, since these features have less impact on the separation, the clusters might overlap more, making it harder to distinguish between different employment statuses.

Assign numeric codes to EmploymentStatus variable:

```
[]: df['EmploymentStatus'] = df['EmploymentStatus'].astype('category').cat.codes
```

Find the most effective feature:

```
[]: # Perform PCA on the dataset pca = PCA(n_components=2)
```

Termd

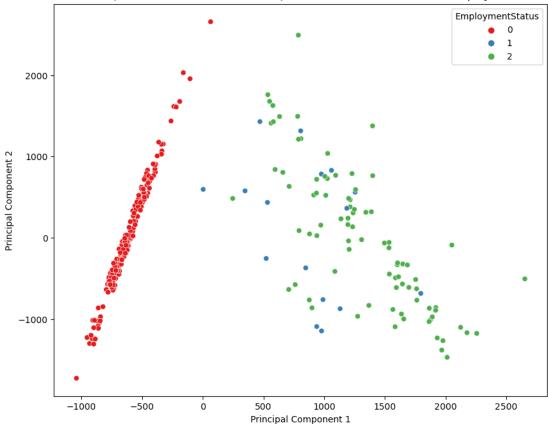
PCA Results (Without Most Effective Feature): Scatter Plot with Color-Coded Employment Status:

```
[]: # Exclude the most effective feature from the dataset
     modified_df = df.drop(columns=df.columns[most_effective_feature])
     # Apply PCA on the modified dataset
     pca_modified = PCA(n_components=2)
     pca_result_modified = pca_modified.fit_transform(modified_df)
     # Visualize the resulting clusters
     plt.figure(figsize=(10, 8))
     sns.scatterplot(data=pd.DataFrame(pca_result_modified, columns=['PC1', 'PC2']),__

¬x='PC1', y='PC2', hue=df['EmploymentStatus'], palette='Set1')

     plt.title('PCA Results (Without Most Effective Feature): Scatter Plot with ⊔
      →Color-Coded Employment Status')
     plt.xlabel('Principal Component 1')
     plt.ylabel('Principal Component 2')
     plt.show()
     # Explore the features most effective in separation (excluding the most _{\sqcup}
     ⇔effective feature)
     remaining_features = modified_df.columns
     print("Remaining Features:", remaining_features)
```





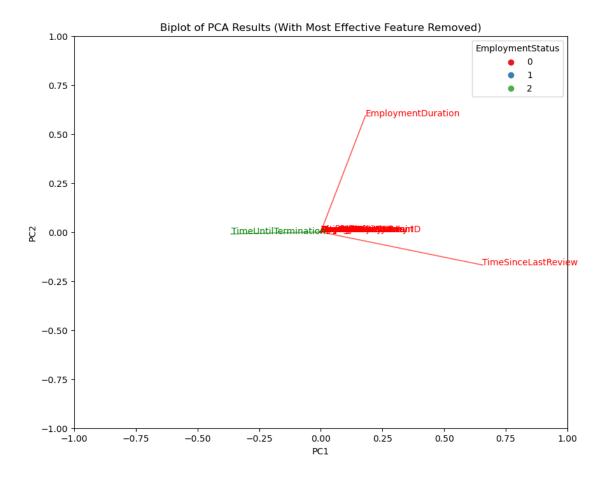
4. Biplot of PCA Results (With Most Effective Feature Removed):

```
[]: # Apply PCA on the modified dataset
pca_modified = PCA(n_components=2)
pca_result_modified = pca_modified.fit_transform(modified_df)

# Create a DataFrame for the modified PCA results
pca_df_modified = pd.DataFrame(data=pca_result_modified, columns=['PC1', 'PC2'])

# Get the explained variance ratio for PC1 and PC2
```

```
explained variance ratio modified = pca modified.explained variance ratio
# Create a biplot (with the most effective feature)
plt.figure(figsize=(10, 8))
# Plot the data points
sns.scatterplot(data=pca_df_modified, x='PC1', y='PC2', __
 ⇔hue=df['EmploymentStatus'], palette='Set1', alpha=0.6)
# Plot the variable vectors
for i, feature in enumerate(modified_df.columns[:-1]):
    arrow_length = np.sqrt(explained_variance_ratio_modified[0]) * pca_modified.
 ⇔components_[0, i]
    arrow_width = np.sqrt(explained_variance_ratio_modified[1]) * pca_modified.
 ⇔components_[1, i]
    plt.arrow(0, 0, arrow_length, arrow_width, color='r', alpha=0.5)
    plt.text(arrow_length, arrow_width, feature, color='r')
# Plot the most effective feature vector
arrow_length = np.sqrt(explained_variance_ratio_modified[0]) * pca_modified.
\hookrightarrowcomponents_[0, -1]
arrow width = np.sqrt(explained variance ratio modified[1]) * pca modified.
 \rightarrowcomponents_[1, -1]
plt.arrow(0, 0, arrow_length, arrow_width, color='g', alpha=0.5)
plt.text(arrow_length, arrow_width, modified_df.columns[-1], color='g')
# Set the limits of the plot
plt.xlim(-1, 1)
plt.ylim(-1, 1)
# Set labels and title
plt.xlabel('PC1')
plt.ylabel('PC2')
plt.title('Biplot of PCA Results (With Most Effective Feature Removed)')
# Show the plot
plt.show()
```



The biplot will provide insights into the relationships between the data points and variables. Here are some aspects to consider when interpreting the biplot:

Proximity of data points: Data points that are close to each other on the biplot are similar in terms of their feature values. Points that are far apart are dissimilar. You can observe the clusters of different employment status groups and identify patterns in their proximity.

Angle and direction of vectors: The vectors represent the variables/features in the biplot. The angle between two vectors indicates the correlation between the corresponding features. Vectors pointing in the same direction or opposite directions indicate positive or negative correlations, respectively.

Variables' contribution: The length of the vectors represents the contribution of the variables to the principal components. Longer vectors indicate higher importance in explaining the data's variance. By examining the lengths and orientations of the vectors, you can identify the variables that have the most significant influence on PC1 and PC2.

By considering these aspects and analyzing the biplot, you can gain insights into the relationships between data points and variables, identify patterns, and understand the variables' contributions to the separation of different employment status groups.

5. Bonus:

```
[]: # Perform PCA on the dataset
     pca = PCA(n_components=2)
     pca_result = pca.fit_transform(df)
     # Calculate the coordinates of each data point in the PCA space
     pca_df = pd.DataFrame(data=pca_result, columns=['PC1', 'PC2'])
     # Calculate the Euclidean distance of each data point from the centroid in the
      →PCA space
     centroid = pca_df.mean()
     distances = np.linalg.norm(pca_df - centroid, axis=1)
     # Define a threshold value to determine outliers
     threshold = np.percentile(distances, 95) # Adjust the percentile as needed
     # Identify the outliers
     outliers = df[distances > threshold]
     # Print the list of outliers
     print(outliers)
     # Explain the findings
     print("Number of outliers:", len(outliers))
     print("Threshold:", threshold)
         MarriedID MaritalStatusID
                                      GenderID DeptID PerfScoreID \
                                                      3
    18
                 0
                                              0
                                                                   3
    42
                  1
                                   1
                                              1
                                                      3
                                                                   3
    55
                  0
                                   0
                                                      5
                                                                   4
                                              1
    76
                                                      3
                  0
                                   0
                                              1
                                                                   4
    96
                  0
                                   0
                                              1
                                                      3
                                                                   4
    108
                  0
                                   0
                                              0
                                                      3
                                                                   3
                                                      6
                                                                   3
    131
                  1
                                   1
                                              0
    150
                  1
                                   1
                                              0
                                                      2
                                                                   3
                                                      3
                                                                   2
    190
                  1
                                   1
                                              1
                                                      4
                  0
                                   0
                                              1
                                                                   4
    212
    239
                                   0
                                                      3
                                                                   3
                  0
                                              0
    240
                 0
                                   0
                                              0
                                                      3
                                                                   3
    243
                  0
                                   0
                                              1
                                                      3
                                                                   3
                 0
                                   2
                                                      3
                                                                   3
    244
                                              1
    292
                 1
                                   1
                                              1
                                                      3
                                                                   3
    308
                 0
                                   0
                                              0
                                                      3
         FromDiversityJobFairID Salary Termd PositionID
                                                              Sex
    18
                               0 110000
                                              1
                                                           8
                                                                0 ...
    42
                               0 110929
                                              0
                                                           5
                                                                1
```

1 ...

0 170500

76 96 108 131 150 190 212 239 240 243 244		0 0 0 0 0 1 1 0 0	138888 178000 114800 180000 250000 157000 108987 120000 150290 140920 148999	0 0 1 0 0 0 0 1 1 0 0	13 12 8 11 16 13 24 29 7 13	1 1 0 0 0 1 1 0 0		
292 308		0	113999 220450	1 0	8 6	1		
18 42 55 76 96 108 131 150 190 212 239 240 243 244 292 308	PerformanceScore 1 1 0 0 0 1 1 1 1 1 1 1 1 1 1 1 1 0		gementSurvey 4.50 4.50 3.70 4.30 5.00 4.60 4.50 4.83 2.39 5.00 3.88 4.94 3.60 4.30 4.33 4.60					
18 42 55 76 96 108 131 150 190 212 239 240 243 244 292	SpecialProjectsCo	int D 5 7 0 5 5 4 0 0 6 3 7 6 7		0 Absences 0 8 0 8 0 15 0 4 0 15 0 10 0 19 0 10 4 13 0 13 0 12 0 17 0 13 0 8 0 9	AgeAt		re 28 56 25 57 330 57 52 443 225 332 441 566 339 53 28	

	${\tt EmploymentDuration}$	TimeSinceLastReview	$ exttt{TimeUntilTermination}$
18	432	3083	240
42	-307	1622	1168
55	2494	1602	1188
76	668	1633	1157
96	1664	1630	1160
108	27	3078	54
131	548	1616	1174
150	1220	1620	1170
190	1358	1584	1206
212	1400	2870	22
239	1405	1958	270
240	-430	1600	1190
243	1018	1588	1202
244	1395	3094	304
292	737	2321	7
308	2034	1585	1205

[16 rows x 26 columns]
Number of outliers: 16

308

Threshold: 39097.90035366955

In this code, df represents your original dataset. The code calculates the Euclidean distances of each data point from the centroid in the PCA space. The threshold is defined as the 95th percentile of the distances, which can be adjusted as per your requirements. The code then identifies the data points that exceed the threshold as outliers and prints the list of outliers.

0

16

30

By analyzing the outliers, you can observe if they have any common characteristics or patterns that differentiate them from the rest of the data points. This analysis can provide insights into potential anomalies or peculiarities in your dataset.

3 Section C (Classification):

Split the dataset into features (X) and target variable (y):

```
[]: X = df.drop('EmploymentStatus', axis=1)
y = df['EmploymentStatus']
```

Split the data into training and testing sets:

3.1 Random Forest Classifier:

Create a Random Forest classifier and perform parameter tuning using GridSearchCV:

```
[]: # Create the Random Forest classifier
    rf = RandomForestClassifier(random_state=42)

# Define the parameter grid for tuning
param_grid = {
        'n_estimators': [100, 200, 300],
        'max_depth': [None, 5, 10],
        'min_samples_split': [2, 5, 10],
        'min_samples_leaf': [1, 2, 4]
}

# Perform grid search to find the best parameters
grid_search = GridSearchCV(rf, param_grid, cv=5)
grid_search.fit(X_train, y_train)

# Get the best parameter values
best_params = grid_search.best_params_
best_params
```

Train the Random Forest classifier with the best parameters:

```
[]: # Create a new Random Forest classifier with the best parameters
rf_best = RandomForestClassifier(random_state=42, **best_params)

# Fit the classifier to the training data
rf_best.fit(X_train, y_train)
```

[]: RandomForestClassifier(min samples split=5, n estimators=200, random_state=42)

Get feature importances using Random Forest and explain their significance::

print(feature_importances)

```
Feature
                             Importance
24
      TimeUntilTermination
                               0.282077
23
       TimeSinceLastReview
                               0.278856
7
                      Termd
                               0.251752
16
          EngagementSurvey
                               0.024534
        EmploymentDuration
22
                               0.022366
13
                 ManagerID
                               0.021174
6
                    Salary
                               0.016125
19
            DaysLateLast30
                               0.013537
14
         RecruitmentSource
                               0.012429
                PositionID
8
                               0.012173
20
                  Absences
                               0.010548
21
                 AgeAtHire
                               0.010124
18
      SpecialProjectsCount
                               0.008907
15
          PerformanceScore
                               0.008434
4
               PerfScoreID
                               0.007148
3
                    DeptID
                               0.004572
1
           MaritalStatusID
                               0.003570
17
           EmpSatisfaction
                               0.003263
0
                 MarriedID
                               0.001891
10
               CitizenDesc
                               0.001606
5
    FromDiversityJobFairID
                               0.001384
12
                  RaceDesc
                               0.001174
2
                  GenderID
                               0.000991
9
                        Sex
                               0.000987
                               0.000378
11
            HispanicLatino
```

Compute accuracy for the model and provide sensitivity and specificity measurements for every class:

```
[]: # Predict the employment status using the trained random forest Classifier
y_pred = rf_best.predict(X_test)

# Compute accuracy
accuracy_rf = accuracy_score(y_test, y_pred)

# Compute sensitivity and specificity for each class
class_labels = rf_best.classes_
sensitivity = {}
specificity = {}

for label in class_labels:
    label_index = np.where(class_labels == label)[0][0]
    true_positive = np.sum((y_test == label) & (y_pred == label))
    false_negative = np.sum((y_test == label) & (y_pred != label))
    true_negative = np.sum((y_test != label) & (y_pred != label))
```

```
false_positive = np.sum((y_test != label) & (y_pred == label))
sensitivity[label] = true_positive / (true_positive + false_negative)
specificity[label] = true_negative / (true_negative + false_positive)

# Print accuracy, sensitivity, and specificity for each class
for label in class_labels:
    print(f"Class: {label}")
    print(f"Accuracy: {accuracy_rf:.4f}")
    print(f"Sensitivity: {sensitivity[label]:.4f}")
    print(f"Specificity: {specificity[label]:.4f}")
    print()
```

Class: 0

Accuracy: 0.9524 Sensitivity: 1.0000 Specificity: 1.0000

Class: 1

Accuracy: 0.9524 Sensitivity: 0.2500 Specificity: 1.0000

Class: 2

Accuracy: 0.9524 Sensitivity: 1.0000 Specificity: 0.9333

The above code performs parameter tuning using GridSearchCV to find the best combination of hyperparameters for the Random Forest classifier. Then, it trains the classifier using the best parameters and computes the feature importances.

The feature importances indicate the relative importance of each feature in predicting the target variable. Higher values indicate greater importance. By analyzing the feature importances, you can identify which features have a stronger influence on the model's predictions. It helps you understand which variables are more relevant in determining the employment status.

3.2 Support Vector Machine:

Create an SVM classifier and perform parameter tuning using GridSearchCV:

```
[]: # Create the SVM classifier
svm = SVC(random_state=42)

# Define the parameter grid for tuning
param_grid = {
    'C': [0.1, 1, 10],
    'kernel': ['linear', 'rbf'],
    'gamma': ['scale', 'auto']
```

```
# Perform grid search to find the best parameters
grid_search = GridSearchCV(svm, param_grid, cv=5)
grid_search.fit(X_train, y_train)

# Get the best parameter values
best_params = grid_search.best_params_
best_params
```

[]: {'C': 0.1, 'gamma': 'scale', 'kernel': 'linear'}

Train the SVM classifier with the best parameters:

```
[]: # Create a new SVM classifier with the best parameters
svm_best = SVC(random_state=42, **best_params)

# Fit the classifier to the training data
svm_best.fit(X_train, y_train)
```

[]: SVC(C=0.1, kernel='linear', random_state=42)

Compute accuracy for the model and provide sensitivity and specificity measurements for every class:

```
[]: # Predict the employment status using the trained sum Classifier
     y_pred = svm_best.predict(X_test)
     # Compute accuracy
     accuracy_svc = accuracy_score(y_test, y_pred)
     # Compute sensitivity and specificity for each class
     class_labels = rf_best.classes_
     sensitivity = {}
     specificity = {}
     for label in class_labels:
         label_index = np.where(class_labels == label)[0][0]
         true_positive = np.sum((y_test == label) & (y_pred == label))
         false_negative = np.sum((y_test == label) & (y_pred != label))
         true_negative = np.sum((y_test != label) & (y_pred != label))
         false_positive = np.sum((y_test != label) & (y_pred == label))
         sensitivity[label] = true_positive / (true_positive + false_negative)
         specificity[label] = true_negative / (true_negative + false_positive)
     # Print accuracy, sensitivity, and specificity for each class
     for label in class_labels:
         print(f"Class: {label}")
```

```
print(f"Accuracy: {accuracy_svc:.4f}")
print(f"Sensitivity: {sensitivity[label]:.4f}")
print(f"Specificity: {specificity[label]:.4f}")
print()
```

Class: 0
Accuracy: 0.9365
Sensitivity: 1.0000
Specificity: 1.0000
Class: 1
Accuracy: 0.9365
Sensitivity: 0.5000
Specificity: 0.9661
Class: 2
Accuracy: 0.9365
Sensitivity: 0.8889
Specificity: 0.9556

The above code performs parameter tuning using GridSearchCV to find the best combination of hyperparameters for the SVM classifier. Then, it trains the classifier using the best parameters and computes accuracy, confusion matrix, and classification report on the test data.

The accuracy score provides an overall measure of how well the model predicts the employment status. The confusion matrix gives a breakdown of the predicted and actual labels, showing true positives, true negatives, false positives, and false negatives. From the confusion matrix, you can calculate sensitivity (true positive rate) and specificity (true negative rate) for each class.

3.3 Gradient Boosting Classifier:

Create a Gradient Boosting Classifier and perform parameter tuning using GridSearchCV:

```
[]: # Create the Gradient Boosting Classifier
gbc = GradientBoostingClassifier(random_state=42)

# Define the parameter grid for tuning
param_grid = {
    'learning_rate': [0.1, 0.5, 1],
    'n_estimators': [50, 100, 200],
    'max_depth': [3, 5, 7]
}

# Perform grid search to find the best parameters
grid_search = GridSearchCV(gbc, param_grid, cv=5)
grid_search.fit(X_train, y_train)

# Get the best parameter values
```

```
best_params = grid_search.best_params_
best_params
```

[]: {'learning_rate': 0.5, 'max_depth': 3, 'n_estimators': 100}

Train the Gradient Boosting Classifier with the best parameters:

```
[]: # Create a new Gradient Boosting Classifier with the best parameters
gbc_best = GradientBoostingClassifier(random_state=42, **best_params)

# Fit the classifier to the training data
gbc_best.fit(X_train, y_train)
```

[]: GradientBoostingClassifier(learning_rate=0.5, random_state=42)

Compute accuracy for the model and provide sensitivity and specificity measurements for every class :

```
[]: # Predict the employment status using the trained Gradient Boosting Classifier
     y_pred = gbc_best.predict(X_test)
     # Compute accuracy
     accuracy_gbc = accuracy_score(y_test, y_pred)
     # Compute sensitivity and specificity for each class
     class_labels = rf_best.classes_
     sensitivity = {}
     specificity = {}
     for label in class_labels:
         label_index = np.where(class_labels == label)[0][0]
         true_positive = np.sum((y_test == label) & (y_pred == label))
         false_negative = np.sum((y_test == label) & (y_pred != label))
         true_negative = np.sum((y_test != label) & (y_pred != label))
         false_positive = np.sum((y_test != label) & (y_pred == label))
         sensitivity[label] = true_positive / (true_positive + false_negative)
         specificity[label] = true_negative / (true_negative + false_positive)
     # Print accuracy, sensitivity, and specificity for each class
     for label in class_labels:
         print(f"Class: {label}")
         print(f"Accuracy: {accuracy_gbc:.4f}")
         print(f"Sensitivity: {sensitivity[label]:.4f}")
         print(f"Specificity: {specificity[label]:.4f}")
         print()
```

Class: 0

Accuracy: 0.9206

Sensitivity: 1.0000 Specificity: 1.0000

Class: 1

Accuracy: 0.9206 Sensitivity: 0.5000 Specificity: 0.9492

Class: 2

Accuracy: 0.9206 Sensitivity: 0.8333 Specificity: 0.9556

Obtain the feature importance from a trained Gradient Boosting Classifier:

TimeSinceLastReview: 0.4540968698912378
TimeUntilTermination: 0.4145000809896252
EngagementSurvey: 0.03190955006893112
EmploymentDuration: 0.026467782003560827

Termd: 0.02381732416831714

Salary: 0.01966346301148855 PositionID: 0.013886897577337064 AgeAtHire: 0.006653270279196991

RecruitmentSource: 0.004805240933436853

Absences: 0.0030264448578457333 ManagerID: 0.0009500512010532898 DeptID: 0.00019079427836132264

 SpecialProjectsCount:
 2.3332388883857895e-05

 HispanicLatino:
 6.43832575253447e-06

 MaritalStatusID:
 9.472581456192514e-07

 EmpSatisfaction:
 9.458217961975859e-07

 DaysLateLast30:
 5.542809330630451e-07

Sex: 1.259239151668275e-08

CitizenDesc: 3.4160603284144587e-11
MarriedID: 2.8205621772115578e-11
PerfScoreID: 9.338868158221302e-12
RaceDesc: 1.940450127019775e-17

GenderID: 0.0

FromDiversityJobFairID: 0.0

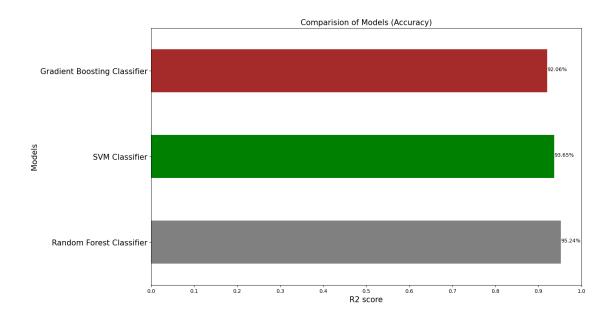
PerformanceScore: 0.0

The above code performs parameter tuning using GridSearchCV to find the best combination of hyperparameters for the Random Forest classifier. Then, it trains the classifier using the best parameters and computes the feature importances.

The feature importances indicate the relative importance of each feature in predicting the target variable. Higher values indicate greater importance. By analyzing the feature importances, you can identify which features have a stronger influence on the model's predictions. It helps you understand which variables are more relevant in determining the employment status.

Comparison of the model in terms of accuracy:

```
[]: scores = pd.Series([accuracy_rf, accuracy_svc, accuracy_gbc, ])
Model_Names = ['Random Forest Classifier','SVM Classifier','Gradient Boosting
Classifier']
```



4 Section D (Regression):

Import the necessary Libraries for Regression:

```
[]: import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LinearRegression
from sklearn.ensemble import RandomForestRegressor, GradientBoostingRegressor
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
from datetime import timedelta
```

Split the dataset into features (X) and target variable (y):

The TimeUntilTermination is already calculate above from DateofTermination and LastPerformanceReview Date.

```
[]: X = df.drop('TimeUntilTermination', axis=1)
y = df['TimeUntilTermination']
```

Split the data into training and testing sets:

```
[]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,_u 
-random_state=42)
```

4.1 Random Forest Regressor:

Create a Random Forest Regressor and perform parameter tuning using GridSearchCV:

```
[]: # Create the Random Forest Regressor
rf = RandomForestRegressor(random_state=42)

# Define the parameter grid for tuning
param_grid = {
        'n_estimators': [100, 200, 300],
        'max_depth': [None, 5, 10],
        'min_samples_split': [2, 5, 10],
        'min_samples_leaf': [1, 2, 4]
}

# Perform grid search to find the best parameters
grid_search = GridSearchCV(rf, param_grid, cv=5)
grid_search.fit(X_train, y_train)

# Get the best parameter values
best_params = grid_search.best_params_
best_params
```

[]: {'max_depth': None,
 'min_samples_leaf': 4,
 'min_samples_split': 10,
 'n_estimators': 300}

Train the Random Forest Regressor with the best parameters:

```
[]: # Create a new Random Forest Regressor with the best parameters
rf_best = RandomForestRegressor(random_state=42, **best_params)

# Fit the Regressor to the training data
rf_best.fit(X_train, y_train)
```

[]: RandomForestRegressor(min_samples_leaf=4, min_samples_split=10, n_estimators=300, random_state=42)

Get feature importances using Random Forest and explain their significance::

```
# Get feature importances
importances = rf_best.feature_importances_

# Create a dataframe to store feature importances
feature_importances = pd.DataFrame({'Feature': X_train.columns, 'Importance':_u
importances})

# Sort the features by importance in descending order
feature_importances = feature_importances.sort_values(by='Importance',_u
ascending=False)
```

```
# Print the feature importances
print(feature_importances)
```

```
Feature
                             Importance
13
          EmploymentStatus
                               0.454860
7
                      Termd
                               0.396037
24
       TimeSinceLastReview
                               0.129885
23
        EmploymentDuration
                               0.006168
14
                 ManagerID
                               0.002254
21
                   Absences
                               0.002114
6
                               0.001865
                     Salary
22
                  AgeAtHire
                               0.001118
          EngagementSurvey
17
                               0.001066
3
                     DeptID
                               0.000683
               CitizenDesc
10
                               0.000647
15
         RecruitmentSource
                               0.000581
8
                PositionID
                               0.000521
18
           EmpSatisfaction
                               0.000506
1
           MaritalStatusID
                               0.000350
12
                   RaceDesc
                               0.000310
5
    FromDiversityJobFairID
                               0.000278
20
            DaysLateLast30
                               0.000252
0
                  MarriedID
                               0.000223
2
                   GenderID
                               0.000078
19
      SpecialProjectsCount
                               0.000062
                               0.000058
9
                        Sex
4
               PerfScoreID
                               0.000054
11
            HispanicLatino
                               0.000015
          PerformanceScore
                               0.000014
```

Evaluate Random Forest Regressor using MAE, RMSE and R2 scores:

```
[]: y_pred = rf_best.predict(X_test)
    mean_absolute_error(y_pred, y_test)
    print('\nMean Absolute Error:', mean_absolute_error(y_pred, y_test))
    print('Root Mean Squared Error:', np.sqrt(mean_squared_error(y_pred, y_test)))
    print('R2 score :', r2_score(y_pred, y_test))
```

Mean Absolute Error: 43.38053964712802 Root Mean Squared Error: 91.50751070428193

R2 score: 0.9620994673655205

5 Linear Regression Model:

Create a Linear Regression Model and train it on training set:

```
[]: # Create the Linear Regression model
Lr = LinearRegression()
model_lr = Lr.fit(X_train, y_train)
y_pred = model_lr.predict(X_test)
```

Evaluate Linear Regression model using MAE, RMSE and R2 scores:

```
[]: y_pred = model_lr.predict(X_test)
    mean_absolute_error(y_pred, y_test)
    print('\nMean Absolute Error:', mean_absolute_error(y_pred, y_test))
    print('Root Mean Squared Error:', np.sqrt(mean_squared_error(y_pred, y_test)))
    print('R2 score :', r2_score(y_pred, y_test))
```

Mean Absolute Error: 66.80676107238526 Root Mean Squared Error: 99.47719653678247

R2 score: 0.9568176276262845

5.1 Gradient Boosting Regressor:

Create a Gradient Boosting regressor and perform parameter tuning using GridSearchCV:

```
[]: # Create the Gradient Boosting Regressor
gbr = GradientBoostingRegressor(random_state=42)

# Define the parameter grid for tuning
param_grid = {
    'learning_rate': [0.1, 0.5, 1],
    'n_estimators': [50, 100, 200],
    'max_depth': [3, 5, 7]
}

# Perform grid search to find the best parameters
grid_search = GridSearchCV(gbr, param_grid, cv=5)
grid_search.fit(X_train, y_train)

# Get the best parameter values
best_params = grid_search.best_params_
best_params
```

[]: {'learning_rate': 0.1, 'max_depth': 3, 'n_estimators': 50}

Train the Gradient Boosting regressor with the best parameters:

```
[]: # Create a new Gradient Boosting Regressor with the best parameters
gbr_best = GradientBoostingRegressor(random_state=42, **best_params)

# Fit the classifier to the training data
```

```
gbr_best.fit(X_train, y_train)
```

[]: GradientBoostingRegressor(n_estimators=50, random_state=42)

Obtain the feature importance from a trained Gradient Boosting regressor:

EmploymentStatus: 0.6707972305145762

Termd: 0.3002432521091373

TimeSinceLastReview: 0.007268161051847526 EmploymentDuration: 0.006547285960554348

DeptID: 0.003737772290699512 ManagerID: 0.0026318126412453106

DaysLateLast30: 0.0014975733931413603

Salary: 0.00147108909464373 Absences: 0.0011478487887006887 CitizenDesc: 0.0009712726619059215 RaceDesc: 0.0009557291830794313

RecruitmentSource: 0.0007111559693962408 MaritalStatusID: 0.0005448672412584865

AgeAtHire: 0.00044450208948630616

EngagementSurvey: 0.00034569345848166995

PositionID: 0.0003418473024510636

PerformanceScore: 0.00010742218492525613

FromDiversityJobFairID: 9.410581457913739e-05

PerfScoreID: 9.184394249165618e-05

SpecialProjectsCount: 4.9417050220901125e-05

MarriedID: 1.1725717787294e-07

GenderID: 0.0

Sex: 0.0

HispanicLatino: 0.0 EmpSatisfaction: 0.0

Evaluate Regression model using MAE, RMSE and R2 scores:

```
[]: y_pred = gbr_best.predict(X_test)
    mean_absolute_error(y_pred, y_test)
    print('\nMean Absolute Error:', mean_absolute_error(y_pred, y_test))
    print('Root Mean Squared Error:', np.sqrt(mean_squared_error(y_pred, y_test)))
    print('R2 score :', r2_score(y_pred, y_test))
```

Mean Absolute Error: 40.03239803442558 Root Mean Squared Error: 77.71043517573587

R2 score : 0.9726115670126408

Post-process the results and find the exact termination date for each employee

```
2022-03-31 12:08:49.982956800
2016-07-01 15:18:04.971715200
2012-09-25 11:07:14.155046400
2022-02-26 21:39:15.687532800
2016-06-07 17:38:38.820710400
2022-03-09 10:50:54.523334400
2022-02-28 10:41:12.283670400
2022-06-14 11:52:15.164025600
2022-04-11 06:21:42.030374400
2022-05-31 18:31:47.443555200
2016-09-15 07:55:35.822294400
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2022-06-18 04:29:27.597753600
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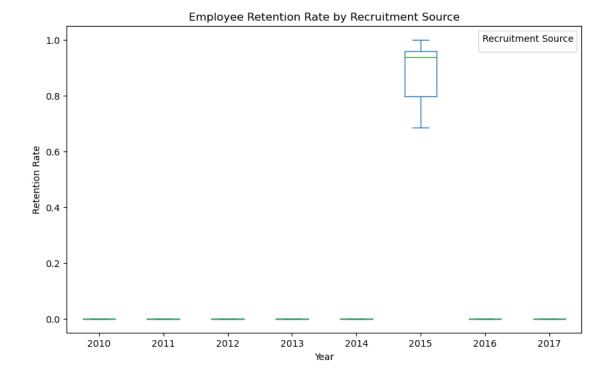
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2016-01-30 11:16:09.764025600
2022-06-03 07:00:20.782684800
2022-04-28 02:47:42.678988800
2022-04-24 06:32:49.100496
```

6 Section E:

Calculate the Employee Retention Rate for each year from 2008 until 2017 for every recruitment source and display it on a suitable graph:

```
[]: df1 = pd.read_csv("Human_Resuorces_Analytics.csv")
[]: import pandas as pd
     import matplotlib.pyplot as plt
     # Assuming you have already loaded the data into a DataFrame called 'df'
     df1['DateofTermination'] = df1['DateofTermination'].
      →fillna(df1['DateofTermination'].mode()[0])
     # Filter the data for the years 2008 to 2017
     df1['Year'] = pd.to datetime(df1['DateofTermination']).dt.year
     filtered_df = df1[df1['Year'].between(2008, 2017)]
     # Calculate the retention rate for each recruitment source and year
     retention_rates = filtered_df.groupby(['RecruitmentSource', 'Year']).
      \rightarrowapply(lambda x: 1 - x['Termd'].mean())
     # Plot the retention rates on a graph
     retention_rates.unstack().plot(kind='box', figsize=(10, 6))
     plt.xlabel('Year')
     plt.ylabel('Retention Rate')
     plt.title('Employee Retention Rate by Recruitment Source')
     plt.legend(title='Recruitment Source')
     plt.show()
```

No artists with labels found to put in legend. Note that artists whose label start with an underscore are ignored when legend() is called with no argument.



Calculate the diversity index for each department based on race, gender, and age and display the department in descending order of their diversity index:

```
[]: import pandas as pd
     # Assuming you have already loaded the data into a DataFrame called 'df'
     # Calculate age at hire
     df1['DOB'] = pd.to_datetime(df1['DOB'])
     df1['DateofHire'] = pd.to_datetime(df1['DateofHire'])
     df1['AgeAtHire'] = (df1['DateofHire'] - df1['DOB']).dt.days // 365
     # Calculate the diversity index
     diversity_df = df1.groupby('Department').agg({'RaceDesc': lambda x: len(x.
      ounique()), 'GenderID': lambda x: len(x.unique()), 'AgeAtHire': lambda x:⊔
      \rightarrowlen(x.unique())})
     diversity_df['DiversityIndex'] = diversity_df['RaceDesc'] +__

diversity_df['GenderID'] + diversity_df['AgeAtHire']

     # Sort departments in descending order of diversity index
     diversity_df = diversity_df.sort_values('DiversityIndex', ascending=False)
     # Display the departments and their diversity index
     print(diversity_df[['DiversityIndex']])
```

DiversityIndex

```
      Department

      Production
      51

      Sales
      28

      IT/IS
      27

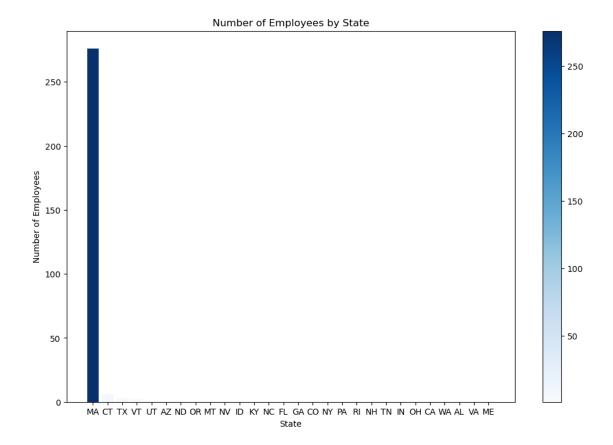
      Software Engineering
      12

      Admin Offices
      11

      Executive Office
      3
```

Create a map visualization that shows the number of employees currently working in the company from each state:

```
[]: import matplotlib.pyplot as plt
     # Count the number of employees from each state
     state_counts = df1['State'].value_counts()
     # Read the state codes and corresponding counts into lists
     state_codes = state_counts.index.tolist()
     employee_counts = state_counts.values.tolist()
     # Create a colormap for coloring the states
     colormap = plt.cm.get_cmap('Blues')
     # Plot the number of employees by state
     fig, ax = plt.subplots(figsize=(12, 8))
     bars = ax.bar(state_codes, employee_counts, color=colormap(employee_counts))
     # Add labels and title
     ax.set_xlabel('State')
     ax.set_ylabel('Number of Employees')
     ax.set_title('Number of Employees by State')
     # Create a colorbar legend
     sm = plt.cm.ScalarMappable(cmap=colormap, norm=plt.
      →Normalize(vmin=min(employee_counts), vmax=max(employee_counts)))
     sm.set array([])
     cbar = plt.colorbar(sm)
     # Show the plot
     plt.show()
```



7 Section F:

Classification Task:

```
from sklearn.metrics import accuracy_score, precision_score, recall_score, u
     ⇒f1_score, roc_auc_score
    # Assuming you have your true labels and predicted labels stored in y_true and_
     \hookrightarrow y_pred, respectively
    # Calculate accuracy
    accuracy = accuracy_score(y_test, y_pred)
    # Calculate precision
    precision = precision_score(y_test, y_pred, average='weighted')
    # Calculate recall
    recall = recall_score(y_test, y_pred, average='weighted')
    # Calculate F1 score
    f1 = f1_score(y_test, y_pred, average='weighted')
    y_pred_prob = rf_best.predict_proba(X_test)
    # Calculate AUC-ROC
    auc_roc = roc_auc_score(y_test, y_pred_prob, multi_class='ovr')
    print('Accuracy: ', accuracy)
    print('Precision: ', precision)
    print('Recall: ', recall)
    print('F1: ', recall)
    print('Auc_Roc: ', auc_roc)
    Accuracy: 0.9523809523809523
    Precision: 0.9487607908660539
    Recall: 0.9523809523809523
    F1: 0.9523809523809523
    Auc_Roc: 0.9704052451698403
    Regression Task:
[]: X = df.drop('TimeUntilTermination', axis=1)
    y = df['TimeUntilTermination']
    ⇒random state=42)
    # Create a new Gradient Boosting Regressor with the best parameters
    gbr_best = GradientBoostingRegressor(random_state=42, **best_params)
    # Fit the classifier to the training data
```

```
gbr_best.fit(X_train, y_train)
# Predict the employment status using the trained model
y_pred = gbr_best.predict(X_test)
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
\# Assuming you have your true values and predicted values stored in y_true and \sqcup
\hookrightarrow y_pred, respectively
# Calculate mean absolute error
mae = mean_absolute_error(y_test, y_pred)
# Calculate mean squared error
mse = mean_squared_error(y_test, y_pred)
# Calculate root mean squared error
rmse = np.sqrt(mse)
# Calculate R-squared
r2 = r2_score(y_test, y_pred)
print('MAE: ', mae)
print('MSE: ', mse)
print('RMSE: ', rmse)
print('R2: ', r2)
```

MAE: 40.03239803442558 MSE: 6038.911735202247 RMSE: 77.71043517573587 R2: 0.9734915883138399

[]: