Problem Statement

Business context

Employee Promotion means the ascension of an employee to higher ranks, this aspect of the job is what drives employees the most. The ultimate reward for dedication and loyalty towards an organization and the HR team plays an important role in handling all these promotion tasks based on ratings and other attributes available.

The HR team in JMD company stored data on the promotion cycle last year, which consists of details of all the employees in the company working last year and also if they got promoted or not, but every time this process gets delayed due to so many details available for each employee - it gets difficult to compare and decide.

Objective

For the upcoming appraisal cycle, the HR team wants to utilize the stored data and leverage machine learning to make a model that will predict if a person is eligible for promotion or not. You, as a data scientist at JMD company, need to come up with the best possible model that will help the HR team to predict if a person is eligible for promotion or not.

Data Description

- employee_id: Unique ID for the employee
- department: Department of employee
- region: Region of employment (unordered)
- education: Education Level
- gender: Gender of Employee
- recruitment channel: Channel of recruitment for employee
- no of trainings: no of other training completed in the previous year on soft skills, technical skills, etc.
- age: Age of Employee
- previous year rating: Employee Rating for the previous year
- length of service: Length of service in years
- awards won: if awards won during the previous year then 1 else 0
- avg training score: Average score in current training evaluations
- is_promoted: (Target) Recommended for promotion

Please read the instructions carefully before starting the project.

This is a commented Jupyter IPython Notebook file in which all the instructions and tasks to be performed are mentioned.

- Blanks '___' are provided in the notebook that needs to be filled with an appropriate code to get the correct result. With every '___' blank, there is a comment that briefly describes what needs to be filled in the blank space.
- Identify the task to be performed correctly, and only then proceed to write the required code.
- Fill the code wherever asked by the commented lines like "# write your code here" or "# complete the code". Running incomplete code may throw error.
- Please run the codes in a sequential manner from the beginning to avoid any unnecessary errors.
- Add the results/observations (wherever mentioned) derived from the analysis in the presentation and submit the same.

Importing necessary libraries

```
# uncomment and run the following line if Google Colab is being used
# !pip install scikit-learn==1.2.2 seaborn==0.13.1 matplotlib==3.7.1 numpy==1.25.2 pandas
==1.5.3 imbalanced-learn==0.10.1 xgboost==2.0.3 -q --user
```

In [4]:

```
# Installing the libraries with the specified version.
# uncomment and run the following lines if Jupyter Notebook is being used
# !pip install scikit-learn==1.2.2 seaborn==0.13.1 matplotlib==3.7.1 numpy==1.25.2 pandas
==1.5.3 imbalanced-learn==0.10.1 xgboost==2.0.3 -q --user
# !pip install --upgrade -q threadpoolctl
```

In [5]:

```
# Libraries to help with reading and manipulating data
import pandas as pd
import numpy as np
# Libaries to help with data visualization
import matplotlib.pyplot as plt
import seaborn as sns
# To tune model, get different metric scores, and split data
from sklearn.metrics import (
   f1 score,
   accuracy score,
   recall score,
   precision score,
   confusion matrix,
   roc auc score,
   ConfusionMatrixDisplay,
from sklearn.model_selection import train_test_split, StratifiedKFold, cross_val_score
# To be used for data scaling and one hot encoding
from sklearn.preprocessing import StandardScaler, MinMaxScaler, OneHotEncoder
# To oversample and undersample data
from imblearn.over sampling import SMOTE
from imblearn.under sampling import RandomUnderSampler
# To do hyperparameter tuning
from sklearn.model selection import RandomizedSearchCV
# To impute missing values
from sklearn.impute import SimpleImputer
# To define maximum number of columns to be displayed in a dataframe
pd.set option("display.max columns", None)
# To supress scientific notations for a dataframe
pd.set option("display.float format", lambda x: "%.3f" % x)
# To help with model building
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import (
   AdaBoostClassifier,
   GradientBoostingClassifier,
   RandomForestClassifier,
   BaggingClassifier,
from xgboost import XGBClassifier
# To suppress scientific notations
pd.set option("display.float format", lambda x: "%.3f" % x)
# To supress warnings
import warnings
warnings.filterwarnings("ignore")
```

Loading the dataset

```
In [6]:
```

```
promotion = pd.read csv("employee promotion.csv")
```

Data Overview

The initial steps to get an overview of any dataset is to:

- observe the first few rows of the dataset, to check whether the dataset has been loaded properly or not
- · get information about the number of rows and columns in the dataset
- find out the data types of the columns to ensure that data is stored in the preferred format and the value of each property is as expected.
- check the statistical summary of the dataset to get an overview of the numerical columns of the data

Checking the shape of the dataset

```
In [7]:
```

```
# Checking the number of rows and columns in the training data promotion.shape ## Complete the code to view dimensions of the train data
```

```
Out[7]:
```

(54808, 13)

In [8]:

```
# let's create a copy of the data
data = promotion.copy()
```

Displaying the first few rows of the dataset

```
In [9]:
```

```
# let's view the first 5 rows of the data data.head() ## Complete the code to view top 5 rows of the data
```

Out[9]:

	employee_id	department	region	education	gender	recruitment_channel	no_of_trainings	age	previous_year_rating	k
0	65438	Sales & Marketing	region_7	Master's & above	f	sourcing	1	35	5.000	
1	65141	Operations	region_22	Bachelor's	m	other	1	30	5.000	
2	7513	Sales & Marketing	region_19	Bachelor's	m	sourcing	1	34	3.000	
3	2542	Sales & Marketing	region_23	Bachelor's	m	other	2	39	1.000	
4	48945	Technology	region_26	Bachelor's	m	other	1	45	3.000	
4										F

```
In [10]:
```

```
# let's view the last 5 rows of the data
data.tail() ## Complete the code to view last 5 rows of the data
```

```
Out[10]:
```

54803	employegg.jd	decartment	reg regien	Baucatips	gender	recruitment chappel	no_of_trainings	age	previous_year_ratir
54804	74592	Operations	region_27	Master's & above	f	other	1	37	2.00
54805	13918	Analytics	region_1	Bachelor's	m	other	1	27	5.00
54806	13614	Sales & Marketing	region_9	NaN	m	sourcing	1	29	1.00
54807	51526	HR	region_22	Bachelor's	m	other	1	27	1.00
4									<u> </u>

Checking the data types of the columns for the dataset

```
In [11]:
```

```
# let's check the data types of the columns in the dataset
data.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 54808 entries, 0 to 54807
Data columns (total 13 columns):
 # Column
                           Non-Null Count Dtype
                            _____
 0 employee id
                            54808 non-null int64
 1 department
                            54808 non-null object
 2 region
                            54808 non-null object
 3 education
                            52399 non-null object
                           54808 non-null object
 4 gender
 5 recruitment channel 54808 non-null object
 6 no_of_trainings 54808 non-null int64
                           54808 non-null int64
   age
 7
    previous_year_rating 50684 non-null float64
 8
9 length_of_service 54808 non-null int64
10 awards_won 54808 non-null int64
11 avg_training_score 52248 non-null float64
12 is_promoted 54808 non-null int64
dtypes: float64(2), int64(6), object(5)
memory usage: 5.4+ MB
```

Checking for duplicate values

```
In [12]:
```

```
# let's check for duplicate values in the data
data.duplicated().sum() ## Complete the code to check duplicate entries in the data
```

```
Out[12]:
```

0

Checking for missing values

```
In [13]:
```

```
# let's check for missing values in the data data.isnull().sum() ## Complete the code to check missing entries in the train data
```

Out[13]:

```
employee_id 0
department 0
region 0
education 2409
gender 0
recruitment_channel 0
no_of_trainings 0
age 0
```

length_of_service 0
awards_won 0
avg_training_score 2560
is_promoted 0
dtype: int64

Statistical summary of the dataset

In [14]:

let's view the statistical summary of the numerical columns in the data
data.describe() ## Complete the code to print the statitical summary of the train data

Out[14]:

	employee_id	no_of_trainings	age	previous_year_rating	length_of_service	awards_won	avg_training_score	is_pr
count	54808.000	54808.000	54808.000	50684.000	54808.000	54808.000	52248.000	54
mean	39195.831	1.253	34.804	3.329	5.866	0.023	63.712	
std	22586.581	0.609	7.660	1.260	4.265	0.150	13.522	
min	1.000	1.000	20.000	1.000	1.000	0.000	39.000	
25%	19669.750	1.000	29.000	3.000	3.000	0.000	51.000	
50%	39225.500	1.000	33.000	3.000	5.000	0.000	60.000	
75%	58730.500	1.000	39.000	4.000	7.000	0.000	77.000	
max	78298.000	10.000	60.000	5.000	37.000	1.000	99.000	
4								Þ

In [15]:

let's view the statistical summary of the numerical columns in the data data.describe().T

Out[15]:

	count	mean	std	min	25%	50%	75%	max
employee_id	54808.000	39195.831	22586.581	1.000	19669.750	39225.500	58730.500	78298.000
no_of_trainings	54808.000	1.253	0.609	1.000	1.000	1.000	1.000	10.000
age	54808.000	34.804	7.660	20.000	29.000	33.000	39.000	60.000
previous_year_rating	50684.000	3.329	1.260	1.000	3.000	3.000	4.000	5.000
length_of_service	54808.000	5.866	4.265	1.000	3.000	5.000	7.000	37.000
awards_won	54808.000	0.023	0.150	0.000	0.000	0.000	0.000	1.000
avg_training_score	52248.000	63.712	13.522	39.000	51.000	60.000	77.000	99.000
is_promoted	54808.000	0.085	0.279	0.000	0.000	0.000	0.000	1.000

Let's check the number of unique values in each column

In [16]:

data.nunique()

Out[16]:

employee id	54808
department	9
region	34
education	3
gender	2
recruitment_channel	3
no_of_trainings	10
200	/1 1

```
aye
                            4 1
                            5
previous year rating
                            35
length_of_service
                            2
awards won
                           59
avg_training_score
                            2
is promoted
dtype: int64
In [17]:
for i in data.describe(include=["object"]).columns:
    print("Unique values in", i, "are :")
    print(data[i].value_counts())
    print("*" * 50)
Unique values in department are :
department
Sales & Marketing 16840
Operations
                     11348
Technology
                      7138
Procurement
                      7138
Analytics
                      5352
                      2536
Finance
HR
                       2418
Legal
                       1039
R&D
Name: count, dtype: int64
***********
Unique values in region are :
region
region 2
            12343
region_22
             6428
region_22 6428
region_7 4843
region_15 2808
region_13 2648
region_26 2260
region_31 1935
region_4 1703
region_27 1659
region_16 1465
region_28 1318
region_11 1315
region_23 1175
region_29 994
              994
region 29
region 32
              945
region 19
              874
region 20
              850
region 14
              827
region 25
              819
region 17
               796
               766
region 5
region 6
               690
region_30
               657
region 8
               655
region 10
               648
              610
region_1
              508
region 24
              500
region_12
region 9
              420
region 21
              411
region 3
              346
              292
region 34
region 33
              269
region_18 31
Name: count, dtype: int64
Unique values in education are :
education
Bachelor's
                     36669
Master's & above 14925
Below Secondary 805
Name: count, dtype: int64
**************
```

```
Unique values in gender are :
gender
   38496
m
f
    16312
Name: count, dtype: int64
*************
Unique values in recruitment channel are :
recruitment channel
          30446
other
          23220
sourcing
referred
          1142
Name: count, dtype: int64
***********
In [18]:
# ID column consists of uniques ID for clients and hence will not add value to the modeli
data.drop(columns="employee id", inplace=True)
In [19]:
data["is promoted"].value_counts(1)
Out[19]:
is promoted
  0.915
   0.085
Name: proportion, dtype: float64
```

Exploratory Data Analysis

The below functions need to be defined to carry out the Exploratory Data Analysis.

```
In [20]:
```

```
# function to plot a boxplot and a histogram along the same scale.
def histogram boxplot(data, feature, figsize=(12, 7), kde=False, bins=None):
   Boxplot and histogram combined
   data: dataframe
   feature: dataframe column
   figsize: size of figure (default (12,7))
   kde: whether to the show density curve (default False)
   bins: number of bins for histogram (default None)
   f2, (ax_box2, ax_hist2) = plt.subplots(
       nrows=2, # Number of rows of the subplot grid= 2
       sharex=True, # x-axis will be shared among all subplots
       gridspec_kw={"height_ratios": (0.25, 0.75)},
       figsize=figsize,
      # creating the 2 subplots
   sns.boxplot(
       data=data, x=feature, ax=ax box2, showmeans=True, color="violet"
     # boxplot will be created and a triangle will indicate the mean value of the colum
n
   sns.histplot(
       data=data, x=feature, kde=kde, ax=ax hist2, bins=bins, palette="winter"
   ) if bins else sns.histplot(
       data=data, x=feature, kde=kde, ax=ax hist2
   ) # For histogram
   ax hist2.axvline(
       data[feature].mean(), color="green", linestyle="--"
   ) # Add mean to the histogram
```

```
ax_hist2.axvline(
    data[feature].median(), color="black", linestyle="-"
) # Add median to the histogram
```

In [21]:

```
# function to create labeled barplots
def labeled barplot(data, feature, perc=False, n=None):
    Barplot with percentage at the top
    data: dataframe
    feature: dataframe column
   perc: whether to display percentages instead of count (default is False)
   n: displays the top n category levels (default is None, i.e., display all levels)
    total = len(data[feature]) # length of the column
    count = data[feature].nunique()
    if n is None:
       plt.figure(figsize=(count + 1, 5))
    else:
       plt.figure(figsize=(n + 1, 5))
    plt.xticks(rotation=90, fontsize=15)
    ax = sns.countplot(
       data=data,
       x=feature,
       palette="Paired",
       order=data[feature].value counts().index[:n].sort values(),
    for p in ax.patches:
       if perc == True:
           label = "{:.1f}%".format(
               100 * p.get height() / total
           ) # percentage of each class of the category
       else:
            label = p.get_height() # count of each level of the category
       x = p.get_x() + p.get_x() / 2 # width_of_the_plot
        y = p.get height() # height of the plot
       ax.annotate(
            label,
            (x, y),
           ha="center",
           va="center",
           size=12,
           xytext=(0, 5),
           textcoords="offset points",
        ) # annotate the percentage
    plt.show() # show the plot
```

In [22]:

```
# function to plot stacked bar chart

def stacked_barplot(data, predictor, target):
    """
    Print the category counts and plot a stacked bar chart

    data: dataframe
    predictor: independent variable
    target: target variable
    """
    count = data[predictor].nunique()
    sorter = data[target].value_counts().index[-1]
```

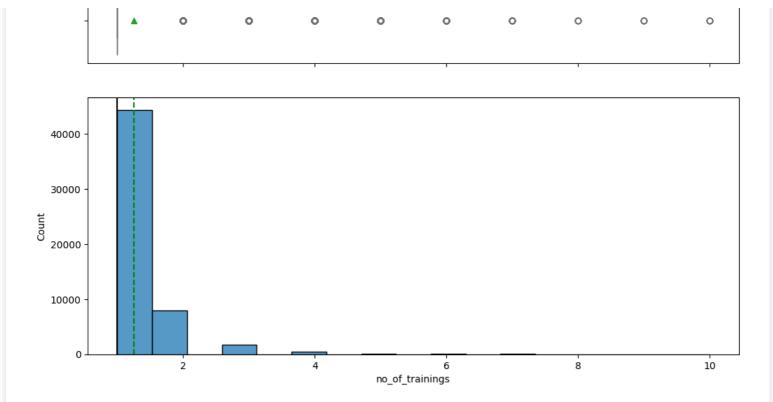
In [23]:

```
### Function to plot distributions
def distribution plot wrt target(data, predictor, target):
    fig, axs = plt.subplots(2, 2, figsize=(12, 10))
    target uniq = data[target].unique()
    axs[0, 0].set title("Distribution of target for target=" + str(target uniq[0]))
    sns.histplot(
       data=data[data[target] == target uniq[0]],
       x=predictor,
       kde=True,
       ax=axs[0, 0],
       color="teal",
    )
    axs[0, 1].set title("Distribution of target for target=" + str(target uniq[1]))
    sns.histplot(
       data=data[data[target] == target uniq[1]],
       x=predictor,
       kde=True,
       ax=axs[0, 1],
        color="orange",
    axs[1, 0].set title("Boxplot w.r.t target")
    sns.boxplot(data=data, x=target, y=predictor, ax=axs[1, 0], palette="gist rainbow")
    axs[1, 1].set title("Boxplot (without outliers) w.r.t target")
    sns.boxplot(
       data=data,
       x=target,
       y=predictor,
       ax=axs[1, 1],
       showfliers=False,
       palette="gist rainbow",
    )
    plt.tight layout()
   plt.show()
```

Univariate analysis

Observations on No. of Trainings

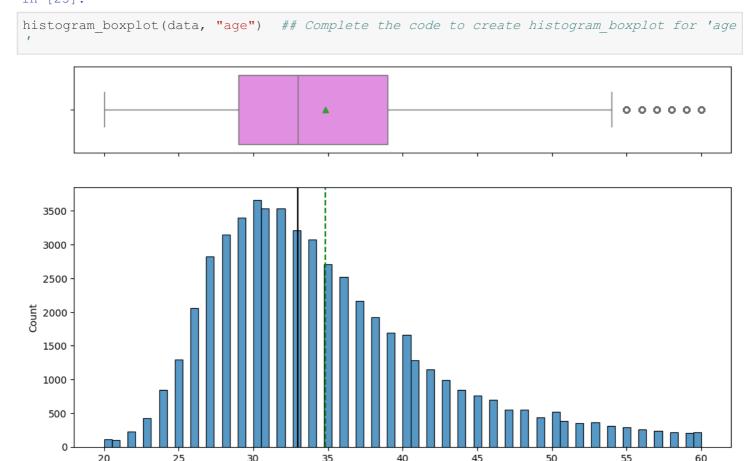
```
In [24]:
histogram_boxplot(data, "no_of_trainings")
```



Let's see the distribution of age of employee

Observations on Age

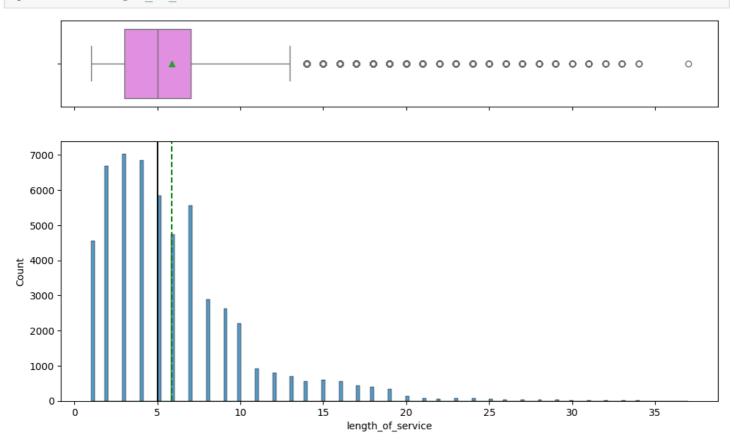
In [25]:



Observations on Length of Service

In [26]:

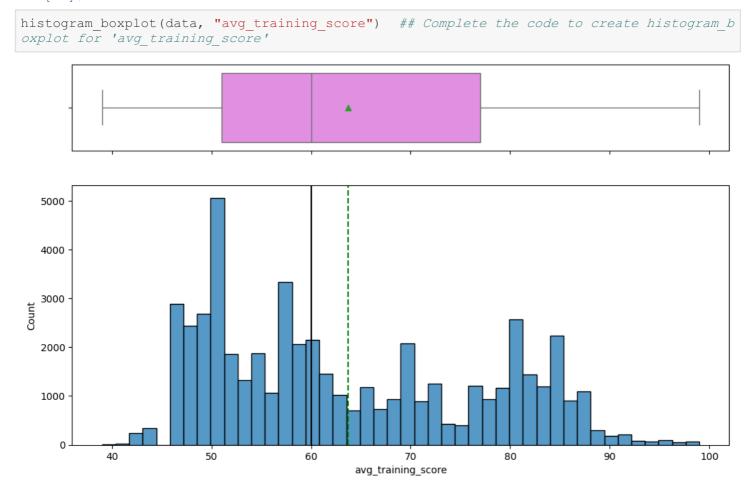




Let's see the distribution of average training score of employee

Observations on Average Training Score

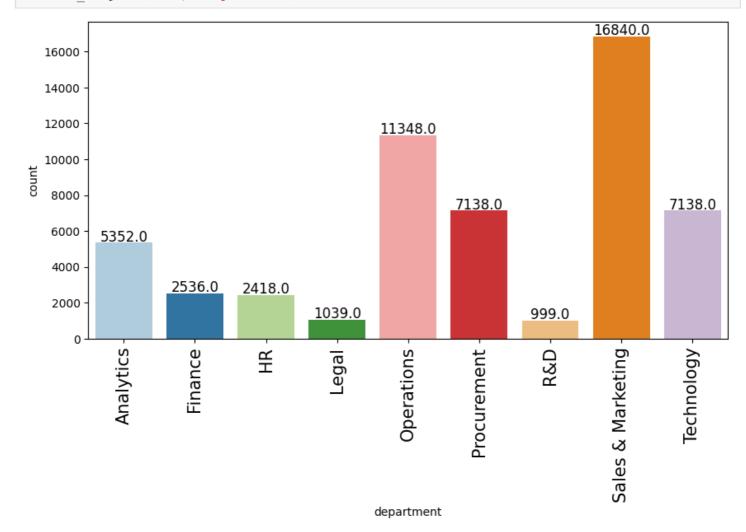
In [27]:



Observations on Department

In [28]:

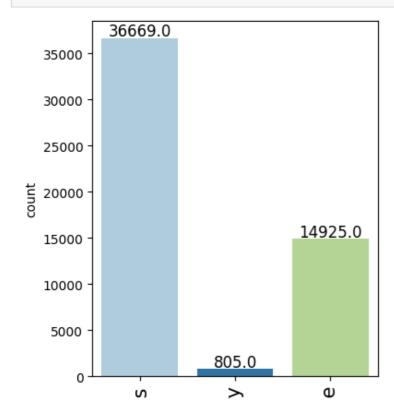
labeled_barplot(data, "department")

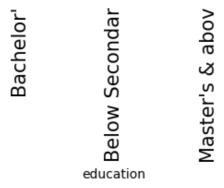


Observations on Education

In [29]:

labeled_barplot(data, "education") ## Complete the code to create labeled_barplot for 'ed
ucation'

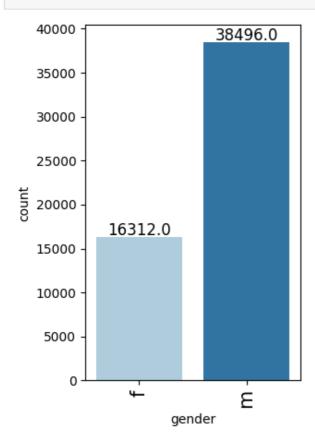




Observations on Gender

In [30]:

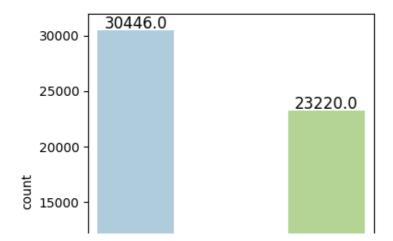
labeled_barplot(data, "gender") ## Complete the code to create labeled_barplot for 'gende
r'

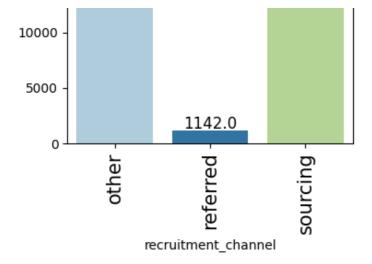


Observations on Recruitment Channel

In [31]:

labeled_barplot(data, "recruitment_channel") ## Complete the code to create labeled_barpl
ot for 'recruitment_channel'

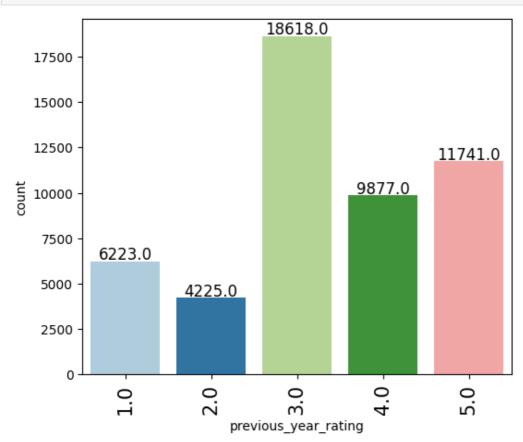




Observations on Previous Year Rating

In [32]:

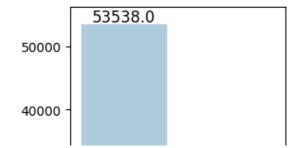
labeled_barplot(data, "previous_year_rating") ## Complete the code to create labeled_barp
lot for 'previous_year_rating'

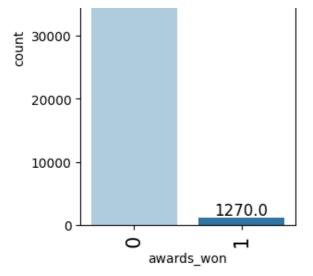


Observations on Awards Won

In [33]:

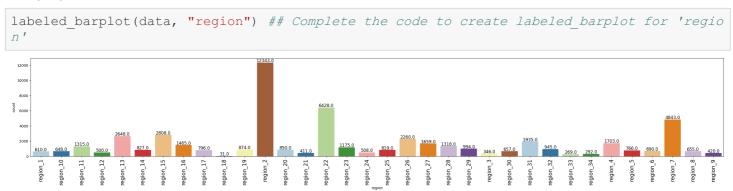
labeled_barplot(data, "awards_won") ## Complete the code to create labeled_barplot for 'a
wards won'





Observations on Region

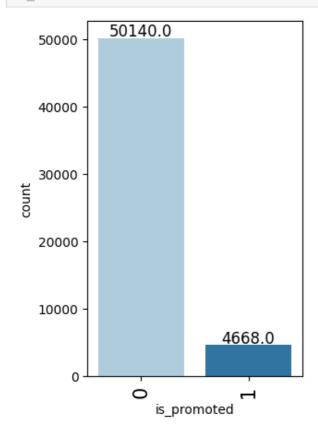
In [34]:



Observations on target variable

In [35]:

labeled_barplot(data, "is_promoted") ## Complete the code to create labeled_barplot for '
is_promoted'



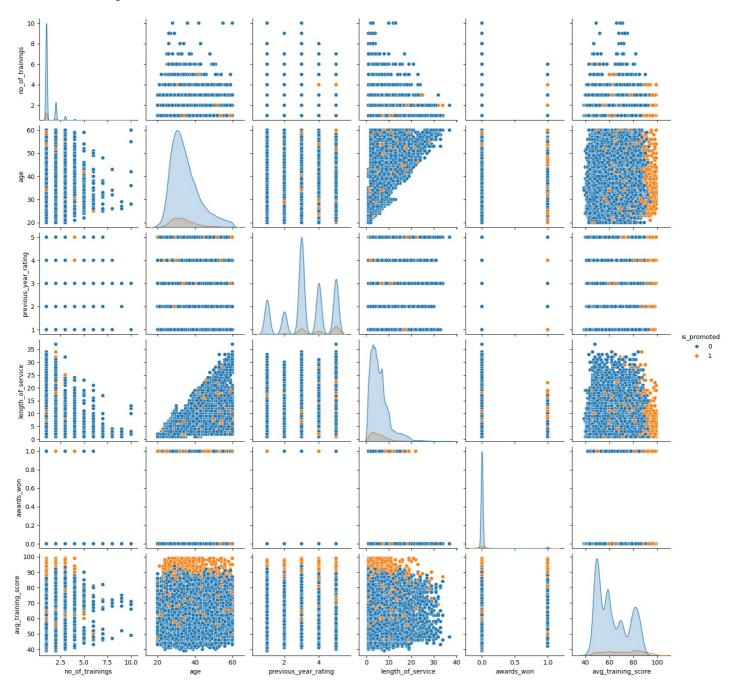
Bivariate Analysis

In [36]:

sns.pairplot(data, hue="is_promoted")

Out[36]:

<seaborn.axisgrid.PairGrid at 0x227014b78e0>

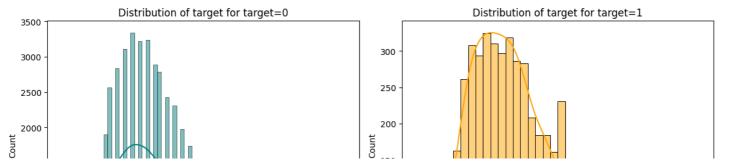


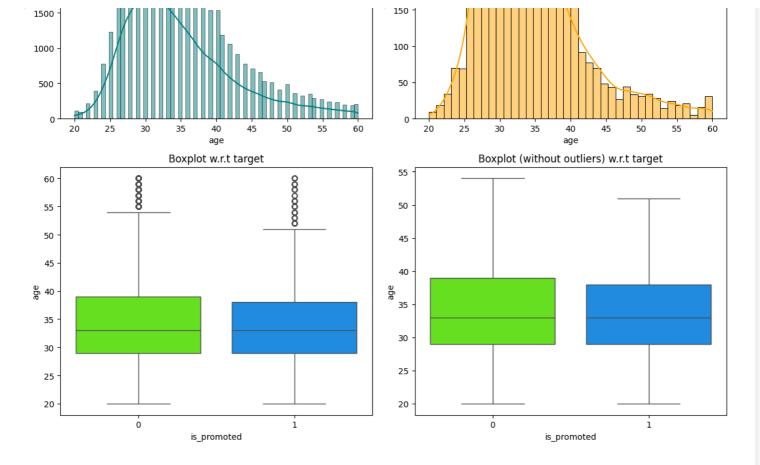
Target variable vs Age

In [37]:

```
distribution_plot_wrt_target(data, "age", "is_promoted")

Distribution of target for target=1
```





Let's see the change in length of service (length_of_service) vary by the employee's promotion status (is_promoted)?

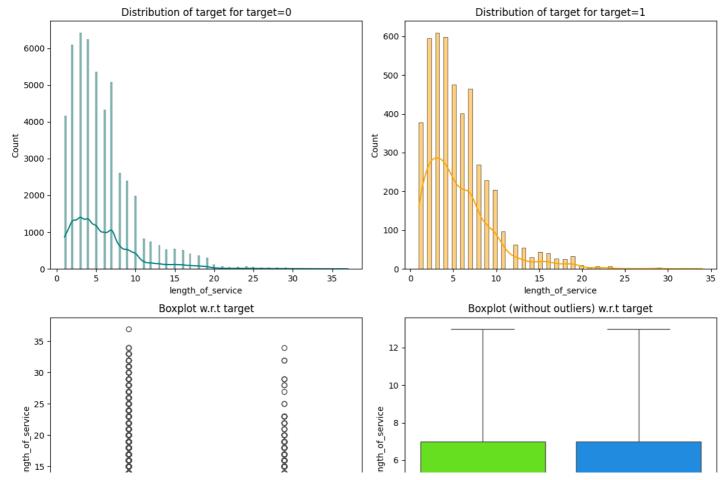
Target variable vs Length of Service

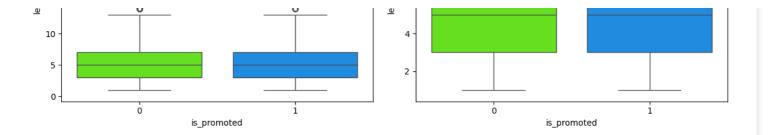
In [38]:

distribution_plot_wrt_target(data, "length_of_service", "is_promoted") ## Complete the co
de to create distribution_plot for length_of_service vs is_promoted

Distribution of target for target=0

Distribution of target for target=1

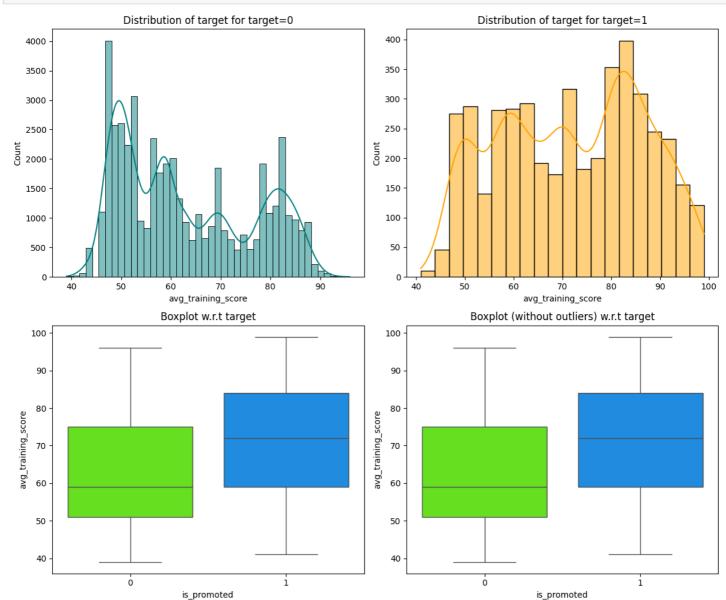




Target variable vs Average Training Score

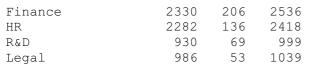
In [39]:

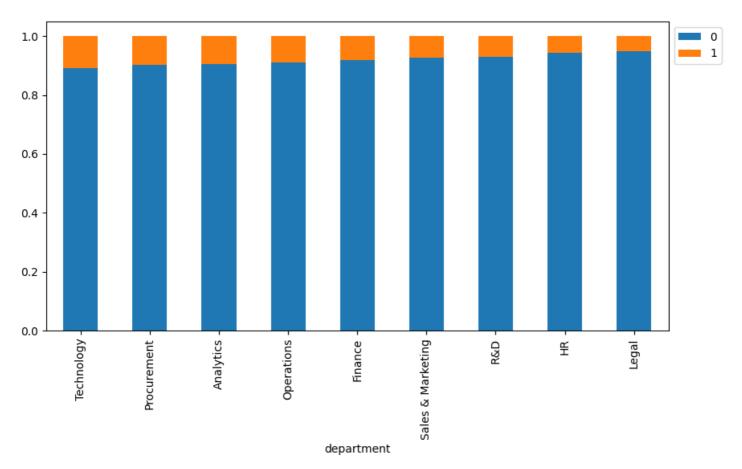
distribution_plot_wrt_target(data, "avg_training_score", "is_promoted") ## Complete the c ode to create distribution_plot for avg_training_score vs is_promoted



Target variable vs Department

In [40]:			
stacked_barplot(da	ata, "de	partme	ent", "i
is_promoted department	0	1	All
All	50140	4668	54808
Sales & Marketing	15627	1213	16840
Operations	10325	1023	11348
Technology	6370	768	7138
Procurement	6450	688	7138
Analytics	4840	512	5352



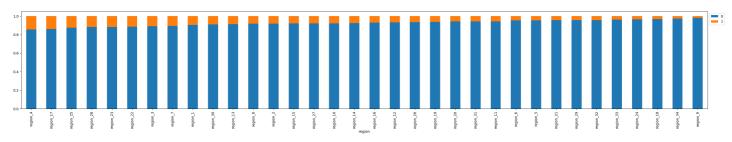


Target variable vs Region

In [41]:

In [41]:					
stacked_barp	olot(dat	a, "re	egion",	"is_promoted")	
is promoted	0	1	All		
region					
All	50140	4668	54808		
region_2	11354	989	12343		
region_22	5694	734	6428		
region_7	4327	516	4843		
region_4	1457	246	1703		
region_13	2418	230	2648		
region_15	2586	222	2808		
region_28	1164	154	1318		
region_26	2117	143	2260		
region_23	1038	137	1175		
region_27	1528	131	1659		
region_31	1825	110	1935		
region_17	687	109	796		
region_25	716	103	819		
region_16	1363	102	1465		
region_11	1241	74	1315		
region_14	765	62	827		
region_30	598	59	657		
region_1	552	58	610		
region_19	821	53	874		
region_8	602	53	655		
region_10	597	51	648		
region_20	801	49	850		
region_29	951	43	994		
region_32	905 309	40 37	945 346		
region_3	309	3 /	346		

region 5	731	35	766
region_12	467	33	500
region_6	658	32	690
region_24	490	18	508
region_21	393	18	411
region_33	259	10	269
region_34	284	8	292
region_9	412	8	420
region_18	30	1	31

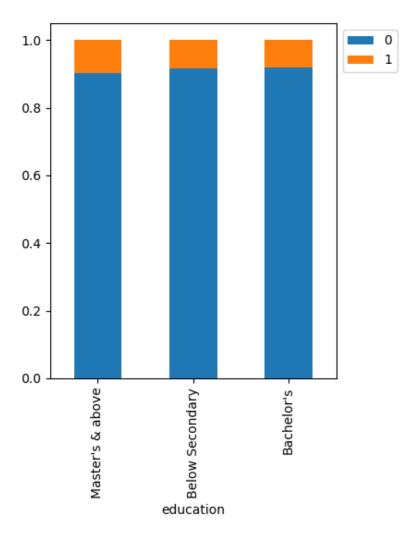


Target variable vs Education

In [42]:

stacked_barplot(data,"education", "is_promoted") ## Complete the code to create distribut
ion_plot for education vs is_promoted

is promoted	0	1	All
education			
All	47853	4546	52399
Bachelor's	33661	3008	36669
Master's & above	13454	1471	14925
Below Secondary	738	67	805

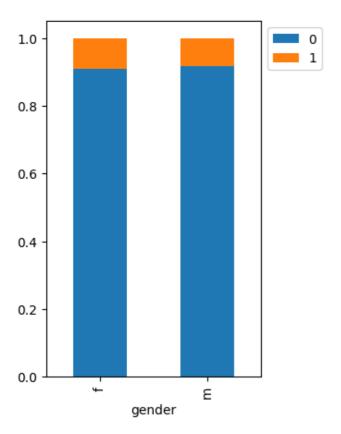


Target variable vs Gender

In [43]:

stacked_barplot(data,"gender", "is_promoted") ## Complete the code to create distribution
_plot for gender vs is_promoted

0	1	All
50140	4668	54808
35295	3201	38496
14845	1467	16312
	35295	35295 3201



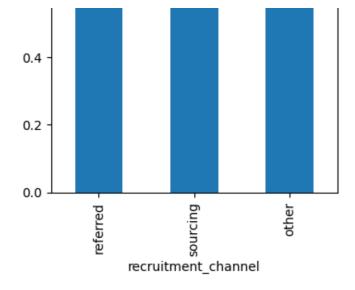
Target variable vs Recruitment Channel

In [44]:

stacked_barplot(data,"recruitment_channel", "is_promoted") ## Complete the code to create
distribution plot for recruitment channel vs is promoted

is_promoted	0	1	All	
recruitment channel				
All	50140	4668	54808	
other	27890	2556	30446	
sourcing	21246	1974	23220	
referred	1004	138	1142	

0.8 -



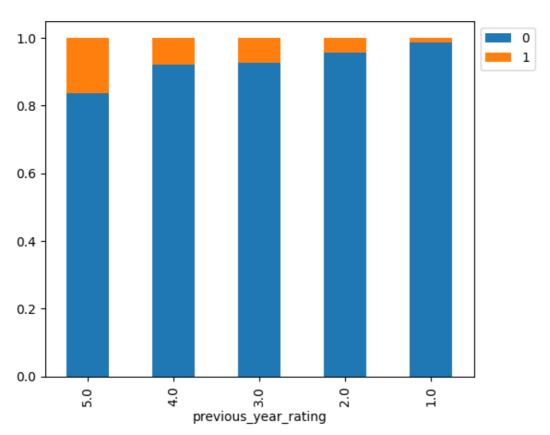
Let's see the previous rating(previous_year_rating) vary by the employee's promotion status (is_promoted)

Target variable vs Previous Year Rating

In [45]:

stacked_barplot(data,"previous_year_rating", "is_promoted") ## Complete the code to creat
e distribution_plot for previous_year_rating vs is_promoted

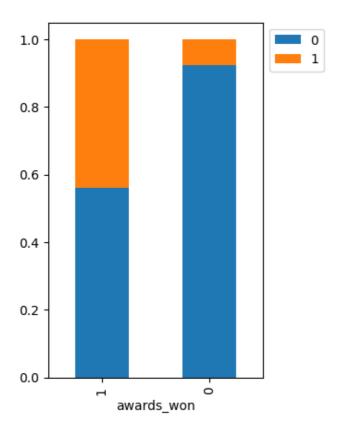
0	1	All
U	Τ	AII
46355	4329	50684
9820	1921	11741
17263	1355	18618
9093	784	9877
4044	181	4225
6135	88	6223
	9820 17263 9093 4044	46355 4329 9820 1921 17263 1355 9093 784 4044 181



In [46]:

stacked_barplot(data,"awards_won", "is_promoted") ## Complete the code to create distribu
tion_plot for awards_won vs is_promoted

is_promoted	0	1	All
awards_won			
All	50140	4668	54808
0	49429	4109	53538
1	711	559	1270

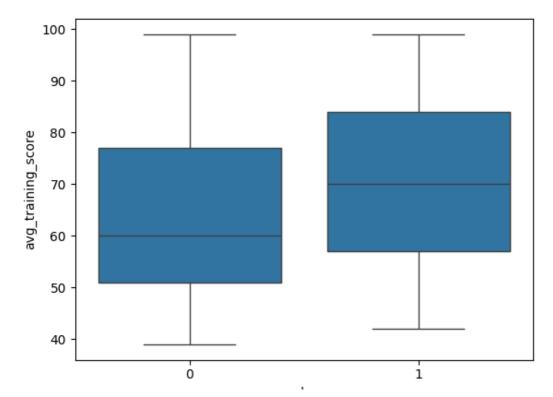


In [47]:

sns.boxplot(data=data, x="awards_won", y="avg_training_score")

Out[47]:

<Axes: xlabel='awards_won', ylabel='avg_training_score'>



Let's see the attributes that have a strong correlation with each other

Correlation Heatmap

```
In [48]:
```

```
# plt.figure(figsize=(15, 7))
# sns.heatmap(data.corr(), annot=True, vmin=-1, vmax=1, fmt=".2f", cmap="Spectral")
# plt.show()

# Create a new DataFrame that only includes numeric columns
numeric_data = data.select_dtypes(include=[np.number])

# Generate the correlation matrix heatmap with only numeric data
plt.figure(figsize=(15, 7))
sns.heatmap(numeric_data.corr(), annot=True, vmin=-1, vmax=1, fmt=".2f", cmap="Spectral")
plt.show()
```



Data Preprocessing

```
In [49]:
```

```
data1 = data.copy()
```

Train-Test Split

```
In [50]:
```

```
X = data1.drop(["is_promoted"], axis=1)
y = data1["is_promoted"]
```

```
In [51]:
```

```
# Splitting data into training and validation set:
```

```
X_train, X_temp, y_train, y_temp = train_test_split(data.drop('is_promoted', axis=1), da
ta['is_promoted'], test_size=0.20, random_state=42) ## Complete the code to split the dat
a into train test in the ratio 80:20

X_test, X_val, y_test, y_val = train_test_split(X_temp, y_temp, test_size=0.25, random_s
tate=42) ## Complete the code to split the data into train test in the ratio 75:25

print(X_train.shape, X_val.shape, X_test.shape)

(43846, 11) (2741, 11) (8221, 11)

Missing value imputation

In [52]:

# Defining the imputers for numerical and categorical variables
imputer_mode = SimpleImputer(strategy="most_frequent")
imputer_median = SimpleImputer(strategy="median")

In [53]:
```

```
# Fit and transform the train data
X_train[["education"]] = imputer_mode.fit_transform(X_train[["education"]])
# Transform the validation data
X_val[["education"]] = imputer_mode.transform(X_val[["education"]]) ## Complete the co
de to impute missing values in X_val

# Transform the test data
X_test[["education"]] = imputer_mode.transform(X_test[["education"]]) ## Complete the co
de to impute missing values in X_test
```

In [54]:

In [55]:

```
# Checking that no column has missing values in train, validation and test sets
print(X_train.isna().sum())
print("-" * 30)
print(X_val.isna().sum())
print("-" * 30)
print(X_test.isna().sum())
```

```
department
                         0
region
                         0
education
gender
recruitment channel
                         0
no_of_trainings
                         0
                         0
age
                         0
previous year rating
                         Ω
length of service
                         0
awards won
avg training score
```

```
dtype: int64
                     Ω
department
                     Ω
region
education
gender
recruitment_channel
no_of_trainings
age
previous_year_rating     0
length_of_service      0
awards_won
                      0
avg_training_score 0
dtype: int64
department
                     0
region
education
gender
recruitment_channel 0
no of_trainings
age
previous_year_rating 0
length_of_service 0
                      0
awards won
avg_training_score 0
dtype: int64
```

Encoding categorical variables

```
In [56]:
```

```
# X_train = pd.get_dummies(X_train, drop_first=True)
# X_val = '____' ## Complete the code to impute missing values in X_val
# X_test = '____' ## Complete the code to impute missing values in X_val
# print(X_train.shape, X_val.shape, X_test.shape)
# print(X_train.shape, X_test.shape)

# Encoding categorical variables for the train, validation, and test data
X_train = pd.get_dummies(X_train, drop_first=True)
X_val = pd.get_dummies(X_val, drop_first=True)
X_test = pd.get_dummies(X_test, drop_first=True)

# Print the shapes of the datasets
print(X_train.shape, X_val.shape, X_test.shape)
```

(43846, 52) (2741, 52) (8221, 52)

Building the model

Model evaluation criterion

Model can make wrong predictions as:

- Predicting an employee should get promoted when he/she should not get promoted
- · Predicting an employee should not get promoted when he/she should get promoted

Which case is more important?

Both cases are important here as not promoting a deserving employee might lead to less productivity and
the company might lose a good employee which affects the company's growth. Further, giving promotion to
a non-deserving employee would lead to loss of monetary resources and giving such employee higher
responsibility might again affect the company's growth.

How to reduce this loss i.e need to reduce False Negatives as well as False Positives?

• Rank would want F1-score to be maximized as both classes are important here. Hence the focus should

be on increasing the F1-score rather than focusing on just one metric i.e. Recall or Precision.

First, let's create two functions to calculate different metrics and confusion matrix, so that we don't have to use the same code repeatedly for each model.

```
In [57]:
```

```
# defining a function to compute different metrics to check performance of a classificati
on model built using sklearn
def model performance classification sklearn (model, predictors, target):
    Function to compute different metrics to check classification model performance
   model: classifier
    predictors: independent variables
    target: dependent variable
    # predicting using the independent variables
    pred = model.predict(predictors)
   acc = accuracy score(target, pred) # to compute Accuracy
    recall = recall score(target, pred) # to compute Recall
    precision = precision score(target, pred) # to compute Precision
    f1 = f1 score(target, pred, average="macro") # to compute F1-score
    # creating a dataframe of metrics
    df perf = pd.DataFrame(
       {
            "Accuracy": acc,
            "Recall": recall,
            "Precision": precision,
            "F1": f1,
        },
        index=[0],
    return df perf
```

In [58]:

```
def confusion_matrix_sklearn(model, predictors, target):
    """
    To plot the confusion_matrix with percentages

    model: classifier
    predictors: independent variables
    target: dependent variable
    """

    y_pred = model.predict(predictors)
    cm = confusion_matrix(target, y_pred)
    labels = np.asarray(
        [
            ["{0:0.0f}".format(item) + "\n{0:.2%}".format(item / cm.flatten().sum())]
            for item in cm.flatten()
        ]
    ).reshape(2, 2)

plt.figure(figsize=(6, 4))
    sns.heatmap(cm, annot=labels, fmt="")
    plt.ylabel("True label")
    plt.xlabel("Predicted label")
```

Model Building - Original Data

```
In [59]:
```

```
# models = [] # Empty list to store all the models
```

```
# # Appending models into the list
# models.append(("Bagging", BaggingClassifier(random state=1)))
# '_____' ## Complete the code to append remaining 4 models in the list models
# results1 = [] # Empty list to store all model's CV scores
# names = [] # Empty list to store name of the models
# # loop through all models to get the mean cross validated score
# print("\n" "Cross-Validation Cost:" "\n")
# for name, model in models:
#
     kfold = StratifiedKFold(
         n_splits=5, shuffle=True, random_state=1
     ) # Setting number of splits equal to 5
     cv result = cross val score(
         estimator=model, X=X train, y=y train, scoring=scorer, cv=kfold
     results1.append(cv result)
     names.append(name)
     print("{}: {}".format(name, cv result.mean()))
# print("\n" "Validation Performance:" "\n")
# for name, model in models:
     model.fit(X_train, y_train)
#
     scores = recall score(y val, model.predict(X val))
#
     print("{}: {}".format(name, scores))
```

In [60]:

```
models = [] # Empty list to store all the models
# Appending models into the list
models.append(("Bagging", BaggingClassifier(random state=1)))
models.append(("RandomForest", RandomForestClassifier(random state=1)))
models.append(("AdaBoost", AdaBoostClassifier(random_state=1)))
models.append(("GradientBoosting", GradientBoostingClassifier(random state=1)))
models.append(("XGBoost", XGBClassifier(random state=1, use label encoder=False, eval me
tric='logloss')))
results1 = [] # Empty list to store all model's CV scores
names = [] # Empty list to store name of the models
# Assuming 'scorer' is previously defined, for example:
scorer = 'accuracy' # You might want to customize this based on your specific evaluation
needs
# loop through all models to get the mean cross validated score
print("\n" "Cross-Validation Cost:" "\n")
for name, model in models:
    kfold = StratifiedKFold(
       n splits=5, shuffle=True, random state=1
      # Setting number of splits equal to 5
    cv result = cross val score(
       estimator=model, X=X train, y=y train, scoring=scorer, cv=kfold
    results1.append(cv result)
    names.append(name)
    print("{}: {}".format(name, cv result.mean()))
print("\n" "Validation Performance:" "\n")
for name, model in models:
   model.fit(X train, y train)
    scores = recall score(y val, model.predict(X val))
   print("{}: {}".format(name, scores))
```

Bagging: 0.9297768871450728 RandomForest: 0.9320120050769705 AdaBoost: 0.925489211047573

GradientBoosting: 0.9367330597199397

XGBoost: 0.9399944632362397

Validation Performance:

Bagging: 0.34051724137931033 RandomForest: 0.28448275862068967 AdaBoost: 0.19827586206896552

GradientBoosting: 0.28448275862068967

XGBoost: 0.34051724137931033

In [61]:

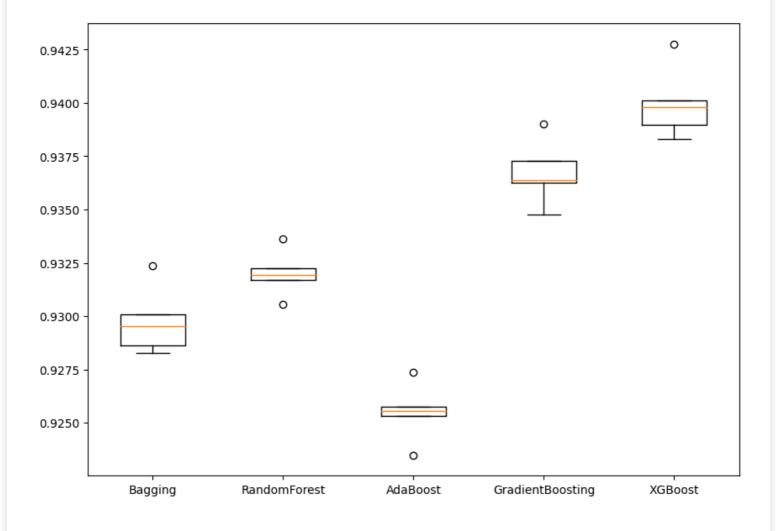
```
# Plotting boxplots for CV scores of all models defined above
fig = plt.figure(figsize=(10, 7))

fig.suptitle("Algorithm Comparison")
ax = fig.add_subplot(111)

plt.boxplot(results1)
ax.set_xticklabels(names)

plt.show()
```

Algorithm Comparison



Model Building - Oversampled Data

In [62]:

```
print("Before Oversampling, counts of label 'Yes': {}".format(sum(y_train == 1)))
print("Before Oversampling, counts of label 'No': {} \n".format(sum(y_train == 0)))
```

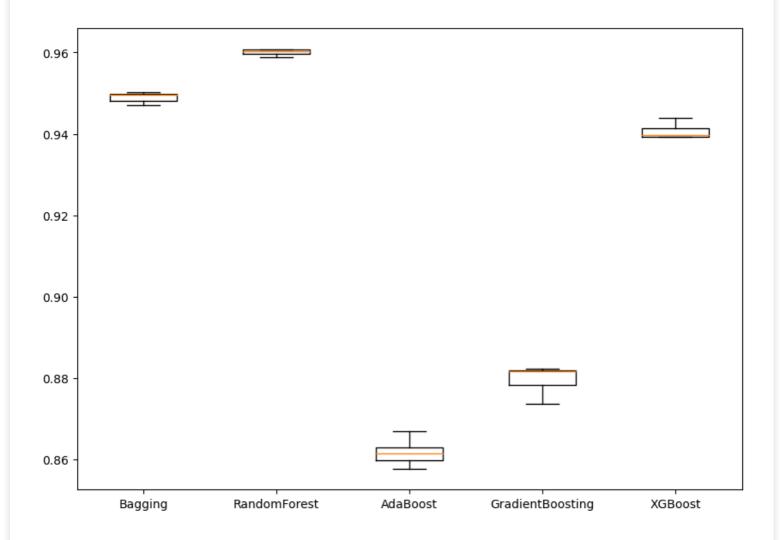
```
sm = SMOTE(
   sampling strategy=1, k neighbors=5, random state=1
) # Synthetic Minority Over Sampling Technique
X train over, y train over = sm.fit resample(X train, y train)
print("After Oversampling, counts of label 'Yes': {}".format(sum(y train over == 1)))
print("After Oversampling, counts of label 'No': {} \n".format(sum(y train over == 0)))
print("After Oversampling, the shape of train X: {}".format(X train over.shape))
print("After Oversampling, the shape of train y: {} \n".format(y train over.shape))
Before Oversampling, counts of label 'Yes': 3760
Before Oversampling, counts of label 'No': 40086
After Oversampling, counts of label 'Yes': 40086
After Oversampling, counts of label 'No': 40086
After Oversampling, the shape of train X: (80172, 52)
After Oversampling, the shape of train y: (80172,)
In [63]:
# ' " ## Complete the code to build models on oversampled data
# ## Note - Take reference from the original models built above
# Assuming 'models' list contains previously initialized models
print("\n" "Validation Performance on Oversampled Data:" "\n")
# Loop to fit each model on the oversampled data and evaluate it
for name, model in models:
   model.fit(X_train_over, y_train_over) # Fit model on oversampled data
    scores = recall_score(y_val, model.predict(X_val)) # Evaluating using the validatio
   print("{}: {}".format(name, scores))
Validation Performance on Oversampled Data:
Bagging: 0.2974137931034483
RandomForest: 0.29310344827586204
AdaBoost: 0.4482758620689655
GradientBoosting: 0.4827586206896552
XGBoost: 0.375
In [64]:
# # Plotting boxplots for CV scores of all models defined above
# '____' ## Write the code to create boxplot to check model performance on oversampled
data
results1_oversampled = [] # Empty list to store all model's CV scores for oversampled da
names oversampled = [] # Empty list to store the names of the models
# Assuming you have a models list set up as follows
models = [
    ("Bagging", BaggingClassifier(random state=1)),
    ("RandomForest", RandomForestClassifier(random state=1)),
    ("AdaBoost", AdaBoostClassifier(random state=1)),
    ("GradientBoosting", GradientBoostingClassifier(random state=1)),
    ("XGBoost", XGBClassifier(random state=1, use label encoder=False, eval metric='logl
oss'))
for name, model in models:
  kfold = StratifiedKFold(n splits=5, shuffle=True, random state=1)
```

```
cv_results = cross_val_score(model, X_train_over, y_train_over, cv=kfold, scoring='a
ccuracy') # Adjust scoring method as needed
    results1_oversampled.append(cv_results)
    names_oversampled.append(name)

# Now plot the boxplot
fig = plt.figure(figsize=(10, 7))
fig.suptitle("Algorithm Comparison on Oversampled Data")
ax = fig.add_subplot(111)

plt.boxplot(results1_oversampled)
ax.set_xticklabels(names_oversampled)
plt.show()
```

Algorithm Comparison on Oversampled Data



Model Building - Undersampled Data

```
In [65]:
```

```
rus = RandomUnderSampler(random_state=1)
X_train_un, y_train_un = rus.fit_resample(X_train, y_train)
```

In [66]:

```
print("Before Under Sampling, counts of label 'Yes': {}".format(sum(y_train == 1)))
print("Before Under Sampling, counts of label 'No': {} \n".format(sum(y_train == 0)))

print("After Under Sampling, counts of label 'Yes': {}".format(sum(y_train_un == 1)))
print("After Under Sampling, counts of label 'No': {} \n".format(sum(y_train_un == 0)))

print("After Under Sampling, the shape of train_X: {}".format(X_train_un.shape))
print("After Under Sampling, the shape of train_y: {} \n".format(y_train_un.shape))
```

Rafora Mindar Camplina counts of labal 'Vac'. 2760

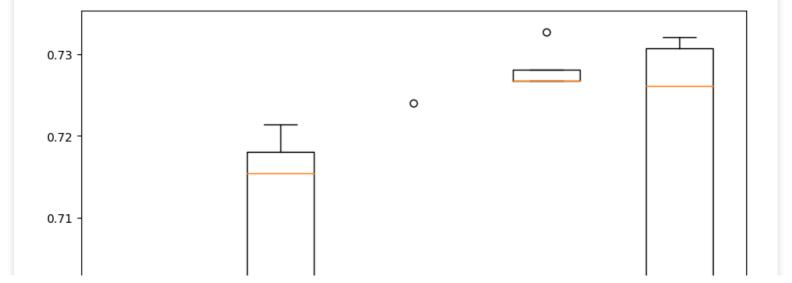
```
After Under Sampling, counts of label 'Yes': 3760
After Under Sampling, counts of label 'No': 3760
After Under Sampling, the shape of train X: (7520, 52)
After Under Sampling, the shape of train y: (7520,)
In [67]:
       ' ## Complete the code to build models on undersampled data
## Note - Take reference from the original models built above
# Loop to fit each model on the undersampled data
for name, model in models:
   model.fit(X train un, y train un) # Fit model on the undersampled training data
    print(f"{name} has been trained on the undersampled data.")
Bagging has been trained on the undersampled data.
RandomForest has been trained on the undersampled data.
AdaBoost has been trained on the undersampled data.
GradientBoosting has been trained on the undersampled data.
XGBoost has been trained on the undersampled data.
In [68]:
# Plotting boxplots for CV scores of all models defined above
# '____' ## Write the code to create boxplot to check model performance on undersample
# Assuming 'models' list exists and 'X_train_un', 'y_train_un' are the undersampled data
results un = [] # List to store cross-validation results for undersampled data
names un = [] # List to store the names of the models
for name, model in models:
   cv results = cross val score(model, X train un, y train un, cv=StratifiedKFold(n spl
its=5), scoring='accuracy')
   results un.append(cv results)
    names un.append(name) # Storing names for x-axis labels in the plot
# Plotting the boxplots
fig = plt.figure(figsize=(10, 7))
fig.suptitle("Algorithm Comparison on Undersampled Data")
ax = fig.add subplot(111)
```

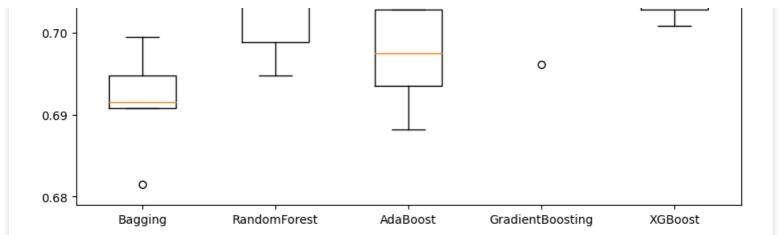
Before Under Sampling, counts of label 'No': 40086

plt.boxplot(results_un)
ax.set xticklabels(names un)

plt.show()

Algorithm Comparison on Undersampled Data





Hyperparameter Tuning

Note

In [70]:

- 1. Sample parameter grid has been provided to do necessary hyperparameter tuning. One can extend/reduce the parameter grid based on execution time and system configuration to try to improve the model performance further wherever needed.
- 2. The models chosen in this notebook are based on test runs. One can update the best models as obtained upon code execution and tune them for best performance.

Tuning AdaBoost using Undersampled data

```
In [69]:
%%time
# defining model
Model = AdaBoostClassifier(random state=1)
# Parameter grid to pass in RandomSearchCV
param grid = {
    "n estimators": np.arange(10, 110, 10),
    "learning rate": [0.1, 0.01, 0.2, 0.05, 1],
    "base estimator": [
        DecisionTreeClassifier(max_depth=1, random_state=1),
        DecisionTreeClassifier(max_depth=2, random_state=1),
        DecisionTreeClassifier(max depth=3, random state=1),
    ],
# Type of scoring used to compare parameter combinations
scorer = 'f1 macro'
#Calling RandomizedSearchCV
randomized cv = RandomizedSearchCV(estimator=Model, param distributions=param grid, n job
s = -1, n iter=50, scoring=scorer, cv=5, random state=1)
#Fitting parameters in RandomizedSearchCV
randomized cv.fit(X train un, y train un)
                                          # This is the line that was needed to complete
the code
 ## Complete the code to fit the model on undersampled data
print("Best parameters are {} with CV score={}:" .format(randomized_cv.best_params_,rand
omized cv.best score ))
Best parameters are {'n estimators': 50, 'learning rate': 1, 'base estimator': DecisionTr
eeClassifier(max depth=2, random state=1)} with CV score=0.7194193297216376:
CPU times: total: 2.86 s
Wall time: 1min 3s
```

```
# # Creating new pipeline with best parameters
# tuned adb1 = AdaBoostClassifier( random state=
# ) ## Complete the code with the best parameters obtained from tuning
# tuned adb1.' ' ## Complete the code to fit the model on undersampled data
# Creating new pipeline with best parameters
tuned adb1 = AdaBoostClassifier(
   random state=1,
   n_estimators=50, # Replace with the best n_estimators found learning_rate=0.1, # Replace with the best learning_rate found
   base_estimator=DecisionTreeClassifier(max_depth=2, random_state=1) # Replace with t
he best max depth found
# Fitting the model on undersampled data
tuned adb1.fit(X train un, y train un)
Out[70]:
          AdaBoostClassifier
 ▶ base estimator: DecisionTreeClassifier
 ▶ DecisionTreeClassifier
In [71]:
# adb1 train
from sklearn.metrics import accuracy_score, classification_report
# Predict on the training set
adb1 train predictions = tuned adb1.predict(X train un)
# Evaluate the predictions using accuracy as an example metric
adb1 train accuracy = accuracy score(y train un, adb1 train predictions)
# You can also use classification report to get a detailed performance report
adb1 train report = classification_report(y_train_un, adb1_train_predictions)
# Store the accuracy in adb1 train and print it
adb1 train = adb1 train accuracy
# Print the accuracy and the detailed report
print("Accuracy on training set:", adb1 train)
print("Detailed classification report:\n", adb1 train report)
Accuracy on training set: 0.7122340425531914
Detailed classification report:
            precision recall f1-score support
         0
                0.67
                        0.85
                                 0.75
                                           3760
                0.79
                        0.57
                                  0.67
                                           3760
   accuracy
                                  0.71
                                           7520
                       0.71
  macro avg
               0.73
                                 0.71
                                           7520
```

In [72]:

weighted avg

0.73

0.71

```
# # Checking model's performance on validation set
# adb1_val = '____' ## Complete the code to check the performance on validation set
# adb1_val
```

7520

0.71

```
from sklearn.metrics import accuracy_score, classification_report

# Predict on the validation set
adb1_val_predictions = tuned_adb1.predict(X_val)

# Evaluate the predictions using accuracy as an example metric
adb1_val_accuracy = accuracy_score(y_val, adb1_val_predictions)

# You can also use classification_report to get a detailed performance report
adb1_val_report = classification_report(y_val, adb1_val_predictions)

# Store the accuracy in adb1_val and print it
adb1_val = adb1_val_accuracy

# Print the accuracy and the detailed report
print("Accuracy on validation set:", adb1_val)
print("Detailed classification report:\n", adb1_val_report)
```

Accuracy on validation set: 0.8314483765049252 Detailed classification report:

	precision	recall	f1-score	support
0	0.96	0.85	0.90	2509
1	0.28	0.62	0.38	232
accuracy			0.83	2741
macro avg weighted avg	0.62 0.90	0.73 0.83	0.64 0.86	2741 2741

Tuning AdaBoost using original data

In [73]:

```
%%time
# defining model
Model = AdaBoostClassifier(random state=1)
# Parameter grid to pass in RandomSearchCV
param grid = {
    "n estimators": np.arange(10, 110, 10),
    "learning rate": [0.1, 0.01, 0.2, 0.05, 1],
    "base estimator": [
        DecisionTreeClassifier(max depth=1, random state=1),
        DecisionTreeClassifier(max depth=2, random state=1),
        DecisionTreeClassifier(max depth=3, random state=1),
    ],
# Type of scoring used to compare parameter combinations
scorer = 'f1 macro'
#Calling RandomizedSearchCV
randomized cv = RandomizedSearchCV(estimator=Model, param distributions=param grid, n job
s = -1, n iter=50, scoring=scorer, cv=5, random state=1)
#Fitting parameters in RandomizedSearchCV
randomized_cv.fit(X_train, y_train) #'_____' ## Complete the code to fit the model on or
iginal data
print("Best parameters are {} with CV score={}:" .format(randomized cv.best params ,rand
omized cv.best score ))
Best parameters are {'n estimators': 90, 'learning rate': 1, 'base estimator': DecisionTr
```

eeClassifier(max depth=2, random state=1)} with CV score=0.7223671392095827:

CPU times: total: 13.2 s Wall time: 5min 17s

```
# # Creating new pipeline with best parameters
# tuned adb2 = AdaBoostClassifier( random_state=___,
# n_estimators= _____, learning_rate= _____, base_estimator= DecisionTreeClassifi
er(max_depth= ____, random_state=1)
# ) ## Complete the code with the best parameters obtained from tuning
# tuned adb2.' ' ## Complete the code to fit the model on original data
# Creating new pipeline with best parameters
tuned adb2 = AdaBoostClassifier(
    random state=1,  # Use the random state from the tuning process
    n_estimators=randomized_cv.best_params_['n_estimators'], # Use the best number of es
timators
   learning rate=randomized cv.best params ['learning rate'], # Use the best learning r
   base estimator=DecisionTreeClassifier(
       max depth=randomized cv.best params ['base estimator'].max depth,
        random state=1
# Fitting the model on the original data
tuned adb2.fit(X train, y train)
Out[74]:
           AdaBoostClassifier
 ▶ base estimator: DecisionTreeClassifier
 ▶ DecisionTreeClassifier
      In [75]:
# adb2 train
from sklearn.metrics import accuracy score, classification report
# Predict on the training set
adb2 train predictions = tuned adb2.predict(X train)
# Evaluate the predictions using accuracy as an example metric
adb2_train_accuracy = accuracy_score(y_train, adb2_train_predictions)
# You can also use classification_report to get a detailed performance report
adb2 train report = classification report(y train, adb2 train predictions)
# Store the accuracy in adb2 train and print it
adb2 train = adb2 train accuracy
# Print the accuracy and the detailed report
print("Accuracy on training set:", adb2 train)
print("Detailed classification report:\n", adb2_train_report)
Accuracy on training set: 0.939652419833052
Detailed classification report:
              precision recall f1-score support

      0.94
      1.00
      0.97

      0.91
      0.33
      0.48

           0
                                               40086
                                                 3760
                                              43846
43846
                                      0.94
   accuracy

      0.92
      0.66
      0.73
      43846

      0.94
      0.94
      0.93
      43846

   macro avg
weighted avg
```

In [76]:

• ولا بالمالية

```
# # Checking model's performance on validation set
\# adb2_val = '_____' \#\# Complete the code to check the performance on validation set
# adb2 val
from sklearn.metrics import accuracy score, classification report
# Predict on the validation set
adb2 val predictions = tuned adb2.predict(X val)
# Evaluate the predictions using accuracy as an example metric
adb2 val accuracy = accuracy score(y val, adb2 val predictions)
# You can also use classification_report to get a detailed performance report
adb2 val report = classification report(y val, adb2 val predictions)
# Store the accuracy in adb2 val and print it
adb2 val = adb2 val accuracy
# Print the accuracy and the detailed report
print("Accuracy on validation set:", adb2 val)
print("Detailed classification report:\n", adb2 val report)
Accuracy on validation set: 0.9398029916089019
Detailed classification report:
```

	precision	recall	f1-score	support
0	0.94	1.00	0.97	2509
1	0.89	0.33	0.48	232
accuracy			0.94	2741
macro avg	0.92	0.66	0.72	2741
weighted avg	0.94	0.94	0.93	2741

Tuning Gradient Boosting using undersampled data

In [77]:

```
%%time
#Creating pipeline
Model = GradientBoostingClassifier(random state=1)
#Parameter grid to pass in RandomSearchCV
param grid = {
   "init": [AdaBoostClassifier(random state=1), DecisionTreeClassifier(random state=1)],
   "n estimators": np.arange(75,150,25),
   "learning rate": [0.1, 0.01, 0.2, 0.05, 1],
   "subsample": [0.5, 0.7, 1],
   "max features": [0.5, 0.7, 1],
# Type of scoring used to compare parameter combinations
scorer = 'f1 macro'
#Calling RandomizedSearchCV
randomized cv = RandomizedSearchCV(estimator=Model, param distributions=param grid, n ite
r=50, scoring=scorer, cv=5, random state=1, n jobs = -1)
#Fitting parameters in RandomizedSearchCV
randomized_cv.fit(X_train_un, y_train_un) ## Complete the code to fit the model on under
sampled data
print("Best parameters are {} with CV score={}:" .format(randomized_cv.best_params_, rand
omized cv.best score ))
```

Best parameters are {'subsample': 0.7, 'n_estimators': 125, 'max_features': 0.7, 'learnin g_rate': 0.2, 'init': AdaBoostClassifier(random_state=1)} with CV score=0.728005306035405 7:
CPU times: total: 2.78 s

Wall time: 47.6 s

```
In [78]:
```

```
# Creating new pipeline with best parameters
# tuned gbm1 = GradientBoostingClassifier(
    max features=
    init=AdaBoostClassifier(random state=1),
#
#
     random state=1,
     learning_rate=__
     n_estimators=____,
     subsample=
# ) ## Complete the code with the best parameters obtained from tuning
# tuned gbm1.fit(X train un, y train un)
# Replace the placeholders with the best parameters obtained from the RandomizedSearchCV
best params = randomized cv.best params
tuned gbm1 = GradientBoostingClassifier(
   max features=best params['max features'],
    init=AdaBoostClassifier(random state=1), # Assuming the best `init` was AdaBoostClas
sifier
   random state=1,
   learning_rate=best_params['learning_rate'],
   n_estimators=best_params['n_estimators'],
   subsample=best_params['subsample']
# Fitting the model on undersampled data
tuned gbm1.fit(X train un, y train un)
# Verify the model is fitted
print("Model is trained with the best parameters from hyperparameter tuning on the unders
ampled data.")
```

Model is trained with the best parameters from hyperparameter tuning on the undersampled data.

In [79]:

```
# gbml_train = '____' ## Complete the code to check the performance on oversampled tra
in set
# gbml_train

from sklearn.metrics import accuracy_score, classification_report

# Predict on the oversampled training set
gbml_train_predictions = tuned_gbml.predict(X_train_over)

# Evaluate the predictions using accuracy as an example metric
gbml_train_accuracy = accuracy_score(y_train_over, gbml_train_predictions)

# You can also use classification_report to get a detailed performance report
gbml_train_report = classification_report(y_train_over, gbml_train_predictions)

# Store the accuracy in gbml_train and print it
gbml_train = gbml_train_accuracy

# Print the accuracy and the detailed report
print("Accuracy on oversampled training set:", gbml_train)
print("Detailed classification report:\n", gbml_train_report)
```

Accuracy on oversampled training set: 0.7931447388115551 Detailed classification report:

	precision	recall	f1-score	support
0	0.79 0.79	0.79 0.79	0.79 0.79	40086 40086
accuracy macro avg	0.79	0.79	0.79	80172 80172

weighted avg U./9 U./9 U./9 8U1/2

```
In [80]:
```

```
# gbml_val = '____' ## Complete the code to check the performance on validation set
# gbml_val

from sklearn.metrics import accuracy_score, classification_report

# Predict on the validation set
gbml_val_predictions = tuned_gbml.predict(X_val)

# Evaluate the predictions using accuracy as an example metric
gbml_val_accuracy = accuracy_score(y_val, gbml_val_predictions)

# You can also use classification_report to get a detailed performance report
gbml_val_report = classification_report(y_val, gbml_val_predictions)

# Store the accuracy in gbml_val and print it
gbml_val = gbml_val_accuracy

# Print the accuracy and the detailed report
print("Accuracy on validation set:", gbml_val)
print("Detailed classification report:\n", gbml_val_report)
```

Accuracy on validation set: 0.7697920466982853

Detailed classification report:

	precision	recall	f1-score	support
0 1	0.97 0.23	0.77 0.72	0.86 0.35	2509 232
accuracy macro avg weighted avg	0.60 0.91	0.75 0.77	0.77 0.60 0.82	2741 2741 2741

Tuning Gradient Boosting using original data

In [81]:

```
%%time
#defining model
Model = GradientBoostingClassifier(random state=1)
#Parameter grid to pass in RandomSearchCV
param grid = {
    "Init": [AdaBoostClassifier(random state=1), DecisionTreeClassifier(random state=1)],
    "n estimators": np.arange(75,150,25),
    "learning rate": [0.1, 0.01, 0.2, 0.05, 1],
    "subsample": [0.5, 0.7, 1],
    "max features": [0.5,0.7,1],
# Type of scoring used to compare parameter combinations
scorer = 'f1_macro'
#Calling RandomizedSearchCV
randomized cv = RandomizedSearchCV(estimator=Model, param distributions=param grid, n ite
r=50, scoring=scorer, cv=5, random state=1, n jobs = -1)
#Fitting parameters in RandomizedSearchCV
randomized_cv.fit(X_train, y_train) ## Complete the code to fit the model on original dat
print("Best parameters are {} with CV score={}:" .format(randomized_cv.best_params_, rand
omized cv.best score ))
```

```
Best parameters are {'subsample': 0.7, 'n estimators': 125, 'max features': 0.7, 'learnin
g rate': 0.2, 'init': AdaBoostClassifier(random state=1)} with CV score=0.721238400832193
7:
CPU times: total: 11.9 s
Wall time: 4min 34s
In [82]:
# # Creating new pipeline with best parameters
# tuned gbm2 = GradientBoostingClassifier(
     max features=
     init=AdaBoostClassifier(random state=1),
#
#
     random state=1,
     learning_rate=____
#
     n_estimators=____,
subsample=___,
#
     subsample=
# ) ## Complete the code with the best parameters obtained from tuning
# tuned gbm2.fit(X train, y train)
# Assuming that randomized cv is the RandomizedSearchCV object and its best params attri
bute contains the optimal values.
best_params = randomized_cv.best_params_
tuned gbm2 = GradientBoostingClassifier(
   max features=best params['max features'],
   init=AdaBoostClassifier(random_state=1), # Assuming the best `init` was AdaBoostClas
sifier; replace if necessary.
   random state=1,
   learning rate=best params['learning rate'],
    n estimators=best params['n estimators'],
    subsample=best_params['subsample']
# Fitting the model on original data
tuned gbm2.fit(X train, y train)
# Verify the model is fitted
print("Model is trained with the best parameters from hyperparameter tuning on the origin
al data.")
Model is trained with the best parameters from hyperparameter tuning on the original data
In [83]:
# gbm2 train
from sklearn.metrics import accuracy score, classification report
# Predict on the original training set
gbm2 train predictions = tuned gbm2.predict(X train)
# Evaluate the predictions using accuracy as an example metric
gbm2_train_accuracy = accuracy_score(y_train, gbm2_train_predictions)
# You can also use classification report to get a detailed performance report
gbm2_train_report = classification_report(y_train, gbm2_train_predictions)
# Store the accuracy in gbm2 train and print it
gbm2 train = gbm2 train accuracy
# Print the accuracy and the detailed report
print("Accuracy on training set:", gbm2 train)
print("Detailed classification report:\n", gbm2 train report)
```

Detailed classification report:

precision recall f1-score support

Accuracy on training set: 0.9409980385896091

```
0.94 1.00
                               0.97
         0
                                        40086
               0.96
                       0.33
                                0.49
                                         3760
                                0.94
   accuracy
                                        43846
                             0.73
           macro avg
                                        43846
weighted avg
                                        43846
In [84]:
# gbm2 val\
from sklearn.metrics import accuracy score, classification report
# Predict on the validation set
gbm2 val predictions = tuned gbm2.predict(X val)
# Evaluate the predictions using accuracy as an example metric
gbm2 val accuracy = accuracy score(y val, gbm2 val predictions)
# You can also use classification_report to get a detailed performance report
gbm2 val report = classification report(y val, gbm2 val predictions)
# Store the accuracy in gbm2 val and print it
gbm2 val = gbm2 val accuracy
# Print the accuracy and the detailed report
print("Accuracy on validation set:", gbm2 val)
print("Detailed classification report:\n", gbm2 val report)
```

Accuracy on validation set: 0.9398029916089019

Detailed classification report:

	precision	recall	f1-score	support
0	0.94	1.00	0.97	2509
1	0.89	0.33	0.48	232
accuracy			0.94	2741
macro avg weighted avg	0.91 0.94	0.66 0.94	0.73 0.93	2741 2741

Model Comparison and Final Model Selection

In [85]:

```
# training performance comparison
# models_train_comp_df = pd.concat(
#
#
        gbm1 train.T,
         gbm2_train.T,
         adb1 train.T,
         adb2 train.T,
     ],
#
     axis=1,
# )
# models train comp df.columns = [
      "Gradient boosting trained with Undersampled data",
#
     "Gradient boosting trained with Original data",
     "AdaBoost trained with Undersampled data",
#
#
      "AdaBoost trained with Original data",
# ]
# print("Training performance comparison:")
# models train comp df
import pandas as pd
```

```
# Assuming gbm1 train, gbm2 train, adb1 train, and adb2 train are scalar values like accu
models train comp df = pd.DataFrame(
        "Gradient boosting trained with Undersampled data": [gbml train],
        "Gradient boosting trained with Original data": [gbm2 train],
        "AdaBoost trained with Undersampled data": [adb1 train],
        "AdaBoost trained with Original data": [adb2 train],
    index=["Training Performance"]
print("Training performance comparison:")
print(models train comp df)
Training performance comparison:
                      Gradient boosting trained with Undersampled data \
Training Performance
                      Gradient boosting trained with Original data \
Training Performance
                      AdaBoost trained with Undersampled data \
Training Performance
                      AdaBoost trained with Original data
Training Performance
                                                     0.940
In [86]:
# validation performance comparison
#' ' ## Write the code to compare the performance on validation set
# Assuming gbm1 val, gbm2 val, adb1 val, and adb2 val are also scalar values
models val comp df = pd.DataFrame(
    {
        "Gradient boosting trained with Undersampled data": [gbml val],
        "Gradient boosting trained with Original data": [gbm2 val],
        "AdaBoost trained with Undersampled data": [adb1 val],
        "AdaBoost trained with Original data": [adb2 val],
    index=["Validation Performance"]
print("Validation performance comparison:")
print(models val comp df)
Validation performance comparison:
                        Gradient boosting trained with Undersampled data \
Validation Performance
                                                                    0.770
                        Gradient boosting trained with Original data \
Validation Performance
                        AdaBoost trained with Undersampled data \
Validation Performance
                                                           0.831
                        AdaBoost trained with Original data
Validation Performance
                                                       0.940
Now we have our final model, so let's find out how our final model is performing on unseen test data.
```

#' ' ## Write the code to check the performance of best model on test data

In [87]:

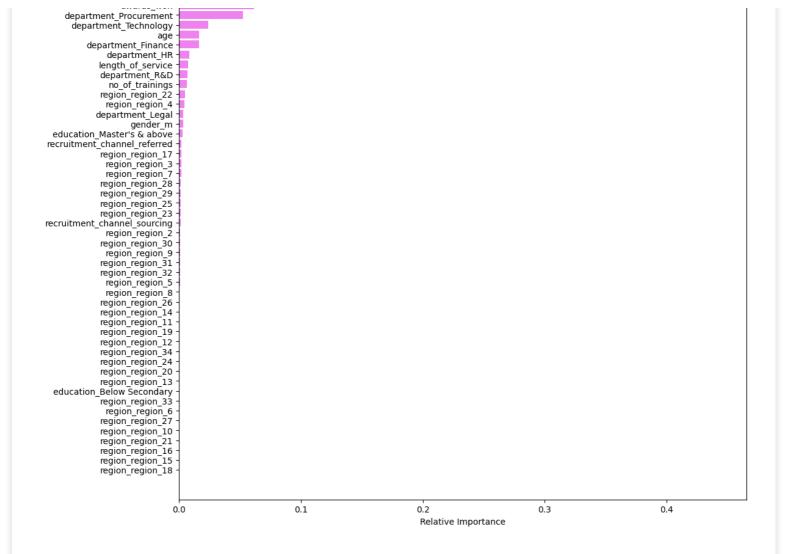
Let's check the performance on test set

```
from sklearn.metrics import accuracy_score, classification_report
# Assuming 'tuned gbm2' is the Gradient Boosting model trained with the original data
# Predict on the test set
test predictions = tuned gbm2.predict(X test)
# Evaluate the predictions using accuracy
test accuracy = accuracy score(y test, test predictions)
# Detailed performance report
test report = classification report(y test, test predictions)
print("Accuracy on test set:", test accuracy)
print("Detailed classification report:\n", test report)
Accuracy on test set: 0.9400316263228318
Detailed classification report:
              precision recall f1-score
                                              support
           0
                   0.94
                           1.00
                                       0.97
                                                 7545
                  0.91
                            0.30
                                       0.45
                                                 676
                                      0.94
                                                 8221
   accuracy
                 0.93
                                      0.71
                            0.65
                                                8221
  macro avg
                  0.94
                             0.94
                                      0.93
                                                8221
weighted avg
```

Feature Importances

```
In [89]:
```

```
# feature names = X train.columns
# importances = '____' ## Complete the code to check the feature importance of the be
st model
# indices = np.argsort(importances)
# plt.figure(figsize=(12, 12))
# plt.title("Feature Importances")
# plt.barh(range(len(indices)), importances[indices], color="violet", align="center")
# plt.yticks(range(len(indices)), [feature names[i] for i in indices])
# plt.xlabel("Relative Importance")
# plt.show()
import numpy as np
import matplotlib.pyplot as plt
# Feature names from the training set
feature names = X train.columns
# Assuming 'tuned gbm2' is the trained Gradient Boosting model
importances = tuned_gbm2.feature_importances_ # This retrieves the feature importances f
rom the model
# Sort the feature importances in ascending order
indices = np.argsort(importances)
# Create a horizontal bar chart to display feature importance
plt.figure(figsize=(12, 12))
plt.title("Feature Importances")
plt.barh(range(len(indices)), importances[indices], color="violet", align="center")
plt.yticks(range(len(indices)), [feature names[i] for i in indices])
plt.xlabel("Relative Importance")
plt.show()
```



Business Insights and Conclusions

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