

# dac-phase-3

October 26, 2023

Date:26/10/2023

Team ID:716

Project Name:Public Health Awareness Campaign Analysis using Data Analytics

## IMPORT DEPENDENCIES

```
[6]: # suppress display of warnings
import warnings
warnings.filterwarnings("ignore")

# 'Pandas' is used for data manipulation and analysis
import pandas as pd

# 'Numpy' is used for mathematical operations on large, multi-dimensional
    ↳ arrays and matrices
import numpy as np

# 'Matplotlib' is a data visualization library for 2D and 3D plots, built on
    ↳ numpy
import matplotlib.pyplot as plt
from matplotlib.colors import ListedColormap

# 'Seaborn' is based on matplotlib; used for plotting statistical graphics
import seaborn as sns

# import 'is_string_dtype' to check if the type of input is string
from pandas.api.types import is_string_dtype

# import various functions to perform classification
from sklearn.naive_bayes import GaussianNB
from sklearn.multiclass import OneVsRestClassifier
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn import metrics
from sklearn.linear_model import LogisticRegression
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import classification_report
```

```

from sklearn.metrics import cohen_kappa_score
from sklearn.metrics import confusion_matrix
from sklearn.metrics import roc_auc_score
from sklearn.metrics import roc_curve
from sklearn import tree
from sklearn.tree import export_graphviz
from sklearn.linear_model import SGDClassifier
from sklearn.model_selection import GridSearchCV
from sklearn.feature_selection import SelectFromModel
from sklearn.svm import SVC
# display all columns of the dataframe
pd.options.display.max_columns = None

```

```

[31]: #setting the plot size using rcParams
plt.rcParams['figure.figsize'] = [10,8]

```

```

[13]: #importing datasets for training and testing the models
df=pd.read_csv("C:/Users/sjana/Downloads/public health survey.csv")

#rwading the first 5 records from the training data set
df.head()

```

```

[13]:

```

	Timestamp	Age	Gender	Country	state	self_employed	\
0	2014-08-27 11:29:31	37	Female	United States	IL	NaN	
1	2014-08-27 11:29:37	44	M	United States	IN	NaN	
2	2014-08-27 11:29:44	32	Male	Canada	NaN	NaN	
3	2014-08-27 11:29:46	31	Male	United Kingdom	NaN	NaN	
4	2014-08-27 11:30:22	31	Male	United States	TX	NaN	

	family_history	treatment	work_interfere	no_employees	remote_work	\
0	No	Yes	Often	6-25	No	
1	No	No	Rarely	More than 1000	No	
2	No	No	Rarely	6-25	No	
3	Yes	Yes	Often	26-100	No	
4	No	No	Never	100-500	Yes	

	tech_company	benefits	care_options	wellness_program	seek_help	\
0	Yes	Yes	Not sure	No	Yes	
1	No	Don't know	No	Don't know	Don't know	
2	Yes	No	No	No	No	
3	Yes	No	Yes	No	No	
4	Yes	Yes	No	Don't know	Don't know	

	anonymity	leave mental_health_consequence	\
0	Yes	Somewhat easy	No
1	Don't know	Don't know	Maybe
2	Don't know	Somewhat difficult	No

3	No	Somewhat difficult	Yes
4	Don't know	Don't know	No

	phys_health_consequence	coworkers	supervisor	mental_health_interview	\
0	No	Some of them	Yes	No	
1	No	No	No	No	
2	No	Yes	Yes	Yes	
3	Yes	Some of them	No	Maybe	
4	No	Some of them	Yes	Yes	

	phys_health_interview	mental_vs_physical	obs_consequence	comments
0	Maybe	Yes	No	NaN
1	No	Don't know	No	NaN
2	Yes	No	No	NaN
3	Maybe	No	Yes	NaN
4	Yes	Don't know	No	NaN

## UNDERSTANDING DATA

```
[14]: #checking the number of rows and columns in the training data set
df.shape
```

```
[14]: (1259, 27)
```

```
[15]: # 'dtypes' gives the data type for each column
df.dtypes
```

```
[15]: Timestamp          object
Age                    int64
Gender                object
Country               object
state                 object
self_employed         object
family_history         object
treatment             object
work_interfere        object
no_employees          object
remote_work           object
tech_company          object
benefits              object
care_options          object
wellness_program      object
seek_help             object
anonymity             object
leave                object
mental_health_consequence object
phys_health_consequence object
```

```
coworkers          object
supervisor         object
mental_health_interview  object
phys_health_interview  object
mental_vs_physical  object
obs_consequence    object
comments           object
dtype: object
```

```
[16]: #Splitting the timestamp feature as it includes both date and time
df[['Date', 'Time']] = df['Timestamp'].str.split(" ", n=1, expand=True)
```

```
[17]: #dropping the timestamp column as we have already created two columns which
      ↪ have date and time
      #axis=1 deletes the entire column
df.drop('Timestamp', axis=1, inplace=True)
```

```
[18]: #converting the datatype of date and time columns
df['Time'] = pd.to_datetime(df['Time'], format='%H:%M:%S')
df['Date'] = pd.to_datetime(df['Date'])
```

```
[19]: #finally checking the columns and their datatypes after alteration
df.dtypes
```

```
[19]: Age          int64
Gender          object
Country         object
state           object
self_employed   object
family_history   object
treatment       object
work_interfere   object
no_employees     object
remote_work      object
tech_company     object
benefits         object
care_options     object
wellness_program object
seek_help        object
anonymity        object
leave           object
mental_health_consequence object
phys_health_consequence object
coworkers        object
supervisor       object
mental_health_interview object
phys_health_interview object
```

```

mental_vs_physical      object
obs_consequence         object
comments               object
Date                   datetime64[ns]
Time                   datetime64[ns]
dtype: object

```

```
[20]: # the describe() returns the statistical summary of the numeric variables
df.describe()
```

```
[20]:
```

	Age	Date \
count	1.259000e+03	1259
mean	7.942815e+07	2014-09-09 10:54:14.011120128
min	-1.726000e+03	2014-08-27 00:00:00
25%	2.700000e+01	2014-08-27 00:00:00
50%	3.100000e+01	2014-08-28 00:00:00
75%	3.600000e+01	2014-08-28 00:00:00
max	1.000000e+11	2016-02-01 00:00:00
std	2.818299e+09	NaN

	Time
count	1259
mean	1900-01-01 13:21:38.166004992
min	1900-01-01 00:02:36
25%	1900-01-01 11:19:59.500000
50%	1900-01-01 13:18:44
75%	1900-01-01 16:13:40
max	1900-01-01 23:59:59
std	NaN

```
[21]: #it gives the statistical summary of all the categorical variables
df.describe(include=object)
```

```
[21]:
```

	Gender	Country	state	self_employed	family_history	treatment \
count	1259	1259	744	1241	1259	1259
unique	49	48	45	2	2	2
top	Male	United States	CA	No	No	Yes
freq	615	751	138	1095	767	637

	work_interfere	no_employees	remote_work	tech_company	benefits \
count	995	1259	1259	1259	1259
unique	4	6	2	2	3
top	Sometimes	6-25	No	Yes	Yes
freq	465	290	883	1031	477

	care_options	wellness_program	seek_help	anonymity	leave \
count	1259	1259	1259	1259	1259

unique	3	3	3	3	5
top	No	No	No	Don't know	Don't know
freq	501	842	646	819	563

	mental_health_consequence	phys_health_consequence	coworkers	\
count	1259	1259	1259	
unique	3	3	3	
top	No	No	Some of them	
freq	490	925	774	

	supervisor	mental_health_interview	phys_health_interview	\
count	1259	1259	1259	
unique	3	3	3	
top	Yes	No	Maybe	
freq	516	1008	557	

	mental_vs_physical	obs_consequence	comments
count	1259	1259	164
unique	3	2	160
top	Don't know	No	* Small family business - YMMV.
freq	576	1075	5

## DATA PREPRATION

```
[22]: #finding the unique values in the column gender
df['Gender'].unique()
```

```
[22]: array(['Female', 'M', 'Male', 'male', 'female', 'm', 'Male-ish', 'maile',
        'Trans-female', 'Cis Female', 'F', 'something kinda male?',
        'Cis Male', 'Woman', 'f', 'Mal', 'Male (CIS)', 'queer/she/they',
        'non-binary', 'Femake', 'woman', 'Make', 'Nah', 'All', 'Enby',
        'fluid', 'Genderqueer', 'Female ', 'Androgynne', 'Agender',
        'cis-female/femme', 'Guy (-ish) ^_^', 'male leaning androgynous',
        'Male ', 'Man', 'Trans woman', 'msle', 'Neuter', 'Female (trans)',
        'queer', 'Female (cis)', 'Mail', 'cis male', 'A little about you',
        'Malr', 'p', 'femail', 'Cis Man',
        'ostensibly male, unsure what that really means'], dtype=object)
```

```
[23]: error={'Female':'F',
        'Male':'M',
        'male':'M',
        'female':'F',
        'm':'M',
        'Male-ish':'M',
        'maile':'M',
        'Trans-female':'T',
        'Cis Female':'F',
```

```

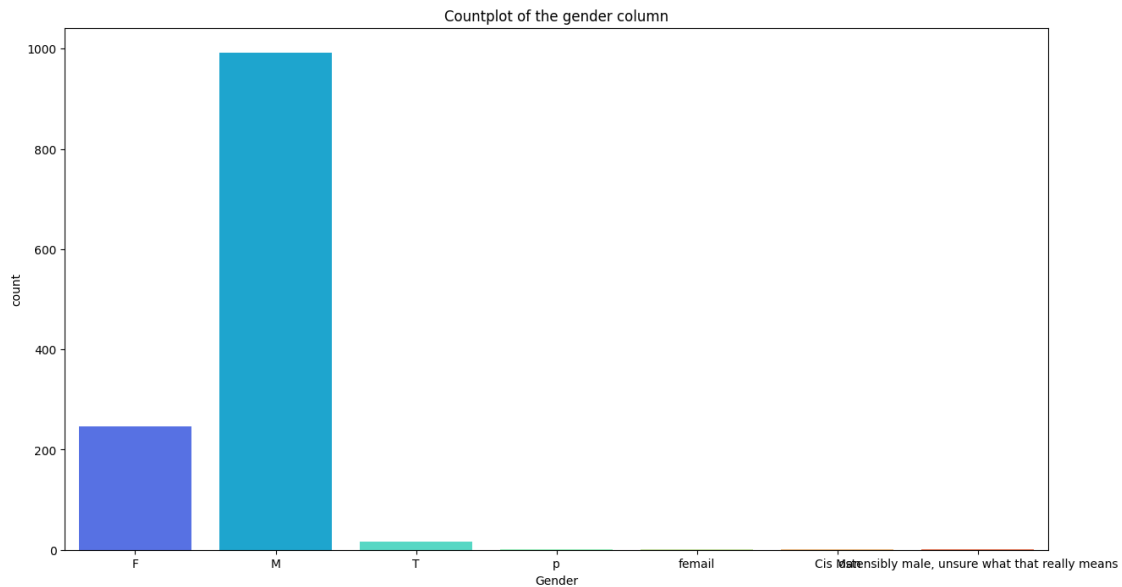
'something kinda male?':'M',
'Cis Male':'M',
'Woman':'F',
'f':'F',
'Mal':'M',
'Male (CIS)':'M',
'queer/she/they':'F',
'non-binary':'T',
'Enby':'T',
'Femake':'F',
'woman':'F',
'Make':'M',
'fluid':'T',
'Malr':'M',
'cis male':'M',
'Female (cis)':'F',
'Guy (-ish) ^_^':'M',
'queer':'T',
'Female (trans)':'T',
'male leaning androgynous':'T',
'Neuter':'T',
'cis-female/femme':'F',
'msle':'M',
'Agender':'T',
'Genderqueer':'T',
'Female':'F',
'Androgyne':'T',
'Nah':'T',
'All':'T',
'Female ':'F',
'Male ':'M',
'Man':'M',
'Trans woman':'T',
'Mail':'M',
'A little about you':'T'}
df['Gender']=df['Gender'].map(error).fillna(df['Gender'])

```

```
[24]: df['Gender'].unique()
```

```
[24]: array(['F', 'M', 'T', 'p', 'femail', 'Cis Man',
          'ostensibly male, unsure what that really means'], dtype=object)
```

```
[25]: #plotting the countplot for the gender column
sns.countplot(df['Gender'],palette='rainbow')
plt.title("Countplot of the gender column")
plt.show()
```



```
[26]: #Checking the number of male,female and transgender in the gender column
df['Gender'].value_counts()
```

```
[26]: Gender
M          992
F          247
T           16
p            1
femail       1
Cis Man       1
ostensibly male, unsure what that really means    1
Name: count, dtype: int64
```

```
[27]: #finding the unique values of age column
df['Age'].unique()
```

```
[27]: array([ 37, 44, 32, 31, 33,
          35, 39, 42, 23, 29,
          36, 27, 46, 41, 34,
          30, 40, 38, 50, 24,
          18, 28, 26, 22, 19,
          25, 45, 21, -29, 43,
          56, 60, 54, 329, 55,
          999999999999, 48, 20, 57, 58,
          47, 62, 51, 65, 49,
          -1726, 5, 53, 61, 8,
          11, -1, 72], dtype=int64)
```



```
[28]: #removing the redundant values from the age column
df=df[df['Age']!=99999999999]
df=df[df['Age']!=-29]
df=df[df['Age']!=329]
df=df[df['Age']!=-1726]
df=df[df['Age']!=5]
df=df[df['Age']!=8]
```

## LABEL ENCODING OF TREATMENT COLUMN

```
[29]: # replace 'no' with zero
df['treatment'] = df['treatment'].replace('No', 0)
# replace 'yes' with one
df['treatment'] = df['treatment'].replace('Yes', 1)

#displaying the first 5 records to check the treatment column after label
↳encoding
df.head()
```

```
[29]:
```

	Age	Gender	Country	state	self_employed	family_history	treatment	\
0	37	F	United States	IL	NaN	No	1	
1	44	M	United States	IN	NaN	No	0	
2	32	M	Canada	NaN	NaN	No	0	
3	31	M	United Kingdom	NaN	NaN	Yes	1	
4	31	M	United States	TX	NaN	No	0	

	work_interfere	no_employees	remote_work	tech_company	benefits	\
0	Often	6-25	No	Yes	Yes	
1	Rarely	More than 1000	No	No	Don't know	
2	Rarely	6-25	No	Yes	No	
3	Often	26-100	No	Yes	No	
4	Never	100-500	Yes	Yes	Yes	

	care_options	wellness_program	seek_help	anonymity	leave	\
0	Not sure	No	Yes	Yes	Somewhat easy	
1	No	Don't know	Don't know	Don't know	Don't know	
2	No	No	No	Don't know	Somewhat difficult	
3	Yes	No	No	No	Somewhat difficult	
4	No	Don't know	Don't know	Don't know	Don't know	

	mental_health_consequence	phys_health_consequence	coworkers	supervisor	\
0	No	No	Some of them	Yes	
1	Maybe	No	No	No	
2	No	No	Yes	Yes	
3	Yes	Yes	Some of them	No	
4	No	No	Some of them	Yes	

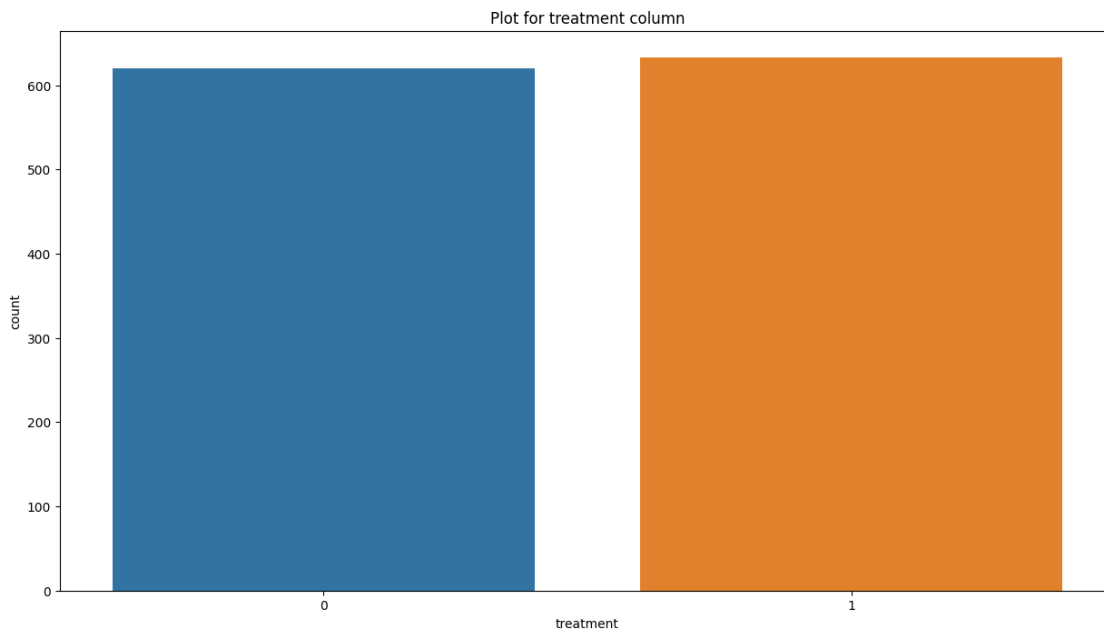
	mental_health_interview	phys_health_interview	mental_vs_physical	\
0	No	Maybe	Yes	
1	No	No	Don't know	
2	Yes	Yes	No	
3	Maybe	Maybe	No	
4	Yes	Yes	Don't know	

	obs_consequence	comments	Date	Time
0	No	NaN	2014-08-27	1900-01-01 11:29:31
1	No	NaN	2014-08-27	1900-01-01 11:29:37
2	No	NaN	2014-08-27	1900-01-01 11:29:44
3	Yes	NaN	2014-08-27	1900-01-01 11:29:46
4	No	NaN	2014-08-27	1900-01-01 11:30:22

```
[30]: #plotting the countplot for treatment column
sns.countplot(df['treatment'])
plt.title("Plot for treatment column")

#checking the count of each class
df['treatment'].value_counts()
```

```
[30]: treatment
1    633
0    620
Name: count, dtype: int64
```



## EXPLORATORY DATA ANALYSIS

```

[32]: # create a list of all categorical variables
# initiate an empty list to store the categorical variables
categorical=[]
z=['Country','state']
# use for loop to check the data type of each variable
for column in df:

    # use 'if' statement with condition to check the categorical type
    if is_string_dtype(df[column]):
        if column!=z:

            # append the variables with 'categorical' data type in the list
            ↪ 'categorical'
            categorical.append(column)

# plot the count plot for each categorical variable
fig, ax = plt.subplots(nrows = 5, ncols = 4, figsize=(25, 30))

# use for loop to plot the count plot for each variable
for variable, subplot in zip(categorical, ax.flatten()):

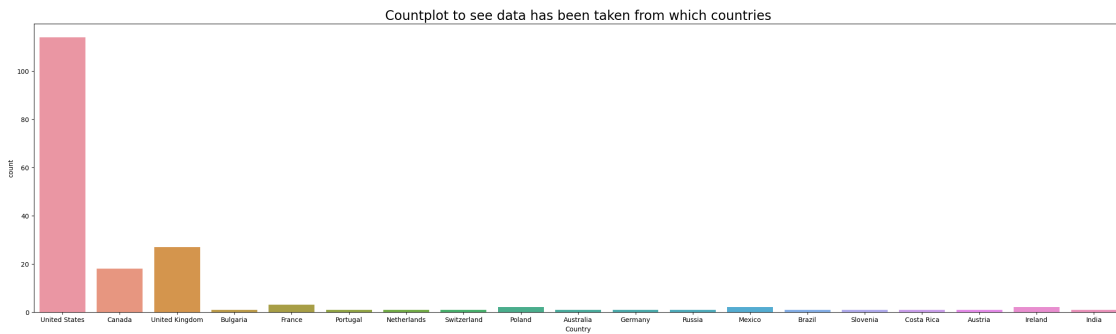
    # use countplot() to plot the graph
    sns.countplot(df[variable], ax = subplot)

# display the plot
plt.show()

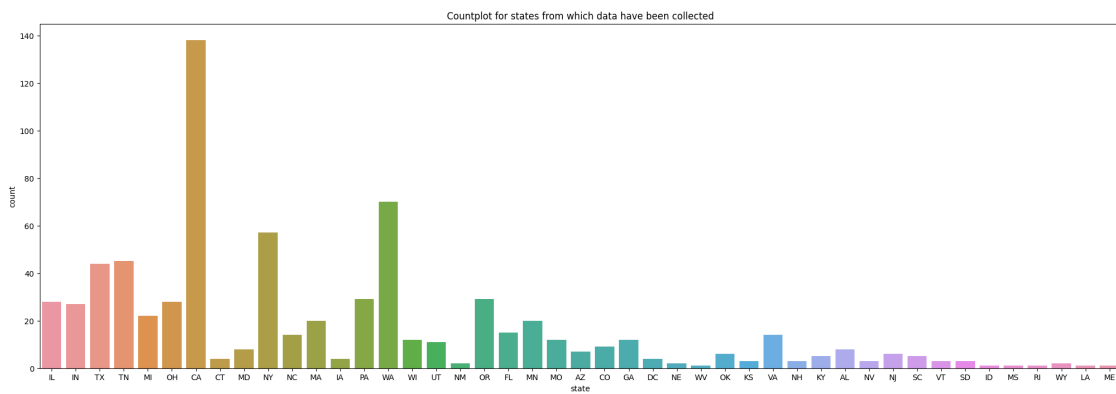
```



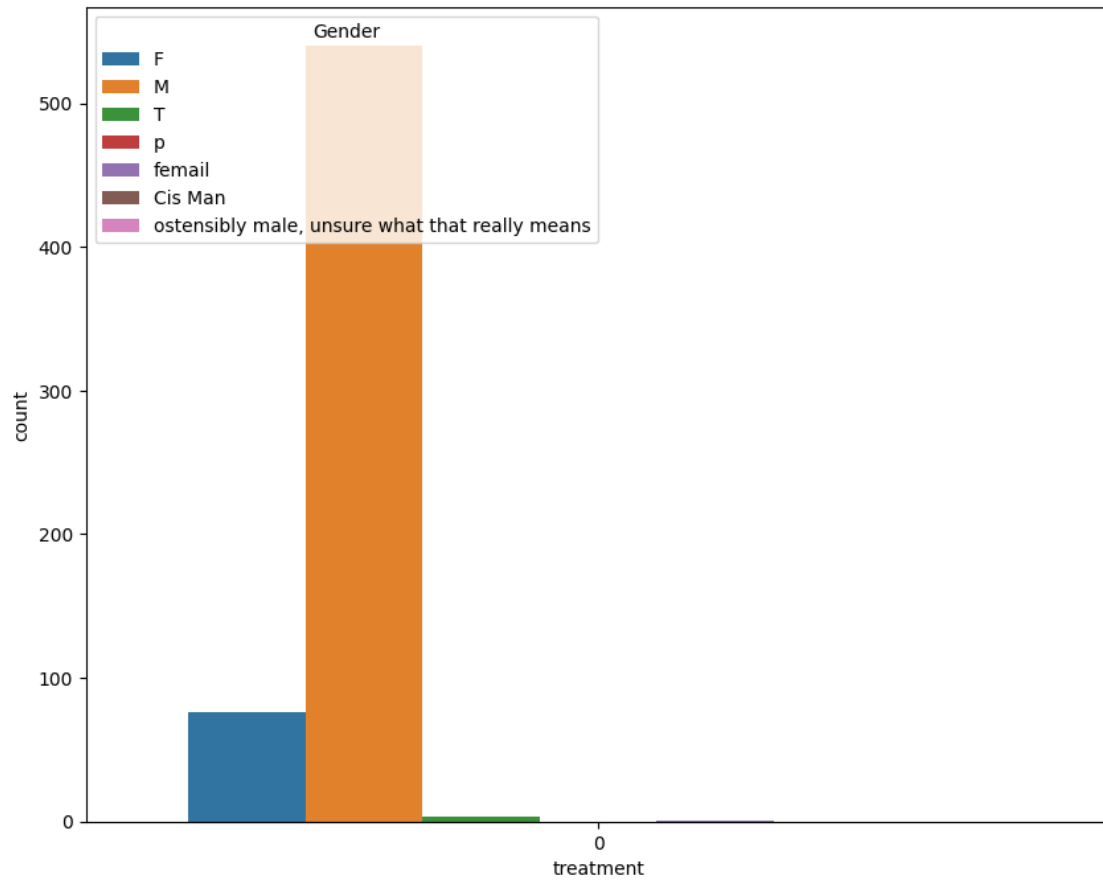
```
[33]: # plotting the counterplot for the country column
#to see
plt.figure(figsize=(30,8))
sns.countplot(df['Country'][:180])
plt.title("Countplot to see data has been taken from which_
countries",fontsize=20)
plt.show()
```



```
[34]: plt.figure(figsize=(25,8))
sns.countplot(df['state'])
plt.title("Countplot for states from which data have been collected")
plt.show()
```

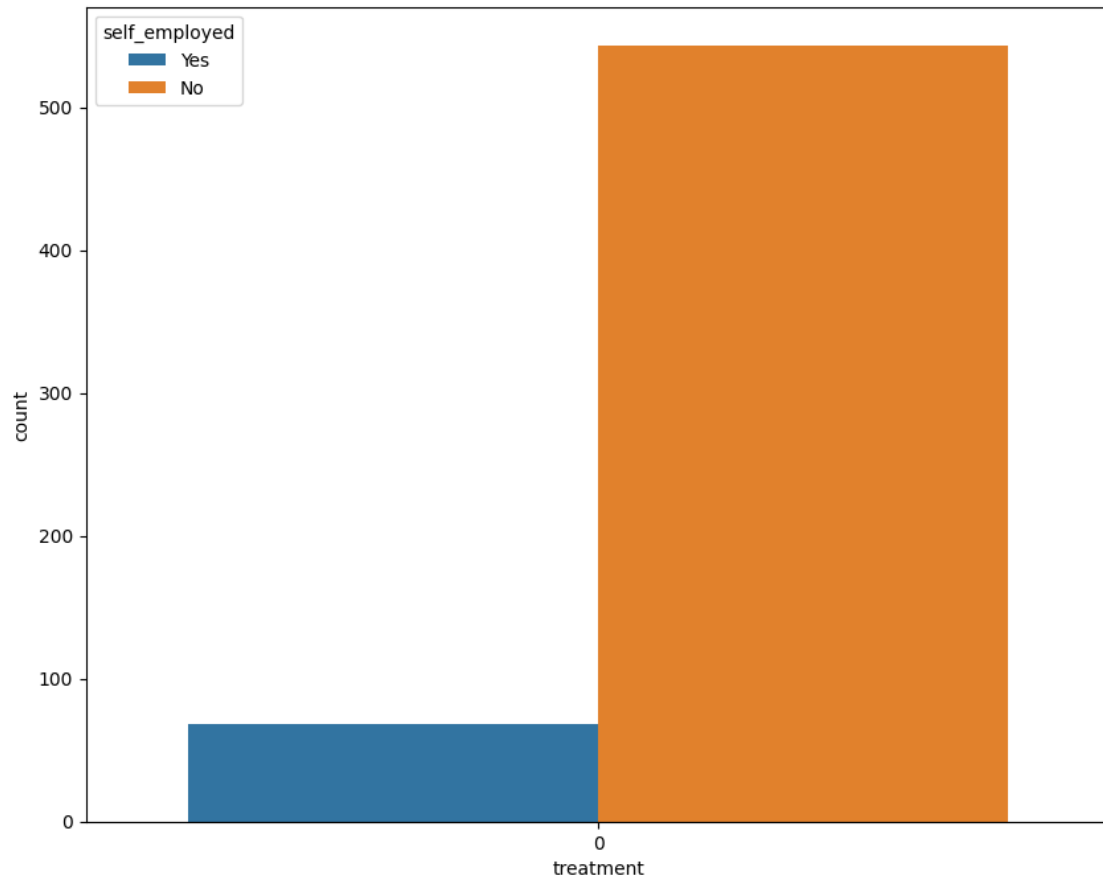


```
[35]: #Checking employees who require treatment are from which gender
sns.countplot(df.treatment, hue=df.Gender, order=df['treatment'].value_counts().
            iloc[1:2].index)
plt.show()
```



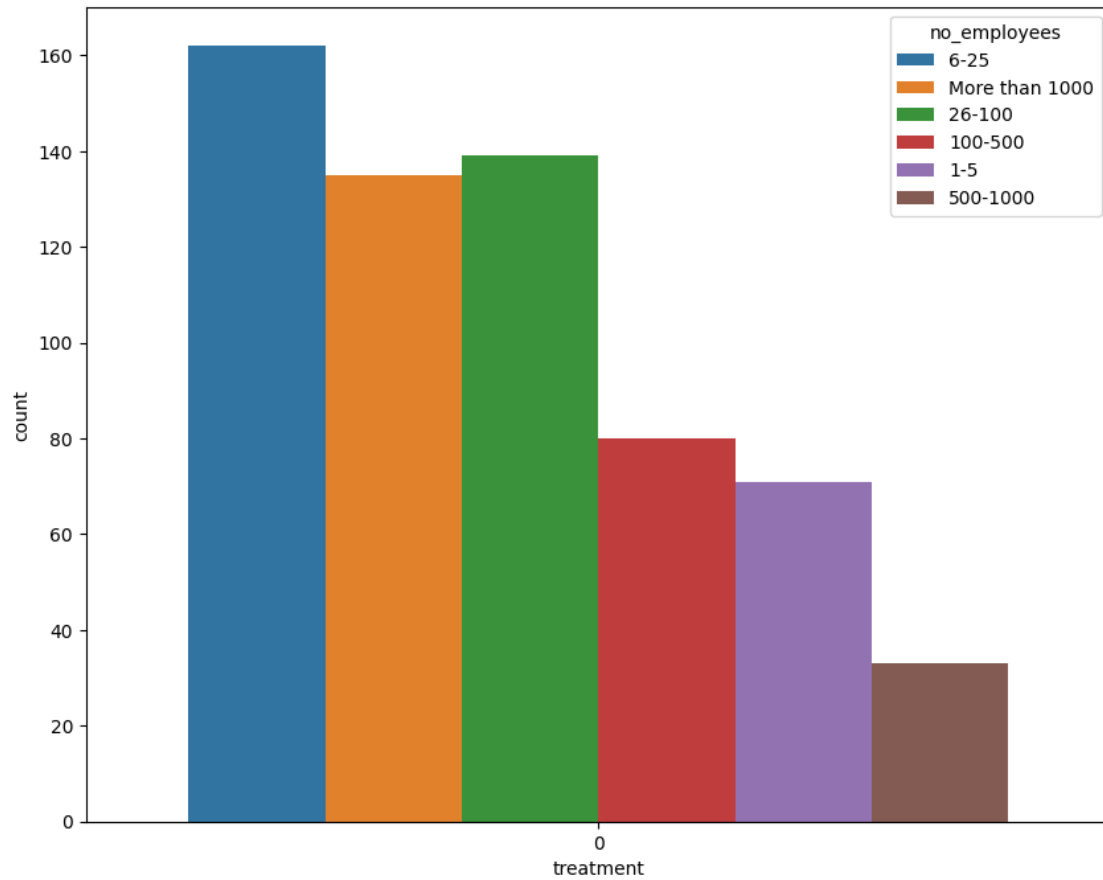
```
[36]: #plotting countplot to see how many self-employed people requires treatment
sns.countplot(df.treatment,hue=df['self_employed'],order=df['treatment'].
↳value_counts().iloc[1:2].index)
```

```
[36]: <Axes: xlabel='treatment', ylabel='count'>
```



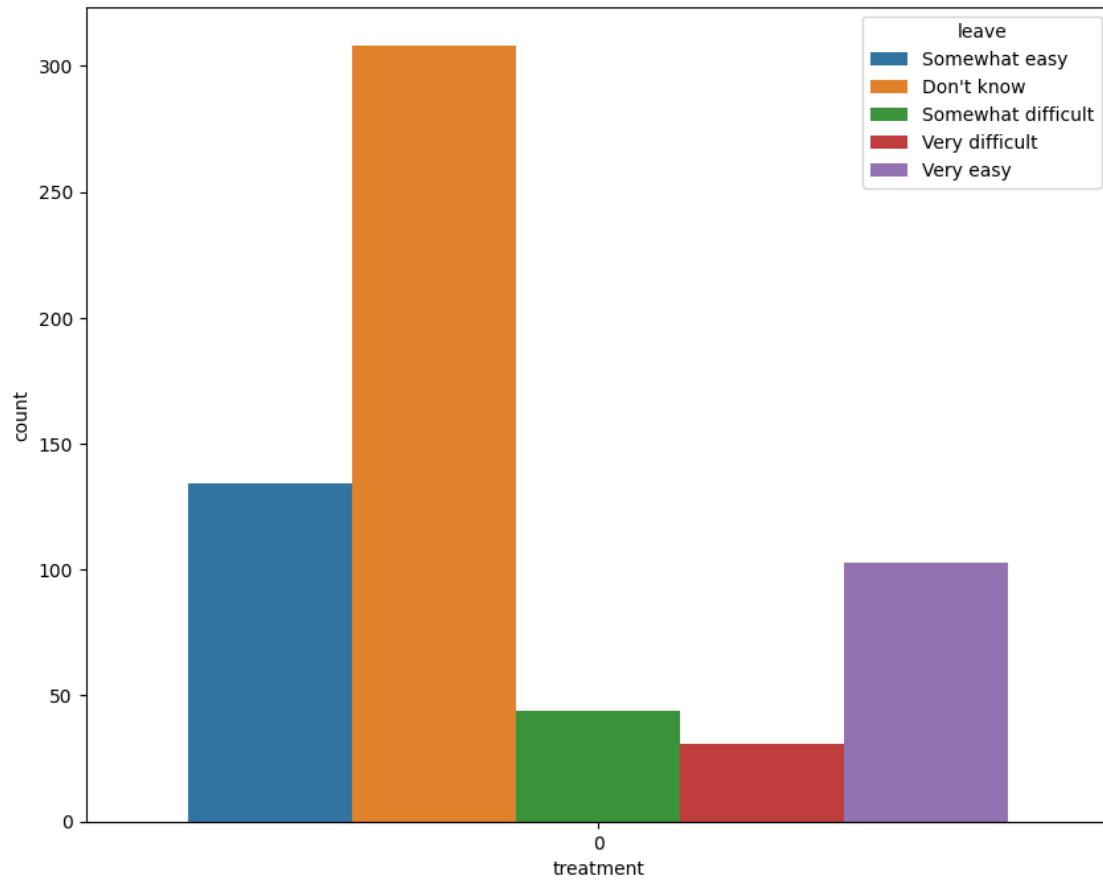
```
[37]: #checking does number of employees in an organisation affects the treatment rate
sns.countplot(df.treatment,hue=df['no_employees'],order=df['treatment'].
↳value_counts().iloc[1:2].index)
```

```
[37]: <Axes: xlabel='treatment', ylabel='count'>
```

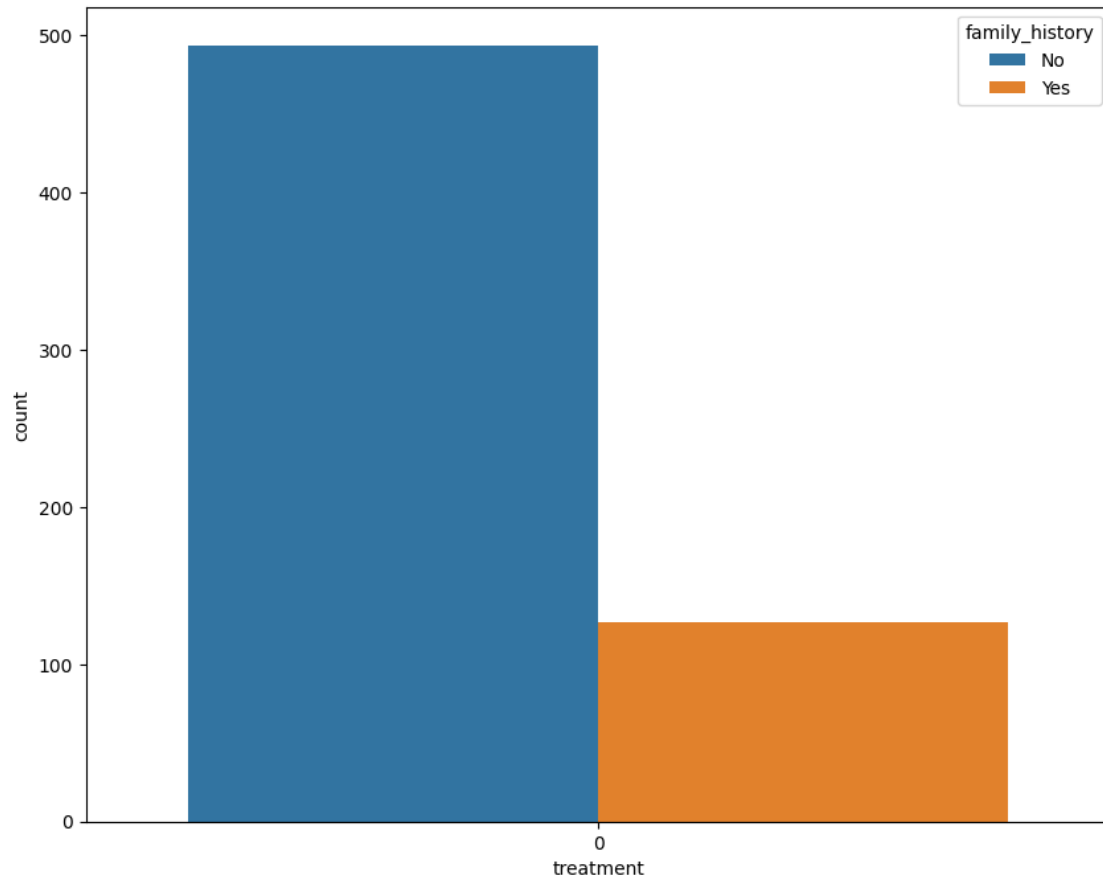


```
[38]: #plot to see does how easy it to take medical leave and its affect on treatment
      ↪requirement
      sns.countplot(df['treatment'],hue=df['leave'],order=df['treatment'].
      ↪value_counts().iloc[1:2].index)
      plt.show()
```

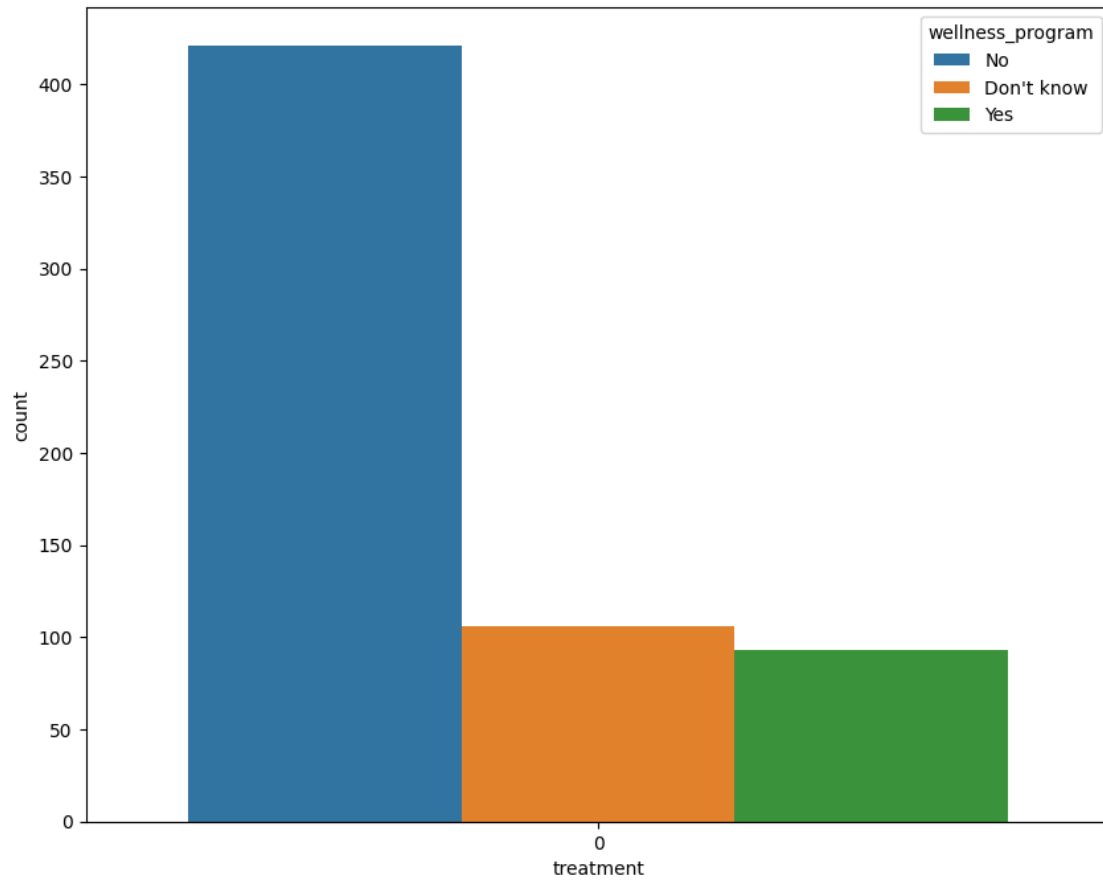




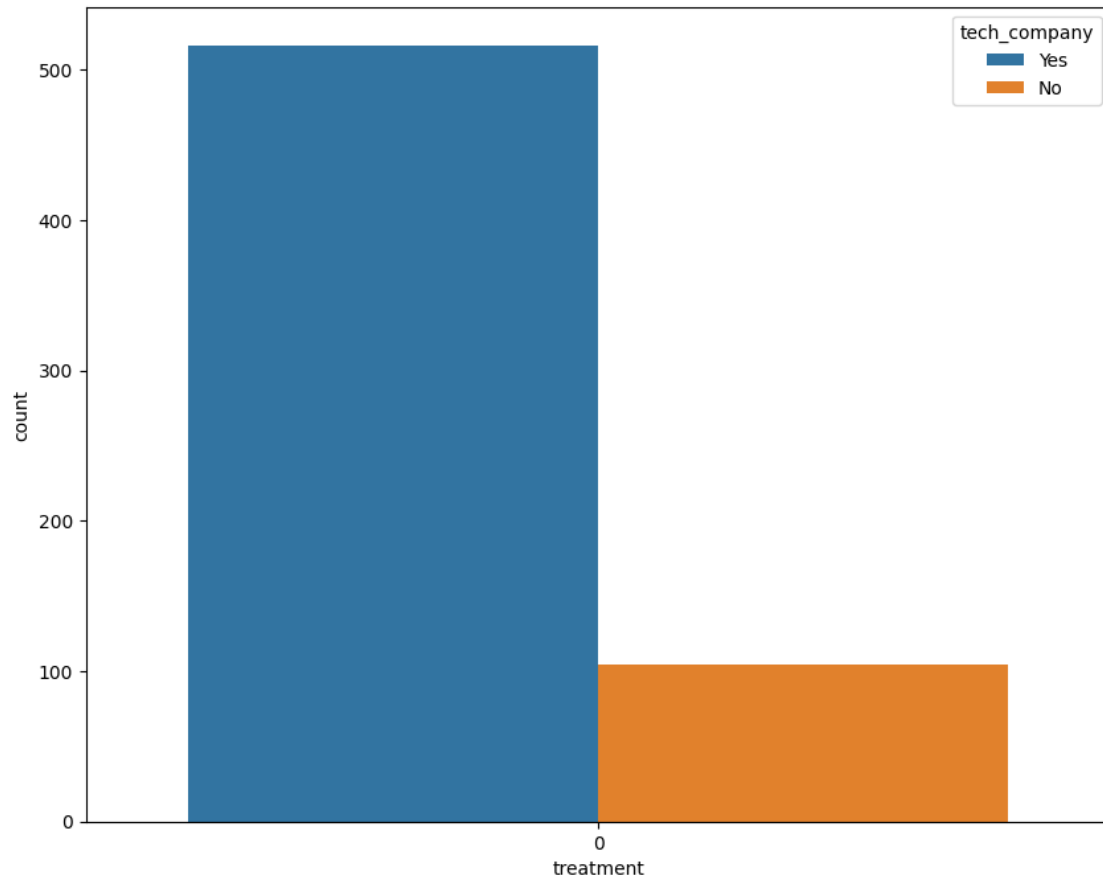
```
[39]: sns.countplot(df.treatment,hue=df['family_history'],order=df['treatment'].  
        ↳value_counts().iloc[1:2].index)  
plt.show()
```



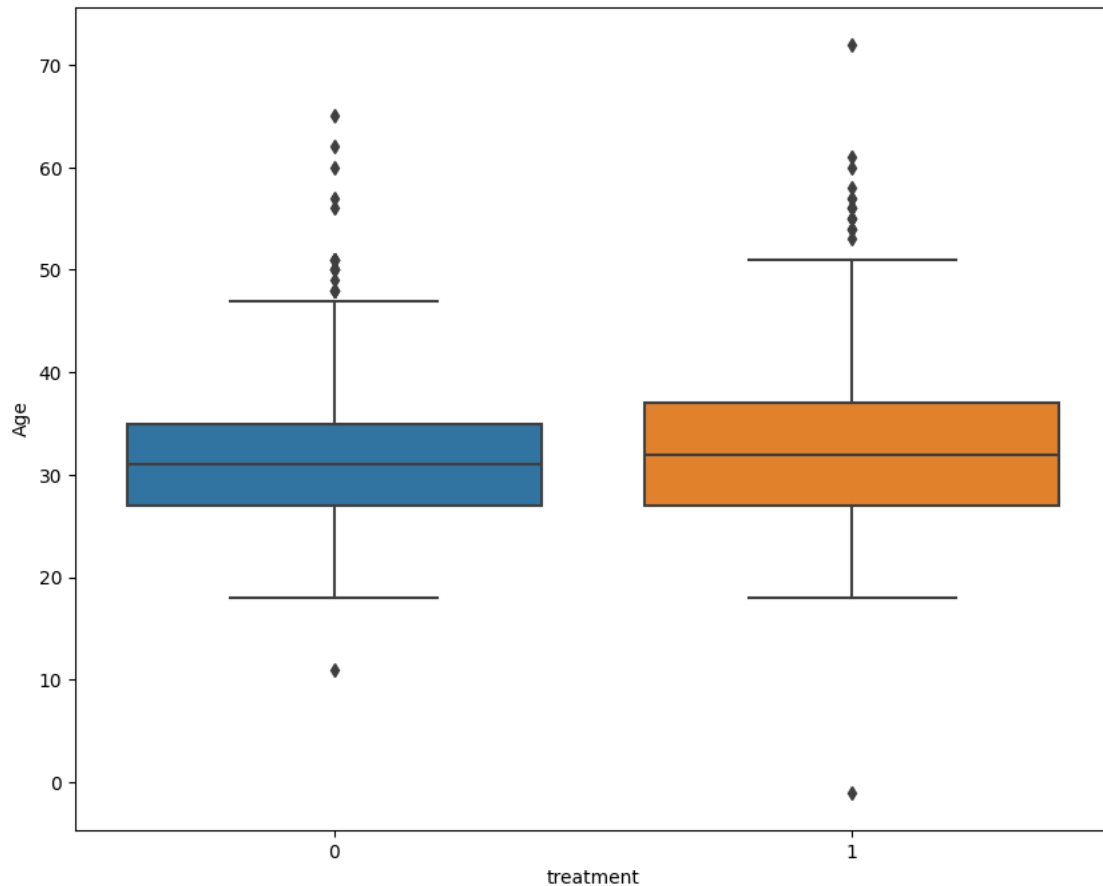
```
[40]: sns.countplot(df['treatment'], hue=df['wellness_program'], order=df['treatment'].  
        ↳ value_counts().iloc[1:2].index)  
plt.show()
```



```
[42]: sns.countplot(df['treatment'], hue=df['tech_company'], order=df['treatment'].  
        ↳ value_counts().iloc[1:2].index)  
plt.show()
```



```
[43]: #plotting the bar plot for age to see if there is any outlier  
sns.boxplot(x=df.treatment,y=df['Age'])  
plt.show()
```



## FINDING THE MISSING VALUES

```
[44]: # sort the variables on the basis of total null values in the variable
# 'isnull().sum()' returns the number of missing values in each variable
Total = df.isnull().sum().sort_values(ascending = False)

# calculate the percentage of missing values
Percent = ((Total*100)/df.isnull().count()).sort_values(ascending = False)

# concat the 'Total' and 'Percent' columns using 'concat' function
missing_data = pd.concat([Total, Percent], axis = 1, keys = ['Total', 'Percentage of Missing Values'])
missing_data
```

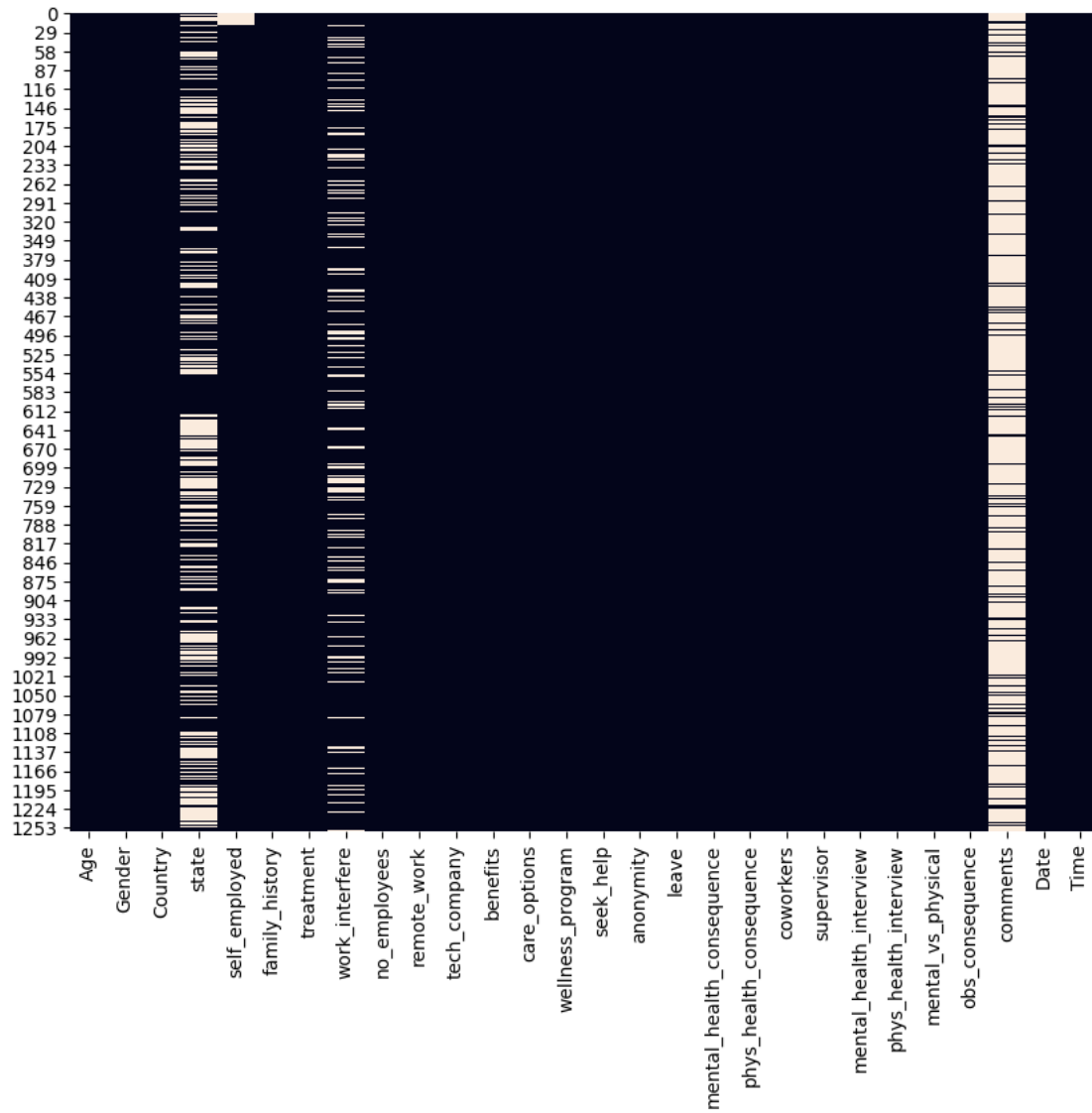
```
[44]:
```

	Total	Percentage of Missing Values
comments	1091	87.071030
state	513	40.941740
work_interfere	262	20.909816
self_employed	18	1.436552

Age	0	0.000000
leave	0	0.000000
Date	0	0.000000
obs_consequence	0	0.000000
mental_vs_physical	0	0.000000
phys_health_interview	0	0.000000
mental_health_interview	0	0.000000
supervisor	0	0.000000
coworkers	0	0.000000
phys_health_consequence	0	0.000000
mental_health_consequence	0	0.000000
seek_help	0	0.000000
anonymity	0	0.000000
Gender	0	0.000000
wellness_program	0	0.000000
care_options	0	0.000000
benefits	0	0.000000
tech_company	0	0.000000
remote_work	0	0.000000
no_employees	0	0.000000
treatment	0	0.000000
family_history	0	0.000000
Country	0	0.000000
Time	0	0.000000

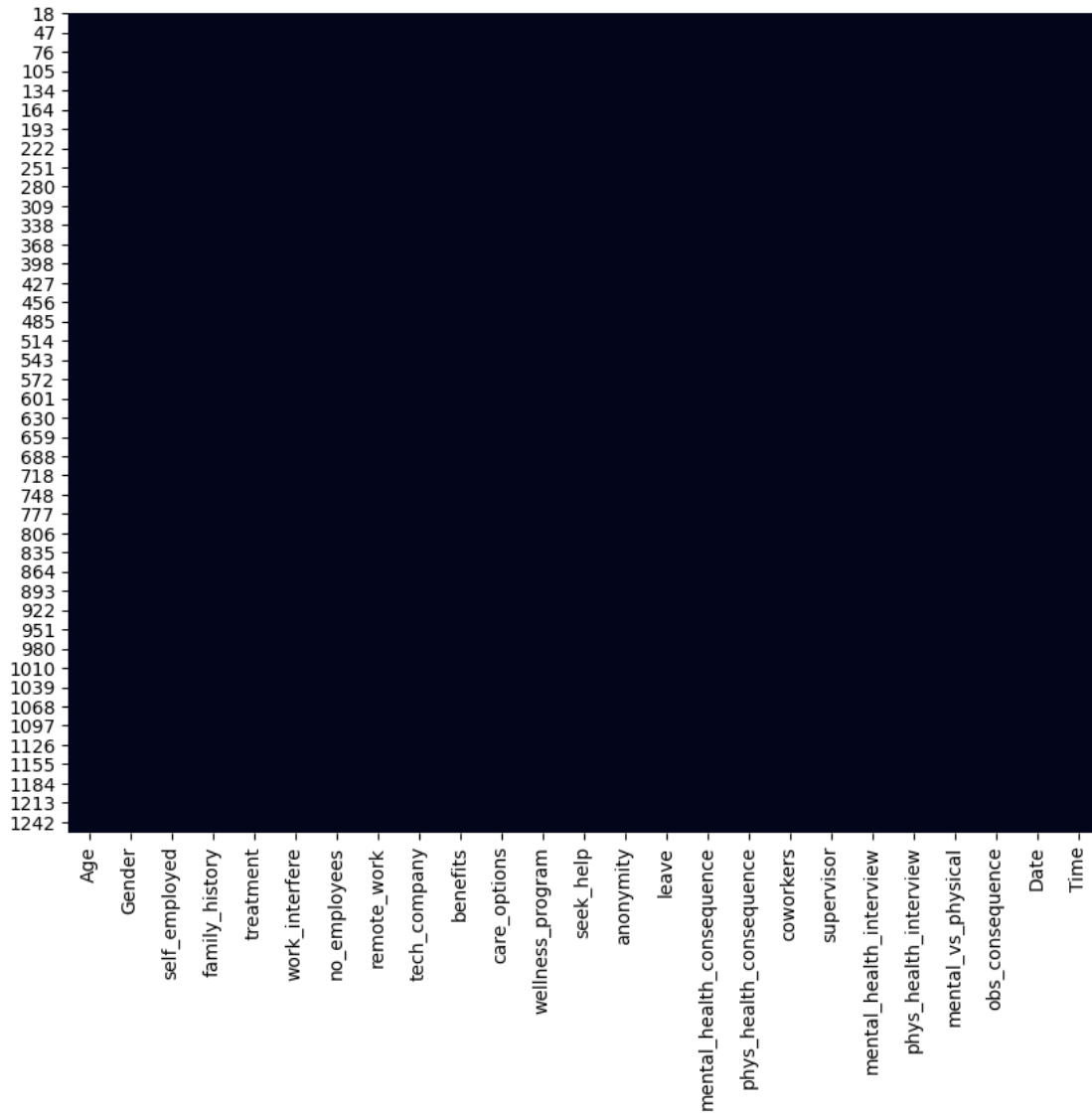
```
[45]: # plot heatmap to check null values
# 'cbar = False' does not show the color axis
sns.heatmap(df.isnull(), cbar=False)

# display the plot
plt.show()
```



## HANDLING THE MISSING VALUES

```
[46]: df.drop('comments',axis=1,inplace=True)
df['work_interfere']=df['work_interfere'].fillna('Not mentioned')
df.drop(['state','Country'],axis=1,inplace=True)
df.dropna(axis=0, inplace=True)
#Checking if all the null values have been handled or not
sns.heatmap(df.isnull(),cbar=False,color='black')
plt.show()
```



## PREPARING THE DATA FOR BUILDING MODEL

```
[47]: #Creating two dataframes df_features and df_target, df_features contains all the
      ↪ important features which we will dummy encode
      #df_target which contains the target variable
      df_features=df.drop(['treatment', 'Age', 'Date', 'Time'],axis=1)
      df_target=df['treatment']
      #dummy encoding the feature(categorical) variables
      df_dummy=pd.get_dummies(df_features,drop_first=True)
      #storing the features in X and the target in y variable
      X=df_dummy
      y=pd.DataFrame(df_target)
```



## CREATING GENERALISED FUNCTIONS

```
[48]: # create a generalized function to calculate the metrics values for test set
def get_test_report(model):

    # return the performace measures on test set
    return(classification_report(y_test, y_pred))
# create a generalized function to calculate the metrics values for test set
def kappa_score(model):

    # return the kappa score on test set
    return(cohen_kappa_score(y_test, y_pred))
# define a to plot a confusion matrix for the model
def plot_confusion_matrix(model):

    # create a confusion matrix
    cm = confusion_matrix(y_test, y_pred)
    conf_matrix = pd.DataFrame(data = cm, columns = ['Predicted:0', 'Predicted:
↪1'], index = ['Actual:0', 'Actual:1'])

    # plot a heatmap to visualize the confusion matrix
    sns.heatmap(conf_matrix, annot = True, fmt = 'd', cmap = _
↪ListedColormap(['lightskyblue']), cbar = False,
                linewidths = 0.1, annot_kws = {'size':25})

    # set the font size of x-axis ticks using 'fontsize'
    plt.xticks(fontsize = 20)

    # set the font size of y-axis ticks using 'fontsize'
    plt.yticks(fontsize = 20)

    # display the plot
    plt.show()
```

```
[51]: # define a function to plot the ROC curve and print the ROC-AUC score
def plot_roc(model):

    # the roc_curve() returns the values for false positive rate, true positive
↪rate and threshold
    fpr, tpr, thresholds = roc_curve(y_test, y_pred)

    # plot the ROC curve
    plt.plot(fpr, tpr)

    # set limits for x and y axes
    plt.xlim([0.0, 1.0])
    plt.ylim([0.0, 1.0])
```

```

# plot the straight line showing worst prediction for the model
plt.plot([0, 1], [0, 1], 'r--')

# add plot and axes labels
# set text size using 'fontsize'
plt.title('ROC Curve', fontsize = 15)
plt.xlabel('False positive rate (1-Specificity)', fontsize = 15)
plt.ylabel('True positive rate (Sensitivity)', fontsize = 15)

# add the AUC score to the plot
plt.text(x = 0.02, y = 0.9, s = ('AUC Score:', round(roc_auc_score(y_test,
↪y_pred), 4)))

# plot the grid
plt.grid(True)

```

```

[52]: # create an empty dataframe to store the scores for various classification
↪algorithms
score_card = pd.DataFrame(columns=['Model', 'AUC Score', 'Precision Score',
↪'Recall Score', 'Accuracy Score',
                                'Kappa Score', 'f1-score'])

def update_score_card(model_name):

    # assign 'score_card' as global variable
    global score_card

    # append the results to the dataframe 'score_card'
    # 'ignore_index = True' do not consider the index labels
    score_card = score_card.append({'Model': model_name,
                                    'AUC Score' : roc_auc_score(y_test, y_pred),
                                    'Precision Score': metrics.
↪precision_score(y_test, y_pred),
                                    'Recall Score': metrics.
↪recall_score(y_test, y_pred),
                                    'Accuracy Score': metrics.
↪accuracy_score(y_test, y_pred),
                                    'Kappa Score': cohen_kappa_score(y_test,
↪y_pred),
                                    'f1-score': metrics.f1_score(y_test,
↪y_pred)}),
                                ignore_index = True)

    return(score_card)

```

```
[53]: # split data into train subset and test subset
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.30,
↳ random_state = 10)

# check the dimensions of the train & test subset using 'shape'
# print dimension of train set
print("X_train",X_train.shape)
print("y_train",y_train.shape)

# print dimension of test set
print("X_test",X_test.shape)
print("y_test",y_test.shape)
```

X\_train (864, 48)

y\_train (864, 1)

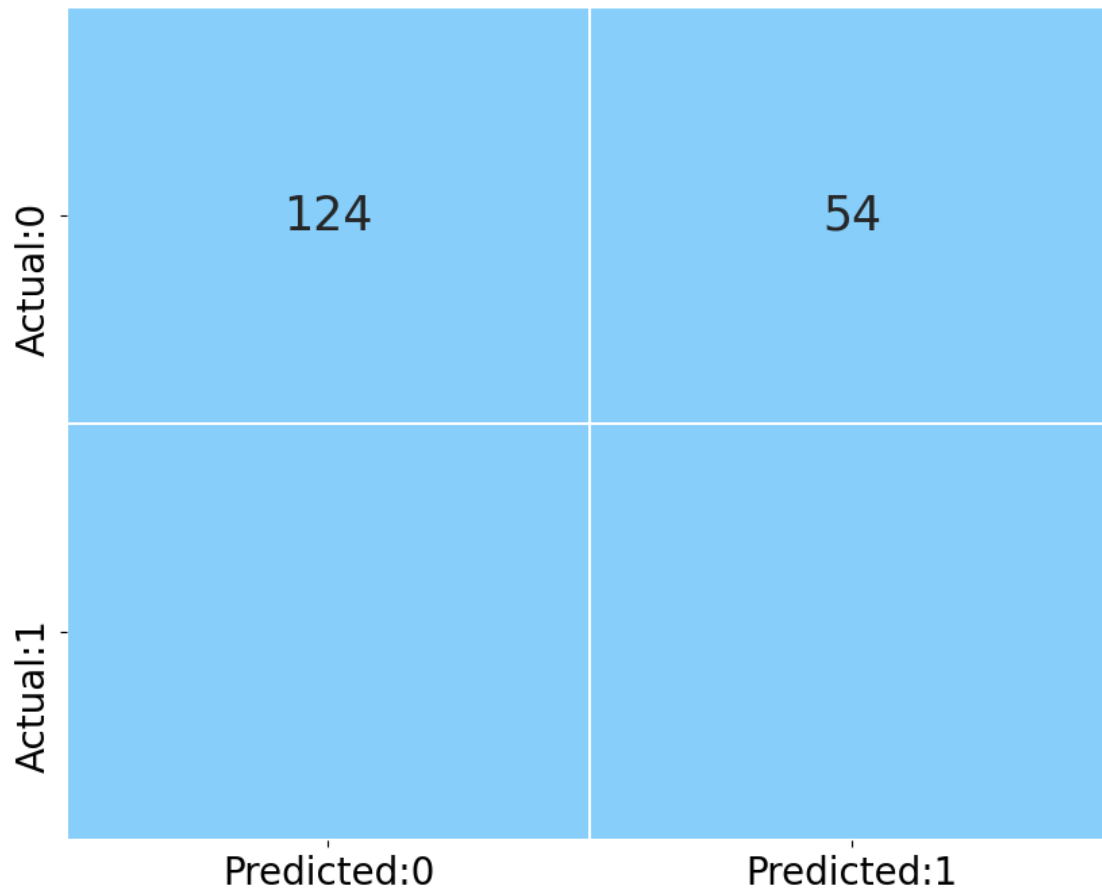
X\_test (371, 48)

y\_test (371, 1)

SUPPORT VECTOR MACHINE

```
[55]: # build the model
svclassifier = SVC(kernel = 'linear')

# fit the model
svc_model=svclassifier.fit(X_train, y_train)
# predict the values
y_pred = svclassifier.predict(X_test)
# call the function to plot the confusion matrix
plot_confusion_matrix(svc_model)
```



```
[56]: # compute the performance measures on test data
test_report = get_test_report(svc_model)

# print the performance measures
print(test_report)
```

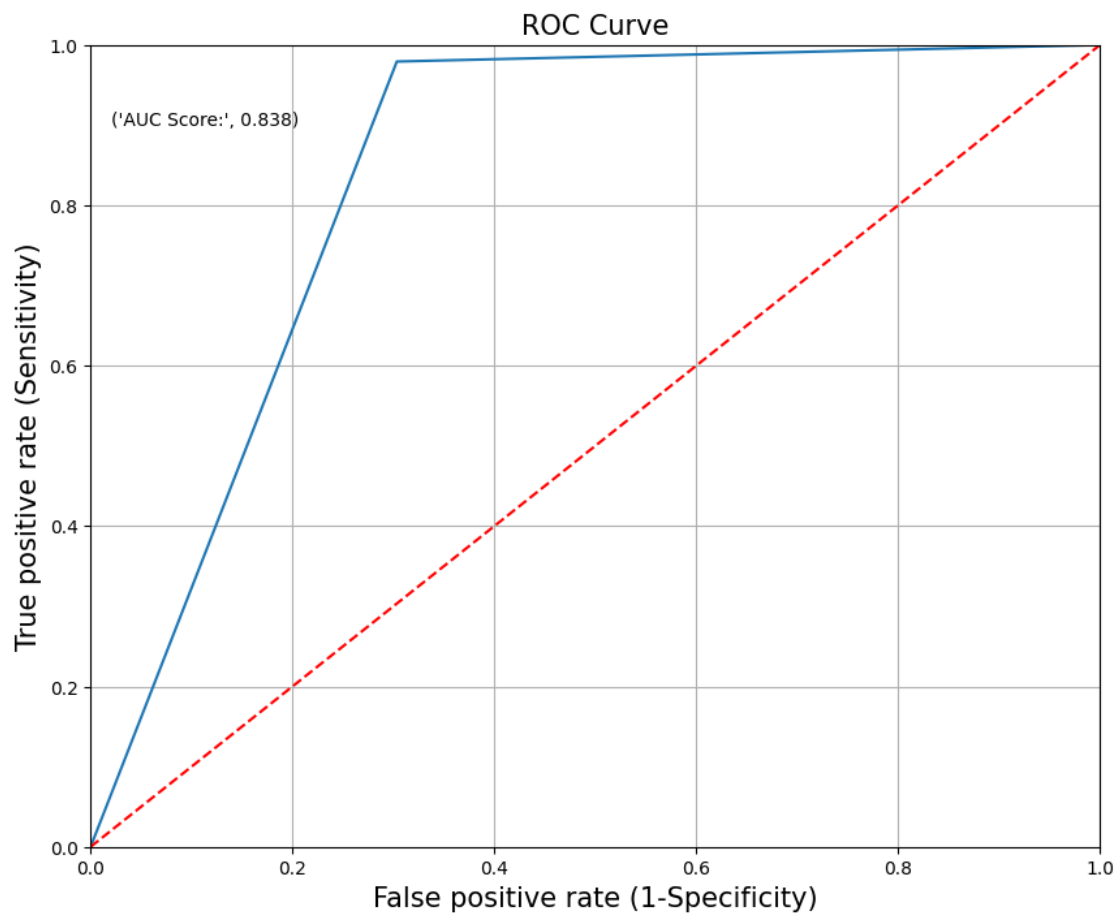
	precision	recall	f1-score	support
0	0.97	0.70	0.81	178
1	0.78	0.98	0.87	193
accuracy			0.84	371
macro avg	0.87	0.84	0.84	371
weighted avg	0.87	0.84	0.84	371

```
[57]: # compute kappa score on test set
kappa_value = kappa_score(svc_model)
```

```
# print the kappa value  
print(kappa_value)
```

0.6833632537743901

```
[71]: plot_roc(svc_model)
```



ACCURACY =0.84

```
[ ]:
```