### DAC\_PHASE5

### DOCUMENTATION AND SUBMISSION

Date: 01-11-2023 Team ID:716

**Project Title: Public Health Awareness Campaign Analysis** 

### **Introduction:**

In this, we are building upon our analysis efforts by utilizing IBM Cognos for data visualization and integrating code, potentially in Python, for advanced data analysis. Our primary objective remains the assessment of the public health awareness campaign's effectiveness and impact. We will design interactive dashboards and reports in IBM Cognos to visually represent campaign reach, awareness levels, and impact metrics, offering valuable insights for stakeholders. Furthermore, we will use code to perform in-depth analysis, including calculating engagement rates, conducting demographic analysis, and running statistical tests, enabling us to provide comprehensive findings that can inform decisions and contribute to the betterment of our communities.

# **Objective:**

The objective of a public health awareness campaign analysis using data analytics is to leverage data-driven insights to assess the effectiveness of the campaign in reaching its target audience, raising awareness, and promoting positive health behaviors. Through data analytics, the campaign aims to identify key performance indicators, assess the impact on public health outcomes, and refine strategies for more impactful and efficient future initiatives.

## **Design Thinking Process:**

#### 1.Data collection

**Data Sources**: Utilize external sources, such as Kaggle, for obtaining datasets containing mental health survey.

**Tools**: Jupyter notebook, IBM cognos tool

## 2.Import necessary libraries

import pandas as

pd import numpy

```
as np import
seaborn as sns
import matplotlib.pyplot as
pltprint('Successfully
imported')
```

This dataset contains the following data:

- \* Timestamp
- \* Age
- \* Gender
- \* Country
- \* state: If you live in the United States, which state or territory do you live in?
- \* self\_employed: Are you self-employed?
- \* family\_history: Do you have a family history of mental illness?
- \* work\_interfere: If you have a mental health condition, do you feel that it interferes with your work?
- \* no\_employees: How many employees does your company or organization have?
- \*remote\_work: Do you work remotely (outside of an office) at least 50% of the time
- \* tech\_company: Is your employer primarily a tech company/organization?
- \*benefits: Does your employer provide mental health benefits? care\_options: Do you know the options for mental health care your employer provides?
- \*wellness\_program: Has your employer ever discussed mental health as part of an employee wellness
- \*seek\_help: Does your employer provide resources to learn more about mental health issues and howto seek help?
  - \* anonymity: Is your anonymity protected if you choose to take advantage of mental health orsubstance abuse treatment resources?
  - \* leave: How easy is it for you to take medical leave for a mental health condition?
  - \* mental\_health\_consequence: Do you think that discussing a mental health issue with your employerwould have negative consequences?
  - \* phys\_health\_consequence: Do you think that discussing a physical health issue with your employerwould have negative consequences?
  - \* coworkers: Would you be willing to discuss a mental health issue with your coworkers?
  - \* supervisor: Would you be willing to discuss a mental health issue with your direct supervisor(s)?
  - \* mental health interview: Would you bring up a mental health issue with a potential

employer in aninterview?

- \* phys\_health\_interview: Would you bring up a physical health issue with a potential employer in aninterview?
- \* mental\_vs\_physical: Do you feel that your employer takes mental health as seriously as physicalhealth?
- \* obs\_consequence: Have you heard of or observed negative consequences for coworkers with mentalhealth conditions in your workplace?
- \* comments: Any additional notes or comments
- \* treatment: Have you sought treatment for a mental health condions.

### 3.Read Dataset

```
data = pd.read_csv('/kaggle/input/mental-health-in-tech-
survey/survey.csv')data.head()
```

### 4. Preprocessing and Cleaning dataset

```
if data.isnull().sum().sum() == 0:
    print ('There is no missing data in our dataset'')

else:
    print('There is {} missing data in our dataset '.format(data.isnull().sum().sum()))

There is 1892 missing data in our dataset

frame = pd.concat([data.isnull().sum(), data.nunique(), data.dtypes], axis = 1,
    sort= False)frame

data['work_interfere'].unique()
    array(['Often', 'Rarely', 'Never', 'Sometimes', nan],
    dtype=object)ax = sns.countplot(data = data , x =
    'work_interfere');

#Add the value of each parametr on the

Plotax.bar_label(ax.containers[0]);
```

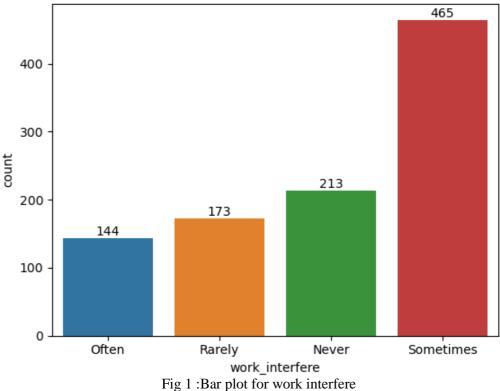
else:

print('There is {} missing data in our dataset '.format(data.isnull().sum().sum()))

There is 1892 missing data in our dataset

frame = pd.concat([data.isnull().sum(), data.nunique(), data.dtypes], axis = 1, sort= False) frame

data['work\_interfere'].unique() array(['Often', 'Rarely', 'Never', 'Sometimes', nan], dtype=object) ax = sns.countplot(data = data , x = 'work\_interfere'); #Add the value of each parametr on the Plot ax.bar\_label(ax.containers[0]);



In this diagram x axis is work interfere and y axis is count this diagram shows sometimes is high.

#For filling nan values i used SimpleImputer but you can use fillnan function toofrom sklearn.impute import SimpleImputer

data = data.drop(columns=['state', 'comments', 'Timestamp', ])# Fill in missing values in work\_interfere column

```
data['work_interfere'] = SimpleImputer(strategy =
'most_frequent').fit_transform(data['work_interfere'].values.reshape(-1,1))
data['self_employed'] = SimpleImputer(strategy =
'most_frequent').fit_transform(data['self_employed'].values.reshape(-1,1))
data.head()
```

ax = sns.countplot(data=data, x='work\_interfere');ax.bar\_label(ax.containers[0]);

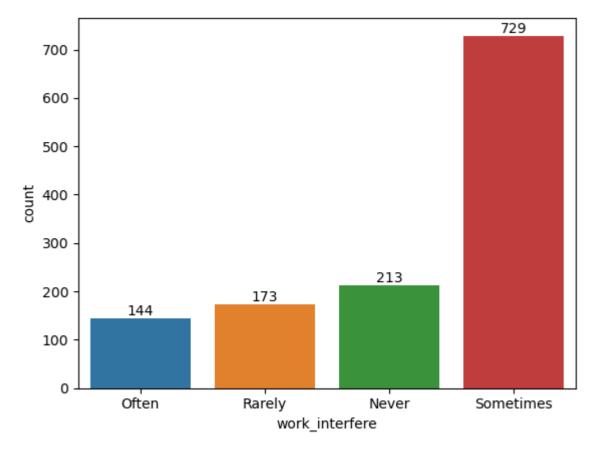


Fig 2:Bar plot for work interfere

In this diagram x axis is work interfere and y axis is count this diagram shows sometimes is high.

```
#Check unique data in gender columns

print(data['Gender'].unique())

print(')

print('-'*75)

print(")

#Check number of unique data too.

print('number of unique Gender in our dataset is :', data['Gender'].nunique())
```

```
['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 ' 'Androgyne' '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']
number of unique Gender in our dataset is : 49
#Gender data contains dictation problems, nonsense answers, and too unique Genders.
#_So Let's clean it and organize it into Male, Female, and other categories
data['Gender'].replace(['Male', 'male', 'M', 'm', 'Male', 'Cis Male',
             'Man', 'cis male', 'Mail', 'Male-ish', 'Male (CIS)',
              'Cis Man', 'msle', 'Malr', 'Mal', 'maile', 'Make',], 'Male', inplace = True)
data['Gender'].replace(['Female', 'female', 'F', 'f', 'Woman', 'Female',
             'femail', 'Cis Female', 'cis-female/femme', 'Femake', 'Female (cis)',
             'woman',], 'Female', inplace = True)
data["Gender"].replace(['Female (trans)', 'queer/she/they', 'non-binary',
             'fluid', 'queer', 'Androgyne', 'Trans-female', 'male leaning androgynous',
              'Agender', 'A little about you', 'Nah', 'All',
              'ostensibly male, unsure what that really means',
              'Genderqueer', 'Enby', 'p', 'Neuter', 'something kinda male?',
              'Guy (-ish) ^_^', 'Trans woman',], 'Other', inplace = True)
print(data['Gender'].unique())
```

```
['Female' 'Male' 'Other']
```

```
#Plot Genders column after cleaning and new categorizing
ax = sns.countplot(data=data, x='Gender');
ax.bar_label(ax.containers[0]);
```

## **Output**

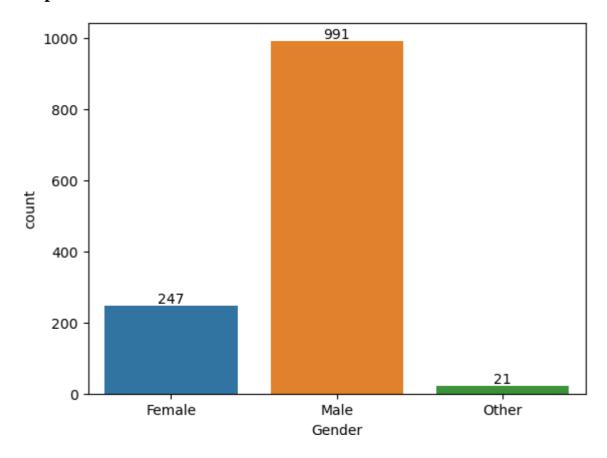


Fig 3:Bar plot for gender and count

In this diagram x axis is gender and y axis is count this diagram shows male is high.

```
#Our data is clean now ? let's see.
if data.isnull().sum().sum() == 0:
    print('There is no missing data')
else:
    print('There is {} missing data'.format(data.isnull().sum().sum()))
```

```
There is no missing data
```

```
#Let's check duplicated data.
if data.duplicated().sum() == 0:
    print('There is no duplicated data:')
else:
    print('Tehre is {} duplicated data:'.format(data.duplicated().sum()))
    #If there is duplicated data drop it.
    data.drop_duplicates(inplace=True)

print('-'*50)
print(data.duplicated().sum())
```

# **Output**

Tehre is 4 duplicated data:

\_\_\_\_\_

0

#Look unique data in Age column data['Age'].unique()

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,	
99	9999999999	,	48,	20,	57,	58,
	47,	62,	51,	65,	49,	
	-1726,	5,	53,	61,	8,	

```
#We had a lot of nonsense answers in the Age column too

#This filtering will drop entries exceeding 100 years and those indicating negative values.

data.drop(data[data['Age']<0].index, inplace = True)

data.drop(data[data['Age']>99].index, inplace = True)

print(data['Age'].unique())

Output

[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 43 56 60 54 55 48 20 57 58 47 62 51 65 49 5 53 61 8 11 72]

#Let's see the Age distribution in this dataset.

plt.figure(figsize = (10,6))

age_range_plot = sns.countplot(data = data, x = 'Age');

age_range_plot.bar_label(age_range_plot.containers[0]);

plt.xticks(rotation=90);
```

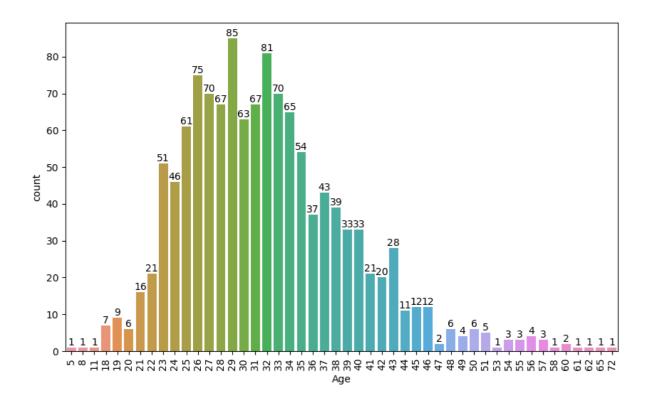


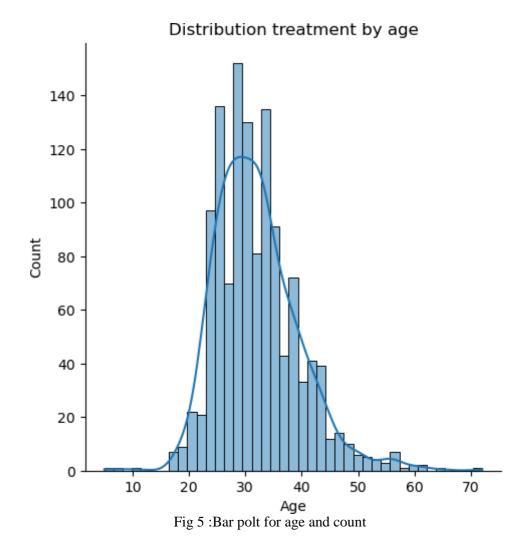
Fig 4 :Range plot for age and count

In this diagram x axis is age and y axis is count

#In this plot moreover on Age distribution we can see treatment distribution by age plt.figure(figsize=(10, 6)); sns.displot(data['Age'], kde = 'treatment'); plt.title('Distribution treatment by age');

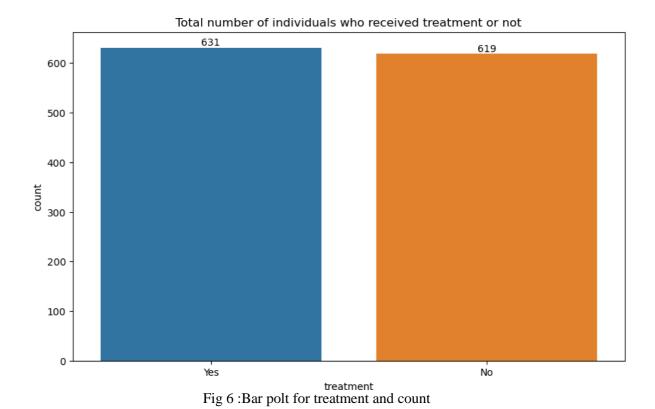
# Output

<Figure size 1000x600 with 0 Axes>



In this diagram x axis is age and y axis is count.

```
#In this plot We can see Total number of individuals who received treatment or not. plt.figure(figsize = (10.6)); treat = sns.countplot(data = data, x = 'treatment'); treat.bar_label(treat.containers[0]); plt.title('Total number of individuals who received treatment or not');
```



In this diagram x axis is treatment and y axis is count.

data.info()

# **Output**

<class 'pandas.core.frame.DataFrame'>

Int64Index: 1250 entries, 0 to 1258

Data columns (total 24 columns):

#	Column	Non-Null Count Dtype
0	Age	1250 non-null int64
1	Gender	1250 non-null object
2	Country	1250 non-null object
3	self_employed	1250 non-null object
4	family_history	1250 non-null object
5	treatment	1250 non-null object
6	work_interfere	1250 non-null object
7	no_employees	1250 non-null object
8	remote_work	1250 non-null object
9	tech_company	1250 non-null object

```
10 benefits
                       1250 non-null object
                         1250 non-null object
11 care_options
12 wellness_program
                            1250 non-null object
13 seek_help
                        1250 non-null object
14 anonymity
                         1250 non-null object
15 leave
                      1250 non-null object
16 mental_health_consequence 1250 non-null object
17 phys_health_consequence 1250 non-null object
18 coworkers
                         1250 non-null object
19 supervisor
                        1250 non-null object
22 mental_vs_physical
                            1250 non-null object
23 obs_consequence
                            1250 non-null object
dtypes: int64(1), object(23)
#Use LabelEncoder to change the Dtypes to 'int'
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
#Make the dataset include all the columns we need to change their dtypes
columns_to_encode = ['Gender', 'Country', 'self_employed', 'family_history', 'treatment',
'work_interfere', 'no_employees',
                 'remote_work', 'tech_company', 'benefits', 'care_options', 'wellness_program',
                 'seek_help', 'anonymity', 'leave', 'mental_health_consequence',
'phys_health_consequence',
                 'coworkers', 'supervisor', 'mental_health_interview', 'phys_health_interview',
                 'mental_vs_physical', 'obs_consequence']
#Write a Loop for fitting LabelEncoder on columns_to_encode
for columns in columns_to_encode:
  data[columns] = le.fit_transform(data[columns])
data.info()
```

<class 'pandas.core.frame.DataFrame'>

Int64Index: 1250 entries, 0 to 1258

Data columns (total 24 columns):

# (	Column	Non-Null Count Dtype			
0	Age	1250 non-null int64			
1 (	Gender	1250 non-null int64			
2	Country	1250 non-null int64			
3 8	self_employed	1250 non-null int64			
4 1	family_history	1250 non-null int64			
5 t	treatment	1250 non-null int64			
6	work_interfere	1250 non-null int64			
7 ı	no_employees	1250 non-null int64			
8 1	remote_work	1250 non-null int64			
9 t	tech_company	1250 non-null int64			
10	benefits	1250 non-null int64			
11	care_options	1250 non-null int64			
12	wellness_program	1250 non-null int64			
13	seek_help	1250 non-null int64			
14	anonymity	1250 non-null int64			
15	leave	1250 non-null int64			
16	mental_health_co	nsequence 1250 non-null int64			
17 phys_health_consequence 1250 non-null int64					
18	coworkers	1250 non-null int64			
19	supervisor	1250 non-null int64			
22	mental_vs_physic	al 1250 non-null int64			
23	obs_consequence	1250 non-null int64			
dtypes: int64(24)					

data.describe()

```
from sklearn.preprocessing import MaxAbsScaler, StandardScaler

data['Age'] = MaxAbsScaler().fit_transform(data[['Age']])

data['Country'] = StandardScaler().fit_transform(data[['Country']])

data['work_interfere'] = StandardScaler().fit_transform(data[['work_interfere']])

data['no_employees'] = StandardScaler().fit_transform(data[['no_employees']])

data['leave'] = StandardScaler().fit_transform(data[['leave']])

data.describe()
```

### Split the data to train and test

```
from sklearn.model_selection import train_test_split

#I wanna work on 'treatment' column.

X = data.drop(columns = ['treatment'])

y = data['treatment']

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.25)

print(X_train.shape, y_train.shape)

print('-'*30)

print('-'*30)
```

## **Output**

```
(937, 23) (937,)
-----
(313, 23) (313,)
```

from sklearn.pipeline import Pipeline

from sklearn.decomposition import PCA

from sklearn.ensemble import RandomForestClassifier as RFC

from sklearn.neighbors import KNeighborsClassifier as KNN

from sklearn.svm import SVC

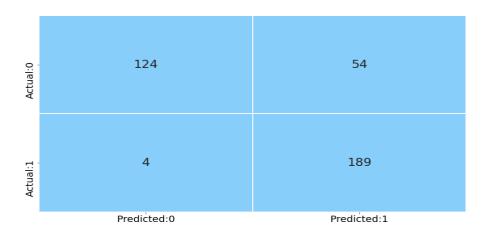
from sklearn.metrics import accuracy\_score

from sklearn.discriminant\_analysis import LinearDiscriminantAnalysis as LDA

# **Support Vector Machine**

A Support Vector Machine (SVM) is a type of supervised machine learning algorithm used for classification and regression tasks. Its primary purpose is to find the optimal hyperplane that best separates data points into different classes or predicts a continuous target variable.

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



 $test\_report = get\_test\_report(svc\_model)$ 

# print the performace 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	0.8	34 3	371	
macro avg	0.87	0.84	0.84	371

```
weighted avg 0.87 0.84 0.84 371
```

# Input

kappa\_value = kappa\_score(svc\_model)

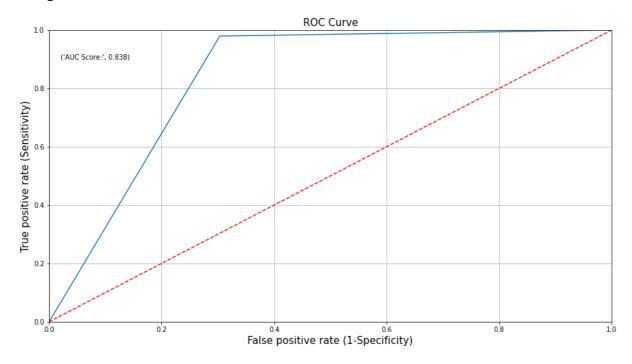
# print the kappa value

print(kappa\_value)

0.6833632537743901

plot\_roc(svc\_model)

## **Output**

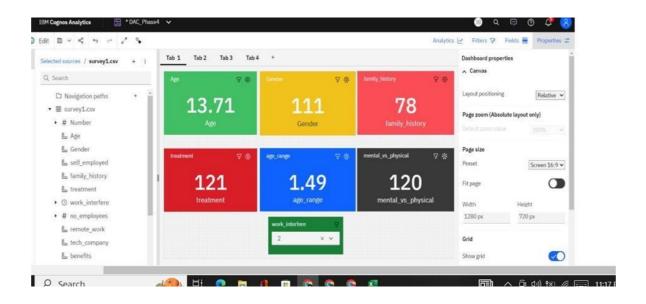


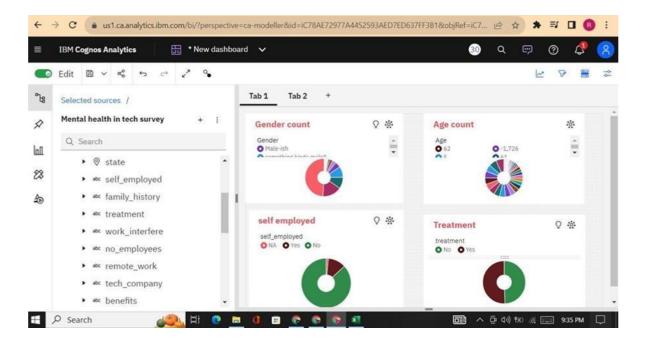
# Accuracy 0.84

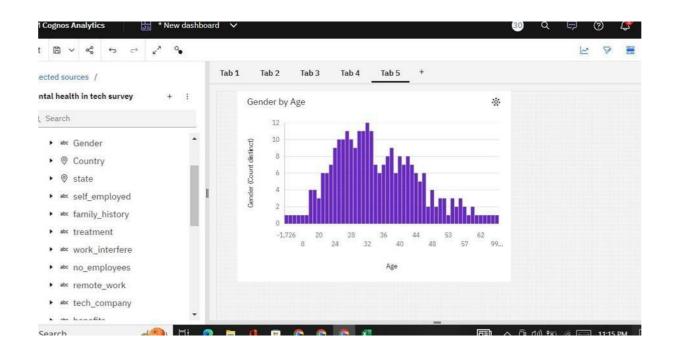
Fig 7: Support Vector Machine

In this diagram x axis is false positive rate and y axis is true positive rate

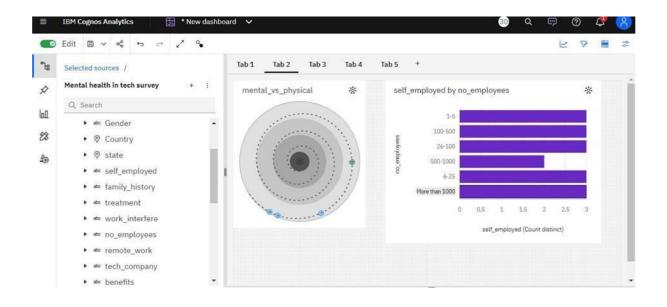
# 5. Visualizations in IBM cognos tool

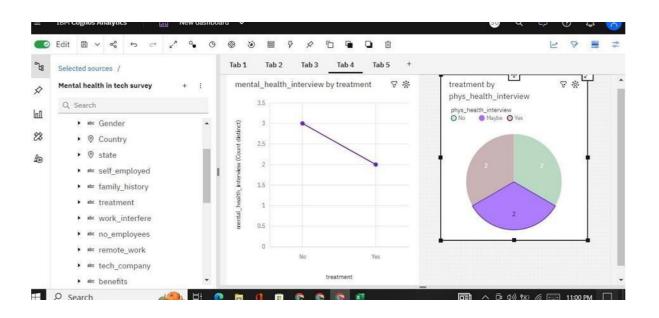


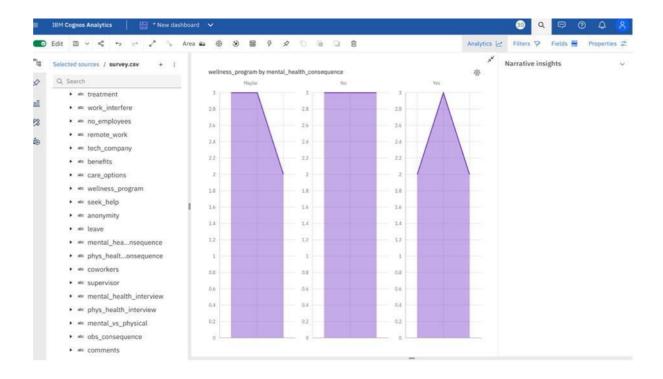






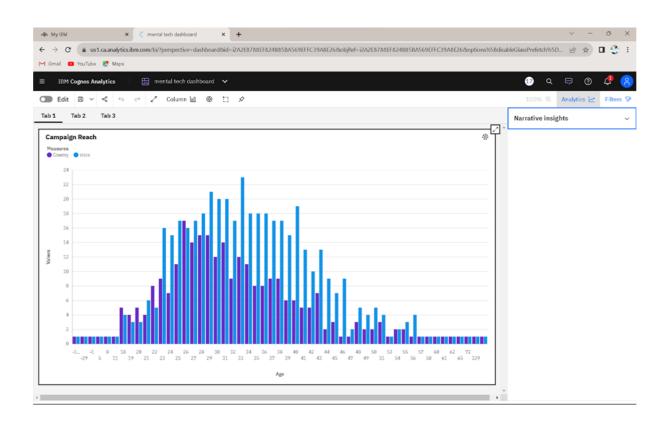




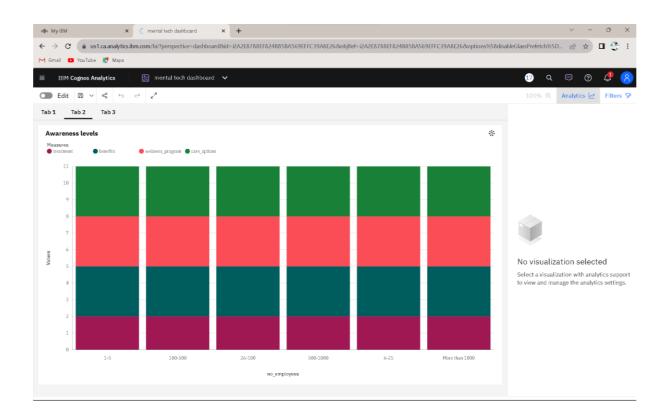


# 6.Dashboard and Report using ibm cognos tool

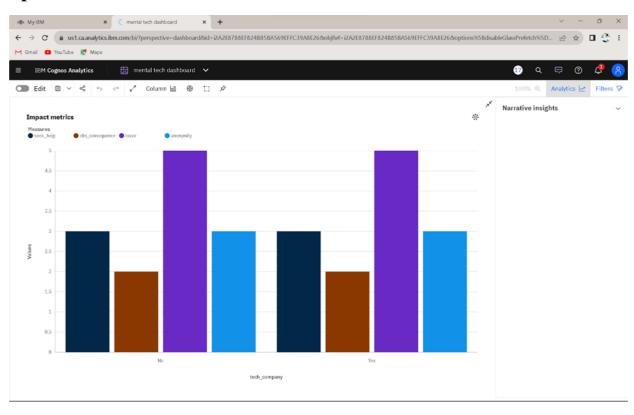
# **Campaign Analysis**



### **Awareness Level**



# **Impact metrics**



## 7.User experience:

- ➤ Data analytics with tools like IBM Cognos can be a game-changer for website owners focused on public health awareness.
- ➤ It provides valuable insights into how people use their websites.
- > These insights can help in several ways: First, by understanding what pages users visit the most and where they leave the site, website owners can make those pages better and fix the weak spots.
- > Second, they can figure out what kinds of information are most interesting to their audience and create more of that.
- Third, they can test different website designs and content to see what works best. Plus, analytics can show if the website is slow or has errors that need fixing.
- ➤ It's like having a map to guide them in making their website more useful and engaging for people interested in public health.

### 8. Conclusion

In conclusion, using data analytics tools like IBM Cognos can be a powerful ally for website owners aiming to enhance the user experience in public health awareness. By understanding how people navigate the website, what content they prefer, and where improvements are needed, owners can tailor their site to be more user-friendly and engaging. It's like having a compass to guide them toward providing vital health information effectively. By continuously analyzing and acting on data-driven insights, they can ensure that their website becomes a valuable resource for promoting public health and well-being, ultimately serving the needs of their audience more efficiently and effectively.