

DEPRESSION, ANXIETY AND STRESS PREDICTION

Collaborators:

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Data info

Q1-Q42 Main questions

- 42 questions with a 4 point rating scale to indicate how often a situation had been true in the past week
- QnA = Stored Response
- QnE = Time taken in milliseconds to answer the question
- QnI = Position on the survey

Recorded durations

- Introelapse = Time spent on introduction/landing page
- Testelapse = Time spent on DASS questions
- Surveyelapse = Time spent on answering the rest of the demographic and survey questions

TIPI = The Ten Item Personality Inventory

- TIPI1: Extraverted-enthusiastic
- TIPI2: Critical-quarrelsome
- TIPI3: Dependable-self_disciplined
- TIPI4: Anxious-easily upset
- TIPI5: Open to new experiences-complex
- TIPI6: Reserved-quiet
- TIPI7: Sympathetic-warm
- TIPI8: Disorganized-careless
- TIPI9: Calm-emotionally_stable
- TIPI10: Conventional-uncreative

-These items were rated "I see myself as:" _____ such that

- 1 = Disagree strongly
- 2 = Disagree moderately
- 3 = Disagree a little
- 4 = Neither agree nor disagree
- 5 = Agree a little
- 6 = Agree moderately
- 7 = Agree strongly

-Score Results

- Extraversion
- Agreeableness
- Conscientiousness
- Emotional Stability
- Openness to Experiences

Gosling, S. D., Rentfrow, P. J., & Swann, W. B., Jr. (2003). A Very Brief Measure of the Big Five Personality Domains. Journal of Research in Personality, 37, 504-528.)

Validity check with list of words Subjects were instructed to check all the words whose definition they knew, where VCL6, VCL9, and VCL12 are not real words and can be used as a validity check.

- VCL1 boat
- VCL2 incoherent
- VCL3 pallid
- VCL4 robot
- VCL5 audible
- VCL6 cuivocal
- VCL7 paucity
- VCL8 epistemology
- VCL9 florted

- VCL10 decide
- VCL11 pastiche
- VCL12 verdid
- VCL13 abysmal
- VCL14 lucid
- VCL15 betray
- VCL16 funny

Demographic and personal information

- education
- marital status
- major
- race
- religion
- age
- sexual orientation

Libraries

```
In [1]: # Basic Libraries
import numpy as np
import pandas as pd
#For regular expressions
import re
# For visuals
from matplotlib import pyplot as plt
import seaborn as sns
```

Data importing and pre-processing

Loading the data

```
In [2]: # Upload data csv file in colab
'''
from google.colab import files
files.upload()
'''
```

```
Out[2]: '\nfrom google.colab import files\nfiles.upload()\n'
```

```
In [3]: # Import the dataset
df=pd.read_csv('data.csv',delimiter='\t')
df.head()
```

```
Out[3]:
```

	Q1A	Q1I	Q1E	Q2A	Q2I	Q2E	Q3A	Q3I	Q3E	Q4A	...	screen	size	uniquen	network	location	hand	religion	orientation	race	voted	married	family	size
0	4	28	3890	4	25	2122	2	16	1944	4	...		1			1	1	12	1	10	2	1		2
1	4	2	8118	1	36	2890	2	35	4777	3	...		2			1	2	7	0	70	2	1		4
2	3	7	5784	1	33	4373	4	41	3242	1	...		2			1	1	4	3	60	1	1		3
3	2	23	5081	3	11	6837	2	37	5521	1	...		2			1	2	4	5	70	2	1		5
4	2	36	3215	2	13	7731	3	5	4156	4	...		2			2	3	10	1	10	2	1		4

5 rows × 172 columns



Characteristics

```
In [4]: print("Dimensions: ", df.shape)
print("Columns:")
print(list(df.columns))
```

```
Dimensions: (39775, 172)
Columns:
['Q1A', 'Q1I', 'Q1E', 'Q2A', 'Q2I', 'Q2E', 'Q3A', 'Q3I', 'Q3E', 'Q4A', 'Q4I', 'Q4E', 'Q5A', 'Q5I', 'Q5E', 'Q6A', 'Q6I', 'Q6E',
'Q7A', 'Q7I', 'Q7E', 'Q8A', 'Q8I', 'Q8E', 'Q9A', 'Q9I', 'Q9E', 'Q10A', 'Q10I', 'Q10E', 'Q11A', 'Q11I', 'Q11E', 'Q12A', 'Q12I',
'Q12E', 'Q13A', 'Q13I', 'Q13E', 'Q14A', 'Q14I', 'Q14E', 'Q15A', 'Q15I', 'Q15E', 'Q16A', 'Q16I', 'Q16E', 'Q17A', 'Q17I', 'Q17E',
'Q18A', 'Q18I', 'Q18E', 'Q19A', 'Q19I', 'Q19E', 'Q20A', 'Q20I', 'Q20E', 'Q21A', 'Q21I', 'Q21E', 'Q22A', 'Q22I', 'Q22E', 'Q23A',
'Q23I', 'Q23E', 'Q24A', 'Q24I', 'Q24E', 'Q25A', 'Q25I', 'Q25E', 'Q26A', 'Q26I', 'Q26E', 'Q27A', 'Q27I', 'Q27E', 'Q28A', 'Q28I',
'Q28E', 'Q29A', 'Q29I', 'Q29E', 'Q30A', 'Q30I', 'Q30E', 'Q31A', 'Q31I', 'Q31E', 'Q32A', 'Q32I', 'Q32E', 'Q33A', 'Q33I', 'Q33E',
'Q34A', 'Q34I', 'Q34E', 'Q35A', 'Q35I', 'Q35E', 'Q36A', 'Q36I', 'Q36E', 'Q37A', 'Q37I', 'Q37E', 'Q38A', 'Q38I', 'Q38E', 'Q39A',
'Q39I', 'Q39E', 'Q40A', 'Q40I', 'Q40E', 'Q41A', 'Q41I', 'Q41E', 'Q42A', 'Q42I', 'Q42E', 'country', 'source', 'introelapse', 'te
stelapse', 'surveyelapse', 'TIPI1', 'TIPI2', 'TIPI3', 'TIPI4', 'TIPI5', 'TIPI6', 'TIPI7', 'TIPI8', 'TIPI9', 'TIPI10', 'VCL1',
'VCL2', 'VCL3', 'VCL4', 'VCL5', 'VCL6', 'VCL7', 'VCL8', 'VCL9', 'VCL10', 'VCL11', 'VCL12', 'VCL13', 'VCL14', 'VCL15', 'VCL16',
'education', 'urban', 'gender', 'engnat', 'age', 'screensize', 'uniquenetworklocation', 'hand', 'religion', 'orientation', 'rac
e', 'voted', 'married', 'familysize', 'major']
```

```
In [5]: # Series data types
series_types = df.dtypes.value_counts()
print("Types of series: ")
print(series_types)

# Series with 'object' data type (strings)
str_series = np.where(df.dtypes == 'object')[0]
str_series = [df.columns[i] for i in str_series]
print("Series with strings: ", str_series)
```

```
Types of series:
int64      170
object      2
dtype: int64
Series with strings: ['country', 'major']
```

There are 170 series with 'int' data types and only 2 series with 'object' types, which is the same as string data type. These 2 series are *country* and *major*.

```
In [6]: # File type
with open("data.csv", "r") as f:
    for i in range(5):
        print(i, "\t", repr(f.readline()))
```

```
0      'Q1A\tQ1I\tQ1E\tQ2A\tQ2I\tQ2E\tQ3A\tQ3I\tQ3E\tQ4A\tQ4I\tQ4E\tQ5A\tQ5I\tQ5E\tQ6A\tQ6I\tQ6E\tQ7A\tQ7I\tQ7E\tQ8A\tQ8I\tQ8
E\tQ9A\tQ9I\tQ9E\tQ10A\tQ10I\tQ10E\tQ11A\tQ11I\tQ11E\tQ12A\tQ12I\tQ12E\tQ13A\tQ13I\tQ13E\tQ14A\tQ14I\tQ14E\tQ15A\tQ15I\tQ15E\tQ
16A\tQ16I\tQ16E\tQ17A\tQ17I\tQ17E\tQ18A\tQ18I\tQ18E\tQ19A\tQ19I\tQ19E\tQ20A\tQ20I\tQ20E\tQ21A\tQ21I\tQ21E\tQ22A\tQ22I\tQ22E\tQ2
3A\tQ23I\tQ23E\tQ24A\tQ24I\tQ24E\tQ25A\tQ25I\tQ25E\tQ26A\tQ26I\tQ26E\tQ27A\tQ27I\tQ27E\tQ28A\tQ28I\tQ28E\tQ29A\tQ29I\tQ29E\tQ30
A\tQ30I\tQ30E\tQ31A\tQ31I\tQ31E\tQ32A\tQ32I\tQ32E\tQ33A\tQ33I\tQ33E\tQ34A\tQ34I\tQ34E\tQ35A\tQ35I\tQ35E\tQ36A\tQ36I\tQ36E\tQ37A
\tQ37I\tQ37E\tQ38A\tQ38I\tQ38E\tQ39A\tQ39I\tQ39E\tQ40A\tQ40I\tQ40E\tQ41A\tQ41I\tQ41E\tQ42A\tQ42I\tQ42E\tcountry\tsource\tintroe
lapse\ttestelapse\tsurveyelapse\tTIPI1\tTIPI2\tTIPI3\tTIPI4\tTIPI5\tTIPI6\tTIPI7\tTIPI8\tTIPI9\tTIPI10\tVCL1\tVCL2\tVCL3\tVCL4
\tVCL5\tVCL6\tVCL7\tVCL8\tVCL9\tVCL10\tVCL11\tVCL12\tVCL13\tVCL14\tVCL15\tVCL16\teducation\turban\tgender\tengnat\tage\tscreens
ize\tuniquenetworklocation\tthand\treligion\torientation\ttrace\tvoted\tmarried\tfamilysize\tmajor'n'
1      '4\t28\t3890\t4\t25\t2122\t2\t16\t1944\t4\t8\t2044\t4\t34\t2153\t4\t33\t2416\t4\t10\t2818\t4\t13\t2259\t2\t21\t5541\t1
\t38\t4441\t4\t31\t2451\t4\t24\t3325\t4\t14\t1416\t4\t37\t5021\t4\t27\t2342\t4\t39\t2480\t3\t6\t2476\t4\t35\t1627\t3\t17\t9050
\t3\t30\t7001\t1\t11\t4719\t4\t20\t2984\t4\t36\t1313\t4\t42\t2444\t4\t1\t9880\t4\t2\t4695\t4\t5\t1677\t3\t4\t6723\t4\t3\t5953\t
2\t26\t8062\t4\t12\t5560\t4\t7\t3032\t2\t29\t3316\t3\t40\t3563\t4\t23\t5594\t4\t41\t1477\t1\t18\t3885\t2\t9\t5265\t4\t19\t1892
\t3\t22\t4228\t4\t32\t1574\t4\t15\t2969\tIN\t2\t19\t167\t166\t1\t5\t7\t7\t7\t7\t5\t1\t1\t1\t0\t0\t1\t1\t0\t1\t0\t0\t1\t0\t0\t0
\t0\t1\t1\t1\t2\t3\t2\t2\t16\t1\t1\t1\t1\t12\t1\t10\t2\t1\t2\t1\t2\t1'n'
2      '4\t2\t8118\t1\t136\t2890\t2\t35\t4777\t3\t28\t3090\t4\t10\t5078\t4\t40\t2790\t3\t18\t3408\t4\t1\t8342\t3\t37\t916\t2\t2
32\t1537\t2\t21\t3926\t2\t25\t3691\t4\t26\t2004\t4\t4\t8888\t3\t27\t4109\t3\t19\t4058\t4\t12\t3692\t2\t6\t3373\t1\t23\t6015\t1
\t16\t3023\t2\t22\t2670\t3\t3\t5727\t1\t39\t3641\t2\t23\t2670\t2\t7\t7649\t3\t11\t2537\t3\t5\t2907\t4\t9\t1685\t3\t41\t4726\t3
\t17\t6063\t2\t20\t3307\t3\t14\t4995\t3\t38\t2505\t2\t34\t2540\t2\t31\t4359\t3\t15\t3925\t4\t13\t4609\t2\t30\t3755\t2\t42\t2323
\t1\t24\t5713\t2\t8\t1334\t2\t29\t5562\tUS\t2\t1\t193\t186\t6\t5\t4\t7\t5\t4\t7\t7\t1\t5\t1\t1\t1\t0\t1\t1\t0\t0\t0\t0\t1\t0\t0\t0
\t1\t1\t1\t1\t2\t3\t2\t1\t16\t2\t1\t2\t7\t0\t70\t2\t1\t2\t1\t4\t4\t1'n'
3      '3\t7\t5784\t1\t33\t4373\t4\t41\t3242\t1\t13\t6470\t4\t11\t3927\t3\t9\t3704\t1\t17\t4550\t3\t5\t3021\t2\t32\t5864\t4\t1
21\t3722\t2\t10\t3424\t1\t36\t3236\t4\t23\t2489\t1\t34\t7290\t4\t12\t6587\t4\t22\t3627\t4\t38\t2905\t2\t18\t2998\t2\t8\t10233\t
1\t16\t4258\t4\t28\t2888\t3\t4\t59592\t2\t3\t11732\t4\t2\t8834\t2\t29\t7358\t1\t30\t4928\t2\t15\t3036\t1\t19\t4127\t2\t37\t3934
\t2\t26\t10782\t4\t1\t8273\t3\t39\t3501\t1\t27\t3824\t4\t25\t2141\t3\t6\t17461\t4\t24\t1557\t4\t40\t4446\t4\t42\t1883\t2\t35\t5
790\t2\t14\t4432\t1\t20\t2203\t4\t31\t5768\tPL\t2\t5\t271\t122\t2\t5\t2\t2\t5\t6\t5\t5\t3\t2\t1\t0\t0\t1\t1\t1\t0\t0\t0\t0\t0\t0\t1\t
0\t0\t1\t1\t1\t1\t2\t3\t2\t2\t17\t2\t1\t1\t4\t3\t60\t1\t1\t3\t1'n'
4      '2\t23\t5081\t3\t11\t6837\t2\t37\t5521\t1\t27\t4556\t3\t28\t3269\t3\t26\t3231\t4\t2\t7138\t2\t19\t3079\t3\t31\t9650\t3
\t17\t4179\t2\t5\t5928\t1\t21\t2838\t1\t20\t2560\t4\t29\t5139\t2\t22\t3597\t2\t35\t3336\t3\t10\t4506\t1\t14\t2695\t1\t25\t8128
\t2\t15\t3125\t1\t6\t4061\t1\t40\t4272\t1\t12\t4029\t1\t9\t5630\t1\t18\t30631\t2\t24\t9870\t4\t4\t2411\t1\t16\t9478\t3\t1\t7618
\t3\t32\t12639\t3\t34\t5378\t1\t41\t8923\t2\t38\t2977\t4\t4\t3\t5620\t1\t7\t16760\t1\t8\t6427\t2\t39\t3760\t1\t13\t4112\t3\t42\t27
69\t4\t33\t4432\t4\t30\t3643\t2\t36\t3698\tUS\t2\t3\t261\t336\t1\t1\t7\t4\t6\t4\t6\t1\t6\t1\t1\t0\t0\t1\t1\t0\t0\t0\t0\t1\t0\t0\t0
\t0\t1\t1\t1\t1\t1\t3\t2\t1\t13\t2\t1\t2\t4\t5\t70\t2\t1\t5\tbiology'n'
```

CSV file, with fields delimited by 't' (tabs) and records delimited by 'n' (new line)

Cleaning the data

```
In [7]: # Removing the people that took answers too quickly or too slowly
df = df[ df['testelapse'] <= df['testelapse'].quantile(0.975)]
df = df[ df['testelapse'] >= df['testelapse'].quantile(0.025)]
df = df[ df['surveyelapse'] <= df['surveyelapse'].quantile(0.975)]
df = df[ df['surveyelapse'] >= df['surveyelapse'].quantile(0.025)]
```

```
In [8]: # Replacing extreme ages
median = df.loc[df['age'] <=80, 'age'].median()
df.loc[df.age > 80, 'age'] = np.nan
df['age'].fillna(median,inplace=True)
```

```
In [9]: # Removing unnecessary columns involving position (QnI) and time (QnE)
position = [i for i in df.iloc[:, 0:126] if 'I' in i]
time = [i for i in df.iloc[:, 0:126] if 'E' in i]
df=df.drop(position, axis=1, errors='ignore')
df=df.drop(time, axis=1, errors='ignore')
# Remove introelapse, testelapse and surveyelapse series
df=df.drop(columns= ["introelapse", "testelapse", "source", "surveyelapse"], axis = 1, errors="ignore")
# Remove engnat, screensize, uniquenetworklocation, hand and voted
df=df.drop(columns= ["engnat", "screensize", "uniquenetworklocation", "hand", "voted"], axis = 1, errors="ignore")
```

```
In [10]: # Removing VCL series
df=df.drop(df.iloc[:,53:69],axis=1)
```

```
In [11]: # Replace the 0's from the categorical variables with 3
df=df.replace(to_replace=0,value=3)
```

```
In [12]: print('New dimensions: ', df.shape)
print("Columns")
print(df.columns)
```

```
New dimensions: (36016, 63)
Columns
Index(['Q1A', 'Q2A', 'Q3A', 'Q4A', 'Q5A', 'Q6A', 'Q7A', 'Q8A', 'Q9A', 'Q10A',
      'Q11A', 'Q12A', 'Q13A', 'Q14A', 'Q15A', 'Q16A', 'Q17A', 'Q18A', 'Q19A',
      'Q20A', 'Q21A', 'Q22A', 'Q23A', 'Q24A', 'Q25A', 'Q26A', 'Q27A', 'Q28A',
      'Q29A', 'Q30A', 'Q31A', 'Q32A', 'Q33A', 'Q34A', 'Q35A', 'Q36A', 'Q37A',
      'Q38A', 'Q39A', 'Q40A', 'Q41A', 'Q42A', 'country', 'TIPI1', 'TIPI2',
      'TIPI3', 'TIPI4', 'TIPI5', 'TIPI6', 'TIPI7', 'TIPI8', 'TIPI9', 'TIPI10',
      'education', 'urban', 'gender', 'age', 'religion', 'orientation',
      'race', 'married', 'familysize', 'major'],
      dtype='object')
```

Canonicalization

It is important to apply a process of canonicalization on the 'major' column since there are many different names that are representing the same type of major, which can represent some problems on the visualization and interpretation that will take place in order to have further understanding of the data.

```
In [13]: # Replace Nan and other values with "No degree" or appropriate string in major feature
df['major'] = df['major'].replace(np.nan, "No Degree")
df['major'] = df['major'].replace(".", "No Degree")
df['major'] = df['major'].replace(",", "No Degree")
df['major'] = df['major'].replace("no", "No Degree")
df['major'] = df['major'].replace("No", "No Degree")
df['major'] = df['major'].replace("a level", "No Degree")
df['major'] = df['major'].replace(" ", "No Degree")
df['major'] = df['major'].replace("None", "No Degree")
df['major'] = df['major'].replace("none", "No Degree")
df['major'] = df['major'].replace(["--", "i do not know ", "???", "-nil-"], "No Degree")
df['major'] = df['major'].replace("-", "No Degree")
df['major'] = df['major'].replace(["thi&#769;t k&#769; ho&#803;a", "&#22810;&#23186;&#39636;&#35373;&#35336;"], "No Degree")
df['major'] = df['major'].replace("t", "IT")
df['major'] = df['major'].replace("yes", "No Degree")
df['major'] = df['major'].replace("undecided", "No Degree")
df['major'] = df['major'].replace("undeclared", "No Degree")
df['major'] = df['major'].replace(["cs", "Computer Programming", "computer sciece", "ca", "game dev", "comp science"], "Computer Science")
df['major'] = df['major'].replace(["it", "college, i.t", "information technology"], "IT")
```

```
In [14]: # Lowercase all the string values
df['major'] = df['major'].str.lower()
```

```
In [15]: # Let's find the most important majors and make sure that they have a standard str
maj = df['major'].value_counts()
print(maj)
```

```
...
mathematics, business, economics    1
commercial art                      1
anthropology and english             1
mechanical engineering...           1
public relation or administrations  1
Name: major, Length: 3804, dtype: int64
```

```
In [16]: # Engineering majors
mask1 = df['major'].str.contains('engineering')
before = [x for x in df[mask1]['major'][:10]]
print(before)
# Engineering majors after canonicalization
df.loc[mask1, 'major'] = 'engineering'
after = [x for x in df[mask1]['major'][:10]]
print(after)
```

[illegible]

```
In [17]: # Psychology majors
mask2 = ( (df['major'].str.startswith('ps')) | (df['major'].str.contains('psychology') == True) | (df['major'].str.contains('behavioral psychology')) )
before = [x for x in df[mask2]['major'][:10]]
print(before)
# Psychology majors after canonicalization
df.loc[mask2, 'major'] = 'psychology'
after = [x for x in df[mask2]['major'][:10]]
print(after)
```

[illegible]

```
In [18]: # English majors
mask3 = ( df['major'].str.startswith("engl") |
          (df['major'].str.contains('lis') == True)
          )
before = [x for x in df[mask3]['major'][:10]]
print(before)
# English majors after canonicalization
df.loc[mask3, 'major'] = 'english'
after = [x for x in df[mask3]['major'][:10]]
print(after)
```

[illegible]

```
In [19]: # Business majors
mask4 = ( (df['major'].str.startswith("b") &
          df['major'].str.endswith("s") &
          (df['major'].str.contains(' ') == False) &
          (df['major'].str.startswith("bio")==False) ) | (df['major'].str.contains('investment') == True) |
          (df['major'].str.contains('admin') == True) | (df['major'].str.contains('business') == True) | (df['major'].str.contains('business'))
before =[x for x in df[mask4]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# Business majors after canonicalization
df.loc[mask4, 'major'] = 'business'
after = [x for x in df[mask4]['major'][:10]
print(after)
```

[illegible]

```
In [20]: #medicine major
mask10 = ( (df['major'].str.contains('medic') == True) | (df['major'].str.contains('nursing') == True)
           | (df['major'].str.contains('dent') == True) | (df['major'].str.contains('health') == True)
           )
before = [x for x in df[mask10]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# medicine majors after canonicalization
df.loc[mask10, 'major'] = 'medicine'
after = [x for x in df[mask10]['major'][:10]
print(after)

['medical technology', 'dentistry', 'nursing', 'medicine', 'medical', 'allied health', 'ba (philosophy) & bmedicine', 'pre-medi
cine', 'nursing', 'medicine ']
['medicine', 'medicine', 'medicine', 'medicine', 'medicine', 'medicine', 'medicine', 'medicine', 'medicine', 'medicine']
```

```
In [21]: #arts major
mask5 = ( (df['major'].str.contains('art') == True) | (df['major'].str.contains('creative') == True) |
           (df['major'].str.contains('illustration') == True) | (df['major'].str.contains('recreation') == True) |
           (df['major'].str.contains('design') == True) | (df['major'].str.contains('music') == True) | (df['major'].str.contains('
           )
before = [x for x in df[mask5]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# arts majors after canonicalization
df.loc[mask5, 'major'] = 'arts'
after = [x for x in df[mask5]['major'][:10]
print(after)

['music', 'art', 'art history', 'art theory', 'italian/history of art', 'animation', 'fine arts', 'culinary arts', 'fine art',
'art and design']
['arts', 'arts', 'arts', 'arts', 'arts', 'arts', 'arts', 'arts', 'arts', 'arts']
```

```
In [22]: #film major
mask6 = ( (df['major'].str.contains('film') == True) | (df['major'].str.contains('cinema') == True) |
           (df['major'].str.contains('theatre') == True) | (df['major'].str.contains('video') == True) )
before = [x for x in df[mask6]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# film majors after canonicalization
df.loc[mask6, 'major'] = 'film'
after = [x for x in df[mask6]['major'][:10]
print(after)

['theatre', 'theatre', 'film', 'film and video production', 'film', 'theatre', 'film studies', 'theatre', 'theatre', 'cinema']
['film', 'film', 'film', 'film', 'film', 'film', 'film', 'film', 'film', 'film']
```

```
In [23]: #law major
mask7 = ( (df['major'].str.contains('law') == True) | (df['major'].str.contains('llb') == True) |
           (df['major'].str.contains('legal') == True) | (df['major'].str.contains('legislation') == True) )
before = [x for x in df[mask7]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# law majors after canonicalization
df.loc[mask7, 'major'] = 'law'
after = [x for x in df[mask7]['major'][:10]
print(after)

['law', 'law', 'law', 'law', 'law', 'law', 'law', 'law', 'law', 'law']
['law', 'law', 'law', 'law', 'law', 'law', 'law', 'law', 'law', 'law']
```

```
In [24]: #accounting and finance major
mask8 = ( (df['major'].str.contains('account') == True) | (df['major'].str.contains('finance') == True) |
           (df['major'].str.contains('econom') == True) | (df['major'].str.contains('bank') == True) | (df['major'].str.contains('c
before = [x for x in df[mask8]['major'][:10] # executing this line, we observe some typos, we'll replace them
print(before)
# accounting majors after canonicalization
df.loc[mask8, 'major'] = 'accounting and finance'
after = [x for x in df[mask8]['major'][:10]
print(after)

['accounting', 'accounting', 'accounting', 'finance', 'economics', 'economy', 'finance', 'account ing', 'accounting', 'accounti
ng']
['accounting and finance', 'accounting and finance', 'accounting and finance', 'accounting and finance', 'accounting and financ
e', 'accounting and finance', 'accounting and finance', 'accounting and finance', 'accounting and finance', 'accounting and fin
ance']
```

```
In [25]: #science major
mask9 = ( (df['major'] == 'science') | (df['major'] == 'chemi') |
          (df['major'] == 'physics') | (df['major'].str.contains('bio') == True) )
before = [x for x in df[mask9]['major']] # executing this line, we observe some typos, we'll replace them
#print(before)
# science majors after canonicalization
df.loc[mask9, 'major'] = 'science'
after = [x for x in df[mask9]['major']]
#print(after)
```

```
In [26]: maj = df['major'].value_counts()
maj.head(60)
```

```
Out[26]: no degree                10130
engineering                3296
business                   2487
accounting and finance     2145
medicine                   1846
psychology                 1795
english                   1568
science                   1309
arts                      1238
law                       716
computer science          493
education                 322
architecture              248
it                        239
pharmacy                  191
chemistry                 190
mathematics               188
communication             143
marketing                 132
history                   131
tourism                   130
film                     112
information technology     102
political science         93
sociology                 90
science                   87
computer science          80
human resource            75
education                 71
architecture              67
culinary                  67
multimedia                66
mass communication        66
social work               61
language                  56
science computer          56
communication             55
math                      54
social science            54
physiotherapy             51
computer                  50
international relations   46
tesl                      43
mathematics               43
human resources           42
chemistry                 41
food science              41
literature                41
hospitality               41
statistics                41
geology                   40
quantity surveying        39
early childhood education 38
teaching                  38
communications            38
pharmacy                  38
philosophy                35
information technology     35
islamic studies           34
applied science           33
Name: major, dtype: int64
```

Data analysis and visualization

Creation of age bins to classify data

The new 'AgeGroup' feature is created to convert the continuous feature of 'Age' into a categorical variable, which allows us to visualize its behavior and calculate its correlation with other features. The chosen categories are the following:

- 0) < 20
- 1) 20-24
- 2) 25-29
- 3) 30-34
- 4) 35-39
- 5) 40-49
- 6) 50-59
- 7) > 60

In [27]: ageGroup = ['< 20', '20-24', '25-29', '30-34', '35-39', '40-49', '50-59', '> 60']

```
def agrp(ag):
    if ag < 20:
        return 0
    elif 20 <= ag <= 24:
        return 1
    elif 25 <= ag <= 29:
        return 2
    elif 30 <= ag <= 34:
        return 3
    elif 35 <= ag <= 39:
        return 4
    elif 40 <= ag <= 49:
        return 5
    elif 50 <= ag <= 59:
        return 6
    else:
        return 7

df['AgeGroup'] = df['age'].apply(agrp)
df.head(3)
```

Out[27]:

	Q1A	Q2A	Q3A	Q4A	Q5A	Q6A	Q7A	Q8A	Q9A	Q10A	...	urban	gender	age	religion	orientation	race	married	familysize	major	AgeGroup
0	4	4	2	4	4	4	4	4	2	1	...	3	2	16.0	12	1	10	1	2	no degree	0
1	4	1	2	3	4	4	3	4	3	2	...	3	2	16.0	7	3	70	1	4	no degree	0
2	3	1	4	1	4	3	1	3	2	4	...	3	2	17.0	4	3	60	1	3	no degree	0

3 rows × 64 columns

DAS Score calculation

We start by filtering the df to separate the 42 questions about mental health from the demographics info

In [28]: qA = df.iloc[:,42]
qA.head(3)

Out[28]:

	Q1A	Q2A	Q3A	Q4A	Q5A	Q6A	Q7A	Q8A	Q9A	Q10A	...	Q33A	Q34A	Q35A	Q36A	Q37A	Q38A	Q39A	Q40A	Q41A	Q42A
0	4	4	2	4	4	4	4	4	2	1	...	2	3	4	4	1	2	4	3	4	4
1	4	1	2	3	4	4	3	4	3	2	...	3	2	2	3	4	2	2	1	2	2
2	3	1	4	1	4	3	1	3	2	4	...	1	4	3	4	4	4	2	2	1	4

3 rows × 42 columns


```
In [29]: demographic = df.iloc[:,43:]
demographic.head(3)
```

```
Out[29]:
```

	TIP1	TIP2	TIP3	TIP4	TIP5	TIP6	TIP7	TIP8	TIP9	TIP10	...	urban	gender	age	religion	orientation	race	married	familysize	major	AgeGroup
0	1	5	7	7	7	7	7	5	1	1	...	3	2	16.0	12	1	10	1	2	no degree	C
1	6	5	4	7	5	4	7	7	1	5	...	3	2	16.0	7	3	70	1	4	no degree	C
2	2	5	2	2	5	6	5	5	3	2	...	3	2	17.0	4	3	60	1	3	no degree	C

3 rows × 21 columns

Each question is scored on a 4-point scale ranging from 0 ("Did not apply to me at all") to 3 ("Applied to me very much, or most of the time"). Thus, we must subtract one to every response, since in the df it is in a scale of 1-4 instead of 0-3.

```
In [30]: qA = qA.subtract(1,axis=1)
qA.head(1)
```

```
Out[30]:
```

	Q1A	Q2A	Q3A	Q4A	Q5A	Q6A	Q7A	Q8A	Q9A	Q10A	...	Q33A	Q34A	Q35A	Q36A	Q37A	Q38A	Q39A	Q40A	Q41A	Q42A
0	3	3	1	3	3	3	3	3	1	0	...	1	2	3	3	0	1	3	2	3	3

1 rows × 42 columns

Now we can assign each question to its corresponding mental illness. The scoring keys are based on the following [DASS-42-Scoring](https://neurocogsystem.com/wp-content/uploads/2021/02/DASS-42-Scoring.pdf) (<https://neurocogsystem.com/wp-content/uploads/2021/02/DASS-42-Scoring.pdf>).

```
In [31]: keys = {"Depression": [3, 5, 10, 13, 16, 17, 21, 24, 26, 31, 34, 37, 38, 42],
               "Anxiety": [2, 4, 7, 9, 15, 19, 20, 23, 25, 28, 30, 36, 40, 41],
               "Stress": [1, 6, 8, 11, 12, 14, 18, 22, 27, 29, 32, 33, 35, 39]}
D, A, S = ([] for i in range(3))
for i in keys["Depression"]:
    D.append('Q'+str(i)+'A')

for i in keys["Anxiety"]:
    A.append('Q'+str(i)+'A')

for i in keys["Stress"]:
    S.append('Q'+str(i)+'A')

depression= qA.filter(D)
anxiety = qA.filter(A)
stress = qA.filter(S)
```

The final scores for each condition are calculated by the sum of their associated questions.

```
In [32]: # We create another column on the three new df with the "Total Score" of the condition
depression["score"] = depression.sum(axis = 1)
anxiety["score"] = anxiety.sum(axis = 1)
stress["score"] = stress.sum(axis = 1)
```

Now, the new dataframes will be joined with the rest of the demographic information to start comparing them.

```
In [33]: Depression = pd.merge(depression,demographic,how='left',left_index=True,right_index=True)
D1 = Depression.pop('score')
Depression['score'] = D1
Depression.head(1)
```

```
Out[33]:
```

	Q3A	Q5A	Q10A	Q13A	Q16A	Q17A	Q21A	Q24A	Q26A	Q31A	...	gender	age	religion	orientation	race	married	familysize	major	AgeGroup	score
0	1	3	0	3	3	2	0	3	3	3	...	2	16.0	12	1	10	1	2	no degree	0	2

1 rows × 36 columns

```
In [34]: Anxiety = pd.merge(anxiety,demographic,how='left',left_index=True,right_index=True)
A1 = Anxiety.pop('score')
Anxiety['score'] = A1
Anxiety.head(1)
```

```
Out[34]:
```

	Q2A	Q4A	Q7A	Q9A	Q15A	Q19A	Q20A	Q23A	Q25A	Q28A	...	gender	age	religion	orientation	race	married	familysize	major	AgeGroup	score	
0	3	3	3	1	3	2	2	3	3	2	...	2	16.0	12		1	10	1	2	no degree	0	34

1 rows × 36 columns

```
In [35]: Stress = pd.merge(stress,demographic,how='left',left_index=True,right_index=True)
S1 = Stress.pop('score')
Stress['score'] = S1
Stress.head(1)
```

```
Out[35]:
```

	Q1A	Q6A	Q8A	Q11A	Q12A	Q14A	Q18A	Q22A	Q27A	Q29A	...	gender	age	religion	orientation	race	married	familysize	major	AgeGroup	score	
0	3	3	3	3	3	3	3	3	3	3	...	2	16.0	12		1	10	1	2	no degree	0	40

1 rows × 36 columns

To interpret the resulting scores we use the following criteria:

	Depression (D)	Anxiety (A)	Stress (S)
Normal	0-9	0-7	0-14
Mild	10-13	8-9	15-18
Moderate	14-20	10-14	19-25
Severe	21-27	15-19	26-33
Extremely Severe	28+	20+	34+

```
In [36]: # Depression results
def resultD(d):
    if d < 10:
        return "Normal"
    elif d >= 10 and d <= 13:
        return "Mild"
    elif d >= 14 and d <= 20:
        return "Moderate"
    elif d >= 21 and d <= 27:
        return "Severe"
    elif d > 27:
        return "Extremely Severe"

Depression['intensity']=Depression['score'].apply(resultD)
Depression['intensity'].value_counts()
```

```
Out[36]:
```

Extremely Severe	12085
Normal	8096
Moderate	6483
Severe	5887
Mild	3465

Name: intensity, dtype: int64

```
Out[37]:
```

Extremely Severe	12626
Normal	8848
Moderate	6419
Severe	5584
Mild	2539

```
Name: intensity, dtype: int64
```

```
Out[38]: Normal      10767
Moderate      7958
Severe        7813
Extremely Severe 4962
Mild          4516
Name: intensity, dtype: int64
```

Visualization of number of answers on each condition level

```
In [39]: level_order = ["Normal", "Mild", "Moderate", "Severe", "Extremely Severe"]
fig, axes = plt.subplots(1,3,figsize=(22, 4), sharey=True)
sns.set_theme(style="whitegrid")
fig.suptitle('DAS count of the Score/Interpretation for each condition')

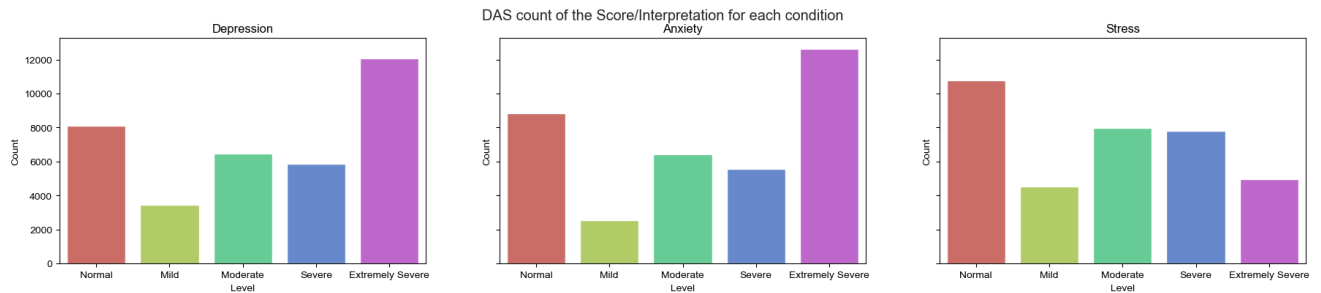
# Visualization of the number of answers on each Level of depression
sns.countplot(ax=axes[0], data=Depression, x='intensity', palette="hls", order = level_order)
#Title
axes[0].set_title('Depression')

# Visualization of the number of answers on each Level of anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='intensity', palette="hls", order = level_order)
#Title
axes[1].set_title('Anxiety')

# Visualization of the number of answers on each Level of stress
sns.countplot(ax=axes[2], data=Stress, x='intensity', palette="hls", order = level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Level', ylabel='Count') for i in range(0,3)]
fig.show()
```

C:\Users\sandr\AppData\Local\Temp\ipykernel_13864\2338039264.py:24: UserWarning: Matplotlib is currently using module://matplotlib_inline.backend_inline, which is a non-GUI backend, so cannot show the figure.
fig.show()



Correlations between categorical features

Most of the features on the three main dataframes are categorical, so they are ready to calculate the correlations between them. However, it is necessary to transform the resulting level of each condition to indexes from 0-4, where:

- 0) Normal
- 1) Mild
- 2) Moderate
- 3) Severe
- 4) Extremely Severe

```
In [40]: def catResults(r):
        if r == 'Normal':
            return 0
        elif r == 'Mild':
            return 1
        elif r == 'Moderate':
            return 2
        elif r == 'Severe':
            return 3
        elif r == 'Extremely Severe':
            return 4

        Depression['cIntensity'] = Depression['intensity'].apply(catResults)
        Anxiety['cIntensity'] = Anxiety['intensity'].apply(catResults)
        Stress['cIntensity'] = Stress['intensity'].apply(catResults)

        Anxiety.head(1)
```

Out[40]:

	Q2A	Q4A	Q7A	Q9A	Q15A	Q19A	Q20A	Q23A	Q25A	Q28A	...	religion	orientation	race	married	familysize	major	AgeGroup	score	intensity	cln
0	3	3	3	1	3	2	2	3	3	2	...	12		1	10	1	2	no degree	0	34	Extremely Severe

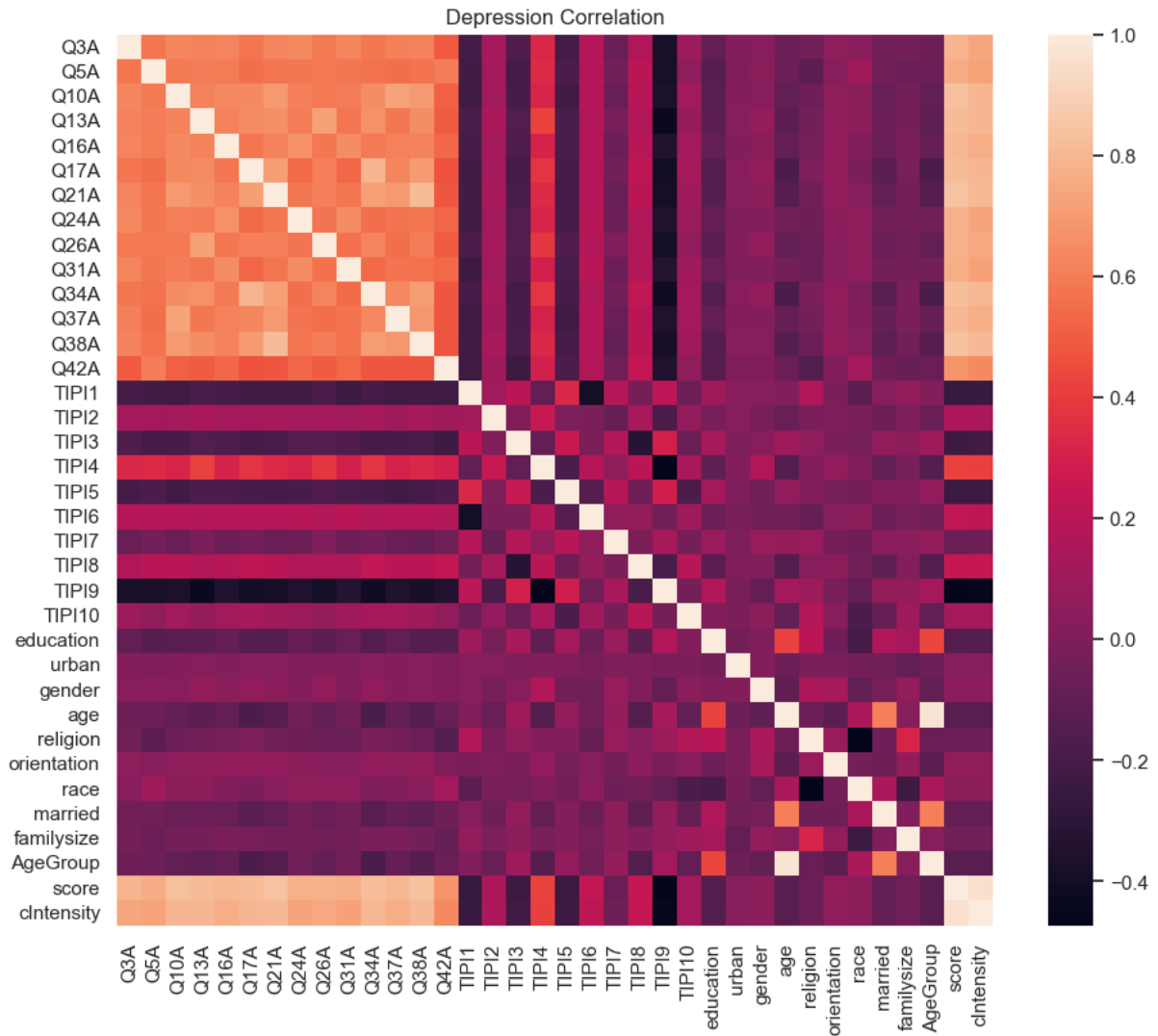
1 rows × 38 columns

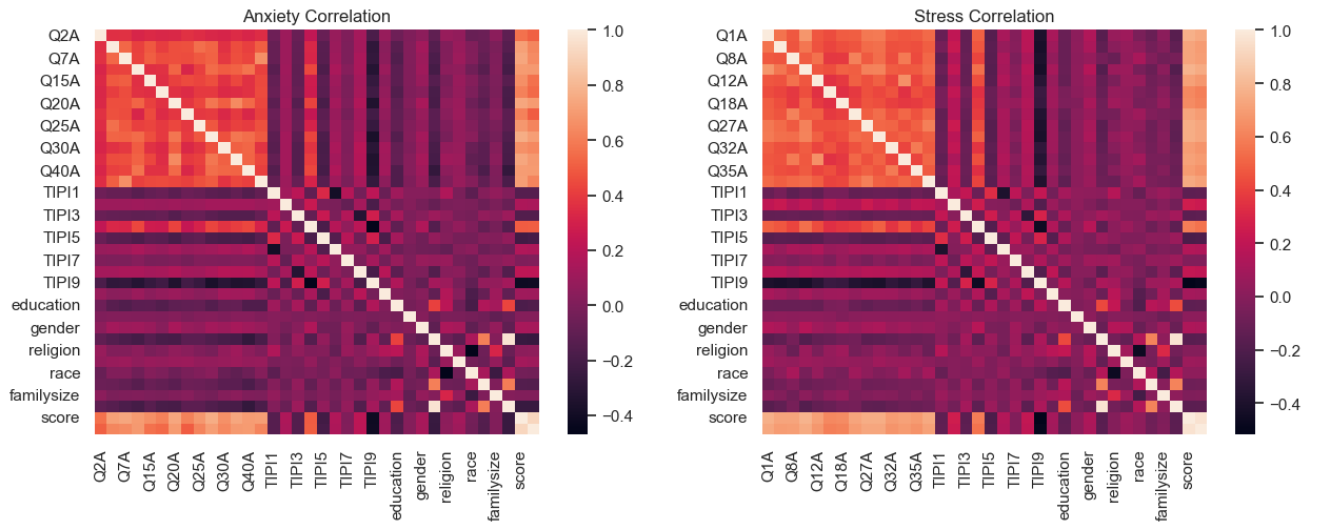
```
In [41]: # Depression df correlation heatmap
plt_1 = plt.figure(figsize=(11, 9))
sns.heatmap(Depression.corr())
plt.title('Depression Correlation')

fig, axes = plt.subplots(1,2,figsize=(15, 5))
# Anxiety df correlation heatmap
sns.heatmap(Anxiety.corr(), ax=axes[0])
axes[0].set_title('Anxiety Correlation')

# Stress df correlation heatmap
sns.heatmap(Stress.corr(), ax=axes[1])
axes[1].set_title('Stress Correlation')

plt.show()
```





It can be seen that TIPI9 (calmness and emotional behavior) has a strong negative correlation with the score, because the most emotionally unstable a person is, the higher their chances of getting a high score.

Now, we only stay with the important demographic information in each dataframe to analyze its behavior.

```
In [42]: Depression0 = Depression.copy()
Depression = Depression.drop(Depression.iloc[:, 0:14], axis=1, errors="ignore")
Depression.head(2)
```

```
Out[42]:
```

	TIPI1	TIPI2	TIPI3	TIPI4	TIPI5	TIPI6	TIPI7	TIPI8	TIPI9	TIPI10	...	religion	orientation	race	married	familysize	major	AgeGroup	score	intensity	cl
0	1	5	7	7	7	7	7	5	1	1	...	12	1	10	1	2	no degree	0	27	Severe	
1	6	5	4	7	5	4	7	7	1	5	...	7	3	70	1	4	no degree	0	24	Severe	

2 rows × 24 columns

```
In [43]: Anxiety0 = Anxiety.copy()
Anxiety = Anxiety.drop(Anxiety.iloc[:, 0:14], axis=1, errors="ignore")
Anxiety.head(2)
```

```
Out[43]:
```

	TIPI1	TIPI2	TIPI3	TIPI4	TIPI5	TIPI6	TIPI7	TIPI8	TIPI9	TIPI10	...	religion	orientation	race	married	familysize	major	AgeGroup	score	intensity	cl
0	1	5	7	7	7	7	7	5	1	1	...	12	1	10	1	2	no degree	0	34	Extremely Severe	
1	6	5	4	7	5	4	7	7	1	5	...	7	3	70	1	4	no degree	0	17	Severe	

2 rows × 24 columns

```
In [44]: Stress0 = Stress.copy()
Stress = Stress.drop(Stress.iloc[:, 0:14], axis=1, errors="ignore")
Stress.head(2)
```

```
Out[44]:
```

	TIPI1	TIPI2	TIPI3	TIPI4	TIPI5	TIPI6	TIPI7	TIPI8	TIPI9	TIPI10	...	religion	orientation	race	married	familysize	major	AgeGroup	score	intensity	cl
0	1	5	7	7	7	7	7	5	1	1	...	12	1	10	1	2	no degree	0	40	Extremely Severe	
1	6	5	4	7	5	4	7	7	1	5	...	7	3	70	1	4	no degree	0	27	Severe	

2 rows × 24 columns

Independant and dependant variables

Dependant

- Score
- Intensity

Independent

- Q1-Q42
- Demographic/Personal info

COMPARISON AND VISUALIZATION OF DEMOGRAPHIC DATA WITH MENTAL DISORDER LEVELS

Create a copy of the df to maintain the numbers for the classification groups that will be used in modeling

```
In [45]: DepressionM = Stress.copy()
AnxietyM = Stress.copy()
StressM = Stress.copy()
```

EDUCATION LEVEL

The 'education' feature has 4 different categories with the following meanings:

- 1) Less than high school
- 2) High school
- 3) University degree
- 4) Graduate degree

```
In [46]: def changeEducationLevelValues(value) -> str:
    if value == 1:
        return '< HighSchool'
    if value == 2:
        return 'HighSchool'
    if value == 3:
        return 'University'
    if value == 4:
        return 'Graduate'

    return value

demographic['education'] = demographic['education'].apply(changeEducationLevelValues)
Depression['education'] = Depression['education'].apply(changeEducationLevelValues)
Anxiety['education'] = Anxiety['education'].apply(changeEducationLevelValues)
Stress['education'] = Stress['education'].apply(changeEducationLevelValues)

print('Count of answers on each education level: ')
print(demographic['education'].value_counts())
```

```
Count of answers on each education level:
University      14433
HighSchool      13542
Graduate         4578
< HighSchool     3463
Name: education, dtype: int64
```



```
In [47]: #education = {1:"Less than high school", 2:"High school", 3:"University Degree", 4:"Graduate Degree"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Education level count associated with each level of intensity of each condition')

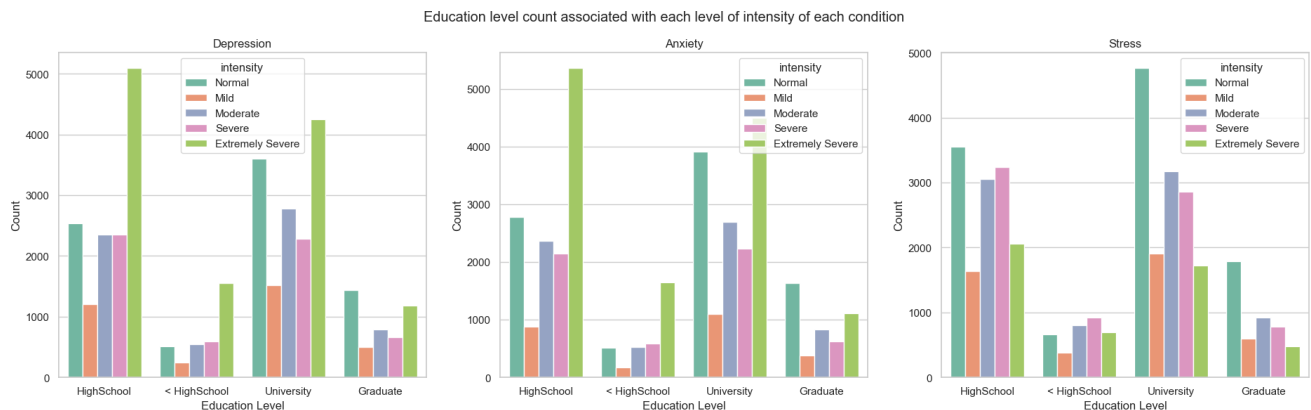
# Education level and depression
sns.countplot(ax=axes[0], data=Depression, x='education', hue=Depression['intensity'], palette="Set2", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Education level and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='education', hue=Anxiety['intensity'], palette="Set2", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Education level and stress
sns.countplot(ax=axes[2], data=Stress, x='education', hue=Stress['intensity'], palette="Set2", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Education Level', ylabel='Count') for i in range(0,3)]
fig.show()
```

C:\Users\sandr\AppData\Local\Temp\ipykernel_13864\54096356.py:23: UserWarning: Matplotlib is currently using module://matplotlib_inline.backend_inline, which is a non-GUI backend, so cannot show the figure.
fig.show()



The graphs show that in the education levels prior to high school, there is not much mental illness and students tend to be more or less relaxed. However, the numbers increase greatly when they enter high school and undergraduate college, and as an example the graph shows that the three analyzed illnesses tend to present the highest number of cases on the 'Extremely Severe' intensity level during this level of education. After high school, the DAS intensity levels start to slowly decrease and the Graduate studies seem to resemble the relaxed intensity levels of the first school years.

GENDER

The 'gender' feature has 3 different categories with the following meanings:

- 1) Male
- 2) Female
- 3) Other

```
In [48]: def changeGenderValues(value) -> str:
    if value == 1:
        return 'Male'
    if value == 2:
        return 'Female'
    if value == 3:
        return 'Other'

    return value

demographic['gender'] = demographic['gender'].apply(changeGenderValues)
Depression['gender'] = Depression['gender'].apply(changeGenderValues)
Anxiety['gender'] = Anxiety['gender'].apply(changeGenderValues)
Stress['gender'] = Stress['gender'].apply(changeGenderValues)

print('Count of answers on each gender: ')
print(demographic['gender'].value_counts().sort_index())
```

```
Count of answers on each gender:
Female    27590
Male      7923
Other      503
Name: gender, dtype: int64
```

```
In [49]: #gen = {1:"Male", 2:"Female", 3:"Other"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Gender count associated with each level of intensity of each condition')

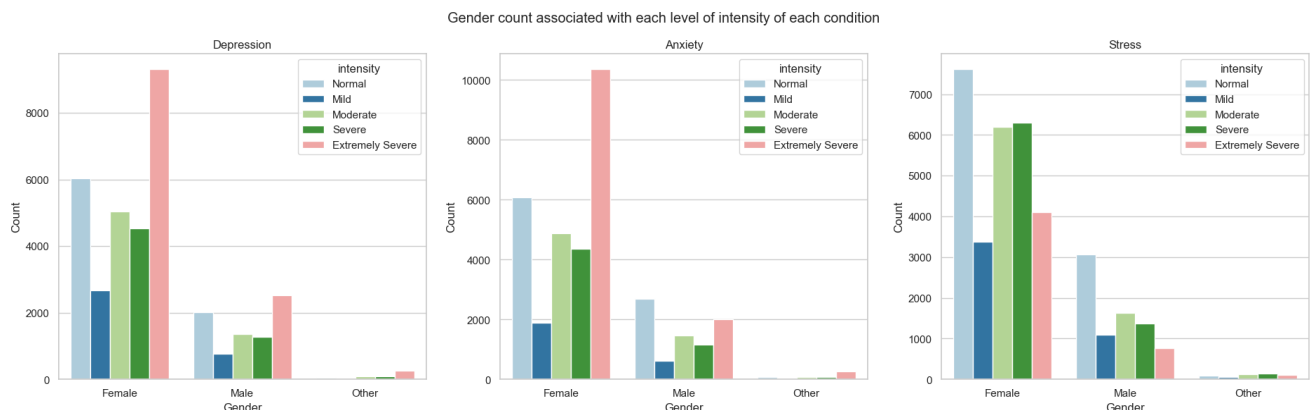
# Gender and depression
sns.countplot(ax=axes[0], data=Depression, x='gender', hue=Depression['intensity'], palette="Paired", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Gender and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='gender', hue=Anxiety['intensity'], palette="Paired", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Gender and stress
sns.countplot(ax=axes[2], data=Stress, x='gender', hue=Stress['intensity'], palette="Paired", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Gender', ylabel='Count') for i in range(0,3)]
fig.show()
```

C:\Users\sandr\AppData\Local\Temp\ipykernel_13864\88411219.py:23: UserWarning: Matplotlib is currently using module://matplotlibl b_inline.backend_inline, which is a non-GUI backend, so cannot show the figure.
fig.show()



The people that took the survey were mostly female and they present a high rate of "Extremely severe" depression and anxiety. Even though there are not many males represented, the graph shows that they tend to have lower scores than women in all three conditions.

AGE GROUP

The 'AgeGroup' feature has 8 different categories with the following meanings:

- 0) < 20
- 1) 20-24
- 2) 25-29

- 3) 30-34
- 4) 35-39
- 5) 40-49
- 6) 50-59
- 7) > 60

```
In [50]: def binAgeValues(value) -> str:
    if value == 0:
        return '< 20'
    if value == 1:
        return '20-24'
    if value == 2:
        return '25-29'
    if value == 3:
        return '30-34'
    if value == 4:
        return '35-39'
    if value == 5:
        return '40-49'
    if value == 6:
        return '50-59'
    if value == 7:
        return '>60'

    return value

demographic['AgeGroup'] = demographic['AgeGroup'].apply(binAgeValues)
Depression['AgeGroup'] = Depression['AgeGroup'].apply(binAgeValues)
Anxiety['AgeGroup'] = Anxiety['AgeGroup'].apply(binAgeValues)
Stress['AgeGroup'] = Stress['AgeGroup'].apply(binAgeValues)

print('Count of answers on each Age Group: ')
print(demographic['AgeGroup'].value_counts().sort_index())
```

Count of answers on each Age Group:

20-24	13620
25-29	4997
30-34	2012
35-39	1034
40-49	1228
50-59	707
< 20	12151
>60	267

Name: AgeGroup, dtype: int64

```
In [51]: #agroup = {0:"<20", 1:"20-24", 2:"25-29", 3:"30-34", 4:"35-39", 5:"40-49", 6:"50-59", 7:">60"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Age group count associated with each level of intensity of each condition')

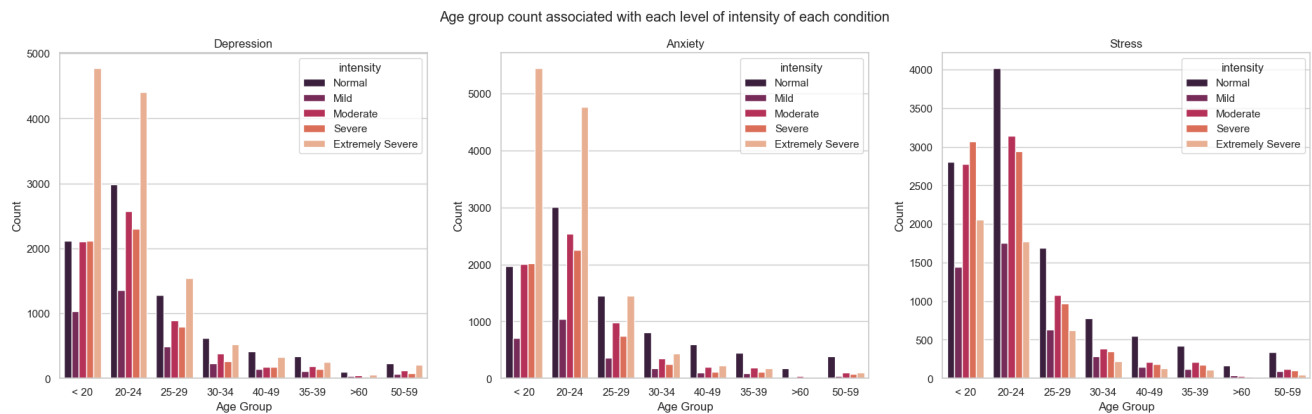
# Age and depression
sns.countplot(ax=axes[0], data=Depression, x='AgeGroup', hue=Depression['intensity'], palette="rocket", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Age and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='AgeGroup', hue=Anxiety['intensity'], palette="rocket", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Age and stress
sns.countplot(ax=axes[2], data=Stress, x='AgeGroup', hue=Stress['intensity'], palette="rocket", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Age Group', ylabel='Count') for i in range(0,3)]
fig.show()
```

C:\Users\sandr\AppData\Local\Temp\ipykernel_13864\3103444475.py:23: UserWarning: Matplotlib is currently using module://matplotlib_inline.backend_inline, which is a non-GUI backend, so cannot show the figure.
fig.show()



The graphs show that the majority of survey respondents are young adult people from around 20 and 24 years old. It also shows that younger people tend to have higher scores (especially with depression and anxiety) and that the scores decrease as people get older.

RELIGION

The 'religion' feature has 12 different categories with the following meanings:

- 1) Agnostic
- 2) Atheist
- 3) Buddhist
- 4) Christian (Catholic)
- 5) Christian (Mormon)
- 6) Christian (Protestant)
- 7) Christian (Other)
- 8) Hindu
- 9) Jewish
- 10) Muslim
- 11) Sikh
- 12) Other

In [52]: *# change 0 value to 12 as it's other value for people who didn't enter value to this field*

```
def updateReligionValue(value):
    if value == 0:
        return 12
    return value

demographic['religion'] = demographic['religion'].apply(updateReligionValue)

def changeReligionValues(value) -> str:
    if (value == 0 or value == 12):
        return 'Other'
    if value == 1:
        return 'Agnostic'
    if value == 2:
        return 'Atheist'
    if value == 3:
        return 'Buddhist'
    if (value == 4 or value == 5 or value == 6 or value == 7):
        return 'Christian'
    if value == 8:
        return 'Hindu'
    if value == 9:
        return 'Jewish'
    if value == 10:
        return 'Muslim'
    if value == 11:
        return 'Sikh'

    return value

demographic['religion'] = demographic['religion'].apply(changeReligionValues)
Depression['religion'] = Depression['religion'].apply(changeReligionValues)
Anxiety['religion'] = Anxiety['religion'].apply(changeReligionValues)
Stress['religion'] = Stress['religion'].apply(changeReligionValues)

print('Count of answers on each religion: ')
print(demographic['religion'].value_counts())
```

Count of answers on each religion:

Muslim	20447
Christian	6294
Atheist	3282
Agnostic	2800
Other	1598
Buddhist	797
Hindu	629
Jewish	111
Sikh	58

Name: religion, dtype: int64

```
In [53]: #rel = {1:"Agnostic", 2:"Atheist", 3:"Buddhist", 4:"Christian (Catholic)", 5:"Christian (Mormon)", 6:"Christian (Protestant)", 7:
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Religion count associated with each level of intensity of each condition')

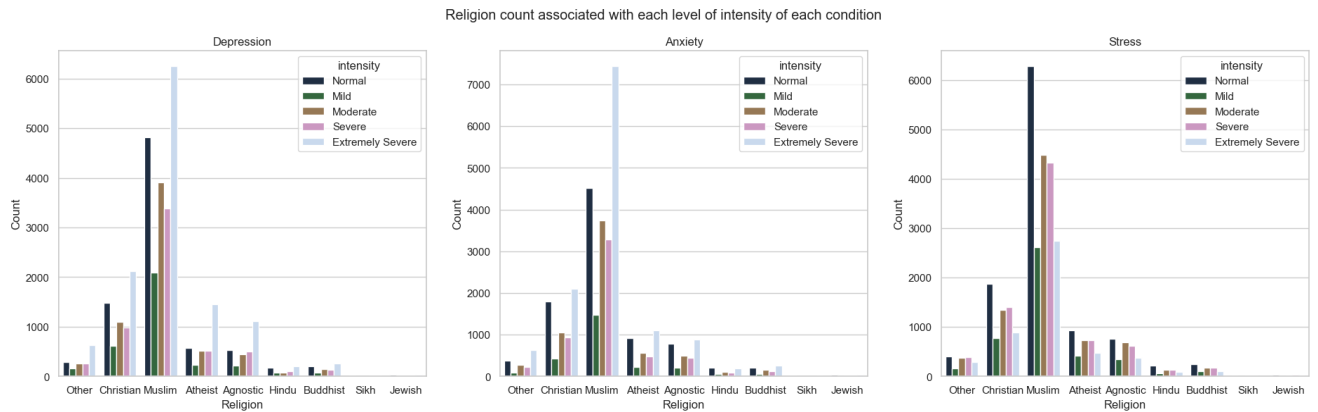
# Religion and depression
sns.countplot(ax=axes[0], data=Depression, x='religion', hue=Depression['intensity'], palette="cubehelix", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Religion and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='religion', hue=Anxiety['intensity'], palette="cubehelix", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Religion and stress
sns.countplot(ax=axes[2], data=Stress, x='religion', hue=Stress['intensity'], palette="cubehelix", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Religion', ylabel='Count') for i in range(0,3)]
```

```
Out[53]: [[Text(0.5, 0, 'Religion'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Religion'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Religion'), Text(0, 0.5, 'Count')]]
```



Most of the people who took the survey were muslim.

SEXUAL ORIENTATION

The 'orientation' feature has 5 different categories with the following meanings:

- 1) Heterosexual
- 2) Bisexual
- 3) Homosexual
- 4) Asexual
- 5) Other

```
In [54]: def changeOrientationValues(value) -> str:
    if value == 1:
        return 'Heterosexual'
    if value == 2:
        return 'Bisexual'
    if value == 3:
        return 'Homosexual'
    if value == 4:
        return 'Asexual'
    if value == 5:
        return 'Other'

    return value

demographic['orientation'] = demographic['orientation'].apply(changeOrientationValues)
Depression['orientation'] = Depression['orientation'].apply(changeOrientationValues)
Anxiety['orientation'] = Anxiety['orientation'].apply(changeOrientationValues)
Stress['orientation'] = Stress['orientation'].apply(changeOrientationValues)

print('Count of answers on each orientation option: ')
print(demographic['orientation'].value_counts())
```

```
Count of answers on each orientation option:
Heterosexual    22133
Homosexual      4615
Bisexual        4375
Other           3102
Asexual         1791
Name: orientation, dtype: int64
```

```
In [55]: #ori = {1:"Heterosexual", 2:"Bisexual", 3:"Homosexual", 4:"Asexual", 5:"Other"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Sexual orientation frequency count associated with each level of intensity of each condition')

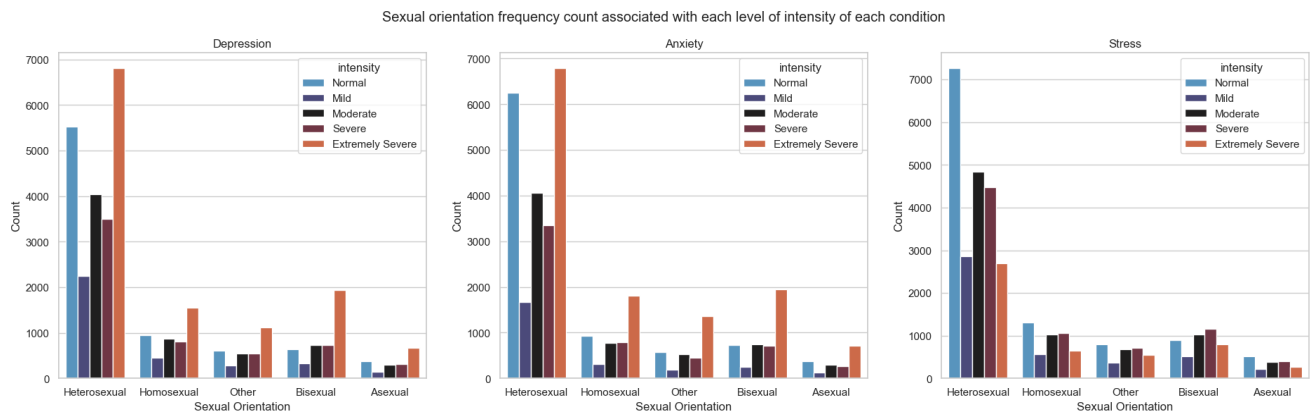
# Orientation and depression
sns.countplot(ax=axes[0], data=Depression, x='orientation', hue=Depression['intensity'], palette="icefire", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Orientation and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='orientation', hue=Anxiety['intensity'], palette="icefire", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Orientation and stress
sns.countplot(ax=axes[2], data=Stress, x='orientation', hue=Stress['intensity'], palette="icefire", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Sexual Orientation', ylabel='Count') for i in range(0,3)]
```

```
Out[55]: [[Text(0.5, 0, 'Sexual Orientation'), Text(0, 0.5, 'Count')],
 [Text(0.5, 0, 'Sexual Orientation'), Text(0, 0.5, 'Count')],
 [Text(0.5, 0, 'Sexual Orientation'), Text(0, 0.5, 'Count')]]
```



The majority of respondents are heterosexual.

RACE

The 'race' feature has 7 different categories with the following meanings:

- 10) Asian
- 20) Arab
- 30) Black
- 40) Indigenous Australian
- 50) Native American
- 60) White
- 70) Other

```
In [56]: def changeRaceValues(value) -> str:
          if value == 10:
              return 'Asian'
          if value == 20:
              return 'Arab'
          if value == 30:
              return 'Black'
          if value == 40:
              return 'Indig. Aust.'
          if value == 50:
              return 'Nat. Am.'
          if value == 60:
              return 'White'
          if value == 70:
              return 'Other'

          return value

demographic['race'] = demographic['race'].apply(changeRaceValues)
Depression['race'] = Depression['race'].apply(changeRaceValues)
Anxiety['race'] = Anxiety['race'].apply(changeRaceValues)
Stress['race'] = Stress['race'].apply(changeRaceValues)

print('Count of answers on each race: ')
print(demographic['race'].value_counts())
```

Count of answers on each race:

Asian	21335
White	9254
Other	4396
Black	542
Arab	289
Nat. Am.	182
Indig. Aust.	18

Name: race, dtype: int64


```
In [57]: #rac = {10:"Asian", 20:"Arab", 30:"Black", 40:"Indigenous Australian", 50:"Native American", 60:"White", 70:"Other"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Race count associated with each level of intensity of each condition')

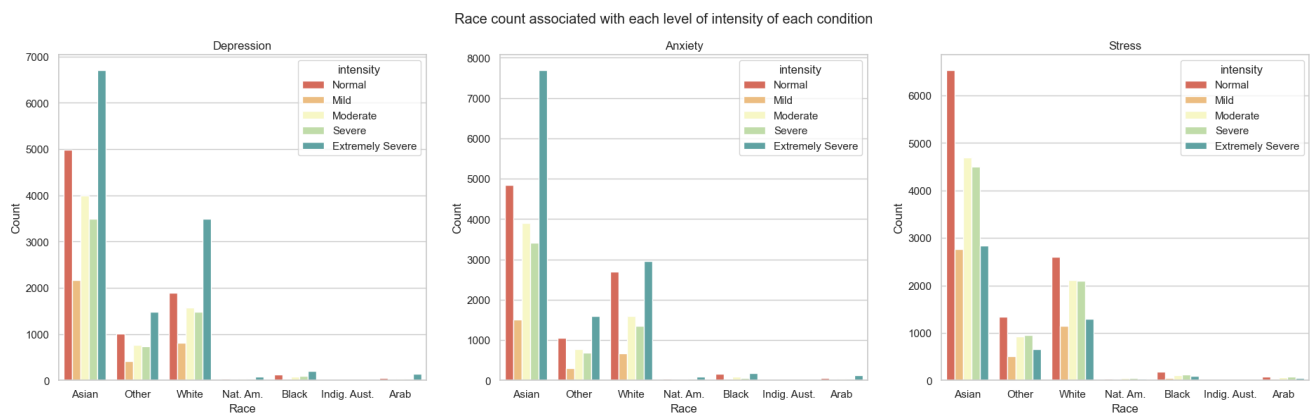
# Race and depression
sns.countplot(ax=axes[0], data=Depression, x='race', hue=Depression['intensity'], palette="Spectral", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Race and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='race', hue=Anxiety['intensity'], palette="Spectral", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Race and stress
sns.countplot(ax=axes[2], data=Stress, x='race', hue=Stress['intensity'], palette="Spectral", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set(xlabel='Race', ylabel='Count') for i in range(0,3)]
```

```
Out[57]: [[Text(0.5, 0, 'Race'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Race'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Race'), Text(0, 0.5, 'Count')]]
```



Most respondents are asian, followed by white and others. Figure 2.6 shows that the predominant categories have similar distributions for the different condition intensities.

MARITAL STATUS

The 'married' feature has 3 different categories with the following meanings:

- 1) Never married
- 2) Currently married
- 3) Previously married

```
In [58]: def changeMaritalStatusValues(value) -> str:
    if value == 1:
        return 'Never Married'
    if value == 2:
        return 'Currently Married'
    if value == 3:
        return 'Previously Married'

    return value

demographic['married'] = demographic['married'].apply(changeMaritalStatusValues)
Depression['married'] = Depression['married'].apply(changeMaritalStatusValues)
Anxiety['married'] = Anxiety['married'].apply(changeMaritalStatusValues)
Stress['married'] = Stress['married'].apply(changeMaritalStatusValues)

print('Count of answers on each marital status: ')
print(demographic['married'].value_counts())
```

```
Count of answers on each marital status:
Never Married      30830
Currently Married   4003
Previously Married  1183
Name: married, dtype: int64
```

```
In [59]: #ms = {1:"Never married", 2:"Currently married", 3:"Previously married"}
fig, axes = plt.subplots(1,3,figsize=(23, 6))
sns.set_theme(style="whitegrid")
fig.suptitle('Marital status count associated with each level of intensity of each condition')

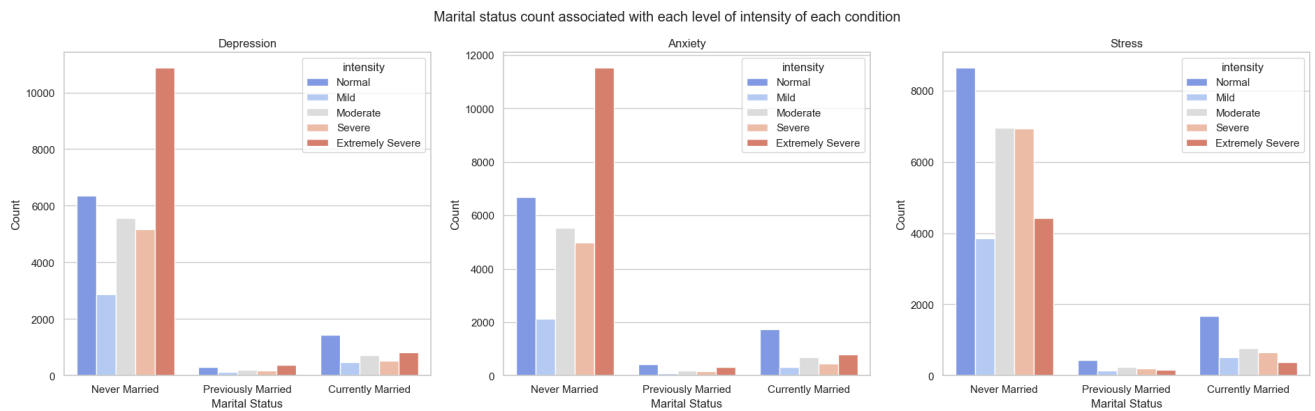
# Marital Status and depression
sns.countplot(ax=axes[0], data=Depression, x='married', hue=Depression['intensity'], palette="coolwarm", hue_order=level_order)
#Title
axes[0].set_title('Depression')

# Marital Status and anxiety
sns.countplot(ax=axes[1], data=Anxiety, x='married', hue=Anxiety['intensity'], palette="coolwarm", hue_order=level_order)
#Title
axes[1].set_title('Anxiety')

# Marital Status and stress
sns.countplot(ax=axes[2], data=Stress, x='married', hue=Stress['intensity'], palette="coolwarm", hue_order=level_order)
#Title
axes[2].set_title('Stress')

#Axis titles
[axes[i].set_xlabel('Marital Status', ylabel='Count') for i in range(0,3)]
```

```
Out[59]: [[Text(0.5, 0, 'Marital Status'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Marital Status'), Text(0, 0.5, 'Count')],
[Text(0.5, 0, 'Marital Status'), Text(0, 0.5, 'Count')]]
```



The graphs show that most respondents are single and that they tend to have higher levels of depression and anxiety than married people.

Creation of a single dataframe

Here, we will create a single DF to use for our model by combining the Depression, Anxiety and Stress DFs and creating one label column (that is used as the ground truth for evaluation in later steps. This label column computed as the sum of the score column in each DF, will be named "DASS" : for Depression, Anxiety and Stress Score

```
In [60]: # Establish some primary keys to perform the inner join
DepressionM['key'] = DepressionM.index
AnxietyM['key'] = AnxietyM.index
StressM['key'] = StressM.index

features = ['key', 'TIPI1', 'TIPI2', 'TIPI3', 'TIPI4', 'TIPI5', 'TIPI6', 'TIPI7', 'TIPI8', 'TIPI9', 'TIPI10', 'education', 'urban',
            'race', 'married', 'familysize', 'major', 'AgeGroup']

dass = pd.merge(DepressionM, AnxietyM,
                how='inner',
                on=features)

dass = pd.merge(dass, StressM,
                how='inner',
                on=features)

dass["DASS"] = dass["score_x"] + dass["score_y"] + dass["score"]
dass = dass.drop(columns=["score_x", "score_y", "score", "key"], axis=1, errors="ignore")
dass = dass.drop(columns=["intensity_x", "intensity_y", "intensity"], axis=1, errors="ignore")
dass = dass.drop(columns=["intensity_x", "intensity_y", "intensity"], axis=1, errors="ignore")
dass = dass.drop(columns=["cIntensity_x", "cIntensity_y", "cIntensity"], axis=1, errors="ignore")
print("DASS df dimensions: ", dass.shape)
dass.head()
```

DASS df dimensions: (36016, 22)

```
Out[60]:
```

	TIPI1	TIPI2	TIPI3	TIPI4	TIPI5	TIPI6	TIPI7	TIPI8	TIPI9	TIPI10	...	gender	age	religion	orientation	race	married	familysize	major	AgeGroup	D
0	1	5	7	7	7	7	7	5	1	1	...	2	16.0	12	1	10	1	2	no degree	0	
1	6	5	4	7	5	4	7	7	1	5	...	2	16.0	7	3	70	1	4	no degree	0	
2	2	5	2	2	5	6	5	5	3	2	...	2	17.0	4	3	60	1	3	no degree	0	
3	1	1	7	4	6	4	6	1	6	1	...	2	13.0	4	5	70	1	5	science	0	
4	2	5	3	6	5	5	5	6	3	3	...	2	19.0	10	1	10	1	4	psychology	0	

5 rows × 22 columns

- 1 -> 25 -> Normal
- 2 -> 26-50 -> Mild
- 3 -> 51-75 -> Moderate
- 4 -> 76-100 -> Severe
- 5 -> 101-126 -> Extremely severe

```
In [61]: # Max and min of DASS score
maxi = dass['DASS'].max()
mini = dass['DASS'].min()
print(maxi, mini)
```

126 0

```
In [62]: # Classify the resulting DASS score
def classifyDASS(x):
    if x < 25:
        return "Normal"
    elif 26 <= x <= 50:
        return "Mild"
    elif 51 <= x <= 75:
        return "Moderate"
    elif 76 <= x <= 100:
        return "Severe"
    else:
        return "Extremely Severe"

dass['cDASS'] = dass['DASS'].apply(classifyDASS)
dass['cDASS'].value_counts().sort_index()
```

```
Out[62]:
```

Extremely Severe	4962
Mild	8203
Moderate	10237
Normal	4801
Severe	7813

Name: cDASS, dtype: int64

```
In [63]: dass = dass.drop('DASS', axis=1, errors="ignore")
n = dass.shape[0] == Depression.shape[0]
print("Does the dass dataset has the same rows as the previous 3 df? ", n)
```

Does the dass dataset has the same rows as the previous 3 df? True

Data Analytics and Modeling

Importing modeling libraries

```
In [64]: from sklearn.preprocessing import MinMaxScaler
from sklearn.model_selection import train_test_split
# Naive bayes
from sklearn.naive_bayes import GaussianNB
# Random forest classifier
from sklearn.ensemble import RandomForestClassifier
# AdaBoost
from sklearn.ensemble import AdaBoostClassifier
# Performance metrics
from sklearn.metrics import r2_score, accuracy_score, precision_score, recall_score, f1_score
from sklearn.metrics import confusion_matrix, plot_confusion_matrix, classification_report
```

DASS dataframe

In this section, we'll analyze the DASS dataframe through modeling with Linear Regression to try to find a good prediction for the dass continuous variable which represents the total score.

Split the data into training and testing set

```
In [65]: # Input data to the model
X = dass.drop(["cDASS", "major"], axis=1, errors="ignore")
# Ground truth Label
y = dass["cDASS"]

# Splitting the data, training set and testing set
X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.80, random_state=100)
scaler = MinMaxScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

# Train and test set dimensions
print('Training Set:', X_train.shape, y_train.shape)
print('Test Set:', X_test.shape, y_test.shape)
```

Training Set: (28812, 20) (28812,)
Test Set: (7204, 20) (7204,)

Gaussian NB

```
In [66]: NB=GaussianNB().fit(X_train_scaled,y_train)
prediction = NB.predict(X_test_scaled)

aNB1=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aNB1)

f1NB1=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1NB1)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

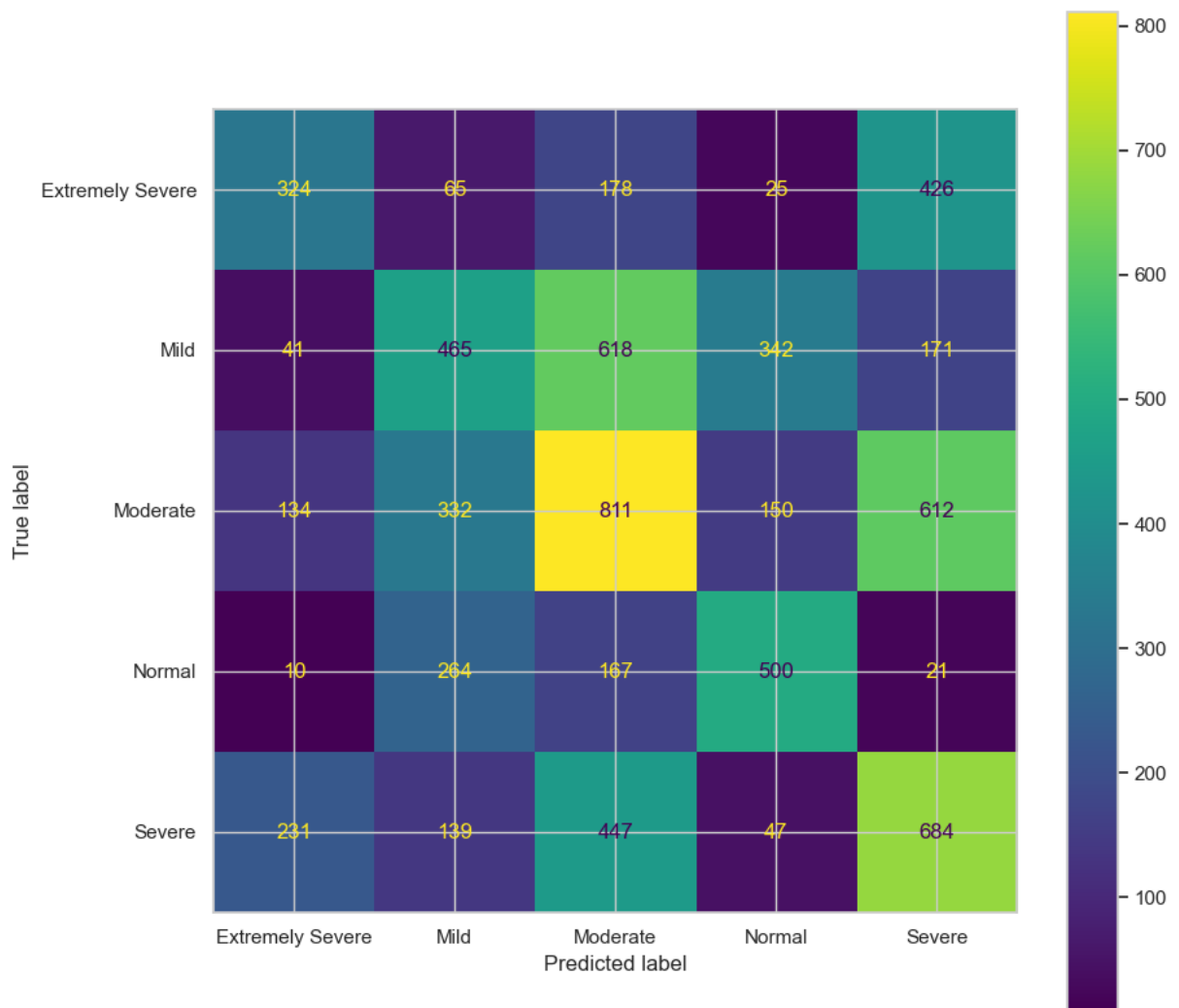
fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(NB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.39
F1_Score: 0.38
Recall_Score: 0.39
Precision_Score: 0.39

		precision	recall	f1-score	support
Extremely	Severe	0.44	0.32	0.37	1018
	Mild	0.37	0.28	0.32	1637
	Moderate	0.37	0.40	0.38	2039
	Normal	0.47	0.52	0.49	962
	Severe	0.36	0.44	0.40	1548
accuracy				0.39	7204
macro avg		0.40	0.39	0.39	7204
weighted avg		0.39	0.39	0.38	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[66]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d5085ef6a0>



Random Forest Classifier

```
In [67]: RF=RandomForestClassifier(n_estimators=200,min_samples_split=3,min_samples_leaf=1,max_depth=160,max_features='auto').fit(X_train,
prediction = RF.predict(X_test_scaled)

aRF1=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aRF1)

f1RF1=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1RF1)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

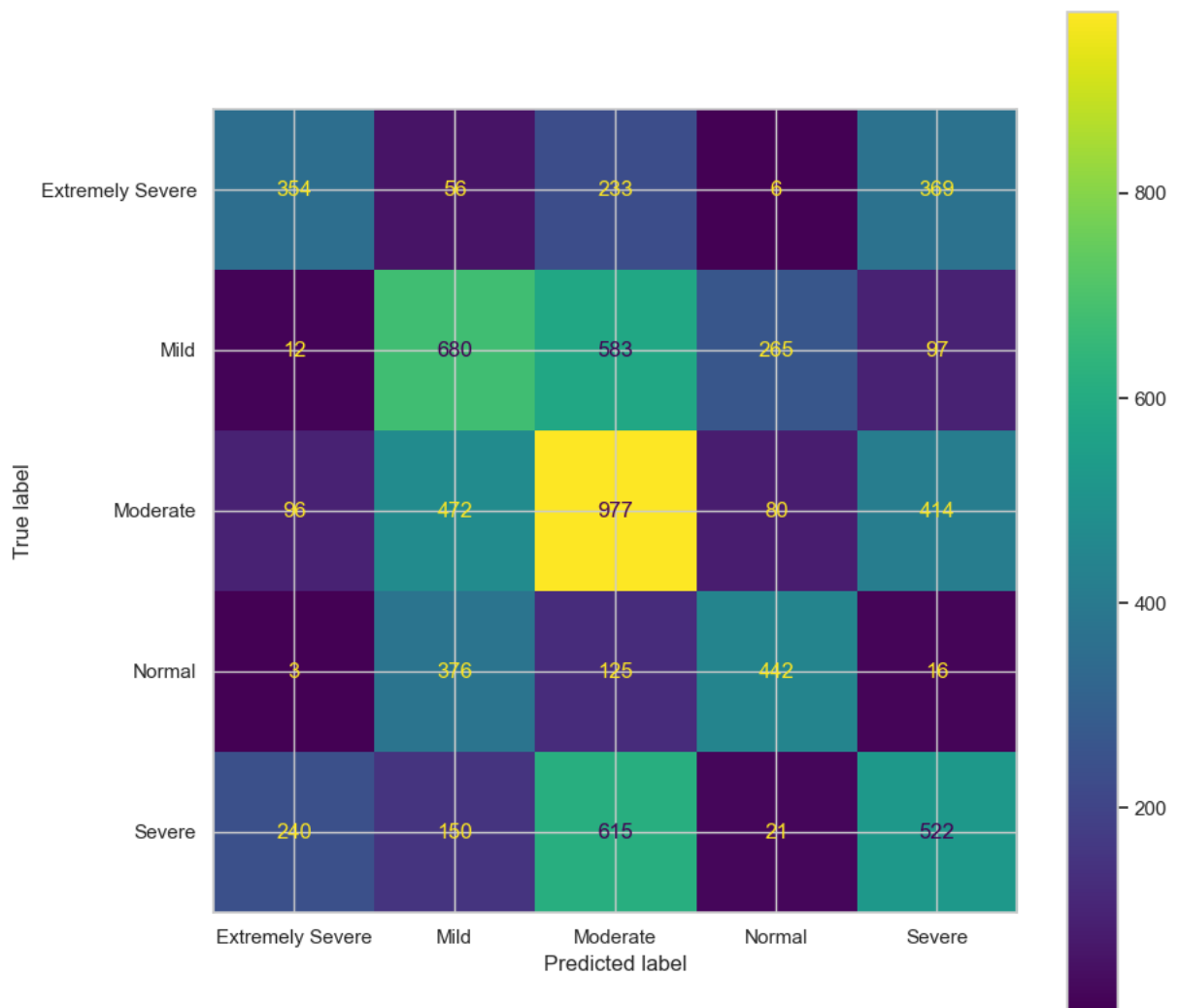
fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(RF,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.41
F1_Score: 0.41
Recall_Score: 0.41
Precision_Score: 0.42

	precision	recall	f1-score	support
Extremely Severe	0.50	0.35	0.41	1018
Mild	0.39	0.42	0.40	1637
Moderate	0.39	0.48	0.43	2039
Normal	0.54	0.46	0.50	962
Severe	0.37	0.34	0.35	1548
accuracy			0.41	7204
macro avg	0.44	0.41	0.42	7204
weighted avg	0.42	0.41	0.41	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[67]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508d24880>



AdaBoost (Ensemble Method)

```
In [68]: AB=AdaBoostClassifier(n_estimators=50,learning_rate=1)
AB.fit(X_train_scaled, y_train)
prediction = AB.predict(X_test_scaled)

aAB1=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aAB1)

f1AB1=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1AB1)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(AB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.42

F1_Score: 0.42

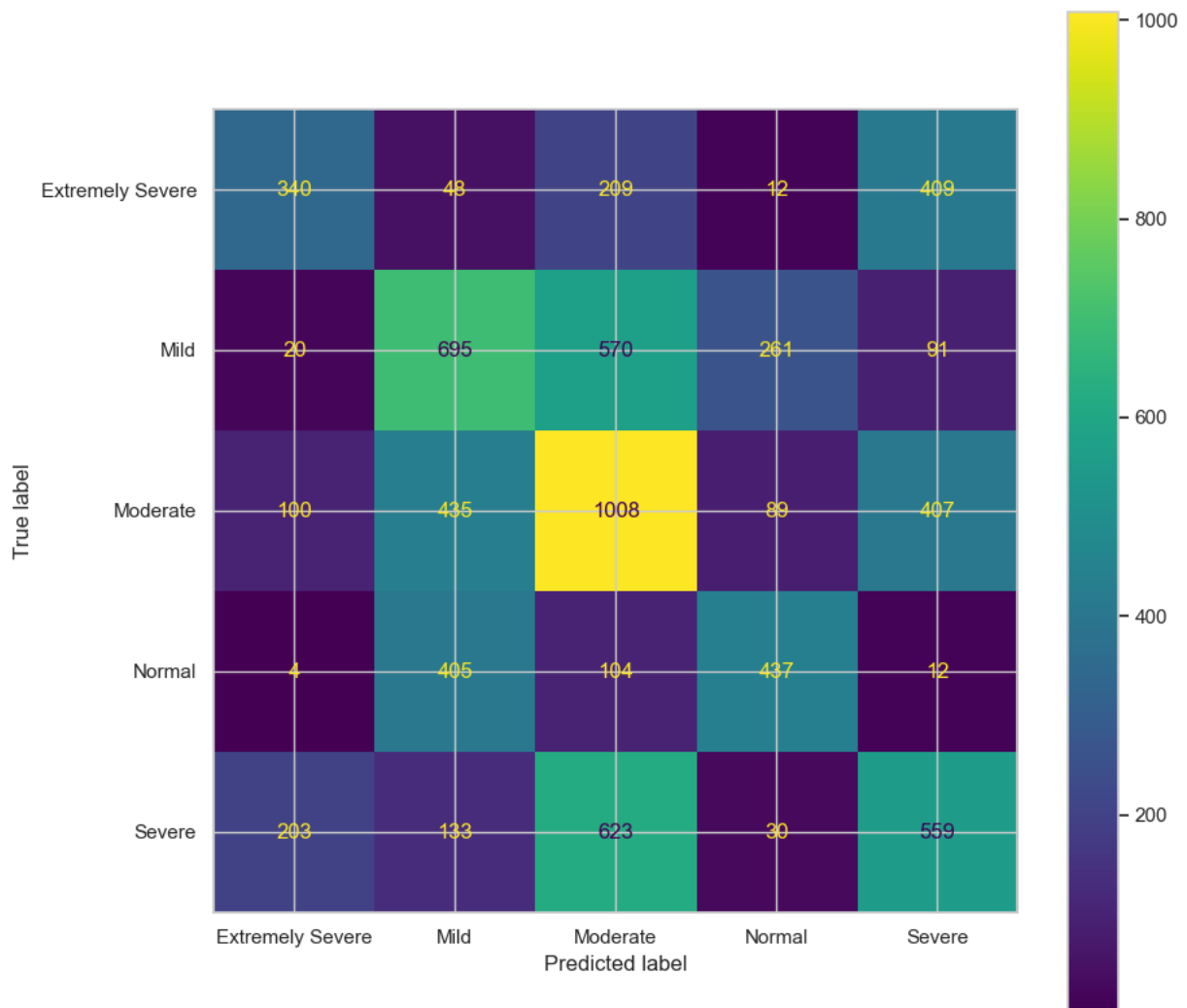
Recall_Score: 0.42

Precision_Score: 0.43

	precision	recall	f1-score	support
Extremely Severe	0.51	0.33	0.40	1018
Mild	0.41	0.42	0.41	1637
Moderate	0.40	0.49	0.44	2039
Normal	0.53	0.45	0.49	962
Severe	0.38	0.36	0.37	1548
accuracy			0.42	7204
macro avg	0.44	0.41	0.42	7204
weighted avg	0.43	0.42	0.42	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[68]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508e1bf40>



Complete DASS score model summary

```
In [69]: summary={
            'Model':['GaussianNB', 'Random-Forest', 'AdaBoost'],
            'Accuracy(%)': [aNB1*100, aRF1*100, aAB1*100],
            'F1_Score(%)': [f1NB1*100, f1RF1*100, f1AB1*100],
        }
summaryDass=pd.DataFrame(summary)
summaryDass['key'] = summaryDass.index
summaryDass
```

Out[69]:

	Model	Accuracy(%)	F1_Score(%)	key
0	GaussianNB	39.0	38.0	0
1	Random-Forest	41.0	41.0	1
2	AdaBoost	42.0	42.0	2

As it can be seen, the use of the whole DAS score gives really low accuracies, with the greatest value being around 45%. Therefore, we will start applying the same models to the other 3 data frames to see if we can get higher accuracies and better predictions.

Depression dataframe

Split the data into training and testing set

```
In [70]: # Input data to the model
X = Depression0.drop(["cIntensity", "intensity", "score", "major"], axis=1, errors="ignore")
# Ground truth label
y = Depression0["intensity"]

# Splitting the data, training set and testing set
X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.80, random_state=100)
scaler = MinMaxScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

# Train and test set dimensions
print('Training Set:', X_train.shape, y_train.shape)
print('Test Set:', X_test.shape, y_test.shape)
```

Training Set: (28812, 34) (28812,)

Test Set: (7204, 34) (7204,)

Gaussian NB

```
In [71]: NB=GaussianNB().fit(X_train_scaled,y_train)
prediction = NB.predict(X_test_scaled)

aNB2=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aNB2)

f1NB2=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1NB2)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(NB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.86

F1_Score: 0.87

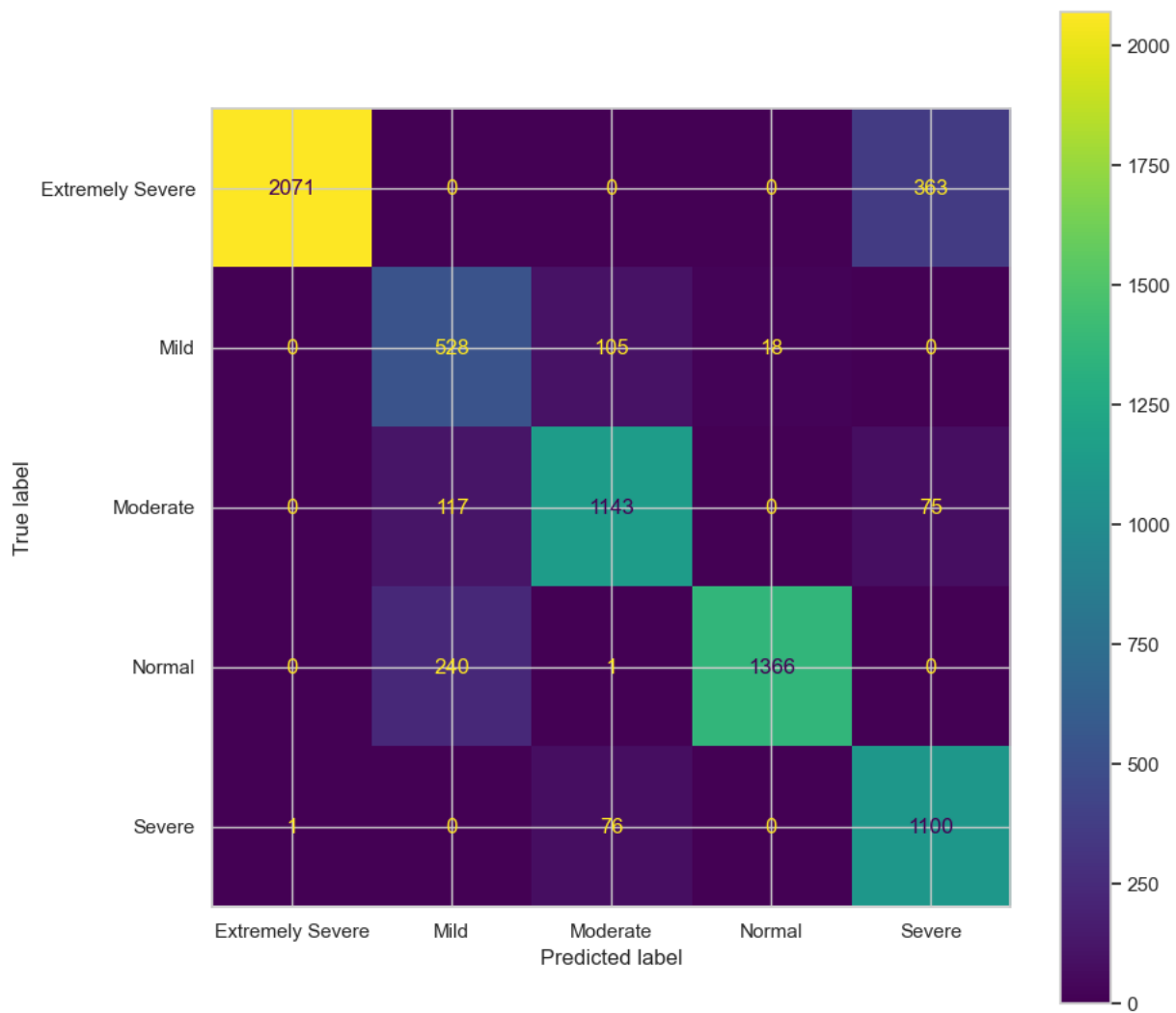
Recall_Score: 0.86

Precision_Score: 0.89

	precision	recall	f1-score	support
Extremely Severe	1.00	0.85	0.92	2434
Mild	0.60	0.81	0.69	651
Moderate	0.86	0.86	0.86	1335
Normal	0.99	0.85	0.91	1607
Severe	0.72	0.93	0.81	1177
accuracy			0.86	7204
macro avg	0.83	0.86	0.84	7204
weighted avg	0.89	0.86	0.87	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[71]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d507b98c10>



Random Forest Classifier

```
In [72]: RF=RandomForestClassifier(n_estimators=200,min_samples_split=3,min_samples_leaf=1,max_depth=160,max_features='auto').fit(X_train,
prediction = RF.predict(X_test_scaled)

aRF2=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aRF2)

f1RF2=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1RF2)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(RF,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.92

F1_Score: 0.92

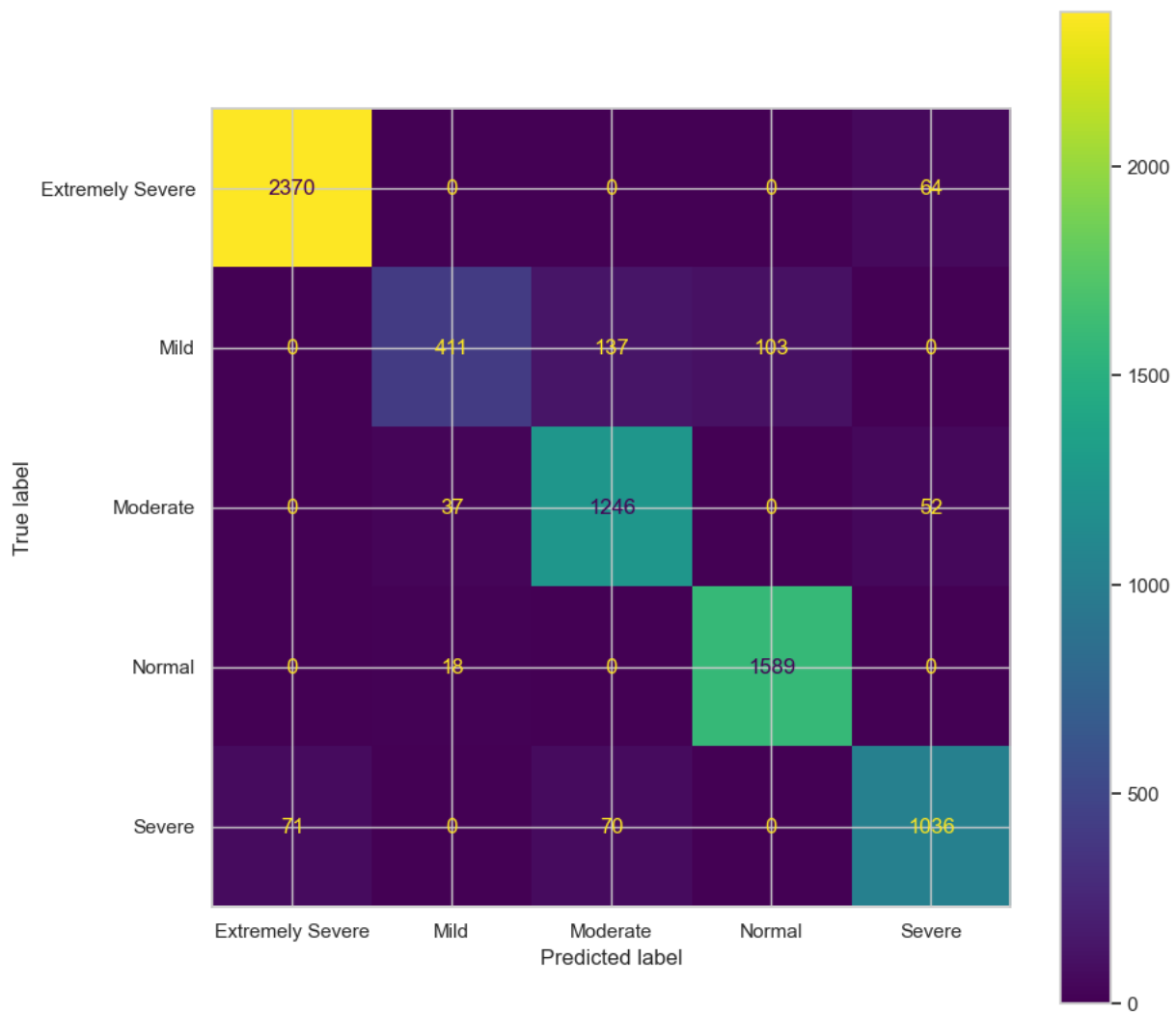
Recall_Score: 0.92

Precision_Score: 0.92

	precision	recall	f1-score	support
Extremely Severe	0.97	0.97	0.97	2434
Mild	0.88	0.63	0.74	651
Moderate	0.86	0.93	0.89	1335
Normal	0.94	0.99	0.96	1607
Severe	0.90	0.88	0.89	1177
accuracy			0.92	7204
macro avg	0.91	0.88	0.89	7204
weighted avg	0.92	0.92	0.92	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[72]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508cfe0a0>



AdaBoost (Ensemble Method)

```
In [73]: AB=AdaBoostClassifier(n_estimators=50,learning_rate=1)
AB.fit(X_train_scaled, y_train)
prediction = AB.predict(X_test_scaled)

aAB2=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aAB2)

f1AB2=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1AB2)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(AB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.75

F1_Score: 0.76

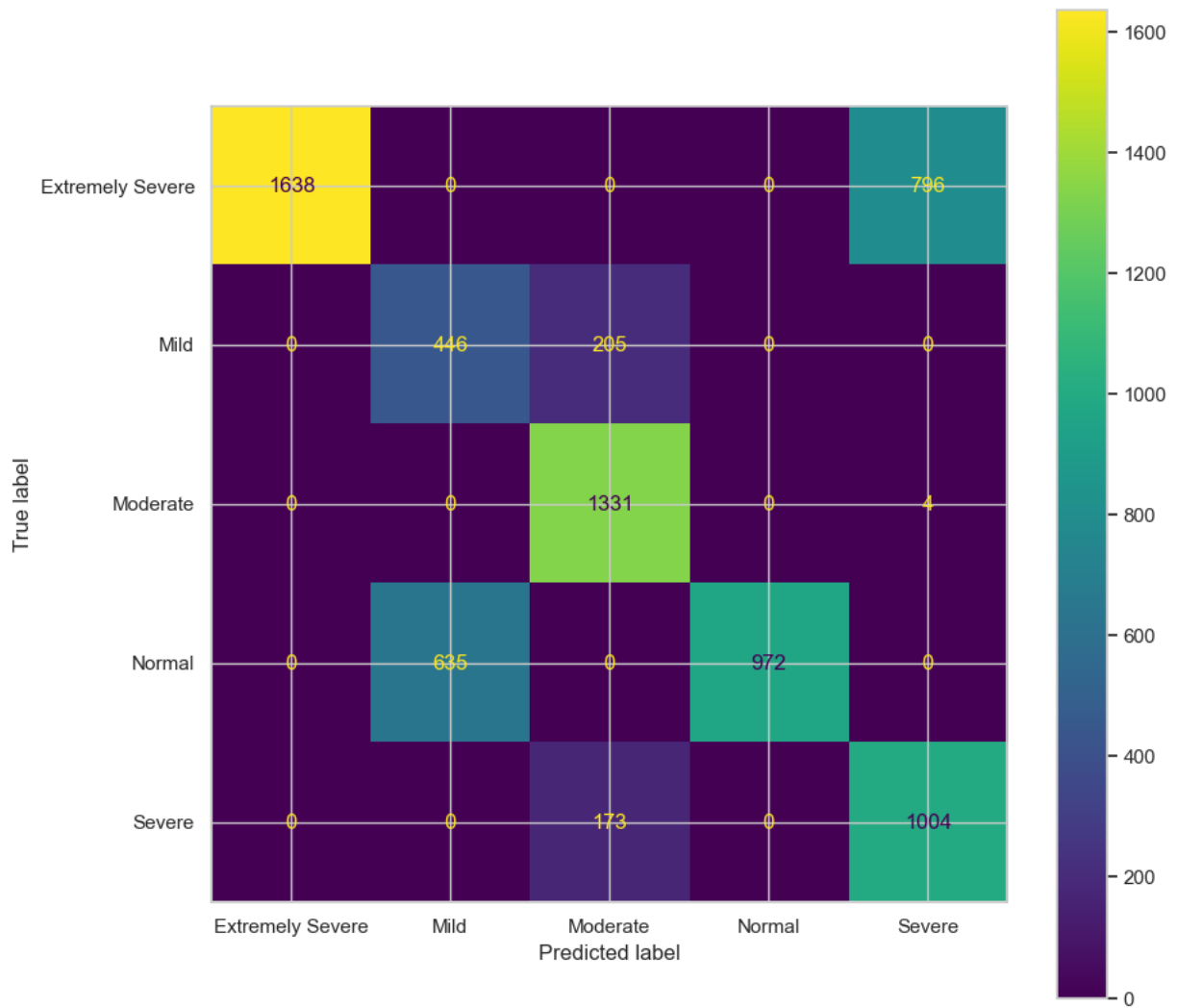
Recall_Score: 0.75

Precision_Score: 0.83

	precision	recall	f1-score	support
Extremely Severe	1.00	0.67	0.80	2434
Mild	0.41	0.69	0.52	651
Moderate	0.78	1.00	0.87	1335
Normal	1.00	0.60	0.75	1607
Severe	0.56	0.85	0.67	1177
accuracy			0.75	7204
macro avg	0.75	0.76	0.72	7204
weighted avg	0.83	0.75	0.76	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[73]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d507f048e0>



Complete Depression modeling summary

```
In [74]: summary={
            'Model': ['GaussianNB', 'Random-Forest', 'AdaBoost'],
            'Accuracy(%)': [aNB2*100, aRF2*100, aAB2*100],
            'F1_Score(%)': [f1NB2*100, f1RF2*100, f1AB2*100],
        }
summaryDepression=pd.DataFrame(summary)
summaryDepression['key'] = summaryDepression.index
summaryDepression
```

Out[74]:

	Model	Accuracy(%)	F1_Score(%)	key
0	GaussianNB	86.0	87.0	0
1	Random-Forest	92.0	92.0	1
2	AdaBoost	75.0	76.0	2

Anxiety dataframe

Split the data into training and testing set

```
In [75]: # Input data to the model
X = Anxiety0.drop(["cIntensity", "intensity", "score", "major"], axis=1, errors="ignore")
# Ground truth label
y = Anxiety0["intensity"]

# Splitting the data, training set and testing set
X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.80, random_state=100)
scaler = MinMaxScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

# Train and test set dimensions
print('Training Set:', X_train.shape, y_train.shape)
print('Test Set:', X_test.shape, y_test.shape)
```

Training Set: (28812, 34) (28812,)

Test Set: (7204, 34) (7204,)

Gaussian NB

```
In [76]: NB=GaussianNB().fit(X_train_scaled,y_train)
prediction = NB.predict(X_test_scaled)

aNB3=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aNB3)

f1NB3=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1NB3)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

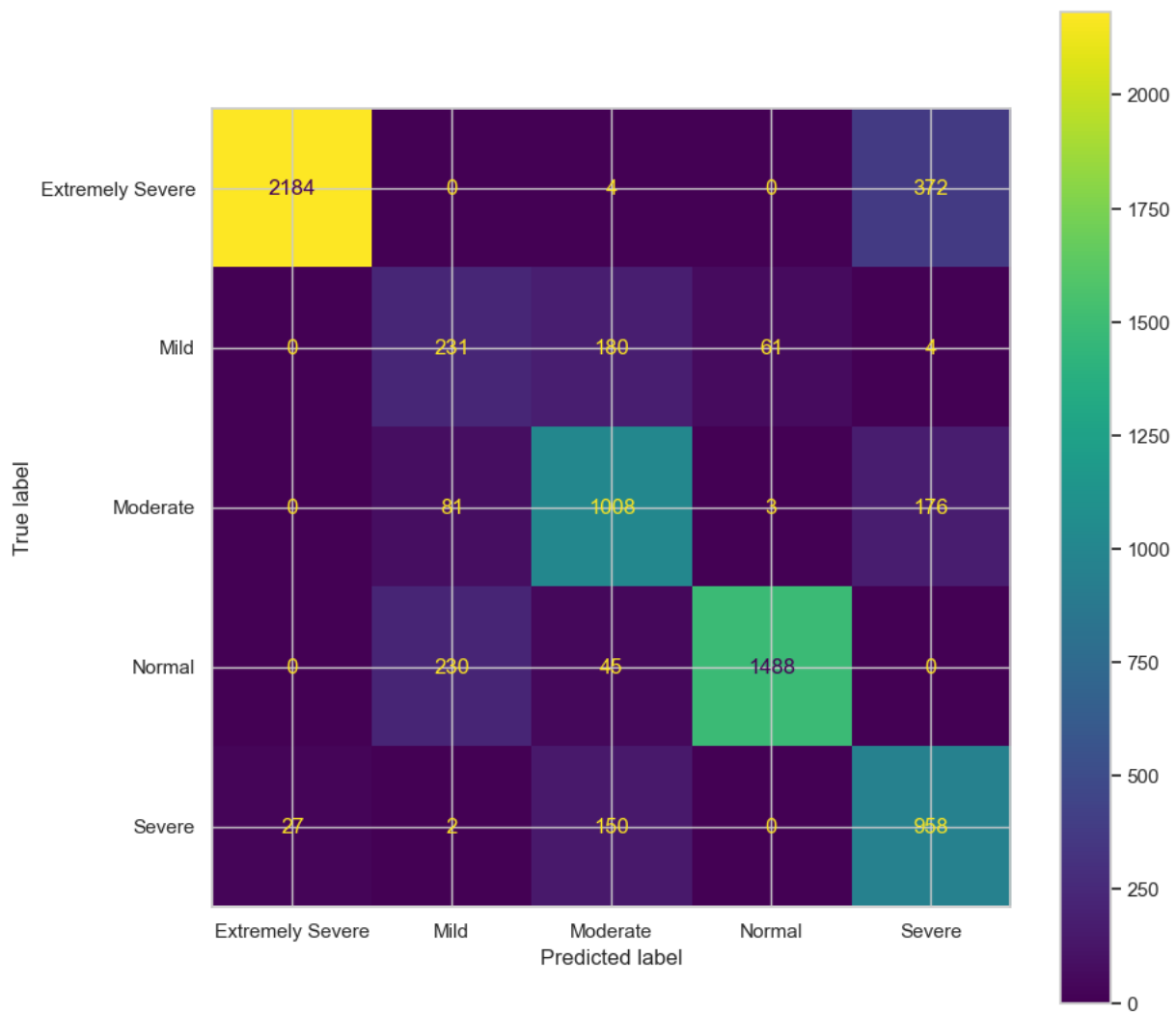
fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(NB,X_test_scaled,y_test,ax=ax)
```

```
Accuracy: 0.81
F1_Score: 0.82
Recall_Score: 0.81
Precision_Score: 0.84
```

		precision	recall	f1-score	support
Extremely	Severe	0.99	0.85	0.92	2560
	Mild	0.42	0.49	0.45	476
	Moderate	0.73	0.79	0.76	1268
	Normal	0.96	0.84	0.90	1763
	Severe	0.63	0.84	0.72	1137
	accuracy			0.81	7204
	macro avg	0.75	0.76	0.75	7204
	weighted avg	0.84	0.81	0.82	7204

```
C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
  warnings.warn(msg, category=FutureWarning)
```

```
Out[76]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508c6b430>
```



Random Forest Classifier

```
In [77]: RF=RandomForestClassifier(n_estimators=200,min_samples_split=3,min_samples_leaf=1,max_depth=160,max_features='auto').fit(X_train,
prediction = RF.predict(X_test_scaled)

aRF3=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aRF3)

f1RF3=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1RF3)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(RF,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.87

F1_Score: 0.84

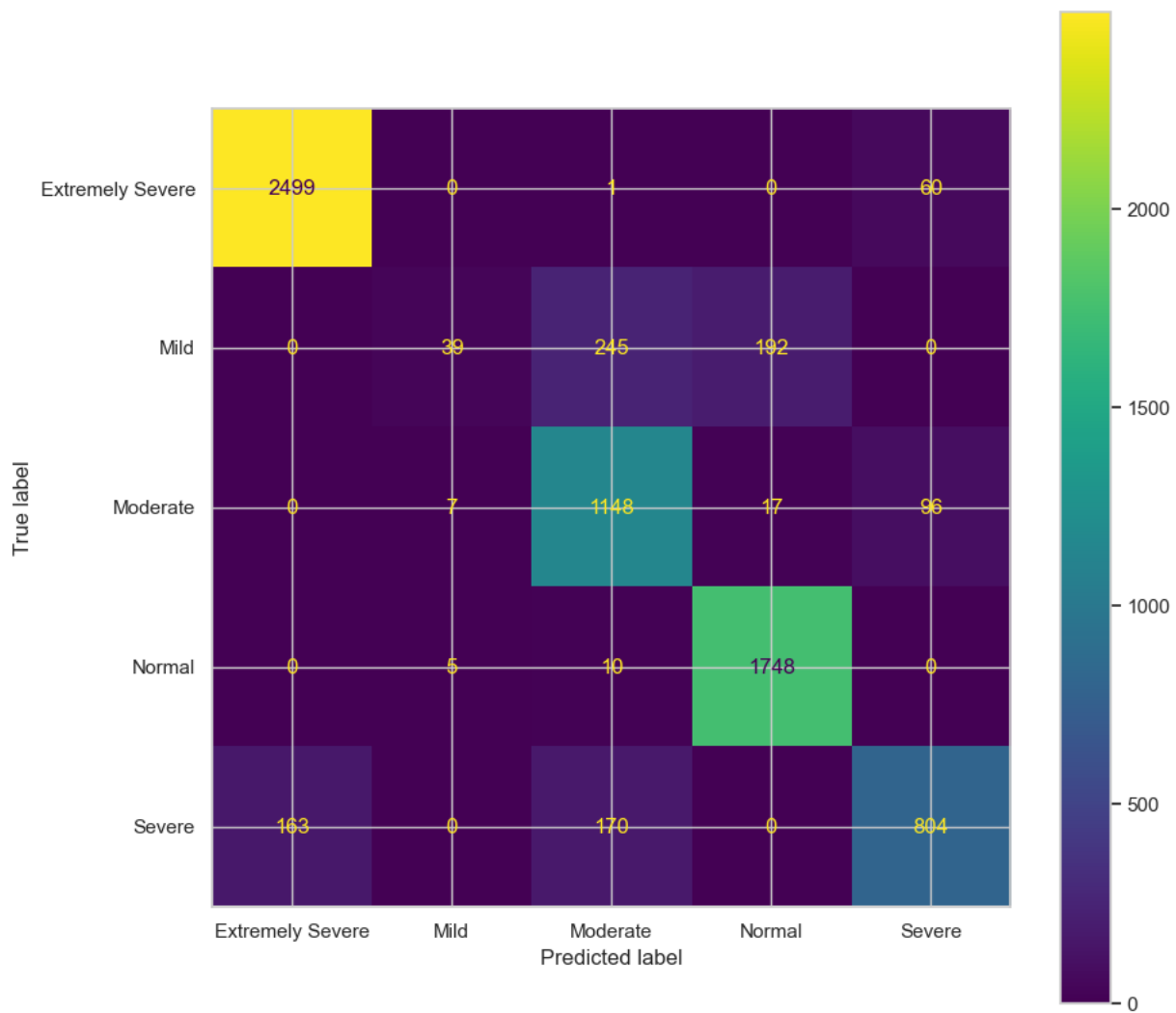
Recall_Score: 0.87

Precision_Score: 0.86

	precision	recall	f1-score	support
Extremely Severe	0.94	0.98	0.96	2560
Mild	0.76	0.08	0.15	476
Moderate	0.73	0.91	0.81	1268
Normal	0.89	0.99	0.94	1763
Severe	0.84	0.71	0.77	1137
accuracy			0.87	7204
macro avg	0.83	0.73	0.72	7204
weighted avg	0.86	0.87	0.84	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[77]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d507ccec0>



AdaBoost (Ensemble Method)

```
In [78]: AB=AdaBoostClassifier(n_estimators=50,learning_rate=1)
AB.fit(X_train_scaled, y_train)
prediction = AB.predict(X_test_scaled)

aAB3=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aAB3)

f1AB3=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1AB3)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(AB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.7

F1_Score: 0.72

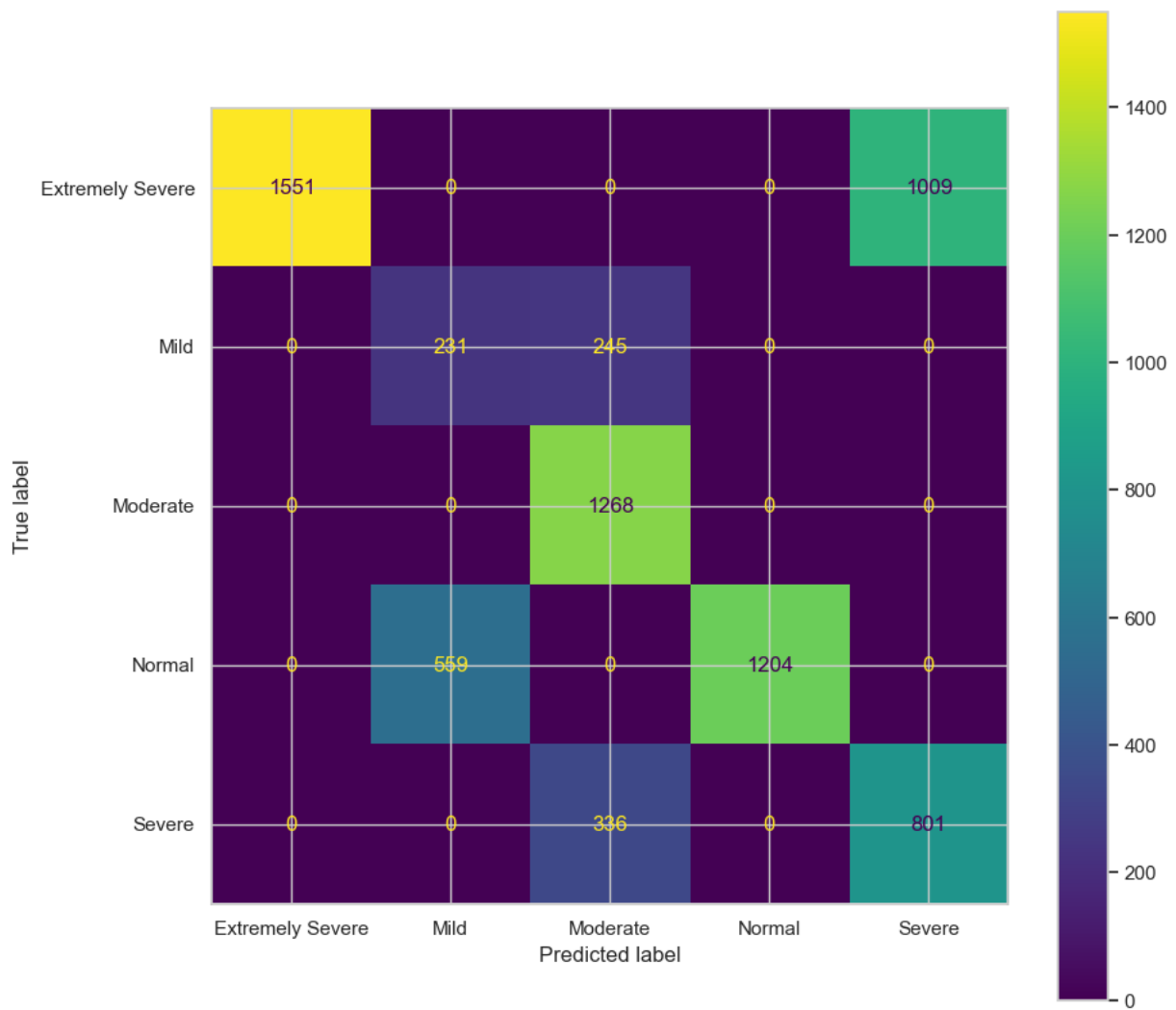
Recall_Score: 0.7

Precision_Score: 0.81

	precision	recall	f1-score	support
Extremely Severe	1.00	0.61	0.75	2560
Mild	0.29	0.49	0.36	476
Moderate	0.69	1.00	0.81	1268
Normal	1.00	0.68	0.81	1763
Severe	0.44	0.70	0.54	1137
accuracy			0.70	7204
macro avg	0.68	0.70	0.66	7204
weighted avg	0.81	0.70	0.72	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[78]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d507db5ee0>



Complete Anxiety modeling summary

```
In [79]: summary={
            'Model': ['GaussianNB', 'Random-Forest', 'AdaBoost'],
            'Accuracy(%)': [aNB3*100, aRF3*100, aAB3*100],
            'F1_Score(%)': [f1NB3*100, f1RF3*100, f1AB3*100],
        }
summaryAnxiety=pd.DataFrame(summary)
summaryAnxiety['key'] = summaryAnxiety.index
summaryAnxiety
```

Out[79]:

	Model	Accuracy(%)	F1_Score(%)	key
0	GaussianNB	81.0	82.0	0
1	Random-Forest	87.0	84.0	1
2	AdaBoost	70.0	72.0	2

Stress dataframe

Split the data into training and testing set

```
In [80]: # Input data to the model
X = Stress0.drop(["cIntensity", "intensity", "score", "major"], axis=1, errors="ignore")
# Ground truth label
y = Stress0["intensity"]

# Splitting the data, training set and testing set
X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.80, random_state=100)
scaler = MinMaxScaler()
X_train_scaled = scaler.fit_transform(X_train)
X_test_scaled = scaler.transform(X_test)

# Train and test set dimensions
print('Training Set:', X_train.shape, y_train.shape)
print('Test Set:', X_test.shape, y_test.shape)
```

Training Set: (28812, 34) (28812,)

Test Set: (7204, 34) (7204,)

Gaussian NB

```
In [81]: NB=GaussianNB().fit(X_train_scaled,y_train)
prediction = NB.predict(X_test_scaled)

aNB4=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aNB4)

f1NB4=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1NB4)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

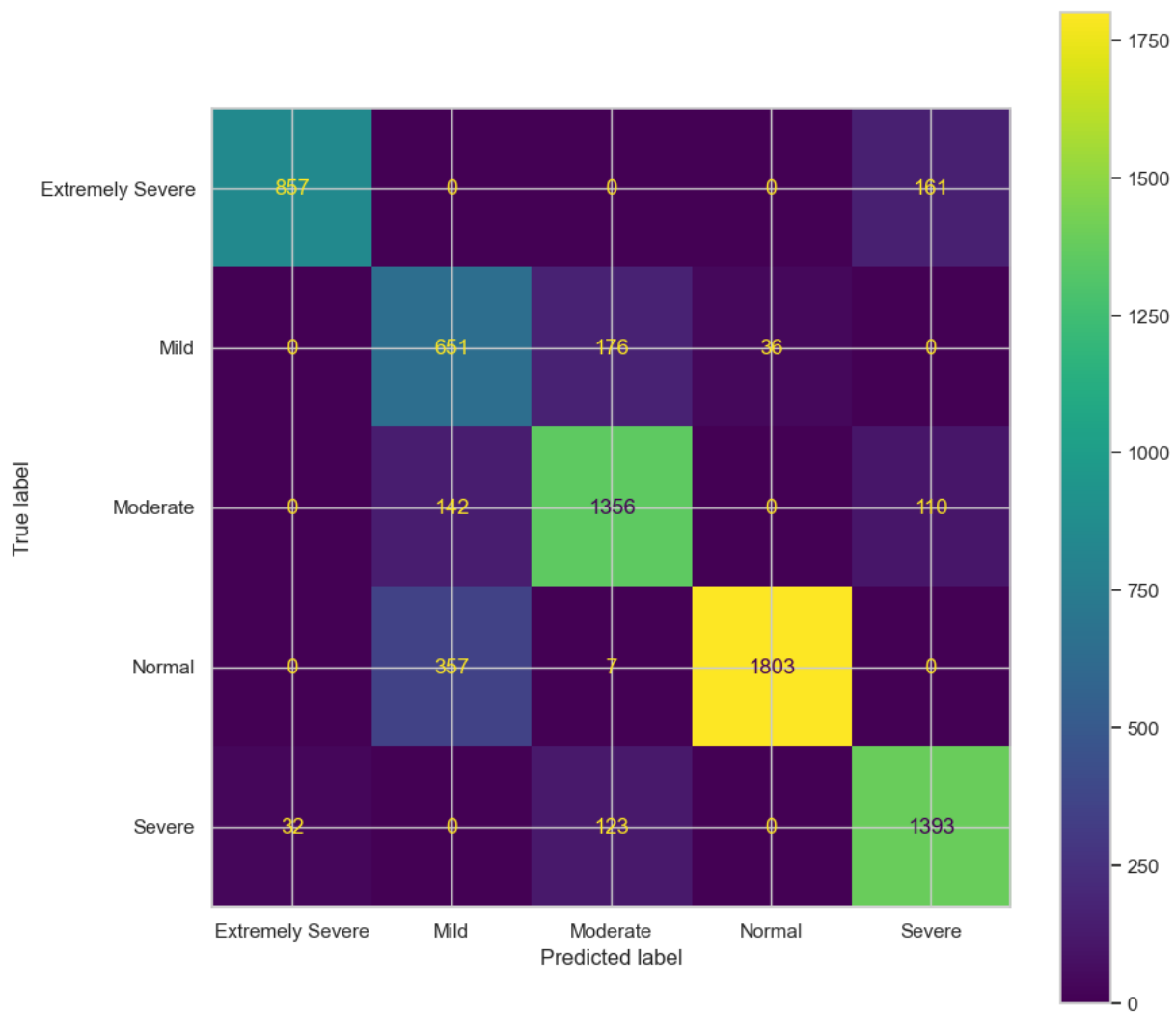
fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(NB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.84
F1_Score: 0.85
Recall_Score: 0.84
Precision_Score: 0.86

		precision	recall	f1-score	support
Extremely	Severe	0.96	0.84	0.90	1018
	Mild	0.57	0.75	0.65	863
	Moderate	0.82	0.84	0.83	1608
	Normal	0.98	0.83	0.90	2167
	Severe	0.84	0.90	0.87	1548
	accuracy			0.84	7204
	macro avg	0.83	0.83	0.83	7204
	weighted avg	0.86	0.84	0.85	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[81]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508708340>



Random Forest Classifier

```
In [82]: RF=RandomForestClassifier(n_estimators=200,min_samples_split=3,min_samples_leaf=1,max_depth=160,max_features='auto').fit(X_train,
prediction = RF.predict(X_test_scaled)

aRF4=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aRF4)

f1RF4=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1RF4)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(RF,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.9

F1_Score: 0.9

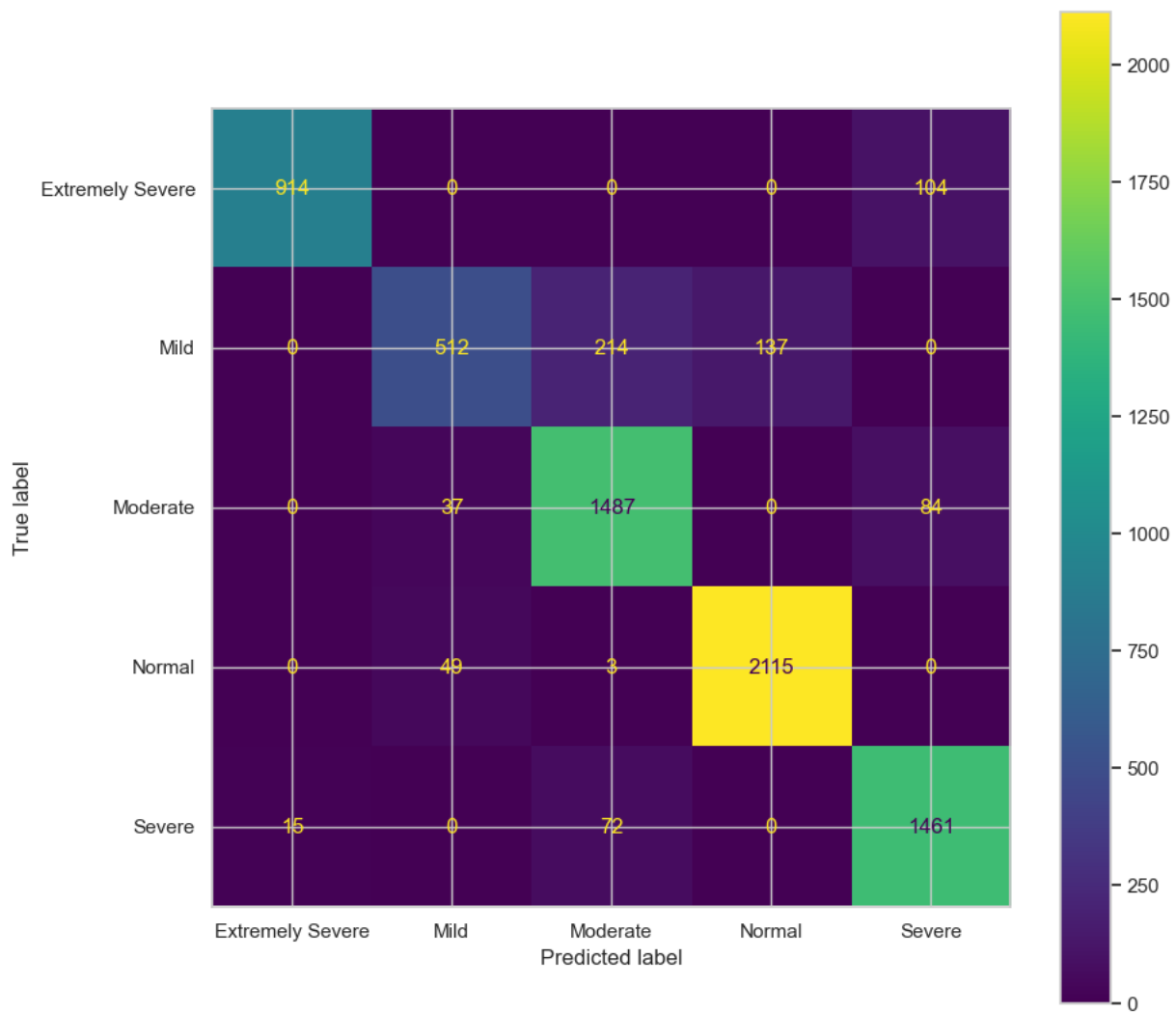
Recall_Score: 0.9

Precision_Score: 0.9

	precision	recall	f1-score	support
Extremely Severe	0.98	0.90	0.94	1018
Mild	0.86	0.59	0.70	863
Moderate	0.84	0.92	0.88	1608
Normal	0.94	0.98	0.96	2167
Severe	0.89	0.94	0.91	1548
accuracy			0.90	7204
macro avg	0.90	0.87	0.88	7204
weighted avg	0.90	0.90	0.90	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[82]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d507c90400>



AdaBoost (Ensemble Method)

```
In [83]: AB=AdaBoostClassifier(n_estimators=50,learning_rate=1)
AB.fit(X_train_scaled, y_train)
prediction = AB.predict(X_test_scaled)

aAB4=round(accuracy_score(y_test,prediction),2)
print('Accuracy:', aAB4)

f1AB4=round(f1_score(y_test,prediction,average='weighted'),2)
print('F1_Score:', f1AB4)

recall=round(recall_score(y_test,prediction,average='weighted'),2)
print('Recall_Score:', recall)

precision=round(precision_score(y_test,prediction,average='weighted'),2)
print('Precision_Score:', precision)

classification=classification_report(
    digits=2,
    y_true=y_test,
    y_pred=prediction)
print(classification)

fig, ax = plt.subplots(figsize=(10, 10))
plot_confusion_matrix(AB,X_test_scaled,y_test,ax=ax)
```

Accuracy: 0.7

F1_Score: 0.71

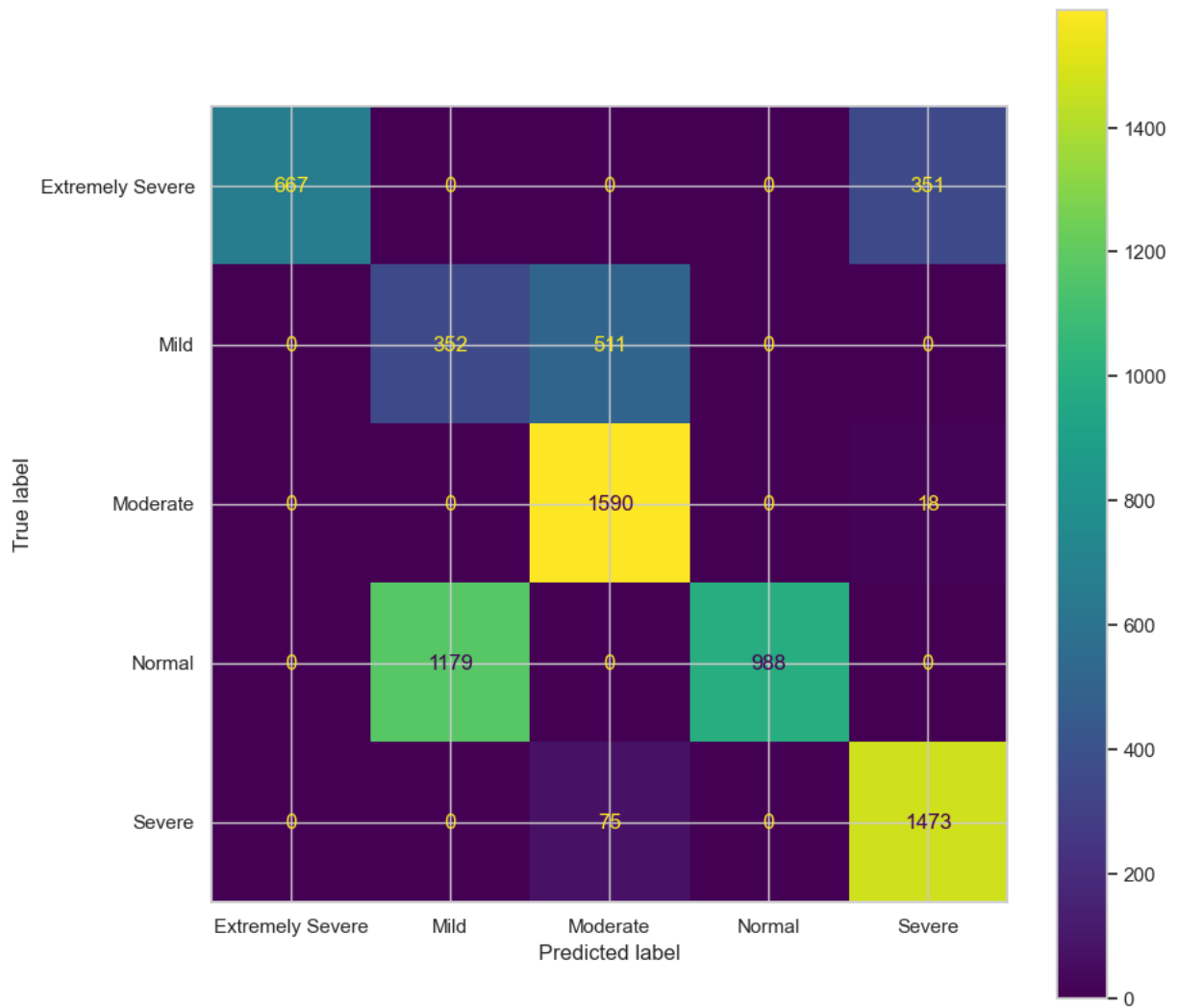
Recall_Score: 0.7

Precision_Score: 0.8

	precision	recall	f1-score	support
Extremely Severe	1.00	0.66	0.79	1018
Mild	0.23	0.41	0.29	863
Moderate	0.73	0.99	0.84	1608
Normal	1.00	0.46	0.63	2167
Severe	0.80	0.95	0.87	1548
accuracy			0.70	7204
macro avg	0.75	0.69	0.68	7204
weighted avg	0.80	0.70	0.71	7204

C:\Users\sandr\anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function 'plot_confusion_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.
warnings.warn(msg, category=FutureWarning)

Out[83]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x1d508911e50>



Complete Stress modeling summary

```
In [84]: summary={
            'Model': ['GaussianNB', 'Random-Forest', 'AdaBoost'],
            'Accuracy(%)': [aNB4*100, aRF4*100, aAB4*100],
            'F1_Score(%)': [f1NB4*100, f1RF4*100, f1AB4*100],
        }
summaryStress=pd.DataFrame(summary)
summaryStress['key'] = summaryStress.index
summaryStress
```

```
Out[84]:
```

	Model	Accuracy(%)	F1_Score(%)	key
0	GaussianNB	84.0	85.0	0
1	Random-Forest	90.0	90.0	1
2	AdaBoost	70.0	71.0	2

Comparison of modeling results on all the dataframes

Accuracy table

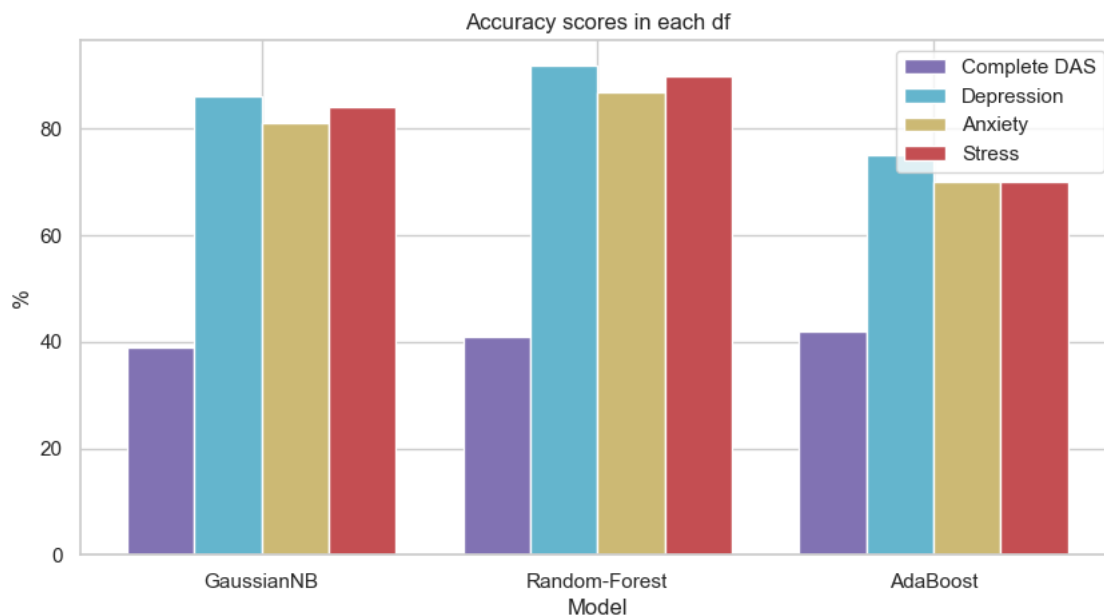
```
In [85]: summary1 = pd.merge(summaryDass,summaryDepression, how='inner', on='key')
summary2 = pd.merge(summaryAnxiety,summaryStress, how='inner', on='key')
summary = pd.merge(summary1,summary2, how='inner', on='key')
summary = summary.drop(columns=["Model_y_x", "key", "Model_x_y", "Model_y_y"], axis=1, errors="ignore")
summary.rename(columns={'Model_x_x': 'Model'}, inplace=True)
summary.rename(columns={'Accuracy(%)_x_x': 'Acc_Das', 'F1_Score(%)_x_x': 'F1_Das'}, inplace=True)
summary.rename(columns={'Accuracy(%)_y_x': 'Acc_Depression', 'F1_Score(%)_y_x': 'F1_Depression'}, inplace=True)
summary.rename(columns={'Accuracy(%)_x_y': 'Acc_Anxiety', 'F1_Score(%)_x_y': 'F1_Anxiety'}, inplace=True)
summary.rename(columns={'Accuracy(%)_y_y': 'Acc_Stress', 'F1_Score(%)_y_y': 'F1_Stress'}, inplace=True)
summaryAccuracy = summary.drop(columns=["F1_Das", "F1_Depression", "F1_Anxiety", "F1_Stress"], axis=1, errors="ignore")
summaryAccuracy
```

Out[85]:

	Model	Acc_Das	Acc_Depression	Acc_Anxiety	Acc_Stress
0	GaussianNB	39.0	86.0	81.0	84.0
1	Random-Forest	41.0	92.0	87.0	90.0
2	AdaBoost	42.0	75.0	70.0	70.0

Accuracy results visualization

```
In [86]: x=['GaussianNB','Random-Forest','AdaBoost']
X=np.arange(len(x))
plt.figure(figsize=(10,5))
bar0=plt.bar(X,summaryDass['Accuracy(%)'],color='m', width = 0.2)
bar1=plt.bar(X+0.2,summaryDepression['Accuracy(%)'],color='c',width = 0.2)
bar2=plt.bar(X+0.4,summaryAnxiety['Accuracy(%)'],color='y',width = 0.2)
bar3=plt.bar(X+0.6,summaryStress['Accuracy(%)'],color='r',width = 0.2)
plt.xticks(X+0.3,x)
plt.legend((bar0, bar1, bar2, bar3),('Complete DAS', 'Depression', 'Anxiety', 'Stress'))
plt.ylabel('%')
plt.xlabel('Model')
plt.title('Accuracy scores in each df')
plt.show()
```



F1 score

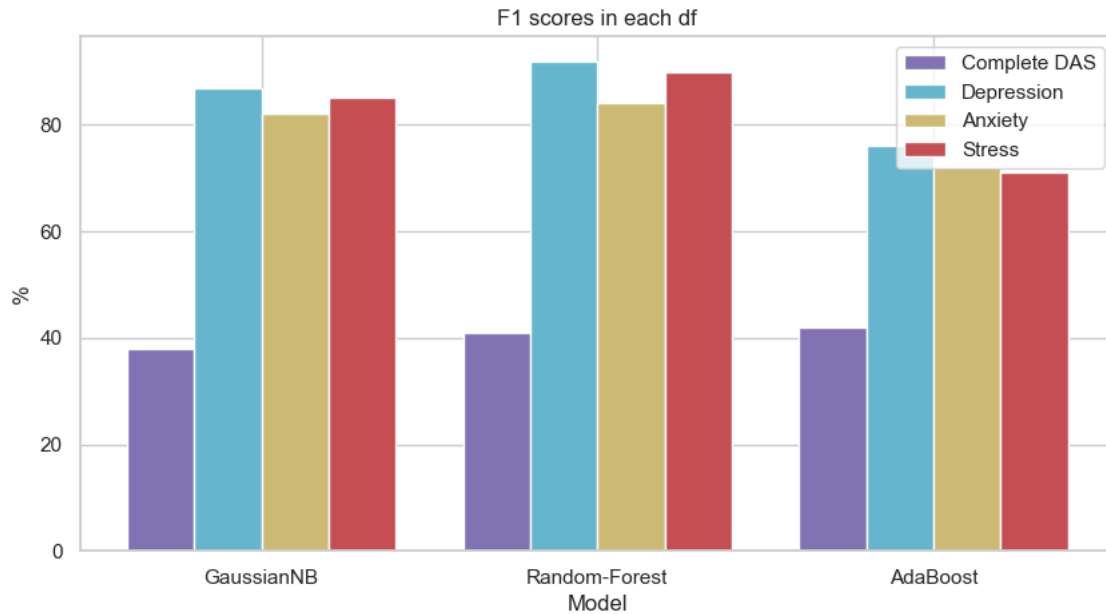
```
In [87]: summaryF1 = summary.drop(columns=["Acc_Das", "Acc_Depression", "Acc_Anxiety", "Acc_Stress"], axis=1, errors="ignore")
summaryF1
```

Out[87]:

	Model	F1_Das	F1_Depression	F1_Anxiety	F1_Stress
0	GaussianNB	38.0	87.0	82.0	85.0
1	Random-Forest	41.0	92.0	84.0	90.0
2	AdaBoost	42.0	76.0	72.0	71.0

F1 score results visualization

```
In [88]: x=['GaussianNB', 'Random-Forest', 'AdaBoost']
X=np.arange(len(x))
plt.figure(figsize=(10,5))
bar0=plt.bar(X,summaryDass['F1_Score(%)'],color='m', width=0.2)
bar1=plt.bar(X+0.2,summaryDepression['F1_Score(%)'],color='c', width=0.2)
bar2=plt.bar(X+0.4,summaryAnxiety['F1_Score(%)'],color='y', width=0.2)
bar3=plt.bar(X+0.6,summaryStress['F1_Score(%)'],color='r', width=0.2)
plt.xticks(X+0.3,x)
plt.legend((bar0, bar1, bar2, bar3),('Complete DAS', 'Depression', 'Anxiety', 'Stress'))
plt.ylabel('%')
plt.xlabel('Model')
plt.title('F1 scores in each df')
plt.show()
```



References

- O. Aran and E. Kapusuz. "Predicting Depression, Anxiety and Stress- EDA". <https://www.kaggle.com/code/orkunaran/predicting-depression-anxiety-and-stress-eda/notebook> (<https://www.kaggle.com/code/orkunaran/predicting-depression-anxiety-and-stress-eda/notebook>), 2021, (accessed Nov. 10, 2022).
- S.H. Lovibond and P.F. Lovibond, "Manual for the Depression Anxiety Stress Scales" *Psychology Foundation*, (2nd ed.), 1995.
- Teju, "DAS PREDICTION". <https://www.kaggle.com/code/teju4405/das-prediction#Races> (<https://www.kaggle.com/code/teju4405/das-prediction#Races>), 2022, (accessed Nov. 10, 2022).