Predict GAD in students

Imports

```
In [27]: import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         from ucimlrepo import fetch_ucirepo
         from sklearn.decomposition import PCA, KernelPCA
         from sklearn.discriminant_analysis import LinearDiscriminantAnalysis as LDA
         from sklearn.preprocessing import StandardScaler
         from sklearn.model_selection import train_test_split
         from sklearn.impute import KNNImputer
         from imblearn.over_sampling import SMOTE, RandomOverSampler
         from imblearn.under_sampling import RandomUnderSampler
         from sklearn.preprocessing import LabelEncoder, OneHotEncoder
         from sklearn.metrics import precision_score, recall_score, f1_score, classification_report, confusion_matrix, roc_curve, roc_auc_score, accuracy_score
         import category_encoders as ce
         import seaborn as sns
         import pandas as pd
         import matplotlib.pyplot as plt
         import seaborn as sns
         # Setting seed to last 3 digits of my student ID as random seed for reproducibility
         seed value = 417
         pd.set_option('display.max_columns', None)
```

Preprocessing

```
In [31]: # fetch dataset
         # Rename cols
         raw df = pd.read csv('dataset.csv')
         df = raw_df.copy()
         col names map = {
           '1. Have you read Participant Information Sheet?': 'c1',
           '2. Do you understand you won\'t be able to withdraw consent because of anonymised data?': 'c2',
           '3. Are you aged 18 years or above?': 'c3',
           '4. Do you agree to take part in questionnaire out of your own free will?': 'c4',
           '5. Do you have a part-time job or full time job?': 'have_job',
           '6. Are you an international student?': 'international student',
           '7. Do you have to support your family financially?': 'support family financially',
           '8. How do you rate your financial worries?': 'financial_worries',
           '9. How satisfied you are with your living arrangements?': 'living_arrangements',
           '10. How do you rate your current level of academic stress?': 'academic_stress',
           '11. How do you feel about your workload this or last semester?': 'workload',
           '12. How often do you feel supported by your friends?': 'supported_by_friends',
           '13. How often do you feel supported by your family?': 'supported_by_family',
           '14. How often do you seek support from faculty or staff?': 'seek staff support',
           '15. How many hours of sleep do you typically get per night?': 'sleep time',
           '16. How often do you exercise per week? (Any kind of exercise counts walking, jogging, gym etc)': 'exercise',
```

```
'17. How would you rate your overall diet?': 'diet',
  '18. What is your degree level?': 'degree level',
  '19. Have you ever faced discriminatory treatment from any of the student or University staff?': 'faced discrimination',
  '20. Are you involved in sports? (Any sports activity at least twice a month)': 'involved_in_sports',
  '21. Do you have a circle of friends you like?': 'friends_circle',
  '22. What are your drinking habits? (Alcoholic drinks)': 'drinking habits',
  '23. What is your Gender?': 'gender',
  '24. Enter your age range?': 'age_range',
  '25. Do you suffer from any chronic Physical health Problem?': 'chronic_health_problem',
  '26. Which study year you are in?': 'study_year',
  '27. How satisfied have you been with your overall well-being over the past two months?': 'overall_well_being',
  '28. During the PAST 7 DAYS, I have...': 'gad_0',
  '28.1. felt moments of sudden terror, fear or fright': 'gad_1',
  '28.2. felt anxious, worried, or nervous': 'gad_2',
  '28.3. had thoughts of bad things happening, such as family tragedy, ill health, loss of a job, or accidents': 'gad_3',
  '28.4. felt a racing heart, sweaty, trouble breathing, faint, or shaky': 'gad_4',
  '28.5. felt tense muscles, felt on edge or restless, or had trouble relaxing or trouble sleeping': 'gad_5',
  '28.6. avoided, or did not approach or enter, situations about which I worry': 'gad_6',
  '28.7. left situations early or participated only minimally due to worries': 'gad 7',
  '28.8. spent lots of time making decisions, putting off making decisions, or preparing for situations, due to worries': 'gad_8',
  '28.9. sought reassurance from others due to worries': 'gad_9',
  '28.10. needed help to cope with anxiety (e.g., alcohol or medication, superstitious objects, or other people)': 'gad_10'
# Rename cols and Remove consent cols
# Overall well being col is dropped
df = df.rename(columns = col names map)
# Apply the strip function to all values in the DataFrame
df = df.applymap(lambda x: x.strip() if isinstance(x, str) else x)
# Drop consent columns and an empty column with no data
df = df.drop(['c1', 'c2', 'c3', 'c4', 'gad_0'], axis=1)
categorical cols = [
  'gender', 'study_year', 'degree_level', 'drinking_habits', 'diet', 'sleep_time', 'age_range']
binary_columns = [
  'have job', 'international student', 'support family financially', 'faced discrimination',
  'involved_in_sports', 'friends_circle', 'chronic_health_problem']
ordinal categorical cols = [
  'financial_worries', 'living_arrangements', 'academic_stress', 'suppported_by_friends',
  'supported_by_family', 'seek_staff_support', 'exercise', 'overall_well_being', 'workload']
```

Remove missing values rows

```
In [32]: # remove missing values rows
print(len(df) - len(df.dropna()), "rows deleted because of missing values.")
df = df.dropna()
```

2 rows deleted because of missing values.

Map GAD column to the respective score and calculate gad score

Then convert the score to respective categories of GAD

```
In [33]: # Gad map
         gad_map = {'Never': 0, 'Occasionally': 1, 'Half of the time': 2, 'Most of the time': 3, 'All of the time': 4}
         for i in range(1,11):
           df[f'gad_{i}'] = df[f'gad_{i}'].map(gad_map)
         # Calculate gad score and covert to classification bins
         df['gad\_score'] = df['gad\_1'] + df['gad\_2'] + df['gad\_3'] + df['gad\_4'] + df['gad\_5'] + df['gad\_6'] + df['gad\_7'] + df['gad\_8'] + df['gad\_9'] + df['gad\_10']
         df.drop(['gad_{}'.format(i) for i in range(1,11)], axis=1, inplace=True)
         # get gad_sum and del GAD columns
         # Define the bins and labels
         bins = [0, 14, 24, 40]
         labels = [0, 1, 2]
         df['gad_scale'] = pd.cut(df['gad_score'], bins=bins, labels=labels, right=True, include_lowest=True).astype(int)
         df.drop(['gad_score'], axis=1, inplace=True)
         df['gad_scale'].value_counts()
         # Create copy for exploratory analysis
         edf = df.copy()
         edf['gad_scale'] = edf['gad_scale'].map({0:'None-Mild', 1: 'Moderate', 2: 'Severe-Extreme'})
         edf['gad_scale'].value_counts()
Out[33]: None-Mild
                            2964
         Moderate
                            884
         Severe-Extreme
                            416
         Name: gad_scale, dtype: int64
In [35]: for column in df.columns:
             print(column, '\t', df[column].unique())
```

```
have_job
                 ['No' 'Yes']
international student
                        ['Yes' 'No']
support family financially
                                 ['Yes' 'No']
                         ['Always' 'Sometimes' 'Often' 'Rarely' 'Never worried']
financial_worries
                         ['Satisfied' 'Not satisfied' 'Very dissatisfied' 'Extremely satisfied']
living_arrangements
                         ['Sometimes stressed' 'Often stressed' 'Always stressed' 'Rarely stressed']
academic stress
                 ['Just right' 'Too heavy' 'Too light']
workload
                       ['Always' 'Often' 'Sometimes' 'Rarely']
suppported_by_friends
                         ['Always' 'Sometimes' 'Often' 'Rarely']
supported_by_family
                         ['Rarely' 'Sometimes' 'Often' 'Always']
seek_staff_support
sleep_time
                 ['6-8 hours' 'Less than 6 hours' 'More than 8 hours']
                 ['0 times' '3-5 times' '1-2 times' '6 or more times']
exercise
        ['Moderately healthy' 'Healthy' 'Unhealthy']
diet
                ['Undergraduate' 'Postgraduate' 'Phd']
degree_level
faced discrimination
                        ['No' 'Yes']
involved_in_sports
                         ['No' 'Yes']
friends_circle ['Yes' 'No']
drinking_habits
                         ['Never' 'Occasionally' 'Regularly' 'Only Weekends']
gender ['Male' 'Female' 'Non-binary' 'Other']
                 ['21-25' '18-20' '26-30' 'Above 30']
age range
chronic_health_problem ['Yes' 'No']
study_year
                 ['Second' 'Third or above' 'First']
                        ['Satisfied' 'Dissatisfied' 'Neutral' 'Very dissatisfied'
overall well being
'Extremely satisfied']
gad_scale
                 [0 1 2]
```

Map binary Variables and ordinal categories

```
In [6]: for col in binary columns:
          df[col] = df[col].map({'Yes': 1, 'No':0})
        ordinal mapping = {
          'financial worries': {'Never worried':0, 'Rarely':1, 'Sometimes':2, 'Often':3, 'Always':4 },
          'living_arrangements': {'Satisfied': 1, 'Extremely satisfied': 0, 'Not satisfied': 2, 'Very dissatisfied': 3},
          'academic_stress': {'Always stressed':3, 'Sometimes stressed':1, 'Often stressed':2, 'Rarely stressed': 0},
          'suppported by friends': {'Always':3, 'Often':2, 'Rarely':1, 'Sometimes': 0},
          'supported_by_family': {'Always':3, 'Often':2, 'Rarely':1, 'Sometimes': 0},
          'seek_staff_support': {'Always':3,'Rarely':0, 'Sometimes':1, 'Often':2},
          'exercise': {'0 times': 0, '3-5 times':2, '1-2 times':1, '6 or more times':3},
          'overall well being': {'Satisfied': 1, 'Extremely satisfied': 0 ,'Neutral': 2, 'Dissatisfied': 3, 'Very dissatisfied': 4},
          'workload': {'Too heavy':2, 'Just right':0, 'Too light': 1}
        for col in ordinal_categorical_cols:
          for key in df[col].unique():
           if key not in ordinal_mapping[col]:
              raise AssertionError('Key {} in not mapped for column {}'.format(key, col))
          df[col] = df[col].map(ordinal_mapping[col])
```

Pedregosa, F., et al. (2011). "Scikit-learn: Machine Learning in Python." Journal of Machine Learning Research, 12, 2825-2830.

This foundational paper on the scikit-learn library explains various preprocessing techniques, including the treatment of ordinal data. It highlights how ordinal encoding can preserve the order of categories and provide meaningful numerical representations for machine learning algorithms.

Encode the categorical columns with one hot encoding

```
In [8]: import pandas as pd

# Apply one-hot encoding
df_encoded = pd.get_dummies(df, columns=categorical_cols)
```

Split in train test

Sample balancing using SMOTE and Random OverSampling

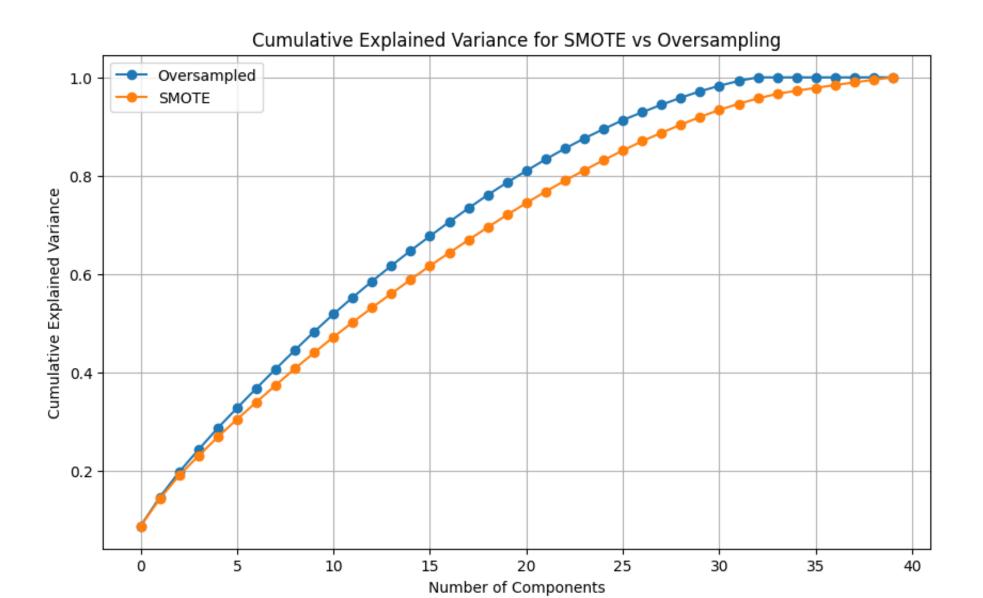
```
In [37]: import pandas as pd
         from sklearn.model_selection import train_test_split
         from imblearn.over sampling import RandomOverSampler, SMOTE
         # Define the features (X) and target (y)
         X = df_encoded.drop('gad_scale', axis=1)
         y = df encoded['gad scale']
         # Split the data into training and testing sets
         X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=seed_value)
         # Apply Random Oversampling
         ros = RandomOverSampler(random_state=seed_value)
         X_train_ros, y_train_ros = ros.fit_resample(X_train, y_train)
         # Apply SMOTE
         smote = SMOTE(random_state=seed_value)
         X_train_smote, y_train_smote = smote.fit_resample(X_train, y_train)
         # Store the balanced datasets in a dictionary
         balanced_datasets = {
             'oversampled': {
                 'X_train': X_train_ros,
                 'y_train': y_train_ros,
                 'X_test': X_test,
                 'y_test': y_test
             },
             'smote': {
                 'X_train': X_train_smote,
                 'y_train': y_train_smote,
                 'X_test': X_test,
                 'y_test': y_test
         # Display the shapes of the resulting DataFrames
         print("Oversampled Data:")
         print(f"X_train shape: {balanced_datasets['oversampled']['X_train'].shape}")
         print(f"y_train shape: {balanced_datasets['oversampled']['y_train'].shape}")
         print("Class distribution in y_train after Random Oversampling:")
         print(balanced_datasets['oversampled']['y_train'].value_counts())
         print("\nSMOTE Data:")
         print(f"X_train shape: {balanced_datasets['smote']['X_train'].shape}")
         print(f"y_train shape: {balanced_datasets['smote']['y_train'].shape}")
         print("Class distribution in y_train after SMOTE:")
         print(balanced_datasets['smote']['y_train'].value_counts())
```

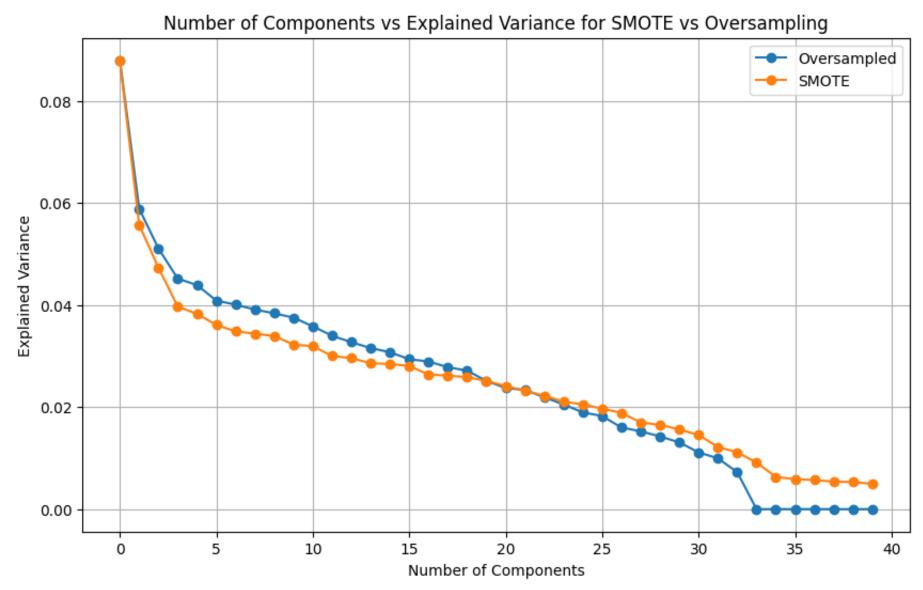
```
Oversampled Data:
X_train shape: (7149, 40)
y_train shape: (7149,)
Class distribution in y_train after Random Oversampling:
    2383
    2383
1
2
    2383
Name: gad_scale, dtype: int64
SMOTE Data:
X_train shape: (7149, 40)
y_train shape: (7149,)
Class distribution in y_train after SMOTE:
    2383
    2383
1
    2383
Name: gad_scale, dtype: int64
```

Standardizd and Apply PCA

```
In [39]: import pandas as pd
         import matplotlib.pyplot as plt
         from sklearn.decomposition import PCA
         from sklearn.preprocessing import StandardScaler
         # Standardize the features before applying PCA
         scaler = StandardScaler()
         X_train_ros_scaled = scaler.fit_transform(balanced_datasets['oversampled']['X_train'])
         X_train_smote_scaled = scaler.fit_transform(balanced_datasets['smote']['X_train'])
         X_test_scaled = scaler.transform(balanced_datasets['smote']['X_test']) # Scaling test data
         # Apply PCA
         pca ros = PCA()
         X_train_ros_pca = pca_ros.fit_transform(X_train_ros_scaled)
         pca_smote = PCA()
         X_train_smote_pca = pca_smote.fit_transform(X_train_smote_scaled)
         # Calculate cumulative explained variance
         cumulative_explained_variance_ros = pca_ros.explained_variance_ratio_.cumsum()
         cumulative_explained_variance_smote = pca_smote.explained_variance_ratio_.cumsum()
         # Plot cumulative explained variance
         plt.figure(figsize=(10, 6))
         plt.plot(cumulative_explained_variance_ros, label='Oversampled', marker='o')
         plt.plot(cumulative explained variance smote, label='SMOTE', marker='o')
         plt.xlabel('Number of Components')
         plt.ylabel('Cumulative Explained Variance')
         plt.title('Cumulative Explained Variance for SMOTE vs Oversampling')
         plt.legend()
         plt.grid(True)
         plt.show()
         # Plot number of components vs explained variance
         plt.figure(figsize=(10, 6))
```

```
plt.plot(pca_ros.explained_variance_ratio_, label='Oversampled', marker='o')
plt.plot(pca_smote.explained_variance_ratio_, label='SMOTE', marker='o')
plt.xlabel('Number of Components')
plt.ylabel('Explained Variance')
plt.title('Number of Components vs Explained Variance for SMOTE vs Oversampling')
plt.legend()
plt.grid(True)
plt.show()
# Decide on the number of components to keep (example: 95% variance)
n_{components} = 0.95
# Determine number of components for 95% variance explained
n_components_ros = (cumulative_explained_variance_ros >= n_components).argmax() + 1
n_components_smote = (cumulative_explained_variance_smote >= n_components).argmax() + 1
# Apply PCA with the chosen number of components
pca_ros_final = PCA(n_components=n_components_ros)
X train ros pca = pca ros final.fit transform(X train ros scaled)
X_test_ros_pca = pca_ros_final.transform(X_test_scaled) # Transform test data
pca_smote_final = PCA(n_components=n_components_smote)
X_train_smote_pca = pca_smote_final.fit_transform(X_train_smote_scaled)
X_test_smote_pca = pca_smote_final.transform(X_test_scaled) # Transform test data
# Store the PCA-transformed datasets in the dictionary
balanced_datasets['oversampled']['X_train_pca'] = X_train_ros_pca
balanced_datasets['oversampled']['X_test_pca'] = X_test_ros_pca
balanced_datasets['smote']['X_train_pca'] = X_train_smote_pca
balanced_datasets['smote']['X_test_pca'] = X_test_smote_pca
# Display the number of components chosen for 95% variance explained
print(f"Number of components for 95% variance (Oversampled): {n_components_ros}")
print(f"Number of components for 95% variance (SMOTE): {n_components_smote}")
```





Number of components for 95% variance (Oversampled): 29 Number of components for 95% variance (SMOTE): 33

Apply all classification algorithms and find best parameters by grid search.

```
In [19]: import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         from sklearn.svm import SVC
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier, ExtraTreesClassifier
         from sklearn.model selection import GridSearchCV, cross val score
         from sklearn.metrics import classification_report, confusion_matrix, roc_curve, auc, roc_auc_score
         import seaborn as sns
         # Define classifiers and their respective parameter grids
         classifiers = {
              'SVM': SVC(probability=True),
             'KNN': KNeighborsClassifier(),
             'RandomForest': RandomForestClassifier(),
             'AdaBoost': AdaBoostClassifier(),
              'ExtraTrees': ExtraTreesClassifier()
         param grids = {
              'SVM': {'C': [0.1, 1, 10], 'kernel': ['linear', 'rbf']},
              'KNN': {'n_neighbors': [3, 5, 7], 'weights': ['uniform', 'distance']},
              'RandomForest': {'n_estimators': [100, 200], 'max_depth': [None, 10, 20]},
             'AdaBoost': {'n_estimators': [50, 100], 'learning_rate': [0.01, 0.1, 1]},
              'ExtraTrees': {'n_estimators': [100, 200], 'max_depth': [None, 10, 20]}
         # Store the best models for each technique
         best models = {'oversampled': {}, 'smote': {}}
         # Hyperparameter tuning with GridSearchCV for each dataset
         print("Best models and their parameters:")
         for technique in ['oversampled', 'smote']:
             for name, clf in classifiers.items():
                 grid_search = GridSearchCV(clf, param_grids[name], cv=10, scoring='accuracy', n_jobs=-1)
                 grid search.fit(balanced datasets[technique]['X train pca'], balanced datasets[technique]['y train'])
                 best models[technique] [name] = grid search.best estimator
                 print(f"{technique} - {name}: {grid_search.best_estimator_.get_params()}")
```

```
Best models and their parameters:
oversampled - SVM: {'C': 10, 'break_ties': False, 'cache_size': 200, 'class_weight': None, 'coef0': 0.0, 'decision_function_shape': 'ovr', 'degree': 3, 'gamma': 'scale',
'kernel': 'rbf', 'max iter': -1, 'probability': True, 'random state': None, 'shrinking': True, 'tol': 0.001, 'verbose': False}
oversampled - KNN: {'algorithm': 'auto', 'leaf_size': 30, 'metric': 'minkowski', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 3, 'p': 2, 'weights': 'distance'}
oversampled - RandomForest: {'bootstrap': True, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'gini', 'max_depth': None, 'max_features': 'auto', 'max_leaf_nodes':
None, 'max samples': None, 'min impurity decrease': 0.0, 'min samples leaf': 1, 'min samples split': 2, 'min weight fraction leaf': 0.0, 'n estimators': 100, 'n jobs': N
one, 'oob_score': False, 'random_state': None, 'verbose': 0, 'warm_start': False}
oversampled - AdaBoost: {'algorithm': 'SAMME.R', 'base_estimator': None, 'learning_rate': 1, 'n_estimators': 100, 'random_state': None}
oversampled - ExtraTrees: {'bootstrap': False, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'gini', 'max_depth': 20, 'max_features': 'auto', 'max_leaf_nodes': No
ne, 'max_samples': None, 'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0, 'n_estimators': 200, 'n_jobs': None
e, 'oob_score': False, 'random_state': None, 'verbose': 0, 'warm_start': False}
smote - SVM: {'C': 10, 'break_ties': False, 'cache_size': 200, 'class_weight': None, 'coef0': 0.0, 'decision_function_shape': 'ovr', 'degree': 3, 'gamma': 'scale', 'kern
el': 'rbf', 'max_iter': -1, 'probability': True, 'random_state': None, 'shrinking': True, 'tol': 0.001, 'verbose': False}
smote - KNN: {'algorithm': 'auto', 'leaf_size': 30, 'metric': 'minkowski', 'metric_params': None, 'n_jobs': None, 'n_neighbors': 5, 'p': 2, 'weights': 'distance'}
smote - RandomForest: {'bootstrap': True, 'ccp_alpha': 0.0, 'class_weight': None, 'criterion': 'gini', 'max_depth': 20, 'max_features': 'auto', 'max_leaf_nodes': None, '
max_samples': None, 'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0, 'n_estimators': 200, 'n_jobs': None, 'o
ob_score': False, 'random_state': None, 'verbose': 0, 'warm_start': False}
smote - AdaBoost: {'algorithm': 'SAMME.R', 'base_estimator': None, 'learning_rate': 1, 'n_estimators': 100, 'random_state': None}
smote - ExtraTrees: {'bootstrap': False, 'ccp alpha': 0.0, 'class weight': None, 'criterion': 'gini', 'max depth': None, 'max features': 'auto', 'max leaf nodes': None,
'max_samples': None, 'min_impurity_decrease': 0.0, 'min_samples_leaf': 1, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0, 'n_estimators': 200, 'n_jobs': None, '
oob_score': False, 'random_state': None, 'verbose': 0, 'warm_start': False}
```


Cross Validation

```
In [20]: # Store results
         results = []
         # Evaluate models and collect results for each technique
         for technique in ['oversampled', 'smote']:
             for name, model in best_models[technique].items():
                 # 10-fold cross-validation scores
                 cross_val_scores = cross_val_score(
                     model,
                     balanced_datasets[technique]['X_train_pca'],
                     balanced_datasets[technique]['y_train'],
                     cv=10,
                     scoring='accuracy'
                 print(f"{technique} - {name} Cross-validation accuracy: {cross_val_scores.mean():.2f} (+/- {cross_val_scores.std() * 2:.2f})")
                 # Evaluation on test set
                 result = evaluate_model(
                     f"{name} ({technique})", model,
                     balanced_datasets[technique]['X_train_pca'],
                     balanced_datasets[technique]['y_train'],
                     balanced_datasets[technique]['X_test_pca'],
                     balanced_datasets[technique]['y_test']
                 results.append(result)
        oversampled - SVM Cross-validation accuracy: 0.98 (+/- 0.01)
```

oversampled - KNN Cross-validation accuracy: 0.92 (+/- 0.02) oversampled - RandomForest Cross-validation accuracy: 0.99 (+/- 0.02) oversampled - AdaBoost Cross-validation accuracy: 0.67 (+/- 0.02) oversampled - ExtraTrees Cross-validation accuracy: 0.99 (+/- 0.02) smote - SVM Cross-validation accuracy: 0.94 (+/- 0.13) smote - KNN Cross-validation accuracy: 0.91 (+/- 0.07) smote - RandomForest Cross-validation accuracy: 0.89 (+/- 0.24) smote - AdaBoost Cross-validation accuracy: 0.74 (+/- 0.18) smote - ExtraTrees Cross-validation accuracy: 0.89 (+/- 0.28)

Visualise Results

```
In [23]: # Function to plot confusion matrix
         def plot_confusion_matrix(cm, class_names, title='Confusion Matrix'):
             plt.figure(figsize=(8, 6))
             sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', xticklabels=class_names, yticklabels=class_names)
             plt.title(title)
             plt.xlabel('Predicted')
             plt.ylabel('True')
             plt.show()
         # Function to plot ROC curve
         def plot_roc_curve(y_test, y_prob, class_names, title='ROC Curve'):
             plt.figure(figsize=(10, 8))
             for i, class_name in enumerate(class_names):
                 fpr, tpr, _ = roc_curve(y_test, y_prob[:, i], pos_label=i)
                 roc auc = auc(fpr, tpr)
                 plt.plot(fpr, tpr, label=f'{class name} (AUC = {roc auc:.2f})')
             plt.plot([0, 1], [0, 1], 'k--')
             plt.xlabel('False Positive Rate')
             plt.ylabel('True Positive Rate')
             plt.title(title)
             plt.legend(loc='lower right')
             plt.grid()
             plt.show()
         # Plot graphs for top 3 performing models
         top_results = sorted(results, key=lambda x: x['accuracy'], reverse=True)[:10]
         for result in top_results:
             model_name = result['model_name']
             y_test = result['y_test']
             y_pred = result['y_pred']
             y_prob = result['y_prob']
             print(f'Performance for {model_name}:')
             print(f'Accuracy: {result["accuracy"]:.2f}')
             print(classification_report(y_test, y_pred, target_names=["None-Mild", "Moderate", "Severe-Extreme"]))
             # Confusion matrix
             cm = confusion_matrix(y_test, y_pred)
             plot_confusion_matrix(cm, class_names=["None-Mild", "Moderate", "Severe-Extreme"], title=f'{model_name} Confusion Matrix')
             # ROC Curve
             if y prob is not None:
                 plot_roc_curve(y_test, y_prob, class_names=["None-Mild", "Moderate", "Severe-Extreme"], title=f'{model_name} ROC Curve')
        Performance for SVM (smote):
        Accuracy: 0.91
                        precision
                                     recall f1-score support
            None-Mild
                             0.91
                                       0.98
                                                 0.95
                                                            581
                                                            181
              Moderate
                             0.93
                                       0.73
                                                 0.82
```

Severe-Extreme

accuracy

macro avq

weighted avg

0.87

0.91

0.91

0.84

0.85

0.91

0.85

0.91

0.87

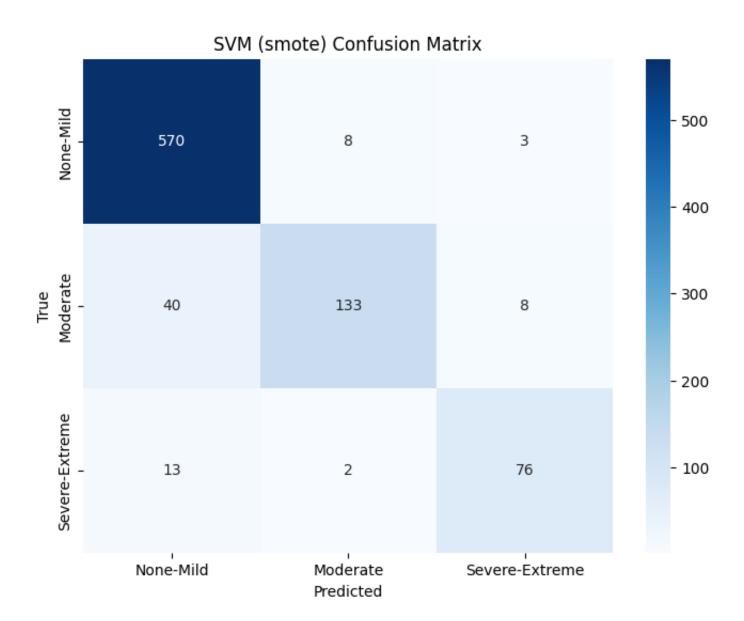
0.91

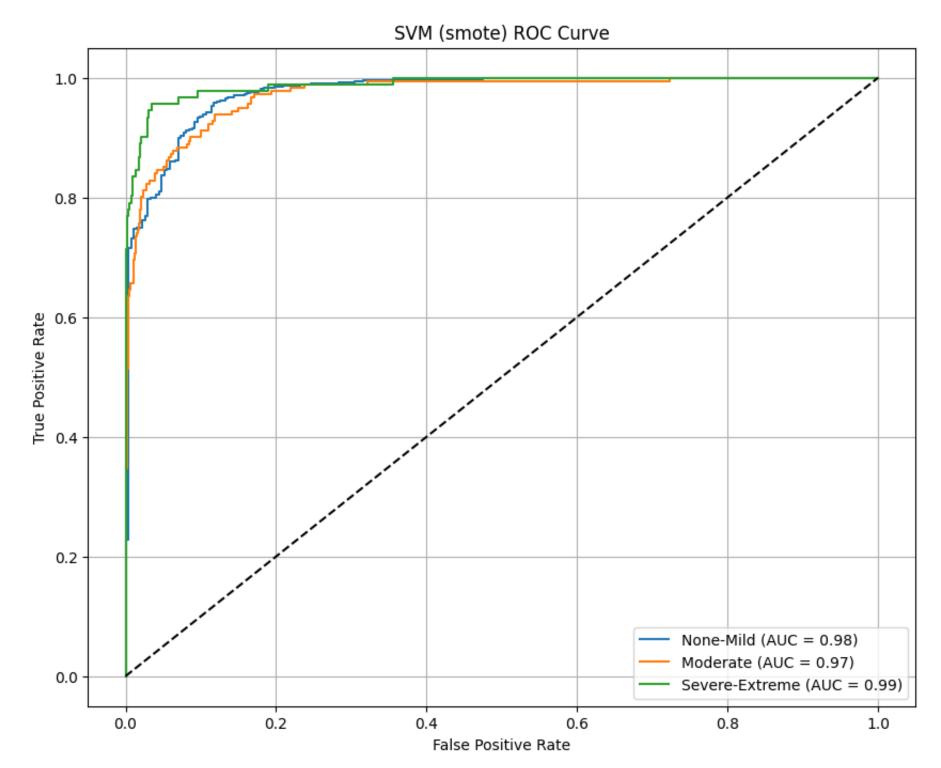
91

853

853

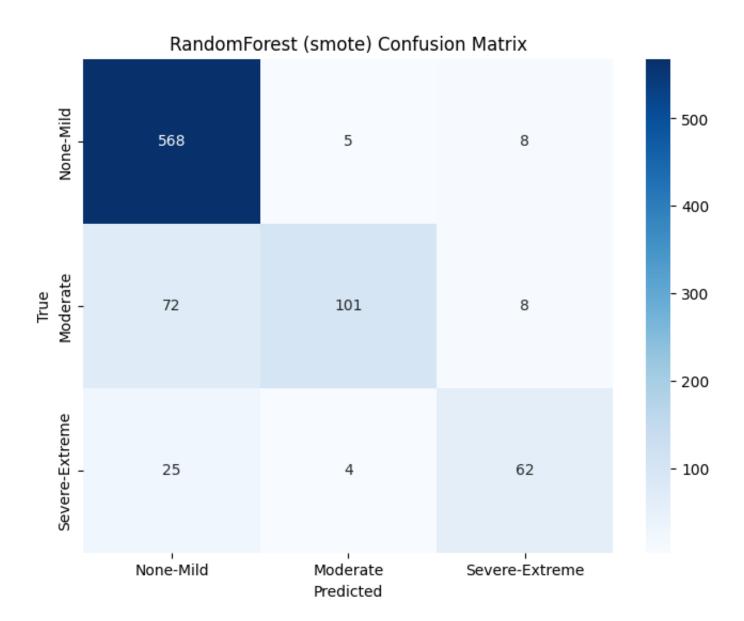
853

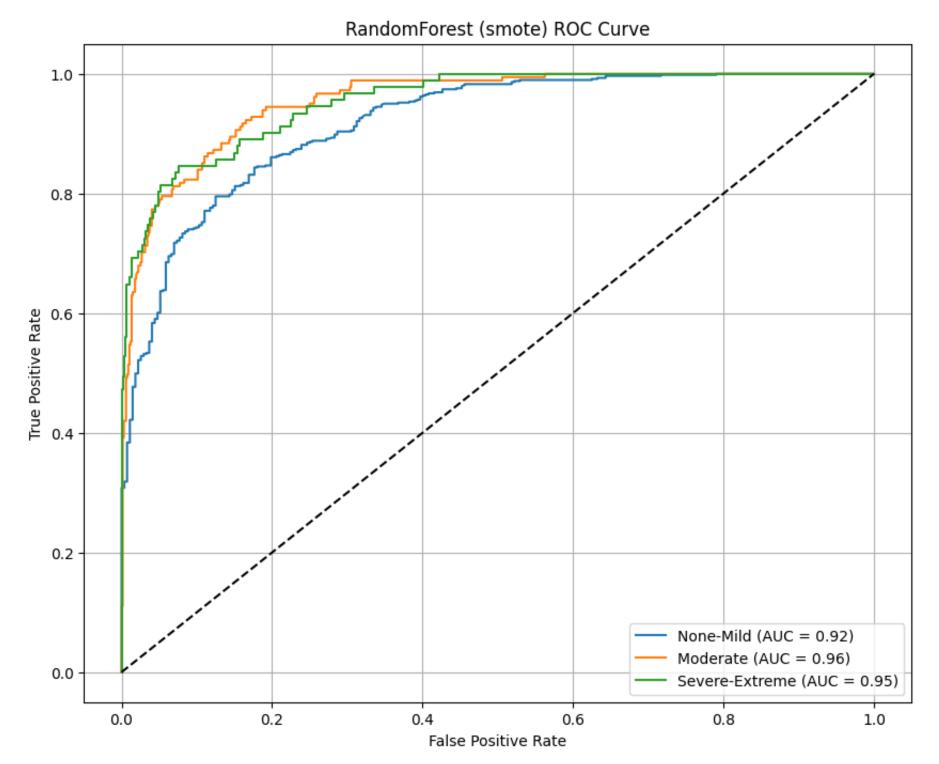




Performance for RandomForest (smote):
Accuracy: 0.86

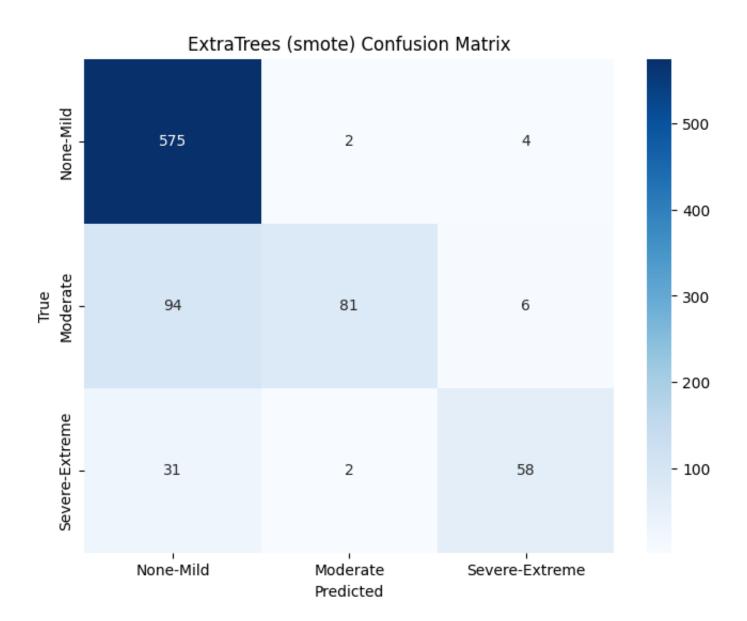
Accuracy: 0.86	precision	recall	f1-score	support
None-Mild Moderate Severe-Extreme	0.85 0.92 0.79	0.98 0.56 0.68	0.91 0.69 0.73	581 181 91
accuracy macro avg weighted avg	0.86 0.86	0.74 0.86	0.86 0.78 0.85	853 853 853

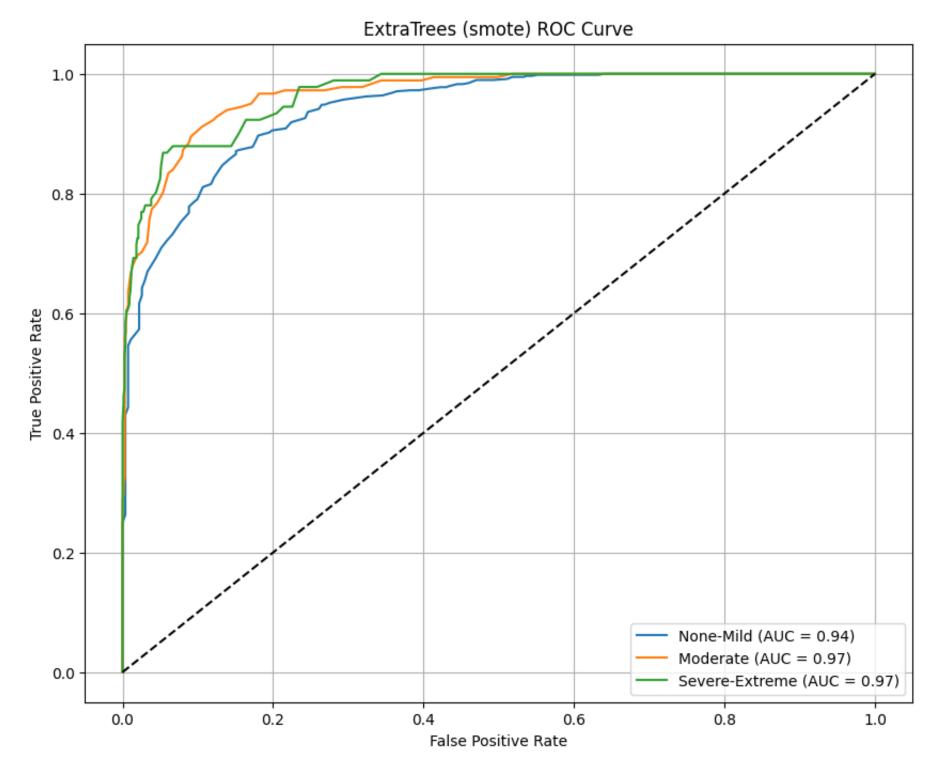




Performance for ExtraTrees (smote):
Accuracy: 0.84

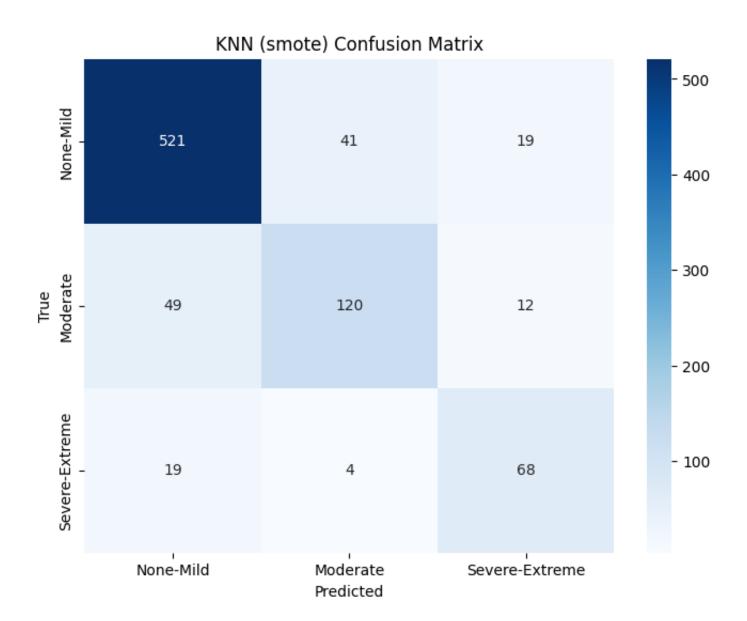
Accuracy: 0.84	precision	recall	f1-score	support
None-Mild Moderate Severe-Extreme	0.82 0.95 0.85	0.99 0.45 0.64	0.90 0.61 0.73	581 181 91
accuracy macro avg weighted avg	0.88 0.85	0.69 0.84	0.84 0.75 0.82	853 853 853

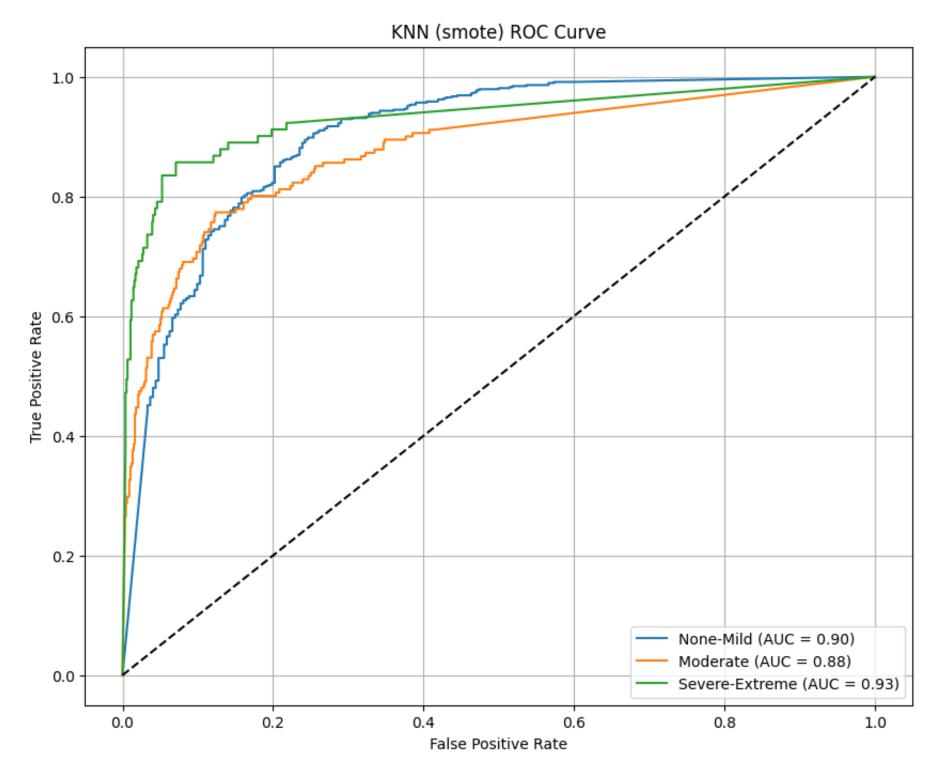




Performance for KNN (smote):
Accuracy: 0.83

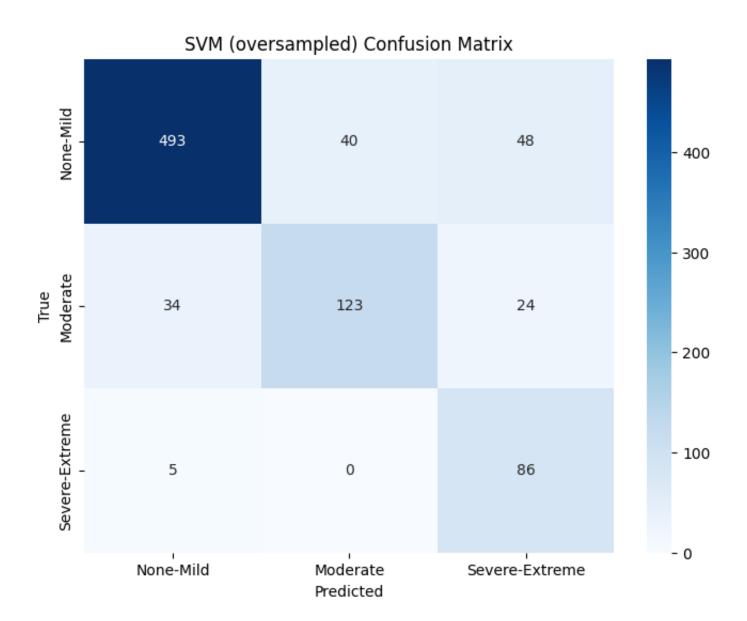
Accuracy: 0.83	precision	recall	f1-score	support
None-Mild Moderate Severe-Extreme	0.88 0.73 0.69	0.90 0.66 0.75	0.89 0.69 0.72	581 181 91
accuracy macro avg weighted avg	0.77 0.83	0.77 0.83	0.83 0.77 0.83	853 853 853

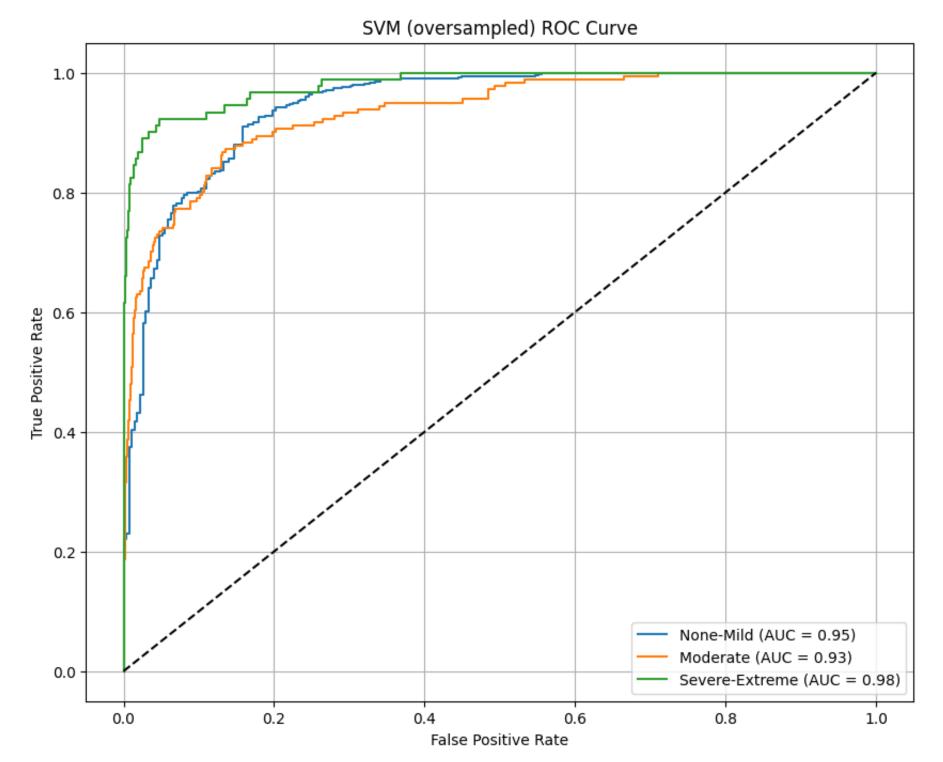




Performance for SVM (oversampled):
Accuracy: 0.82

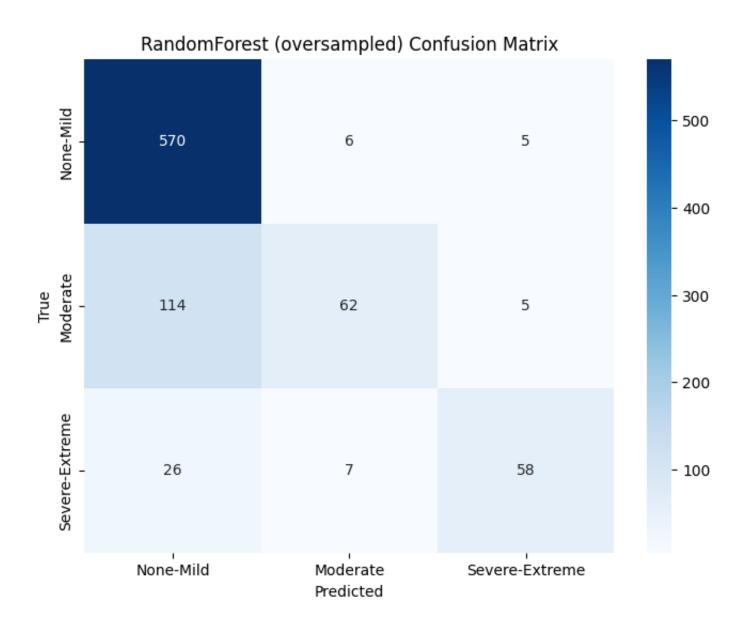
Accuracy: 0.82	precision	recall	f1-score	support
None-Mild Moderate Severe-Extreme	0.93 0.75 0.54	0.85 0.68 0.95	0.89 0.72 0.69	581 181 91
accuracy macro avg weighted avg	0.74 0.85	0.82 0.82	0.82 0.76 0.83	853 853 853





Performance for RandomForest (oversampled): Accuracy: 0.81

Accuracy: 0.81	precision	recall	f1-score	support
None-Mild	0.80	0.98	0.88	581
Moderate	0.83	0.34	0.48	181
Severe-Extreme	0.85	0.64	0.73	91
accuracy			0.81	853
macro avg	0.83	0.65	0.70	853
weighted avg	0.81	0.81	0.78	853



RandomForest (oversampled) ROC Curve 1.0 0.8 True Positive Rate 0.2 None-Mild (AUC = 0.89) Moderate (AUC = 0.90) 0.0 Severe-Extreme (AUC = 0.94) 0.8 0.2

0.4

False Positive Rate

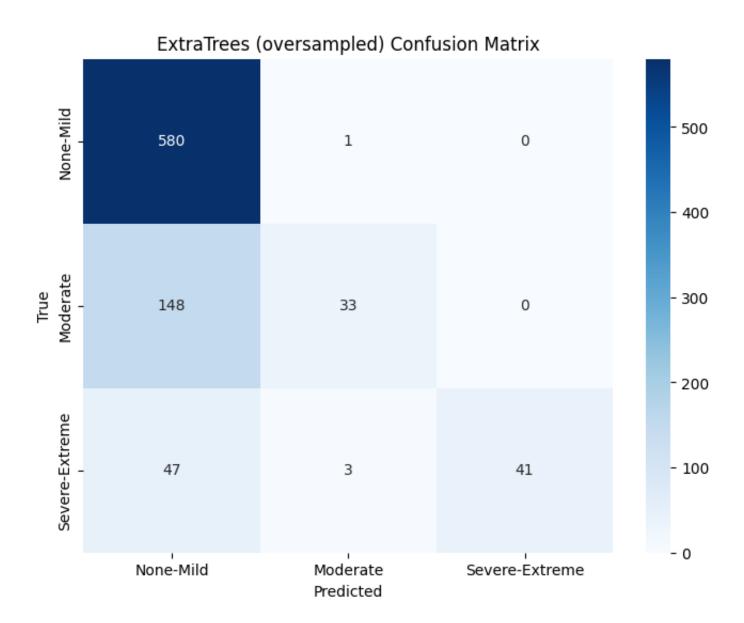
0.6

1.0

Performance for ExtraTrees (oversampled): Accuracy: 0.77

0.0

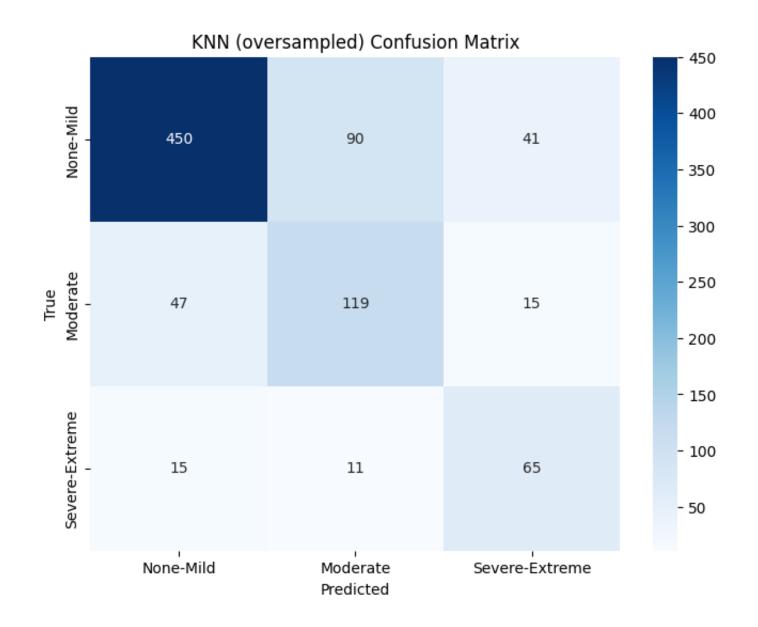
Accuracy: 0.77	precision	recall	f1-score	support
None-Mild Moderate Severe-Extreme	0.75 0.89 1.00	1.00 0.18 0.45	0.86 0.30 0.62	581 181 91
accuracy macro avg weighted avg	0.88 0.81	0.54 0.77	0.77 0.59 0.71	853 853 853

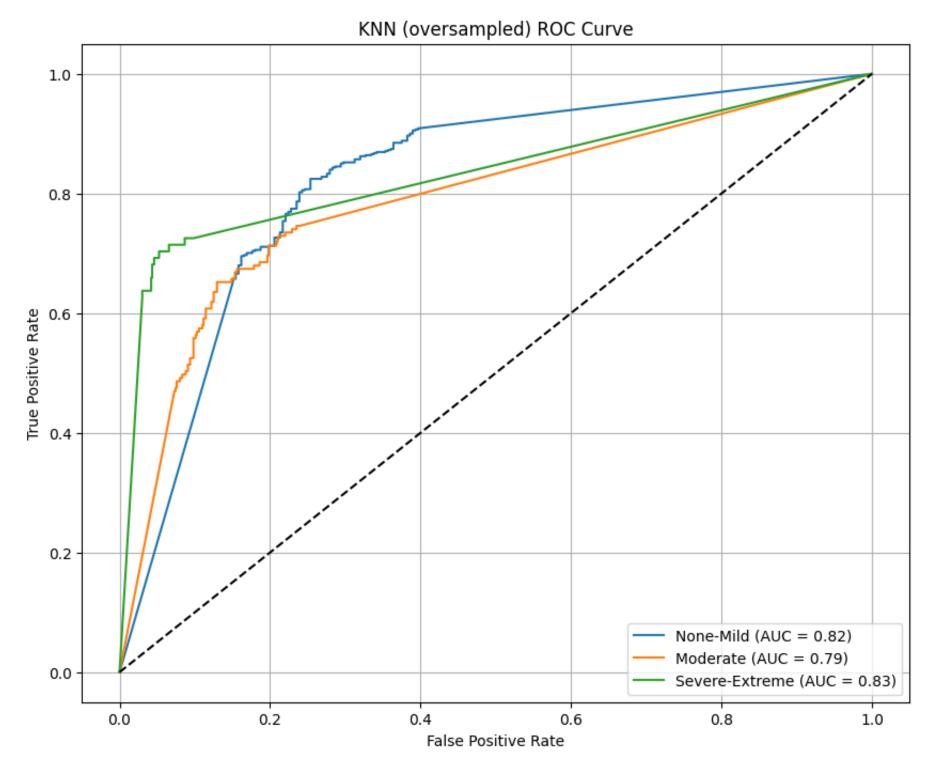


ExtraTrees (oversampled) ROC Curve 1.0 0.8 True Positive Rate 0.2 None-Mild (AUC = 0.90) Moderate (AUC = 0.93) 0.0 Severe-Extreme (AUC = 0.95) 0.8 0.2 0.6 1.0 0.0 0.4 False Positive Rate

Performance for KNN (oversampled):

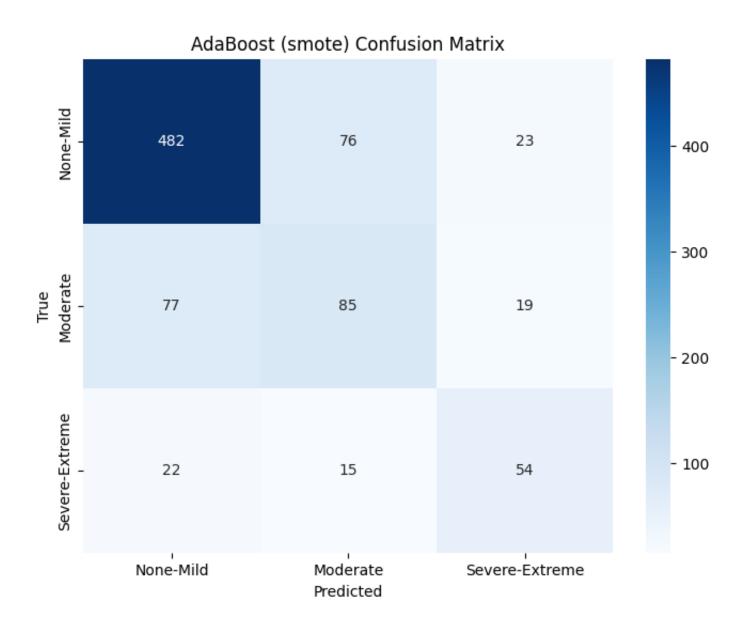
Accuracy: 0.74				
	precision	recall	f1–score	support
None-Mild	0.88	0.77	0.82	581
Moderate	0.54	0.66	0.59	181
Severe-Extreme	0.54	0.71	0.61	91
accuracy			0.74	853
macro avg	0.65	0.72	0.68	853
weighted avg	0.77	0.74	0.75	853

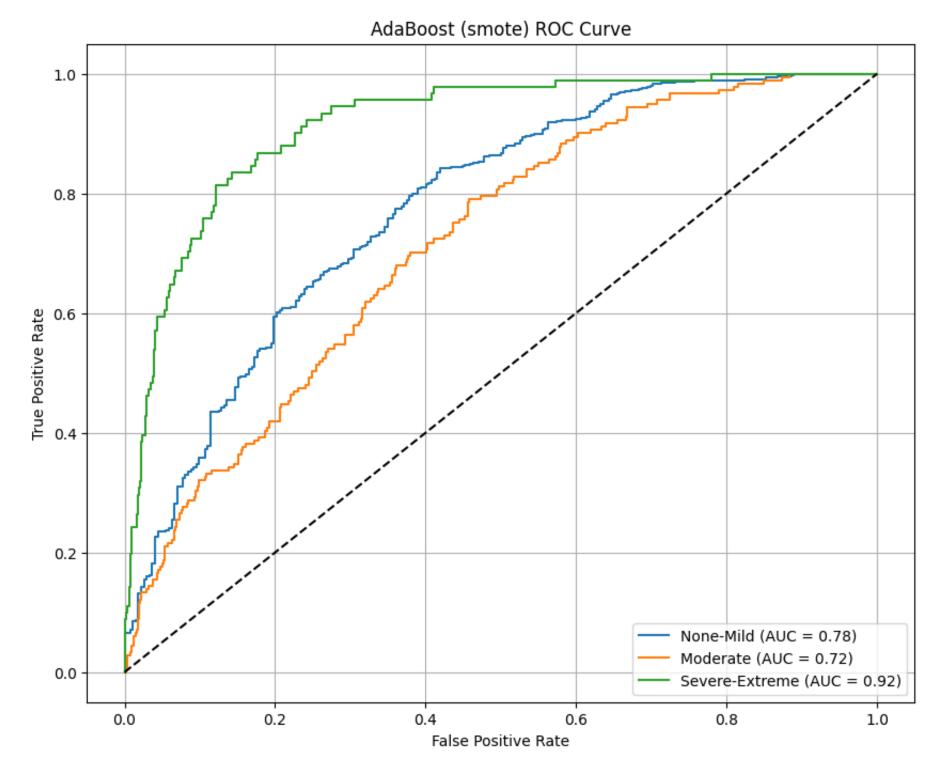




Performance for AdaBoost (smote): Accuracy: 0.73

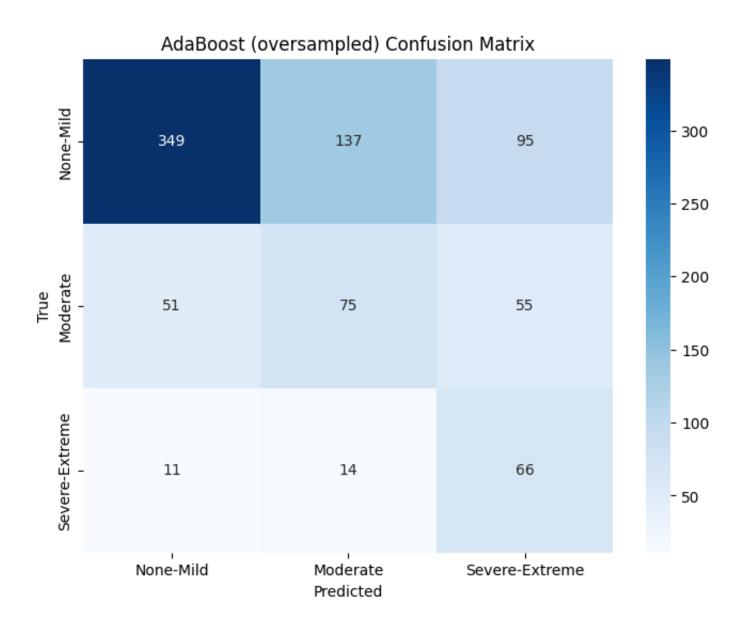
Accuracy: 0.73	precision	recall	f1-score	support
None-Mild Moderate	0.83 0.48	0.83 0.47	0.83 0.48	581 181
Severe-Extreme	0.56	0.59	0.58	91
accuracy macro avg weighted avg	0.63 0.73	0.63 0.73	0.73 0.63 0.73	853 853 853

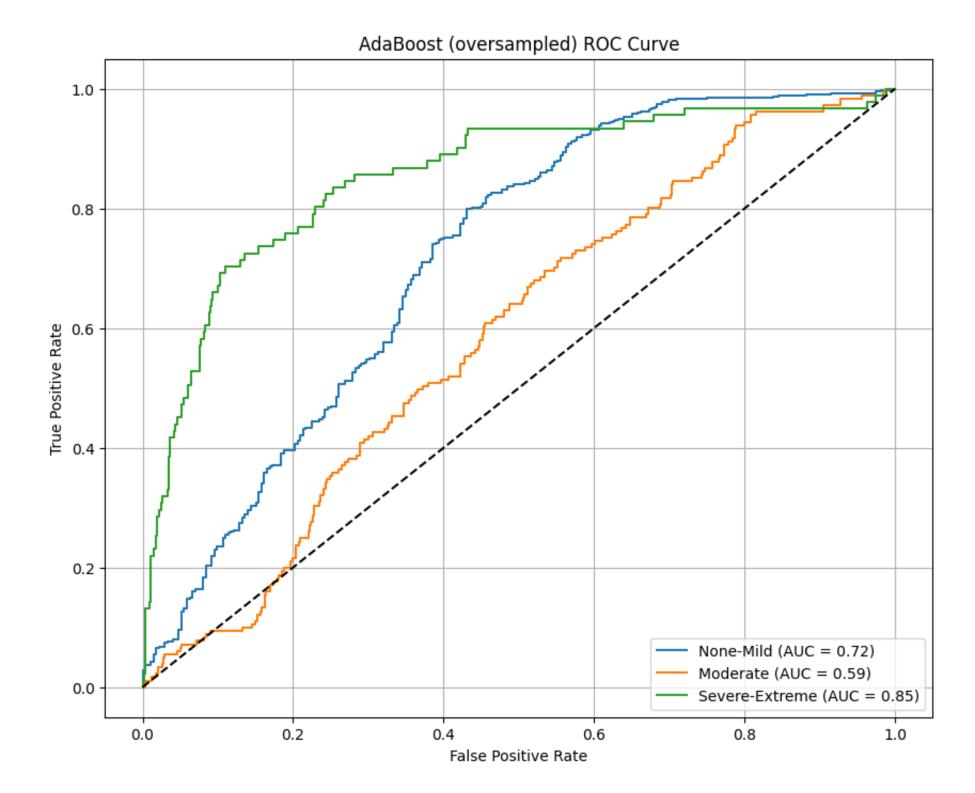




Performance for AdaBoost (oversampled): Accuracy: 0.57

Accuracy: 0.5/				
	precision	recall	f1-score	support
None-Mild	0.85	0.60	0.70	581
Moderate	0.33	0.41	0.37	181
Severe-Extreme	0.31	0.73	0.43	91
accuracy			0.57	853
macro avg	0.50	0.58	0.50	853
weighted avg	0.68	0.57	0.60	853



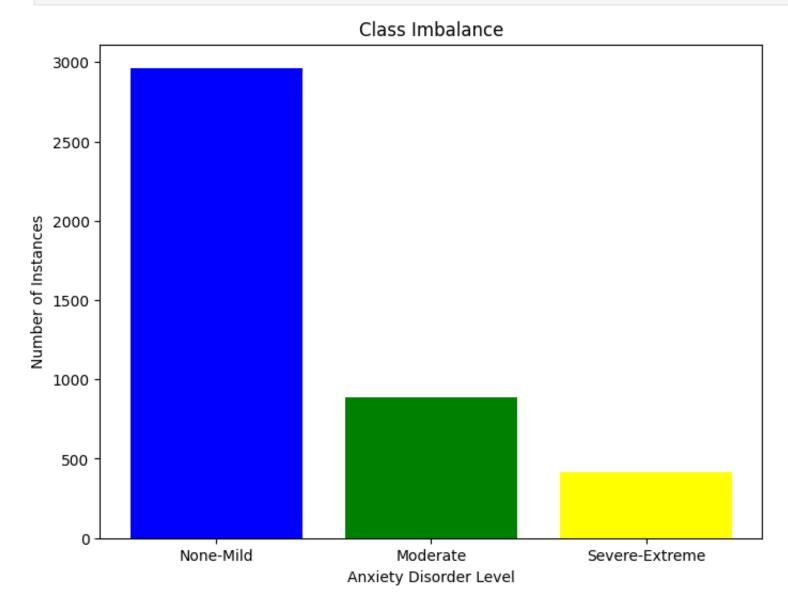


Creating model, predicting, calculating performance measures and drawing graphs

```
import matplotlib.pyplot as plt

# Class labels and counts
class_labels = ['None-Mild', 'Moderate', 'Severe-Extreme']
class_counts = [2964, 884, 416]

# Plotting the class imbalance
plt.figure(figsize=(8, 6))
plt.bar(class_labels, class_counts, color=['blue', 'green', 'yellow', 'orange', 'red'])
plt.xlabel('Anxiety Disorder Level')
plt.ylabel('Number of Instances')
plt.title('Class Imbalance')
plt.show()
```

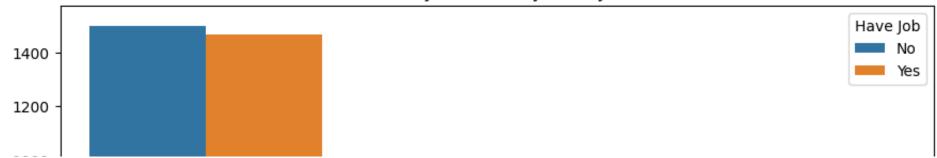


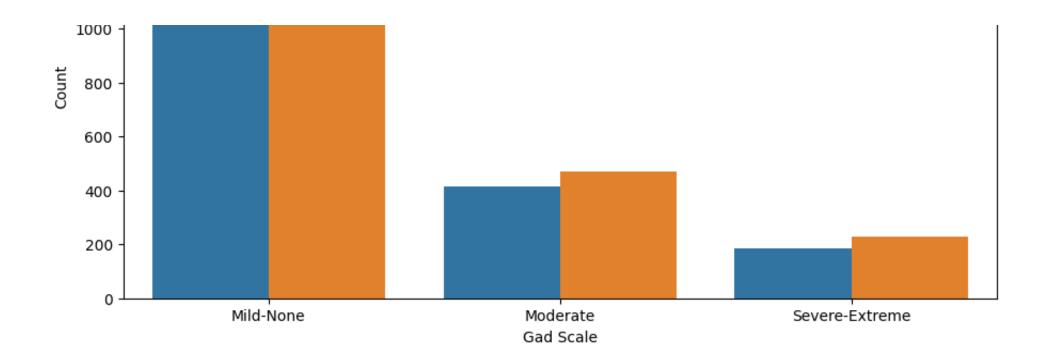
Graphs for EDA

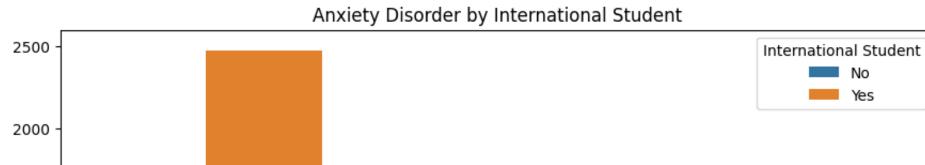
```
In [41]: df = edf.copy()
df['gad_scale'] = df['gad_scale'].map({'None-Mild': 'Mild-None', 'Moderate': 'Moderate', 'Severe-Extreme': 'Severe-Extreme'})
```

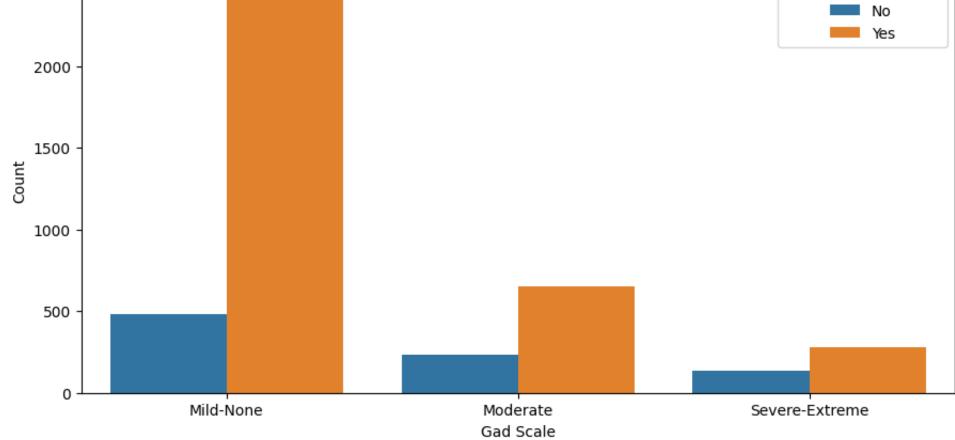
```
In [48]: import pandas as pd
         import matplotlib.pyplot as plt
         import seaborn as sns
         # Assuming the dataframe is already loaded and named df
         # Function to plot actual count plots and print values
         def plot_actual_countplot(data, x, hue, title, ax):
             # Compute the actual count
             counts = data.groupby([x, hue]).size().rename('count').reset_index()
             sns.barplot(data=counts, x=x, y='count', hue=hue, ax=ax)
             ax.set_title(title)
             ax.set_ylabel('Count')
             ax.set_xlabel(x.replace('_', ' ').title())
             ax.legend(title=hue.replace('_', ' ').title())
         # List of columns to analyze
         columns_to_analyze = ['have_job', 'international_student', 'support_family_financially',
                 'financial_worries', 'living_arrangements', 'academic_stress',
                 'workload', 'suppported_by_friends', 'supported_by_family',
                'seek_staff_support', 'sleep_time', 'exercise', 'diet', 'degree_level',
                'faced_discrimination', 'involved_in_sports', 'friends_circle',
                'drinking_habits', 'gender', 'age_range', 'chronic_health_problem',
                 'study_year'
         # Create subplots
         fig, axes = plt.subplots(len(columns_to_analyze), 1, figsize=(10, 5 * len(columns_to_analyze)))
         fig.tight_layout(pad=5.0)
         # Plot each variable
         for ax, column in zip(axes, columns_to_analyze):
             plot_actual_countplot(df, 'gad_scale', column, f'Anxiety Disorder by {column.replace("_", " ").title()}', ax)
         plt.show()
         # Save a summary of these visualizations in a multi-page PDF
         from matplotlib.backends.backend_pdf import PdfPages
         with PdfPages('anxiety_level_trends_actual.pdf') as pdf:
             for column in columns_to_analyze:
                 fig, ax = plt.subplots(figsize=(10, 6))
                 plot_actual_countplot(df, 'gad_scale', column, f'Anxiety Disorder by {column.replace("_", " ").title()}', ax)
                 pdf.savefig()
                 plt.close()
```

Anxiety Disorder by Have Job

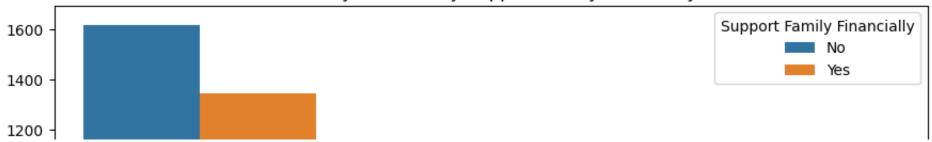


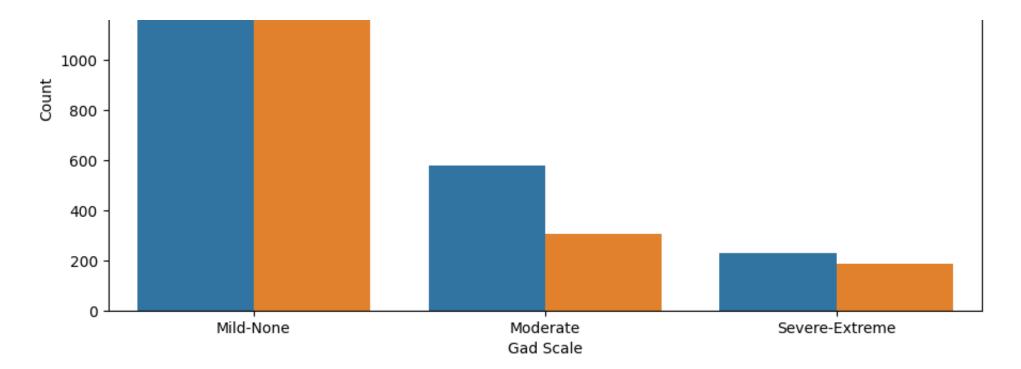


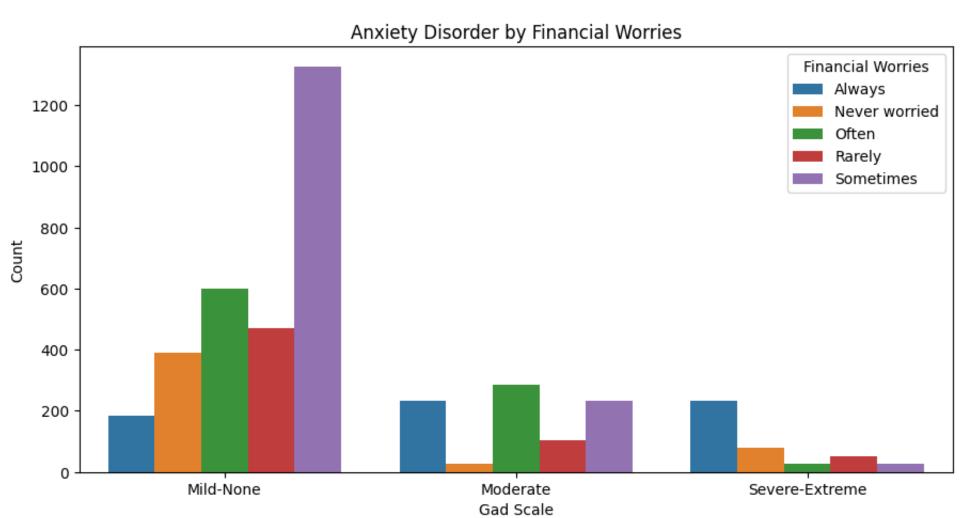


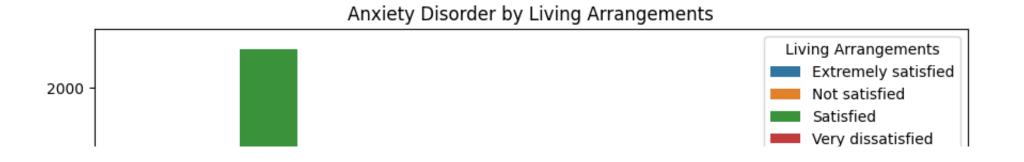


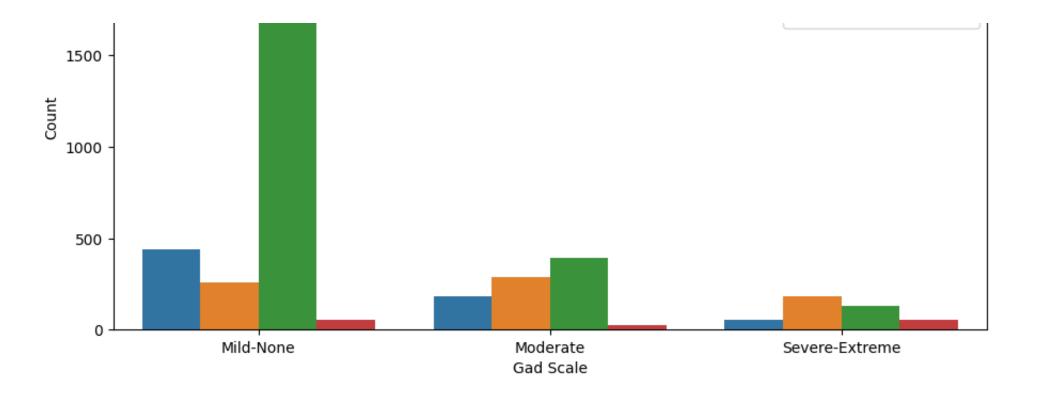


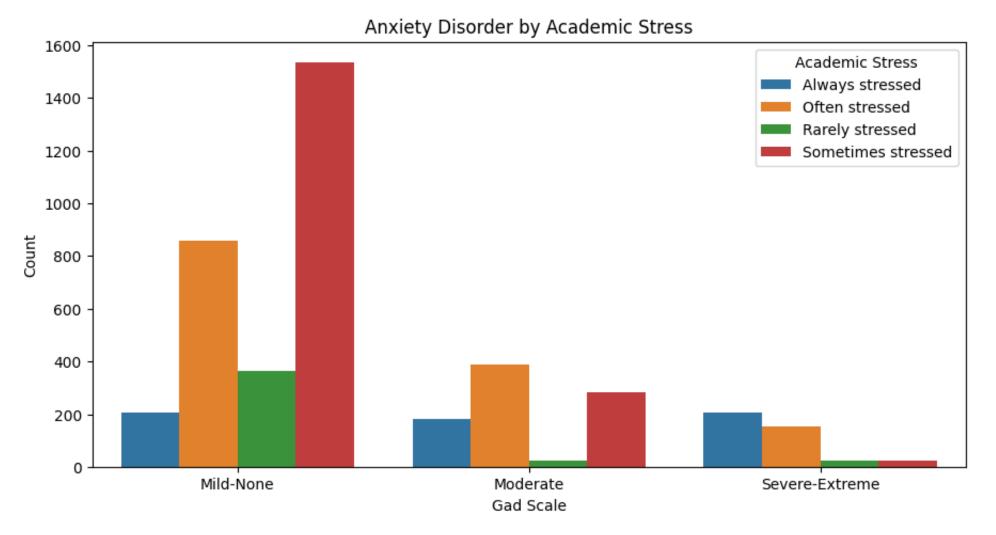


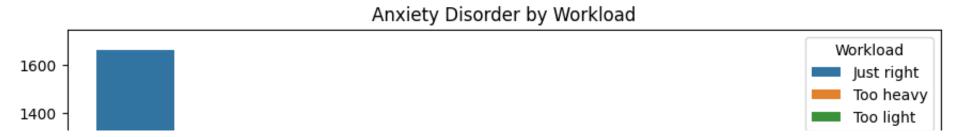


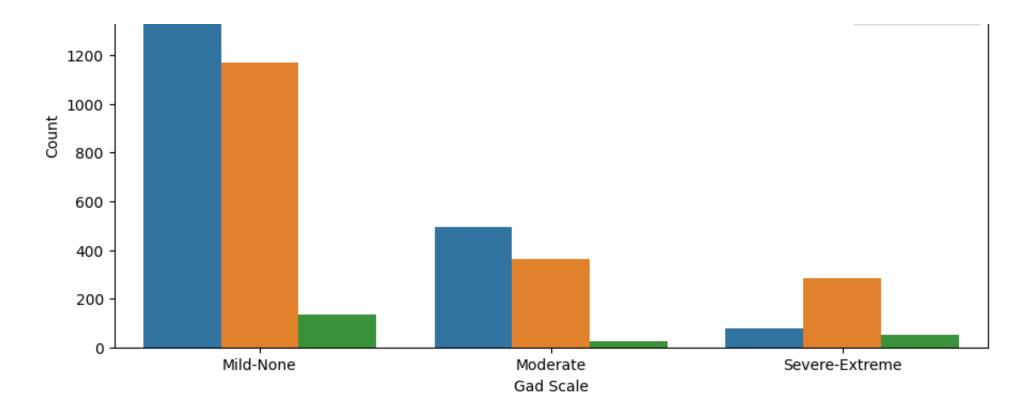




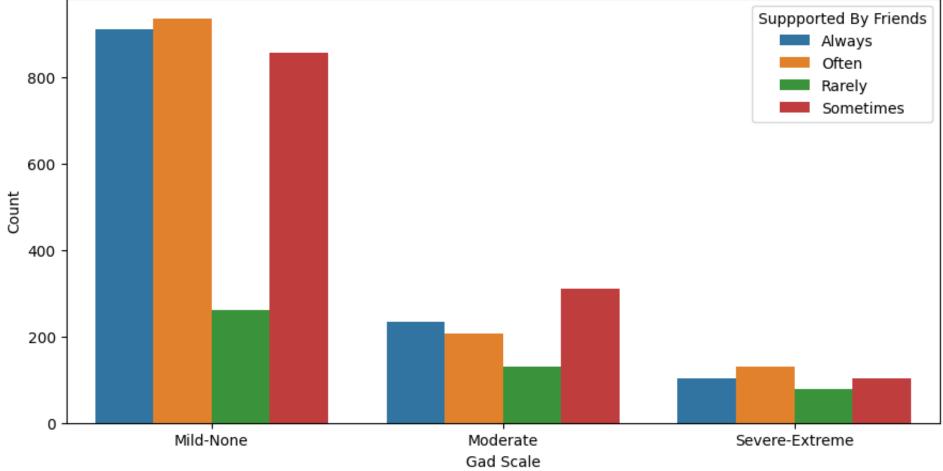




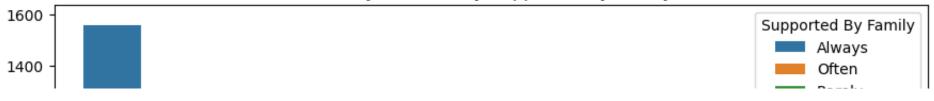


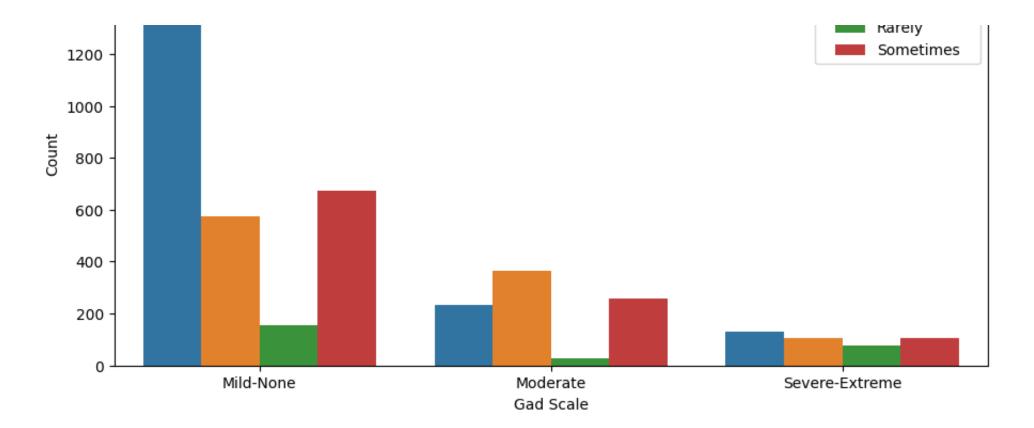


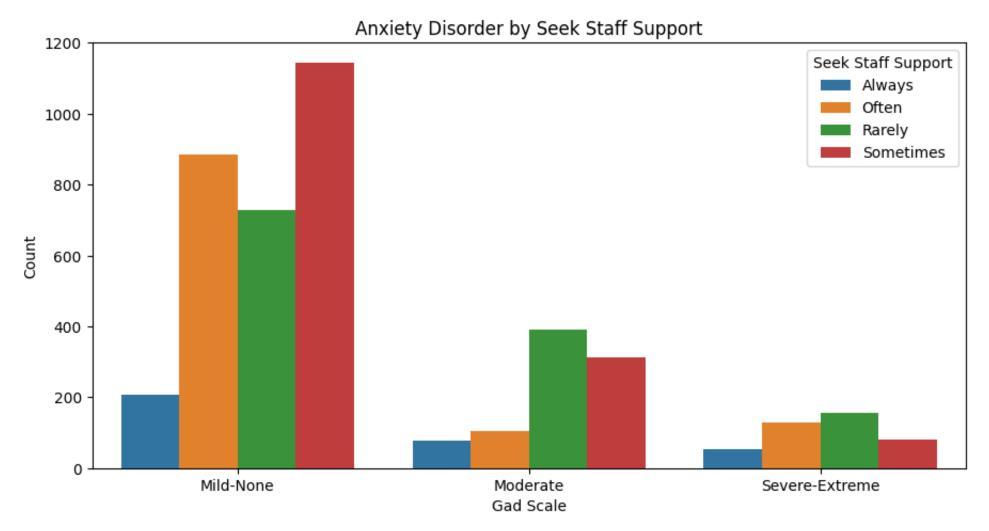


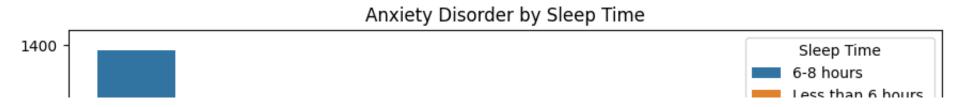


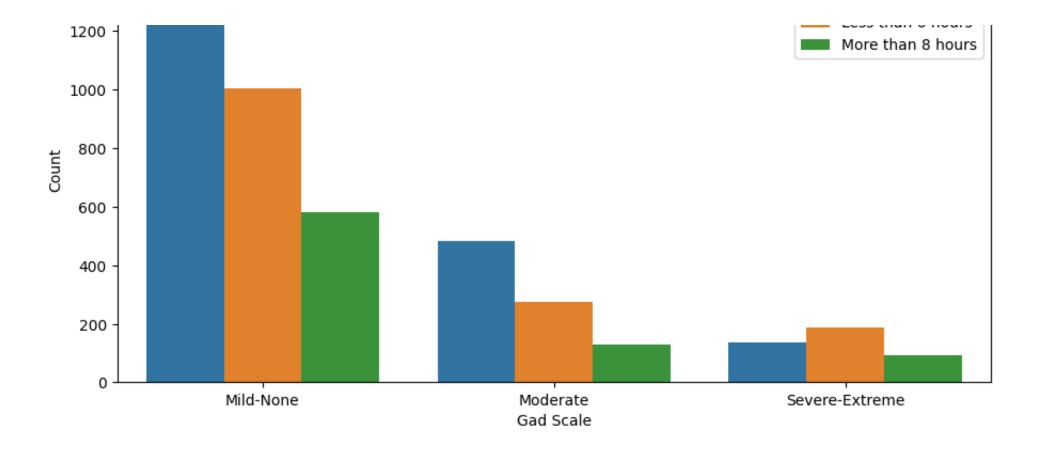
Anxiety Disorder by Supported By Family

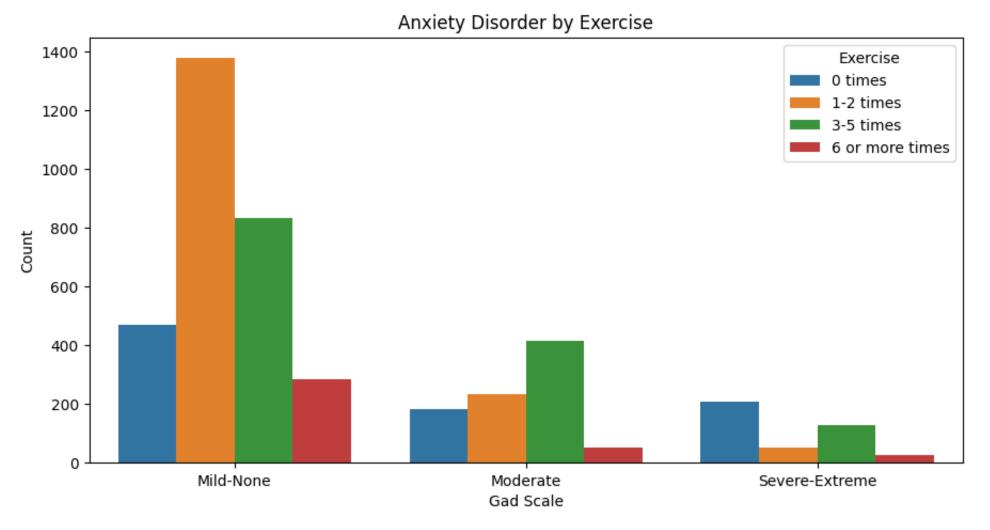


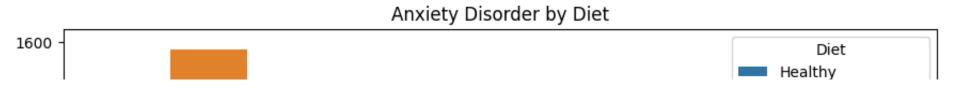


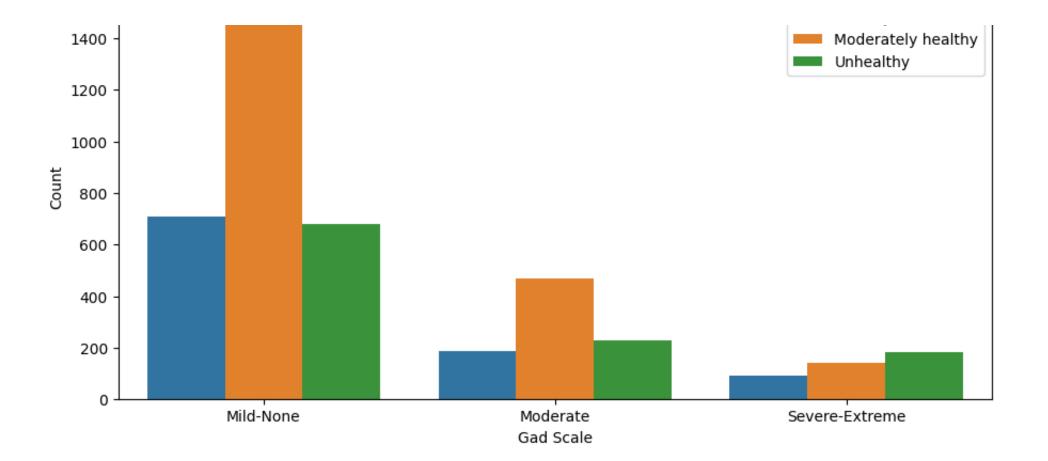


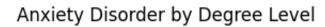


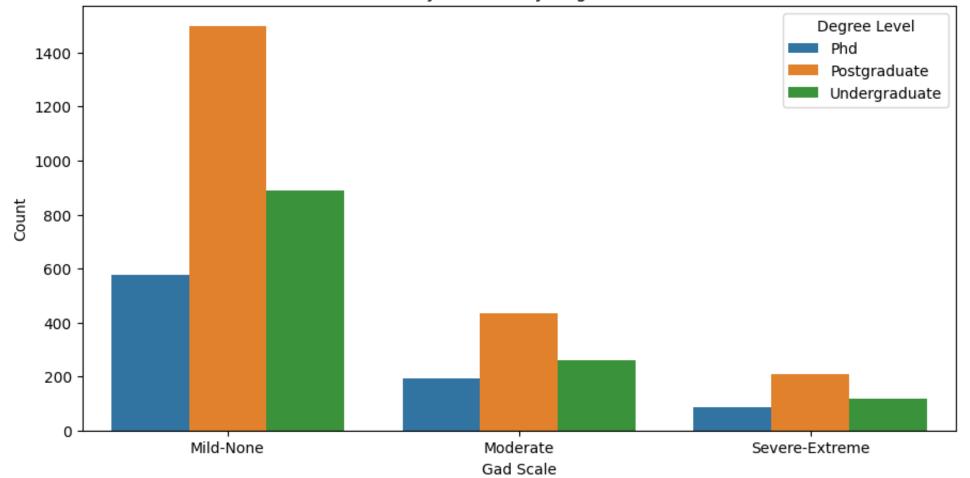




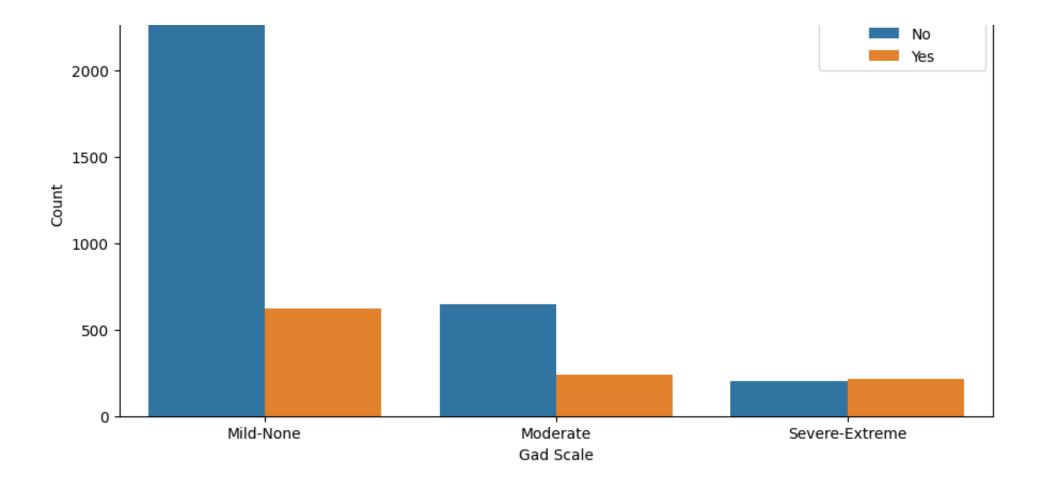




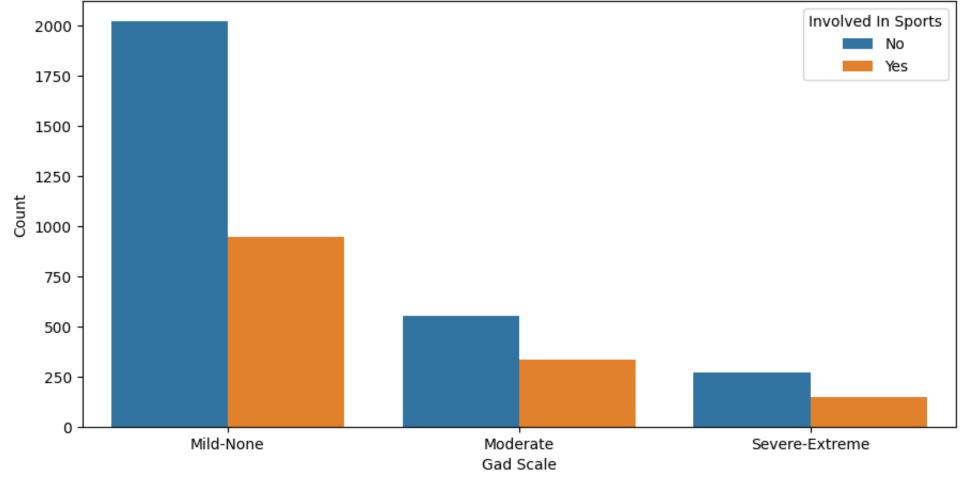




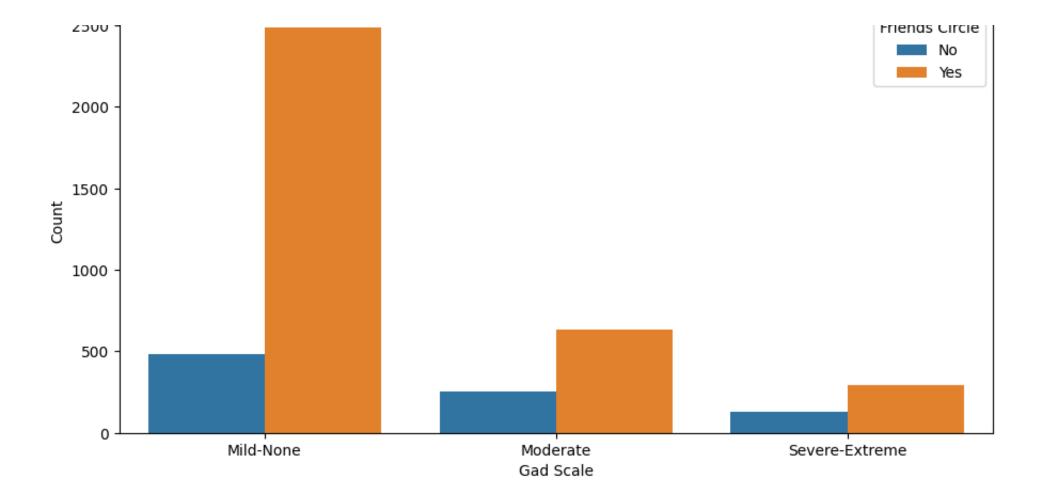
Anxiety Disorder by Faced Discrimination

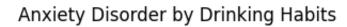


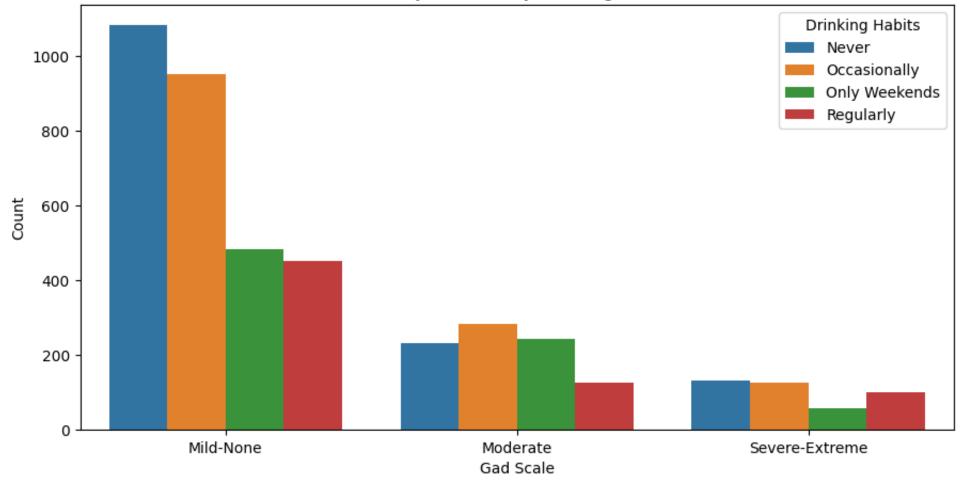




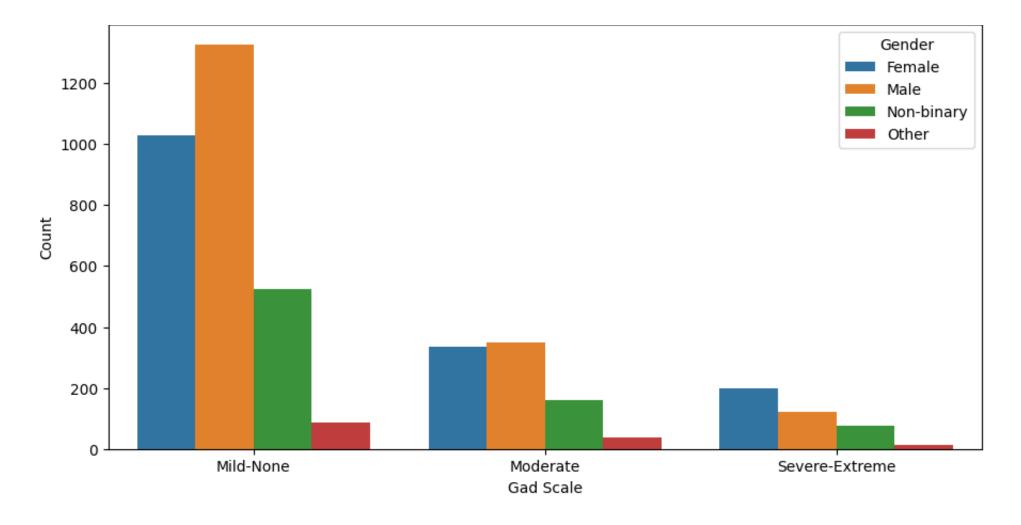
Anxiety Disorder by Friends Circle

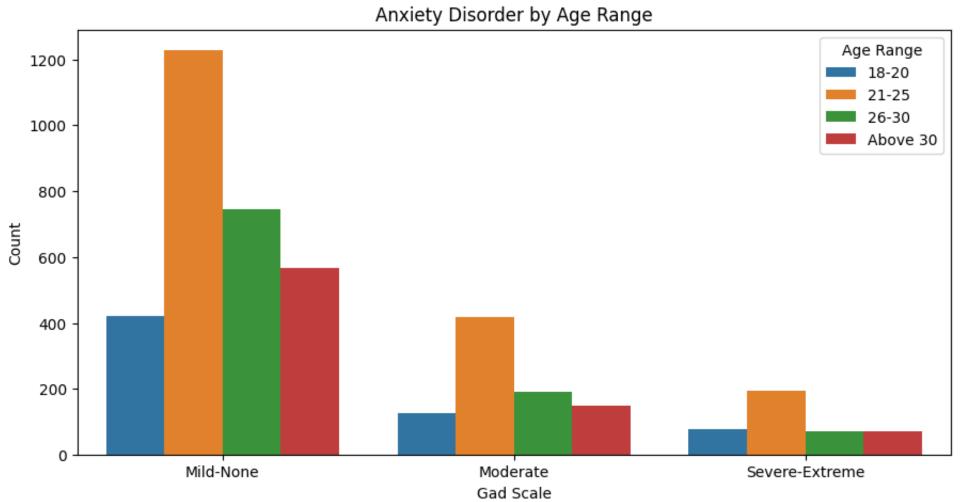




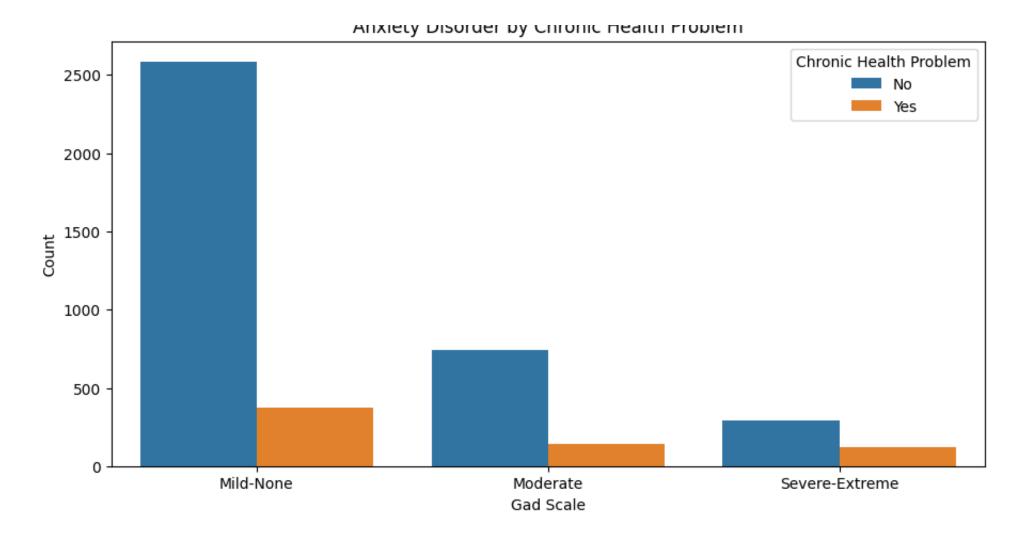


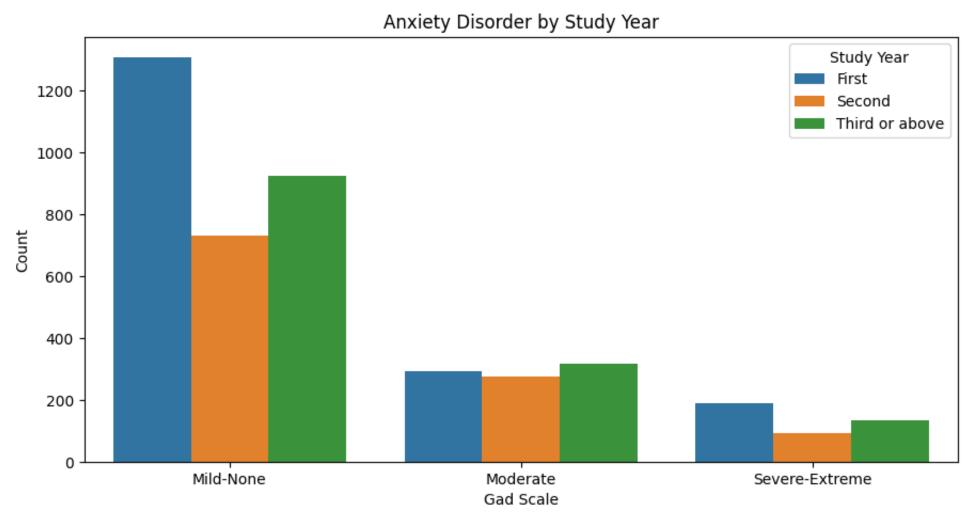
Anxiety Disorder by Gender



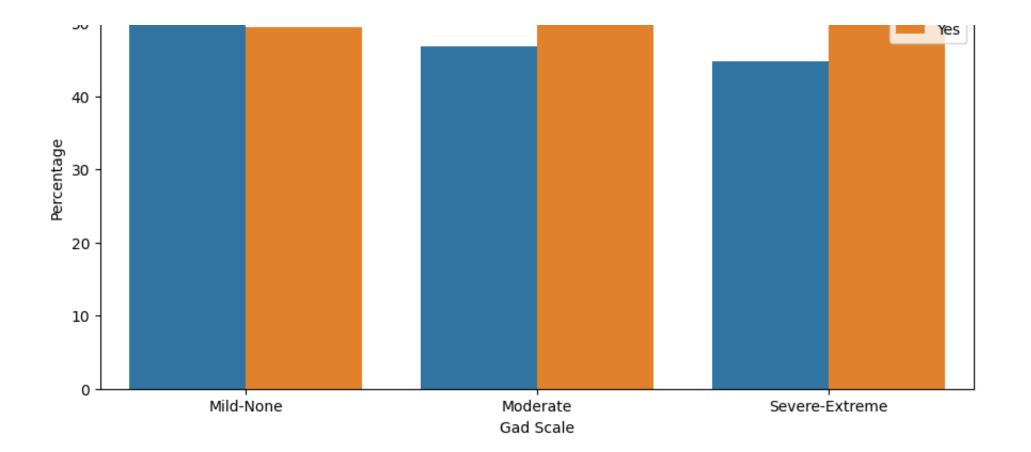


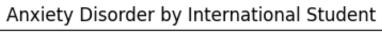
Anxiety Disarder by Chronic Health Droblem

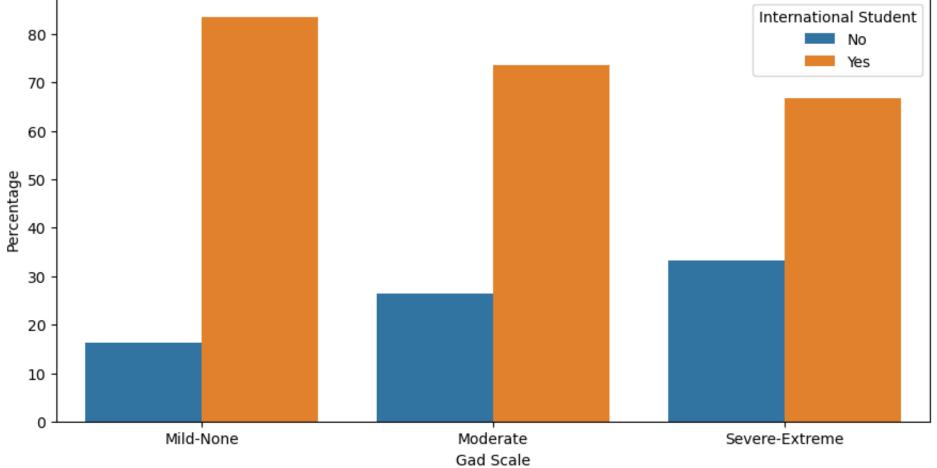




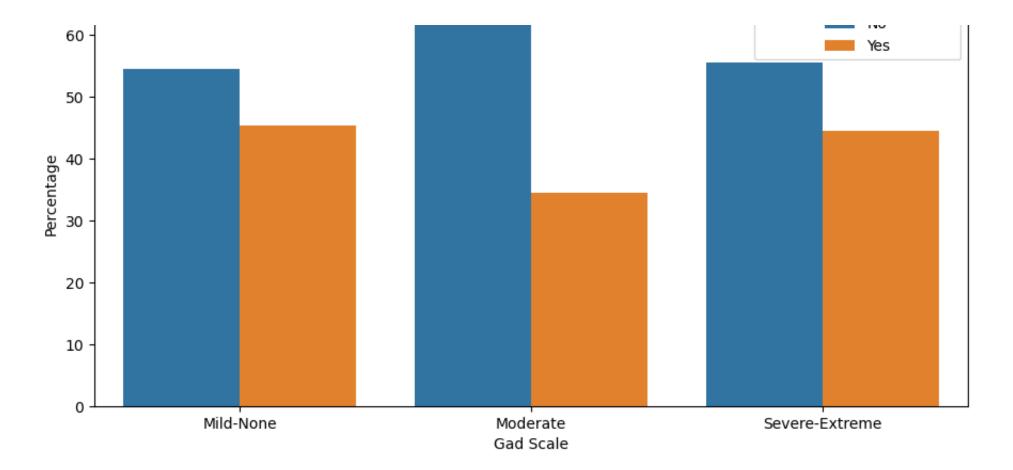
```
In [47]: import pandas as pd
         import matplotlib.pyplot as plt
         import seaborn as sns
         # Assuming the dataframe is already loaded and named df
         # Function to plot percentage count plots and print values
         def plot_percentage_countplot(data, x, hue, title, ax):
             # Compute the actual count
             counts = data.groupby([x, hue]).size().rename('count').reset_index()
             # Calculate the percentage
             total_counts = counts.groupby(x)['count'].transform('sum')
             counts['percentage'] = (counts['count'] / total_counts) * 100
             # Plot the data
             sns.barplot(data=counts, x=x, y='percentage', hue=hue, ax=ax)
             ax.set_title(title)
             ax.set_ylabel('Percentage')
             ax.set_xlabel(x.replace('_', ' ').title())
             ax.legend(title=hue.replace('_', '').title())
         # List of columns to analyze
         columns_to_analyze = ['have_job', 'international_student', 'support_family_financially',
                 'financial_worries', 'living_arrangements', 'academic_stress',
                 'workload', 'suppported_by_friends', 'supported_by_family',
                'seek_staff_support', 'sleep_time', 'exercise', 'diet', 'degree_level',
                'faced_discrimination', 'involved_in_sports', 'friends_circle',
                'drinking_habits', 'gender', 'age_range', 'chronic_health_problem',
                'study_year'
         # Create subplots
         fig, axes = plt.subplots(len(columns_to_analyze), 1, figsize=(10, 5 * len(columns_to_analyze)))
         fig.tight_layout(pad=5.0)
         # Plot each variable
         for ax, column in zip(axes, columns_to_analyze):
             plot_percentage_countplot(df, 'gad_scale', column, f'Anxiety Disorder by {column.replace("_", " ").title()}', ax)
         plt.show()
         # Save a summary of these visualizations in a multi-page PDF
         from matplotlib.backends.backend pdf import PdfPages
         with PdfPages('anxiety_level_trends_actual.pdf') as pdf:
             for column in columns_to_analyze:
                 fig, ax = plt.subplots(figsize=(10, 6))
                 plot_percentage_countplot(df, 'gad_scale', column, f'Anxiety Disorder by {column.replace("_", " ").title()}', ax)
                 pdf.savefig()
                 plt.close()
```

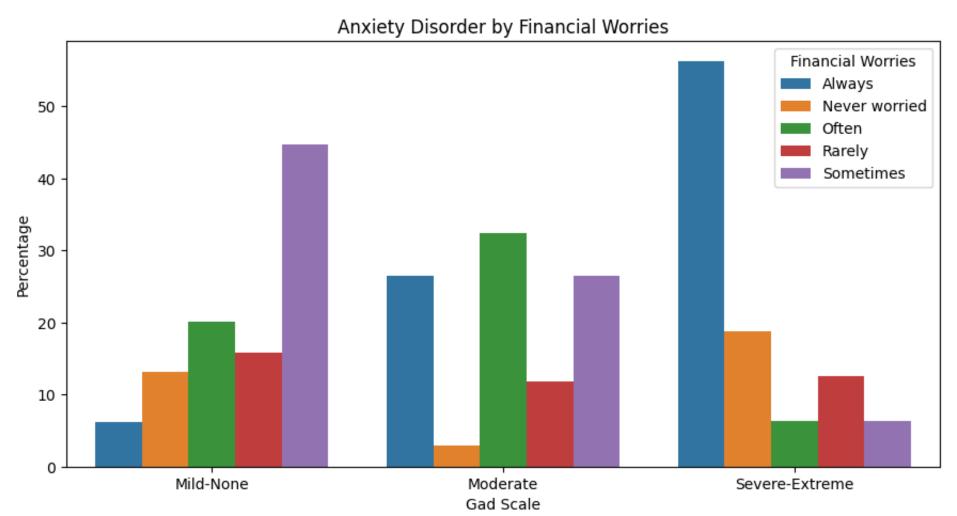






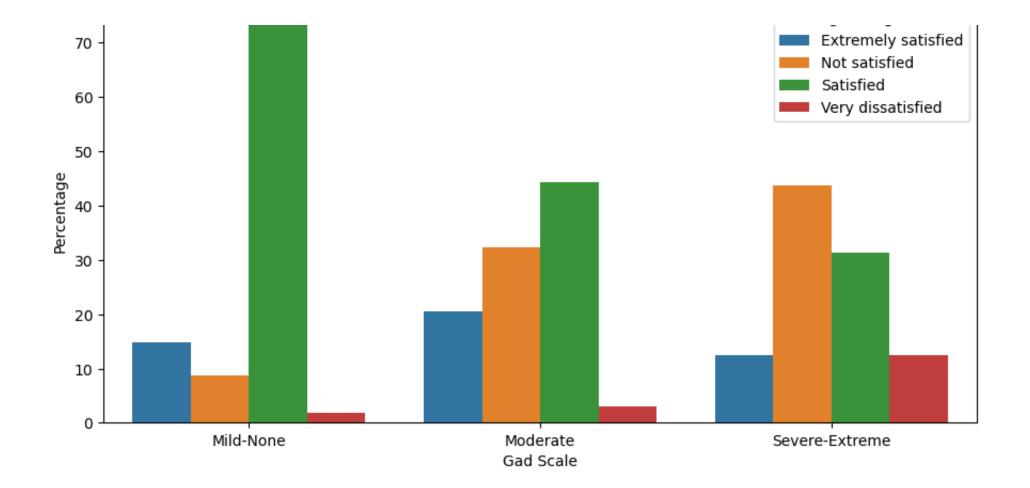
Anxiety Disorder by Support Family Financially

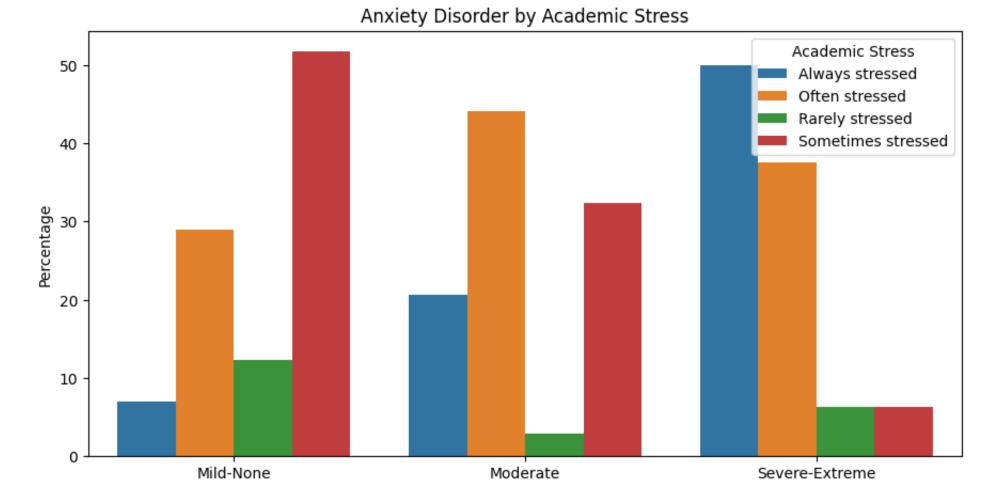




Anxiety Disorder by Living Arrangements

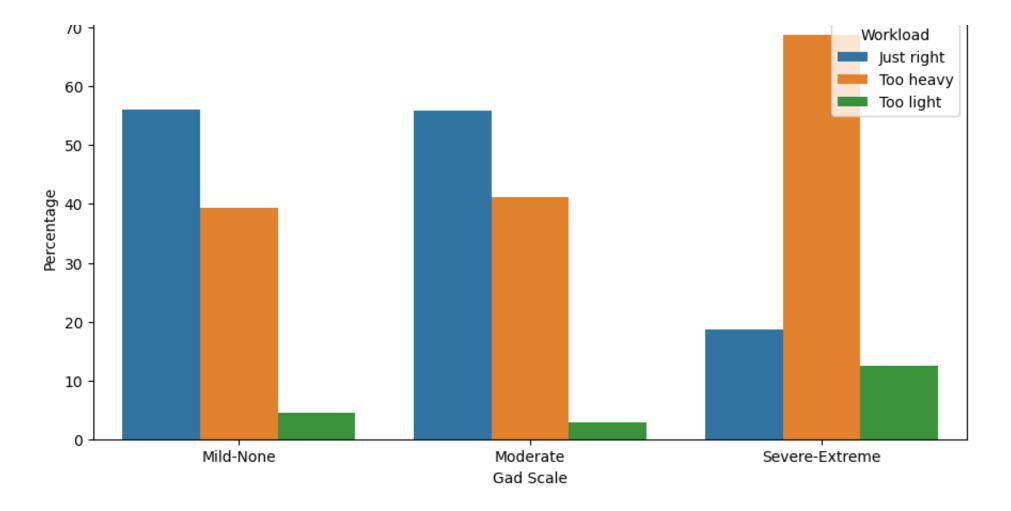
Living Arrangements

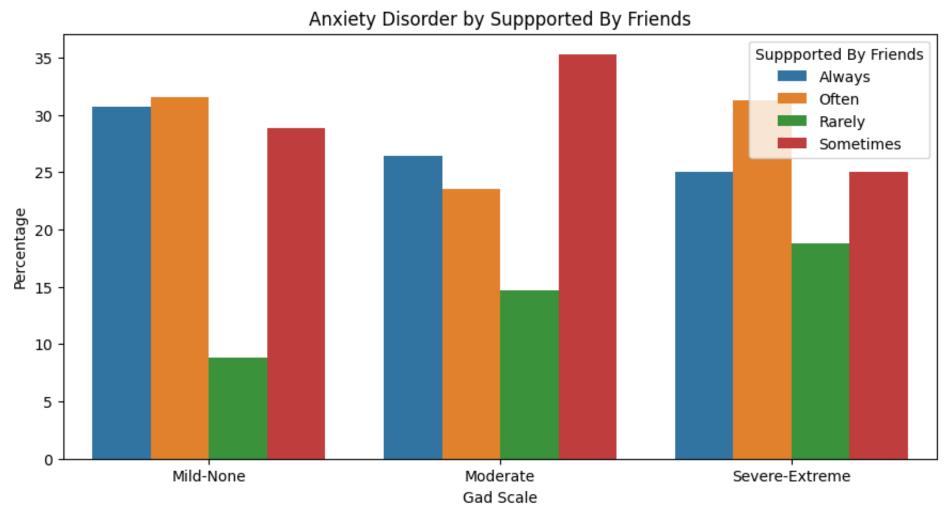




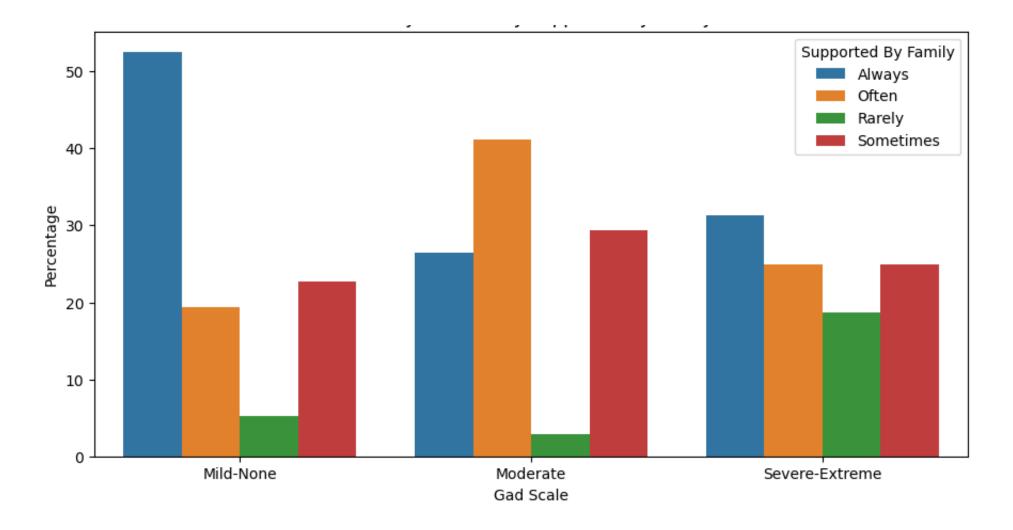
Anxiety Disorder by Workload

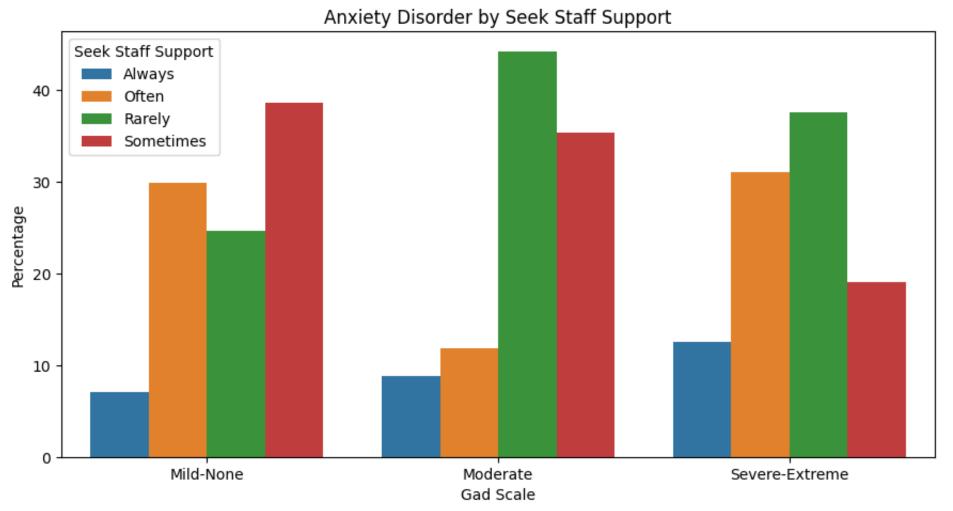
Gad Scale

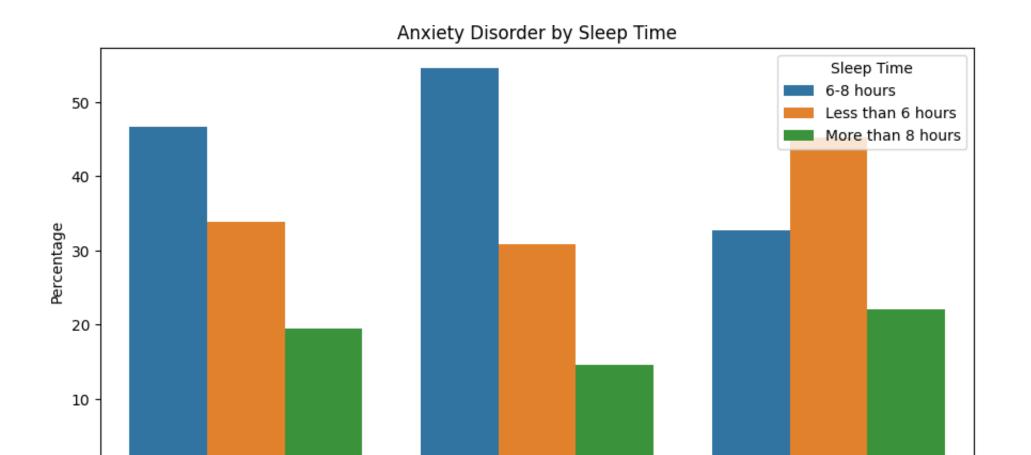




Anxiety Disorder by Supported By Family





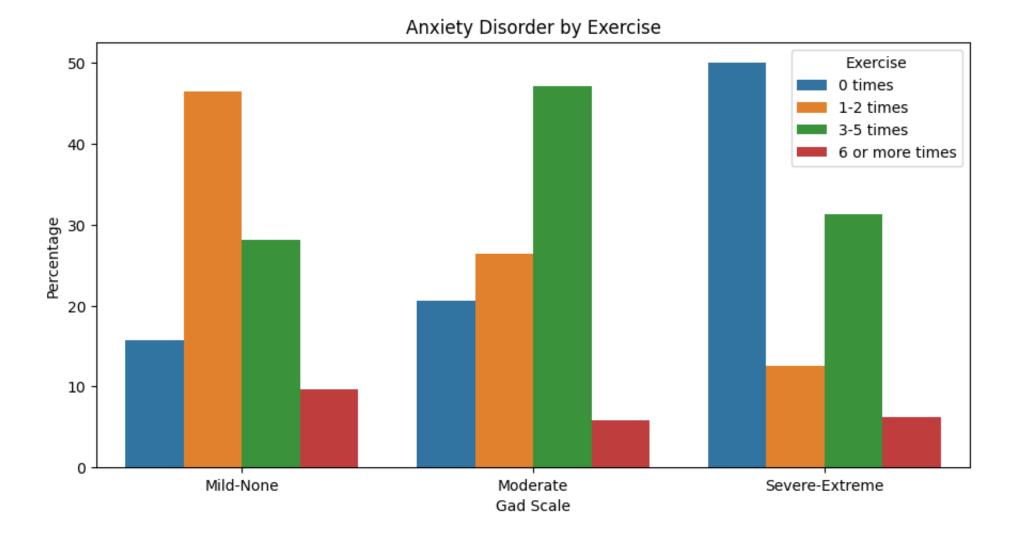


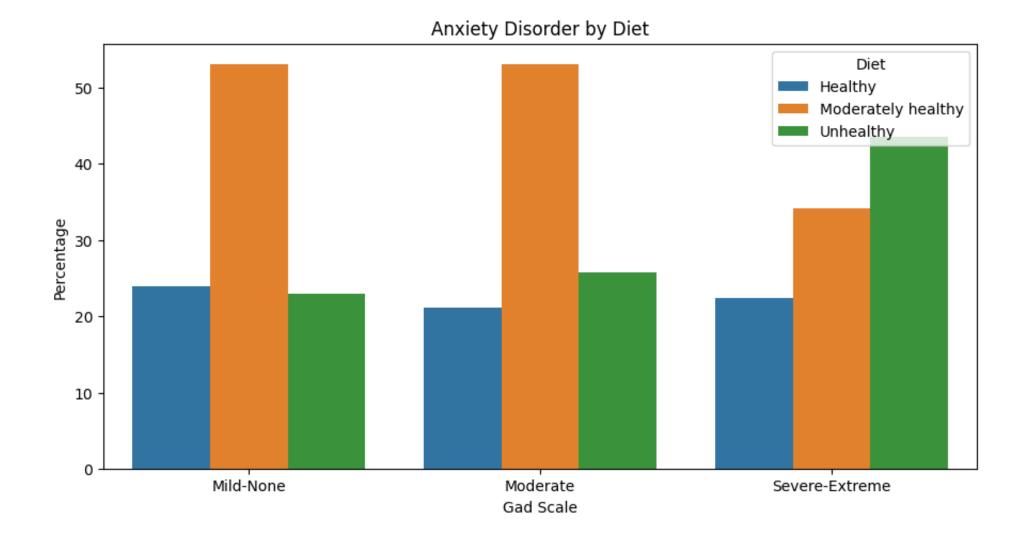
Moderate

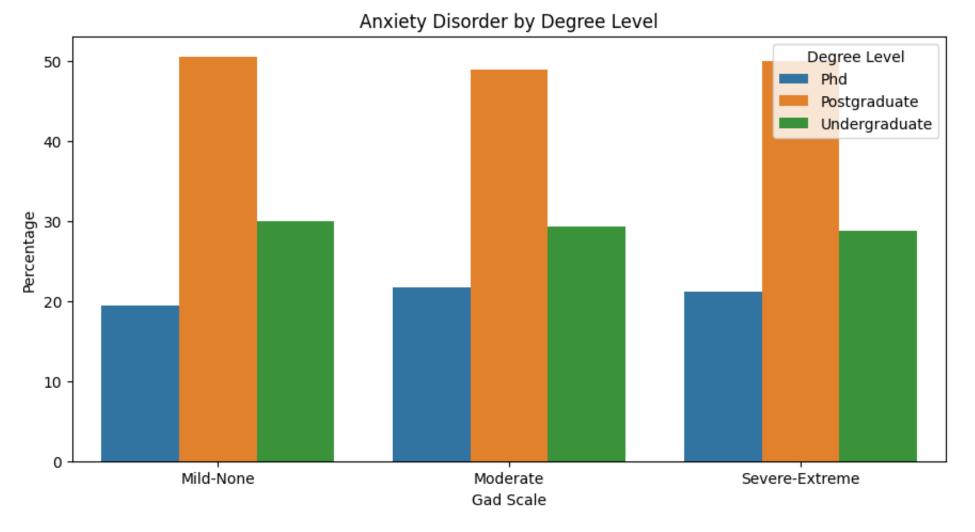
Gad Scale

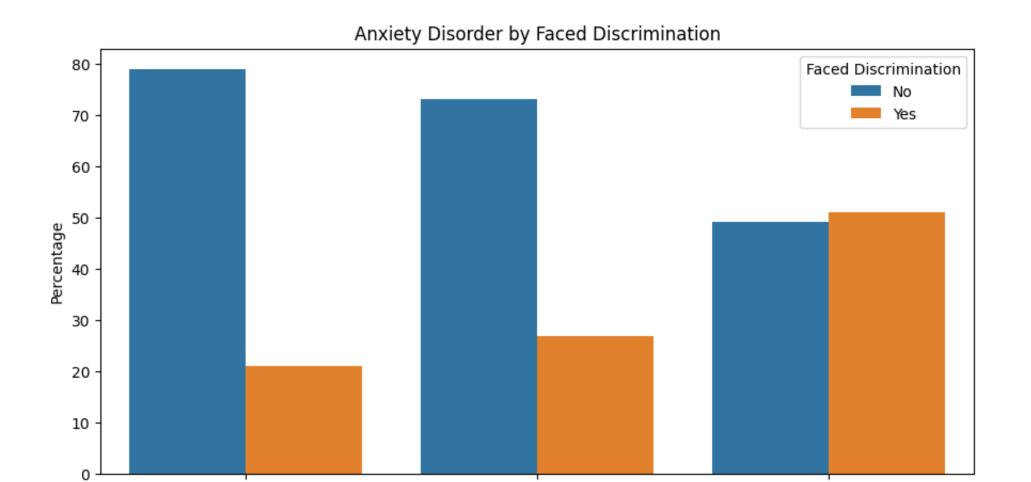
Severe-Extreme

Mild-None







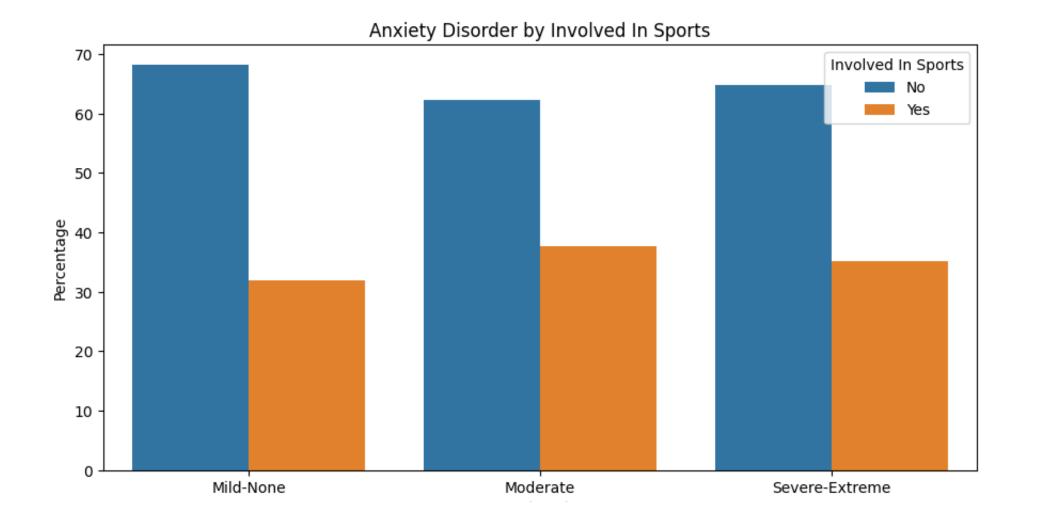


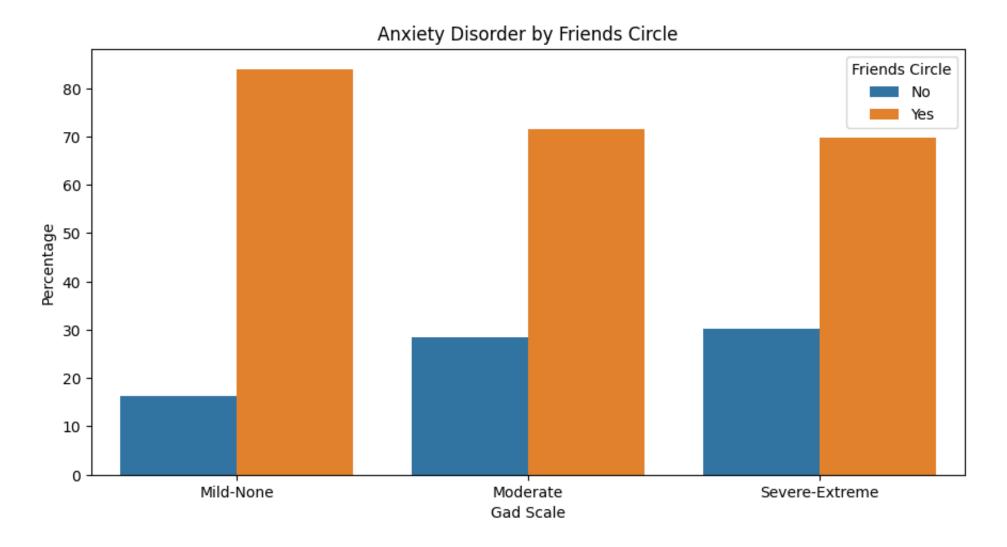
Moderate

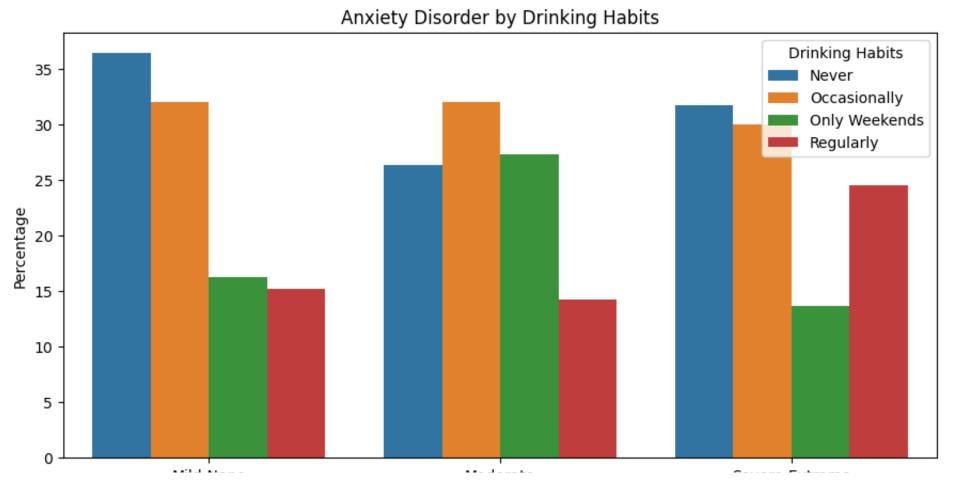
Gad Scale

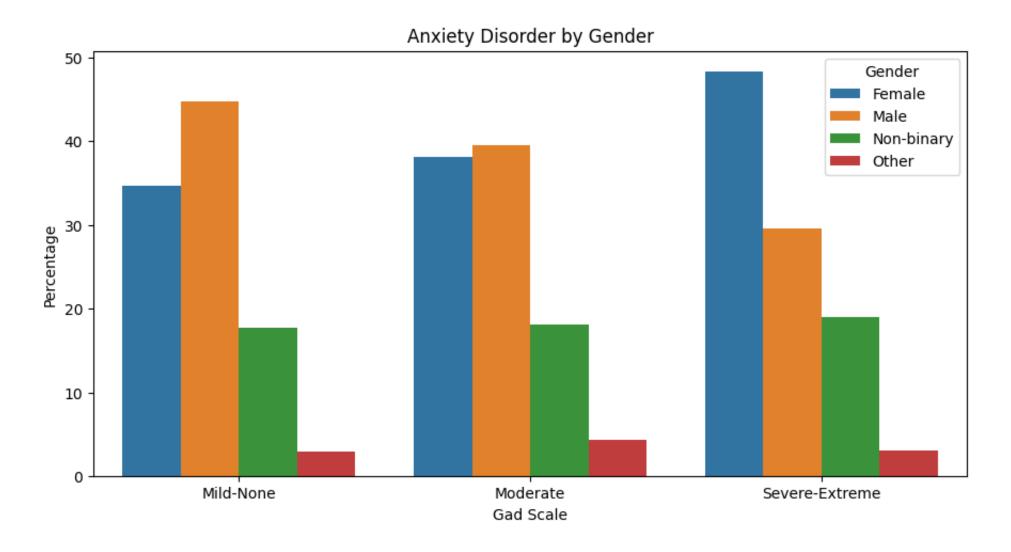
Severe-Extreme

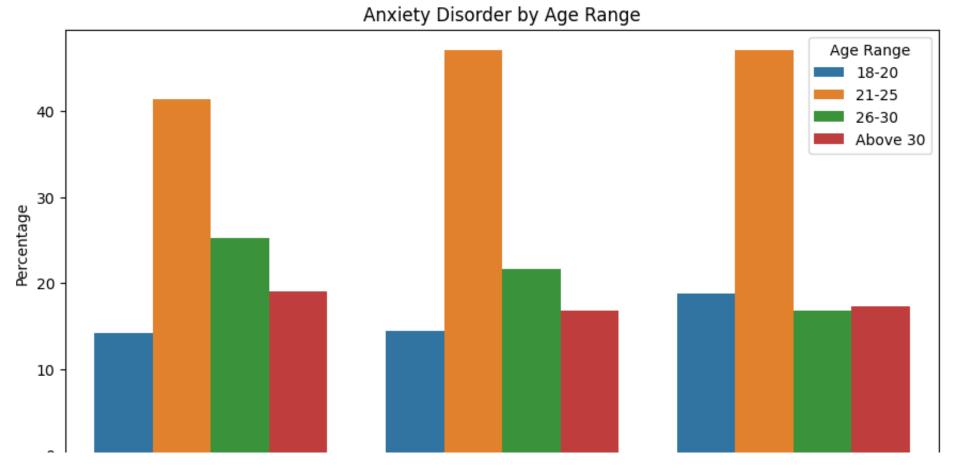
Mild-None



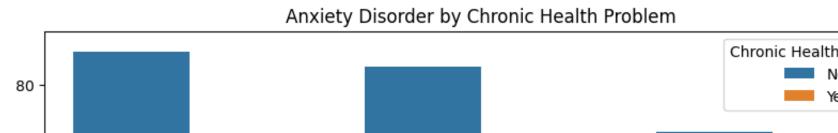


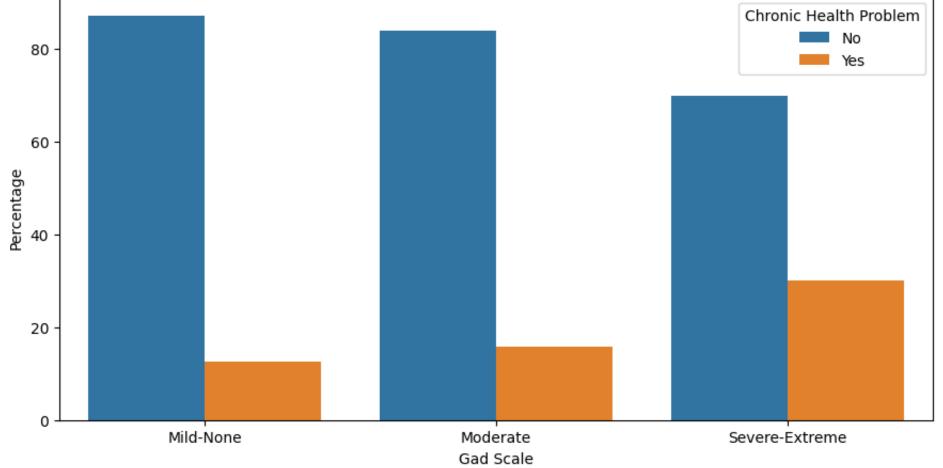




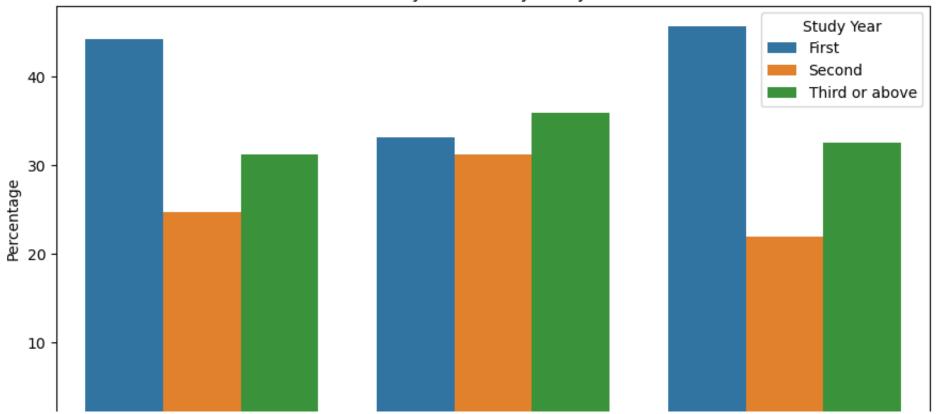


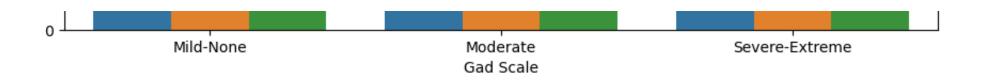












In []: