

# Asthma Disease Prediction using Machine Learning



In [1]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
import plotly.express as px
import plotly.graph_objects as go
```

In [2]:

```
import warnings
warnings.filterwarnings('ignore')
```

In [3]:

```
df = pd.read_csv("asthama.csv")
```

In [4]:

```
df.head()
```

Out[4]:

	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	None_Sympton	Pains	Nasal-Congestion	Runny-Nose	None_E
0	1	1	1	1	0	1	1	1	
1	1	1	1	1	0	1	1	1	
2	1	1	1	1	0	1	1	1	
3	1	1	1	1	0	1	1	1	
4	1	1	1	1	0	1	1	1	

In [5]:

```
df.tail()
```

Out[5]:

	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	None_Sympton	Pains	Nasal-Congestion	Runny-Nose	N
316795	0	0	0	0	1	0	0	0	
316796	0	0	0	0	1	0	0	0	
316797	0	0	0	0	1	0	0	0	
316798	0	0	0	0	1	0	0	0	
316799	0	0	0	0	1	0	0	0	

In [6]:

```
df.shape
```

Out[6]:

(316800, 19)

In [7]:

```
df.columns
```

Out[7]:

```
Index(['Tiredness', 'Dry-Cough', 'Difficulty-in-Breathing', 'Sore-Throat',  
      'None_Sympton', 'Pains', 'Nasal-Congestion', 'Runny-Nose',  
      'None_Experiencing', 'Age_0-9', 'Age_10-19', 'Age_20-24', 'Age_25-59',  
      'Age_60+', 'Gender_Female', 'Gender_Male', 'Severity_Mild',  
      'Severity_Moderate', 'Severity_None'],  
      dtype='object')
```

In [8]:

```
df.duplicated().sum()
```

Out[8]:

```
311040
```

In [9]:

```
df.isnull().sum()
```

Out[9]:

```
Tiredness          0  
Dry-Cough          0  
Difficulty-in-Breathing  0  
Sore-Throat        0  
None_Sympton       0  
Pains              0  
Nasal-Congestion   0  
Runny-Nose         0  
None_Experiencing  0  
Age_0-9            0  
Age_10-19          0  
Age_20-24          0  
Age_25-59          0  
Age_60+            0  
Gender_Female      0  
Gender_Male        0  
Severity_Mild      0  
Severity_Moderate  0  
Severity_None      0  
dtype: int64
```

In [10]:

```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 316800 entries, 0 to 316799
Data columns (total 19 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   Tiredness                            316800 non-null  int64
1   Dry-Cough                           316800 non-null  int64
2   Difficulty-in-Breathing              316800 non-null  int64
3   Sore-Throat                         316800 non-null  int64
4   None_Sympton                        316800 non-null  int64
5   Pains                               316800 non-null  int64
6   Nasal-Congestion                    316800 non-null  int64
7   Runny-Nose                          316800 non-null  int64
8   None_Experiencing                   316800 non-null  int64
9   Age_0-9                             316800 non-null  int64
10  Age_10-19                           316800 non-null  int64
11  Age_20-24                           316800 non-null  int64
12  Age_25-59                           316800 non-null  int64
13  Age_60+                             316800 non-null  int64
14  Gender_Female                       316800 non-null  int64
15  Gender_Male                         316800 non-null  int64
16  Severity_Mild                       316800 non-null  int64
17  Severity_Moderate                   316800 non-null  int64
18  Severity_None                       316800 non-null  int64
dtypes: int64(19)
memory usage: 45.9 MB
```

In [11]:

```
df.describe()
```

Out[11]:

	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	None_Sympton	
count	316800.000000	316800.000000	316800.000000	316800.000000	316800.000000	316800.000000
mean	0.500000	0.562500	0.500000	0.312500	0.062500	0.500000
std	0.500001	0.496079	0.500001	0.463513	0.242062	0.500001
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000
25%	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000
50%	0.500000	1.000000	0.500000	0.000000	0.000000	0.000000
75%	1.000000	1.000000	1.000000	1.000000	0.000000	1.000000
max	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000

In [12]:

```
df.nunique()
```

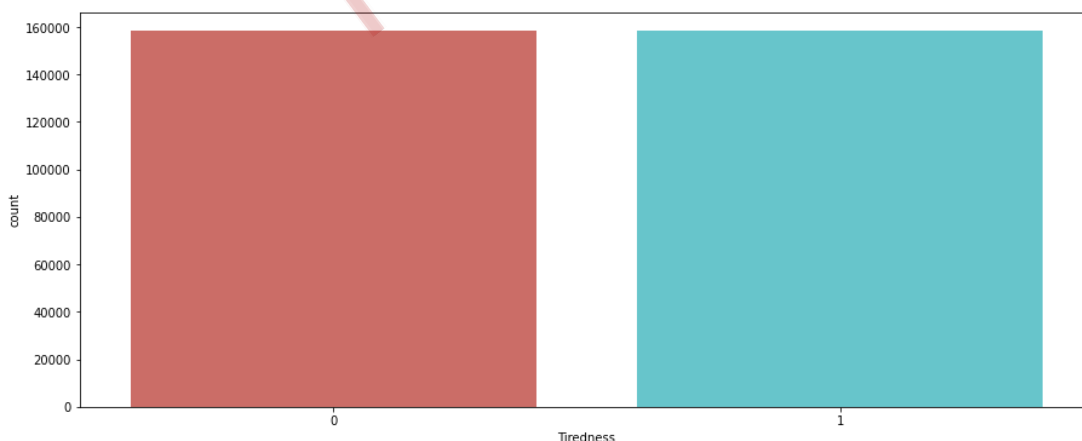
Out[12]:

```
Tiredness                2
Dry-Cough                2
Difficulty-in-Breathing  2
Sore-Throat              2
None_Sympton             2
Pains                    2
Nasal-Congestion         2
Runny-Nose               2
None_Experiencing        2
Age_0-9                  2
Age_10-19                2
Age_20-24                2
Age_25-59                2
Age_60+                  2
Gender_Female            2
Gender_Male              2
Severity_Mild             2
Severity_Moderate        2
Severity_None            2
dtype: int64
```

In [13]:

```
for i in df.columns:
    print('Countplot for:', i)
    plt.figure(figsize=(15,6))
    sns.countplot(x = df[i], data = df, palette = 'hls')
    plt.show()
    print('\n')
```

Countplot for: Tiredness

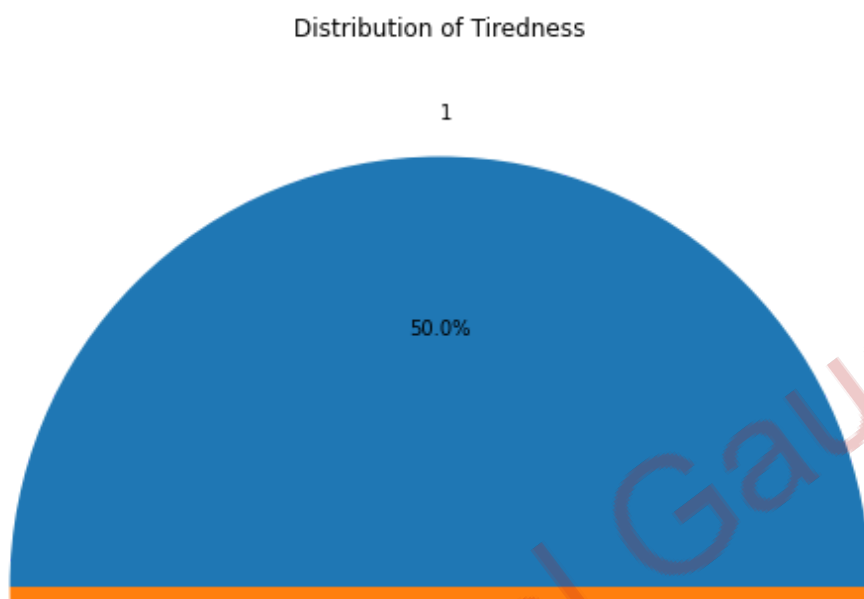


Countplot for: Dry-Cough

In [14]:

```
for i in df.columns:  
    print('Pie plot for:', i)  
    plt.figure(figsize=(20, 10))  
    df[i].value_counts().plot(kind='pie', autopct='%1.1f%%')  
    plt.title('Distribution of ' + i)  
    plt.ylabel('')  
    plt.show()  
    print('\n')
```

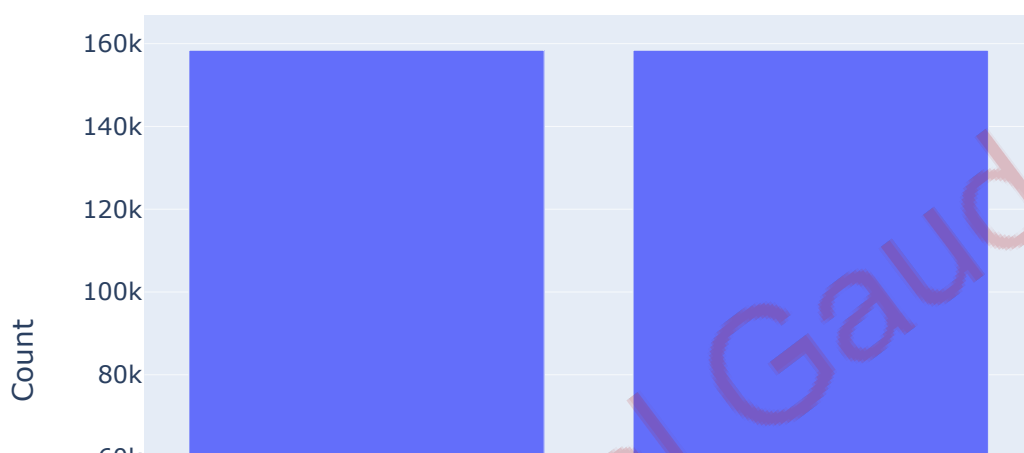
Pie plot for: Tiredness



In [15]:

```
for i in df.columns:
    fig = go.Figure(data=[go.Bar(x=df[i].value_counts().index,
                                y=df[i].value_counts())])
    fig.update_layout(
        title=i,
        xaxis_title=i,
        yaxis_title="Count")
    fig.show()
```

## Tiredness

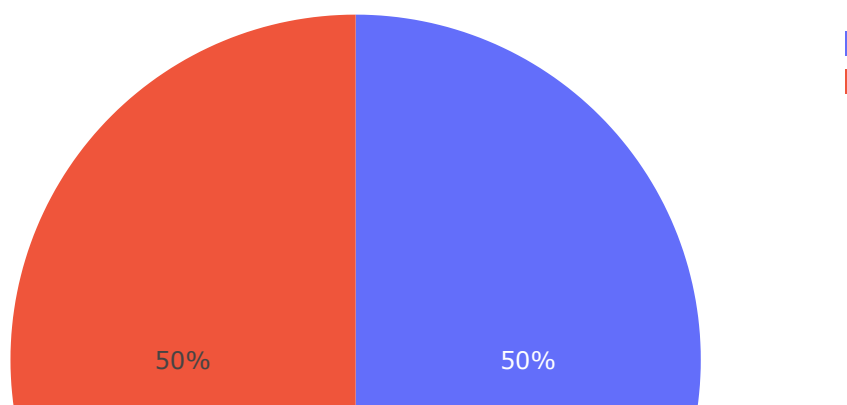


In [16]:

```
for i in df.columns:
    print('Pie plot for:', i)
    fig = px.pie(df, names=i, title='Distribution of ' + i)
    fig.show()
    print('\n')
```

Pie plot for: Tiredness

## Distribution of Tiredness



In [17]:

```
df = df.drop(['Severity_None'], axis = 1)
```

In [18]:

```
df['Asthma_Severity'] = df[['Severity_Mild', 'Severity_Moderate']].idxmax(axis=1)
```

In [19]:

```
severity_mapping = {  
    'Severity_Mild': 'Mild',  
    'Severity_Moderate': 'Moderate'  
}
```

In [20]:

```
df['Asthma_Severity'] = df['Asthma_Severity'].map(severity_mapping)
```

In [21]:

```
df['Asthma_Severity']
```

Out[21]:

```
0      Mild  
1      Mild  
2      Mild  
3  Moderate  
4  Moderate  
...  
316795  Mild  
316796  Mild  
316797  Mild  
316798  Mild  
316799  Mild  
Name: Asthma_Severity, Length: 316800, dtype: object
```

In [22]:

```
severity_mapping = {  
    'Mild': 0,  
    'Moderate': 1  
}
```

In [23]:

```
df['Asthma_Severity'] = df['Asthma_Severity'].map(severity_mapping)
```



In [24]:

```
df['Asthma_Severity'].value_counts()
```

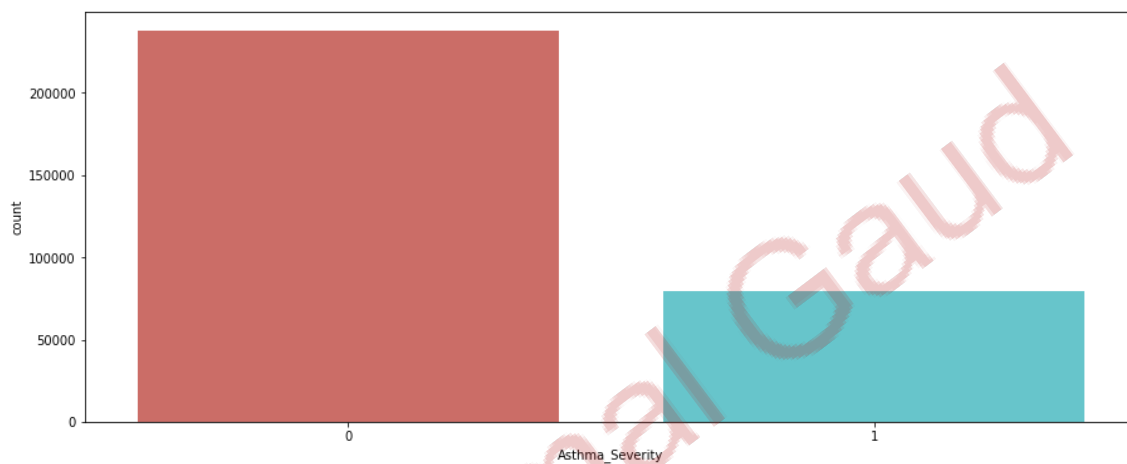
Out[24]:

```
0    237600
1     79200
Name: Asthma_Severity, dtype: int64
```

In [25]:

```
print('Countplot for:', 'Asthma_Severity')
plt.figure(figsize=(15,6))
sns.countplot(x = df['Asthma_Severity'], data = df, palette = 'hls')
plt.show()
```

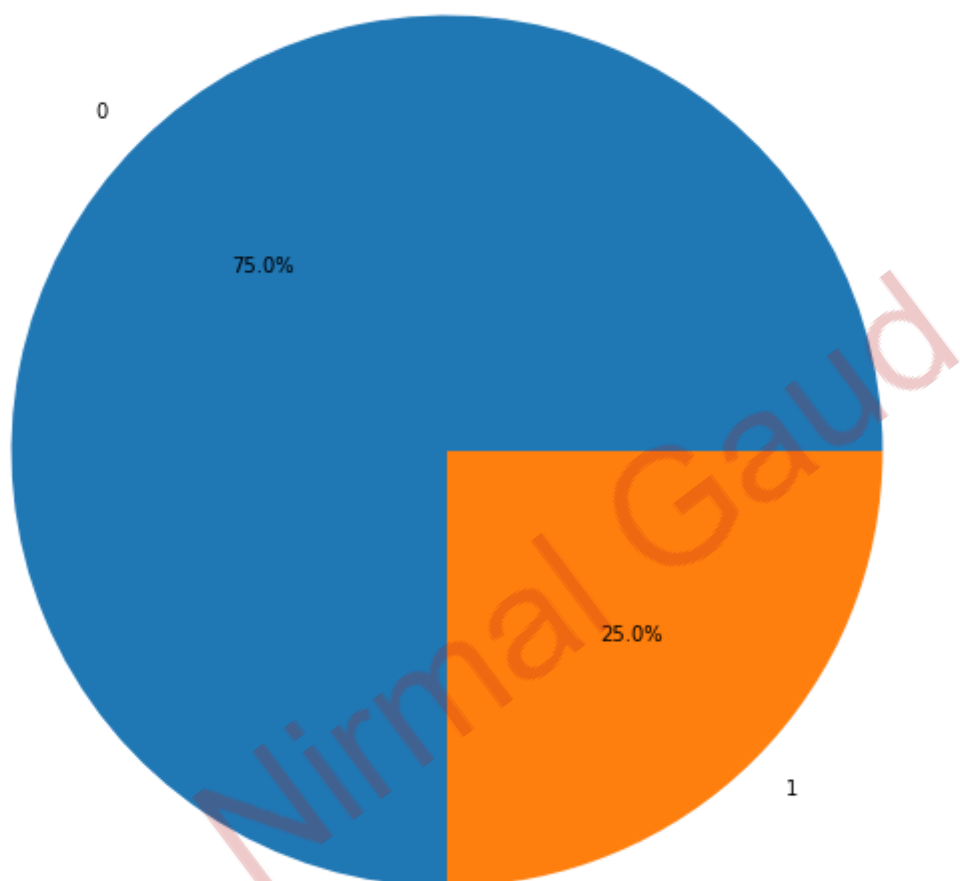
Countplot for: Asthma\_Severity



In [26]:

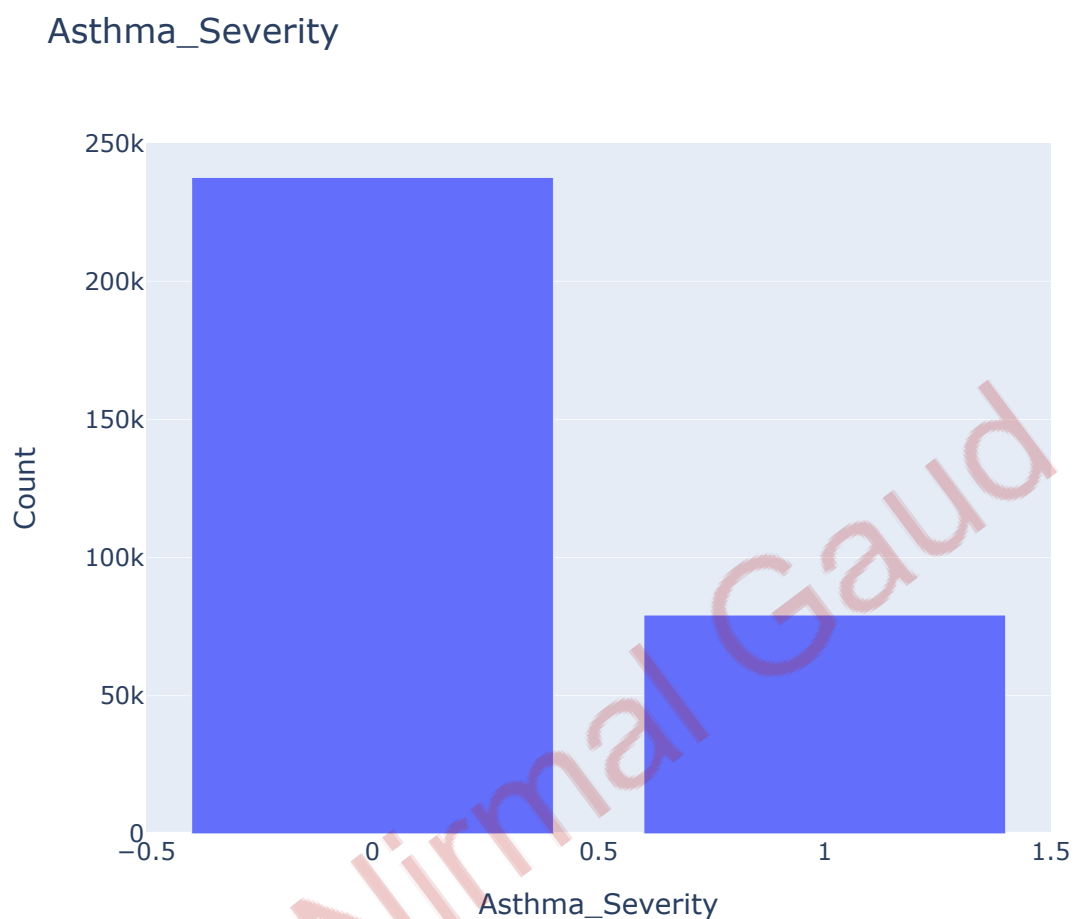
```
print('Pie plot for:', 'Asthma_Severity')
plt.figure(figsize=(20, 10))
df['Asthma_Severity'].value_counts().plot(kind='pie', autopct='%1.1f%%')
plt.ylabel('')
plt.show()
```

Pie plot for: Asthma\_Severity



In [27]:

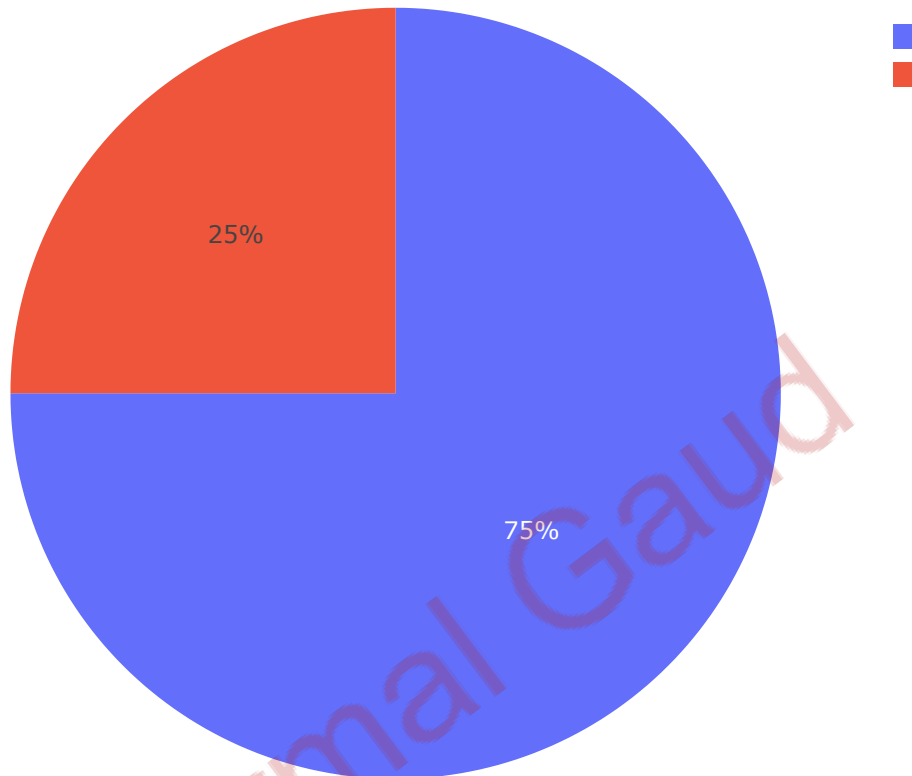
```
fig = go.Figure(data=[go.Bar(x=df['Asthma_Severity'].value_counts().index,  
                             y=df['Asthma_Severity'].value_counts())])  
fig.update_layout(title='Asthma_Severity', xaxis_title='Asthma_Severity',  
                  yaxis_title="Count")  
fig.show()
```



In [28]:

```
print('Pie plot for:', 'Asthma_Severity')
fig = px.pie(df, names='Asthma_Severity')
fig.show()
```

Pie plot for: Asthma\_Severity



In [29]:

```
df.columns
```

Out[29]:

```
Index(['Tiredness', 'Dry-Cough', 'Difficulty-in-Breathing', 'Sore-Throat',
      'None_Sympton', 'Pains', 'Nasal-Congestion', 'Runny-Nose',
      'None_Experiencing', 'Age_0-9', 'Age_10-19', 'Age_20-24', 'Age_25-59',
      'Age_60+', 'Gender_Female', 'Gender_Male', 'Severity_Mild',
      'Severity_Moderate', 'Asthma_Severity'],
      dtype='object')
```

In [30]:

```
severity_levels = ['Severity_Mild', 'Severity_Moderate']  
severity_distribution = df[severity_levels].sum()
```

In [31]:

```
severity_distribution
```

Out[31]:

```
Severity_Mild      79200  
Severity_Moderate  79200  
dtype: int64
```

In [32]:

```
df = df.drop(['Severity_Mild', 'Severity_Moderate'], axis = 1)
```

In [33]:

```
symptoms = ['Tiredness', 'Dry-Cough', 'Difficulty-in-Breathing', 'Sore-Throat', 'Pains',  
symptom_counts = df[symptoms].sum()
```

In [34]:

```
symptom_counts
```

Out[34]:

```
Tiredness      158400  
Dry-Cough      178200  
Difficulty-in-Breathing  158400  
Sore-Throat     99000  
Pains          115200  
Nasal-Congestion  172800  
Runny-Nose      172800  
dtype: int64
```

In [35]:

```
age_groups = ['Age_0-9', 'Age_10-19', 'Age_20-24', 'Age_25-59', 'Age_60+']  
age_distribution = df[age_groups].sum()
```

In [36]:

```
age_distribution
```

Out[36]:

```
Age_0-9      63360
Age_10-19    63360
Age_20-24    63360
Age_25-59    63360
Age_60+      63360
dtype: int64
```

In [37]:

```
gender_groups = ['Gender_Female', 'Gender_Male']
gender_distribution = df[gender_groups].sum()
```

In [38]:

```
gender_distribution
```

Out[38]:

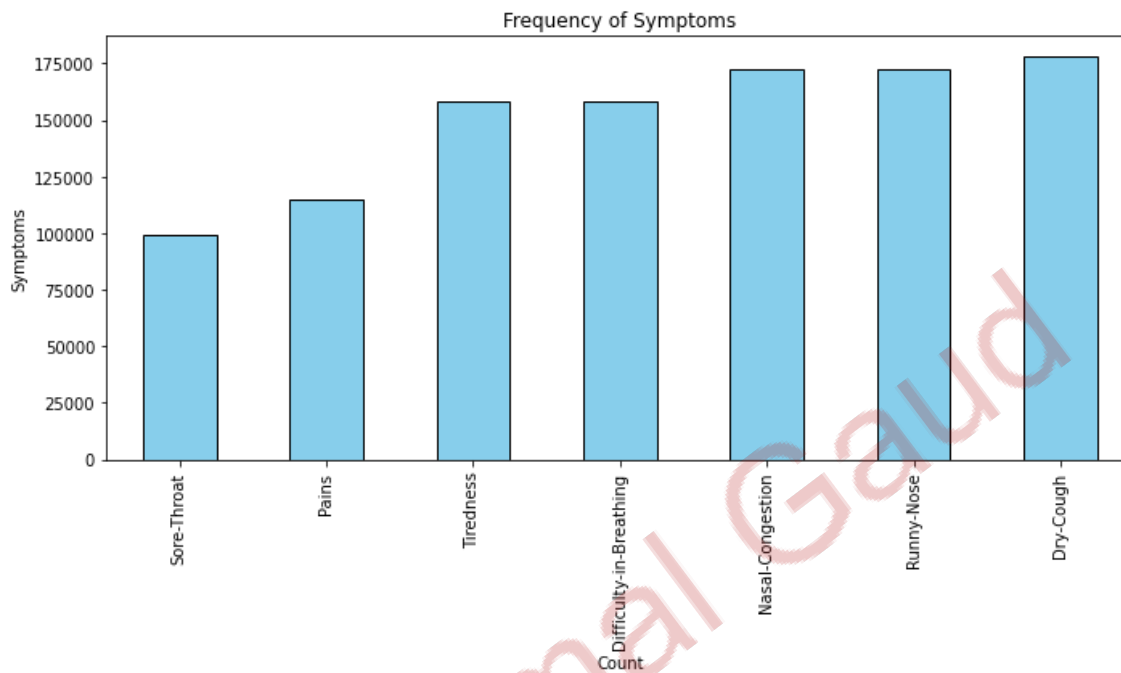
```
Gender_Female  105600
Gender_Male    105600
dtype: int64
```

In [39]:

```
fig, ax = plt.subplots(figsize=(10, 6))

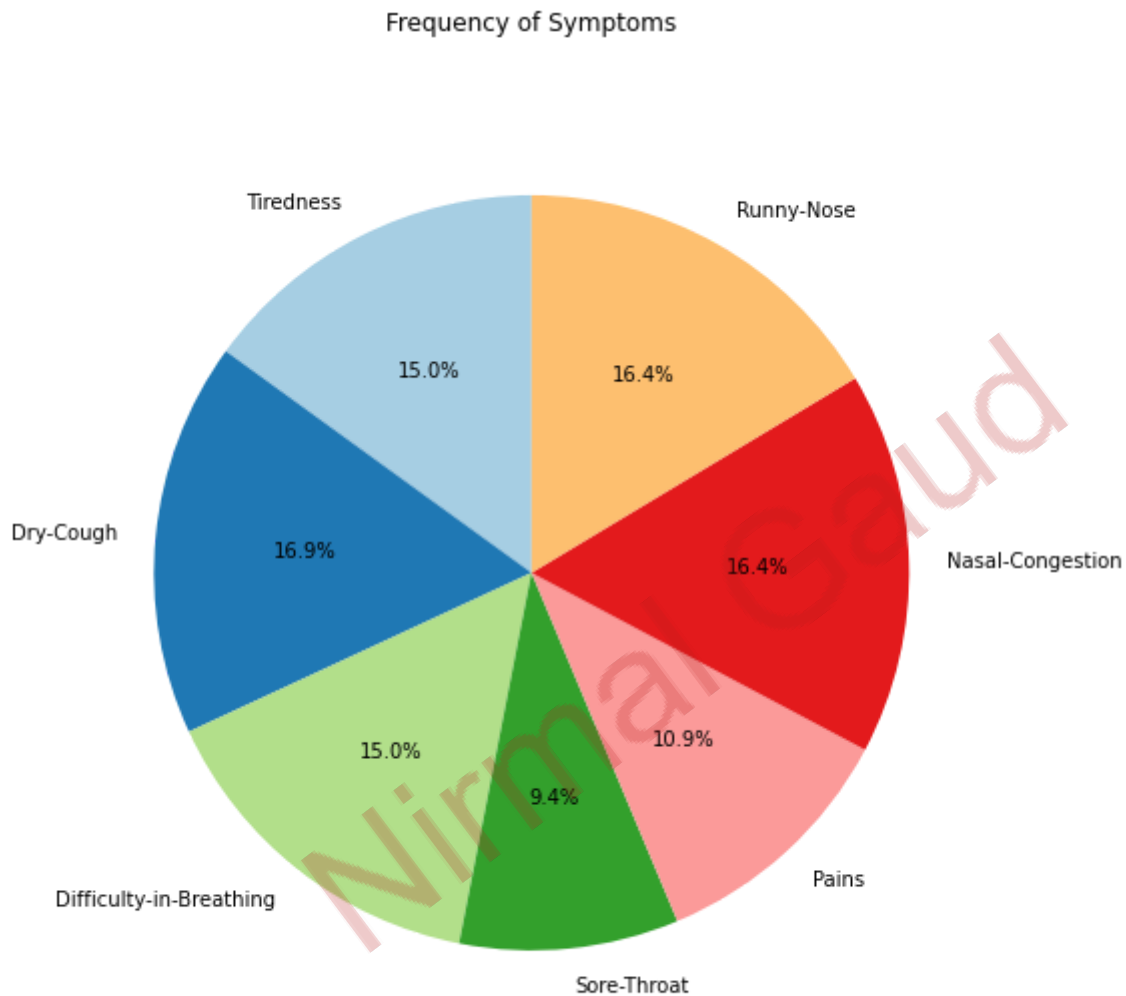
symptom_counts.sort_values().plot(kind='bar', ax=ax, color='skyblue',
                                   edgecolor='black')
ax.set_title('Frequency of Symptoms')
ax.set_xlabel('Count')
ax.set_ylabel('Symptoms')

plt.tight_layout()
plt.show()
```



In [40]:

```
fig, ax = plt.subplots(figsize=(8, 8))
ax.pie(symptom_counts, labels=symptom_counts.index, autopct='%1.1f%%',
      startangle=90,
      colors=plt.cm.Paired(range(len(symptom_counts))))
ax.set_title('Frequency of Symptoms')
plt.axis('equal')
plt.tight_layout()
plt.show()
```

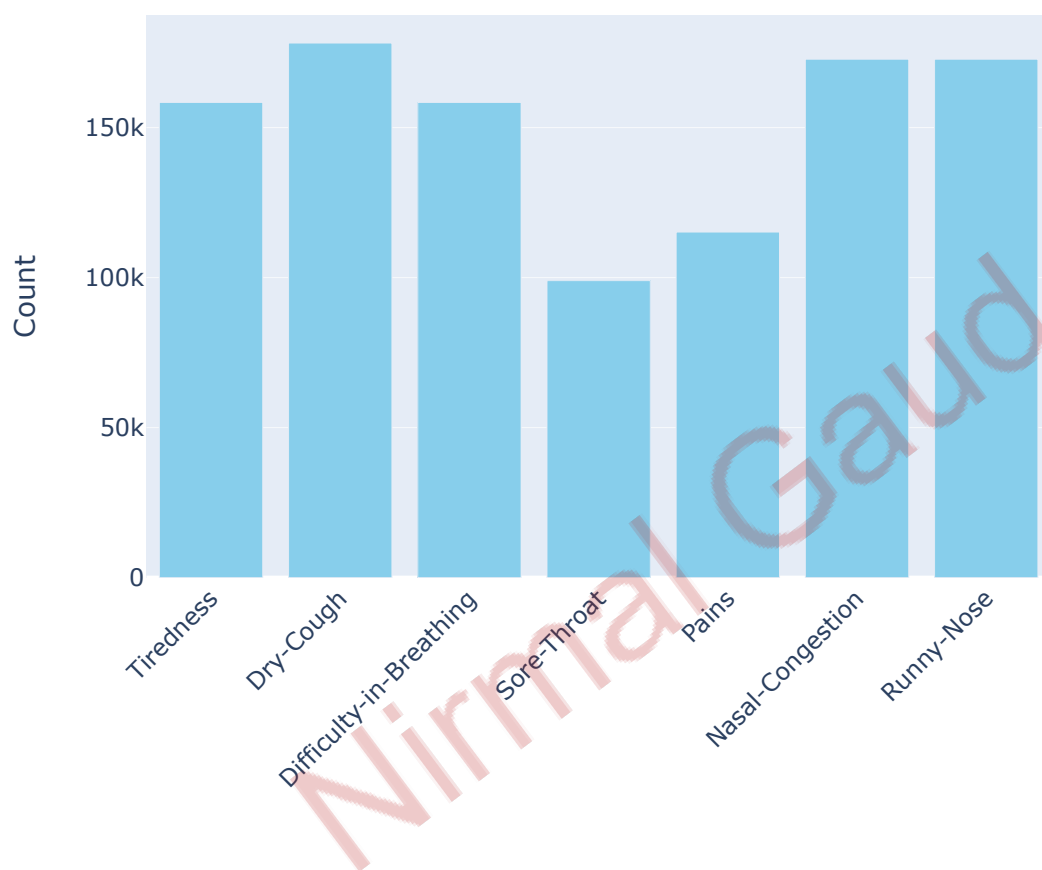




In [41]:

```
fig = px.bar(symptom_counts, x=symptom_counts.index, y=symptom_counts.values,  
             labels={'x': 'Symptoms', 'y': 'Count'},  
             title='Frequency of Symptoms', color_discrete_sequence=['skyblue'])  
fig.update_layout(xaxis_tickangle=-45)  
fig.show()
```

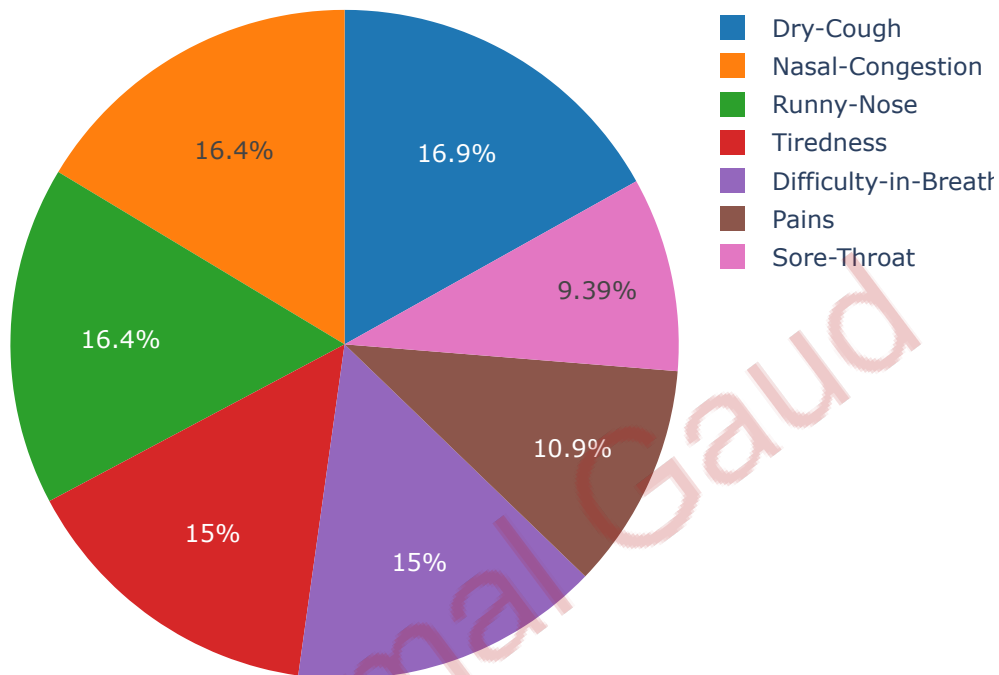
Frequency of Symptoms



In [42]:

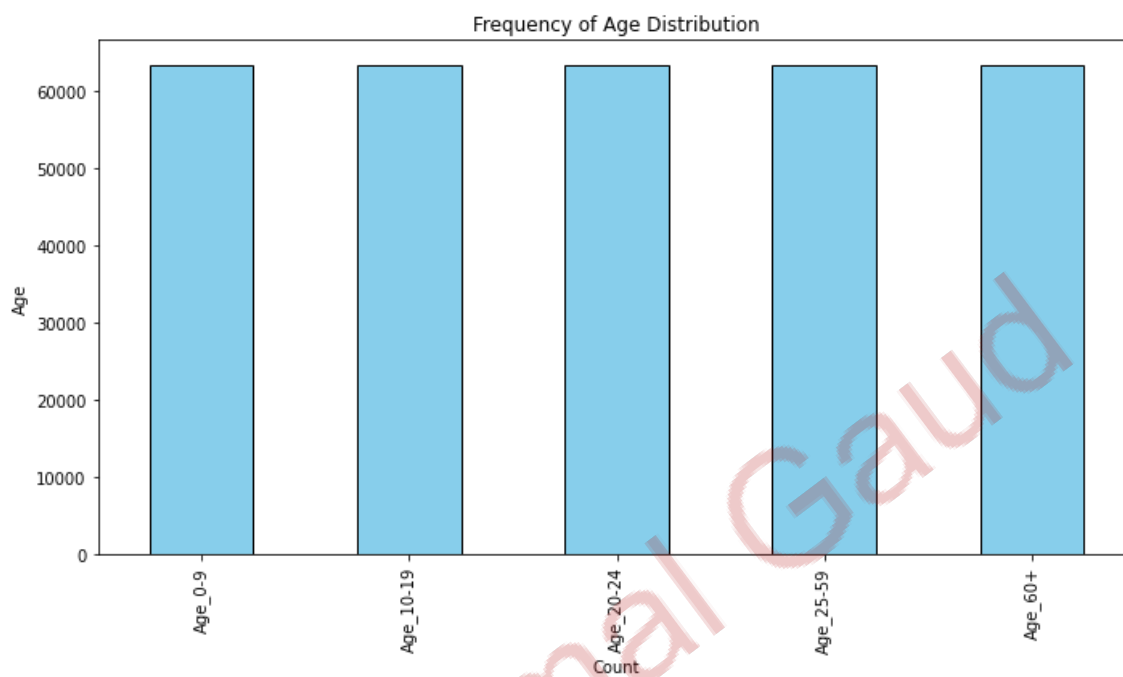
```
custom_colors = ['#1f77b4', '#ff7f0e', '#2ca02c', '#d62728', '#9467bd', '#8c564b', '#e377c2']  
fig = px.pie(symptom_counts, values=symptom_counts.values, names=symptom_counts.index,  
             title='Frequency of Symptoms', color_discrete_sequence=custom_colors)  
fig.show()
```

Frequency of Symptoms



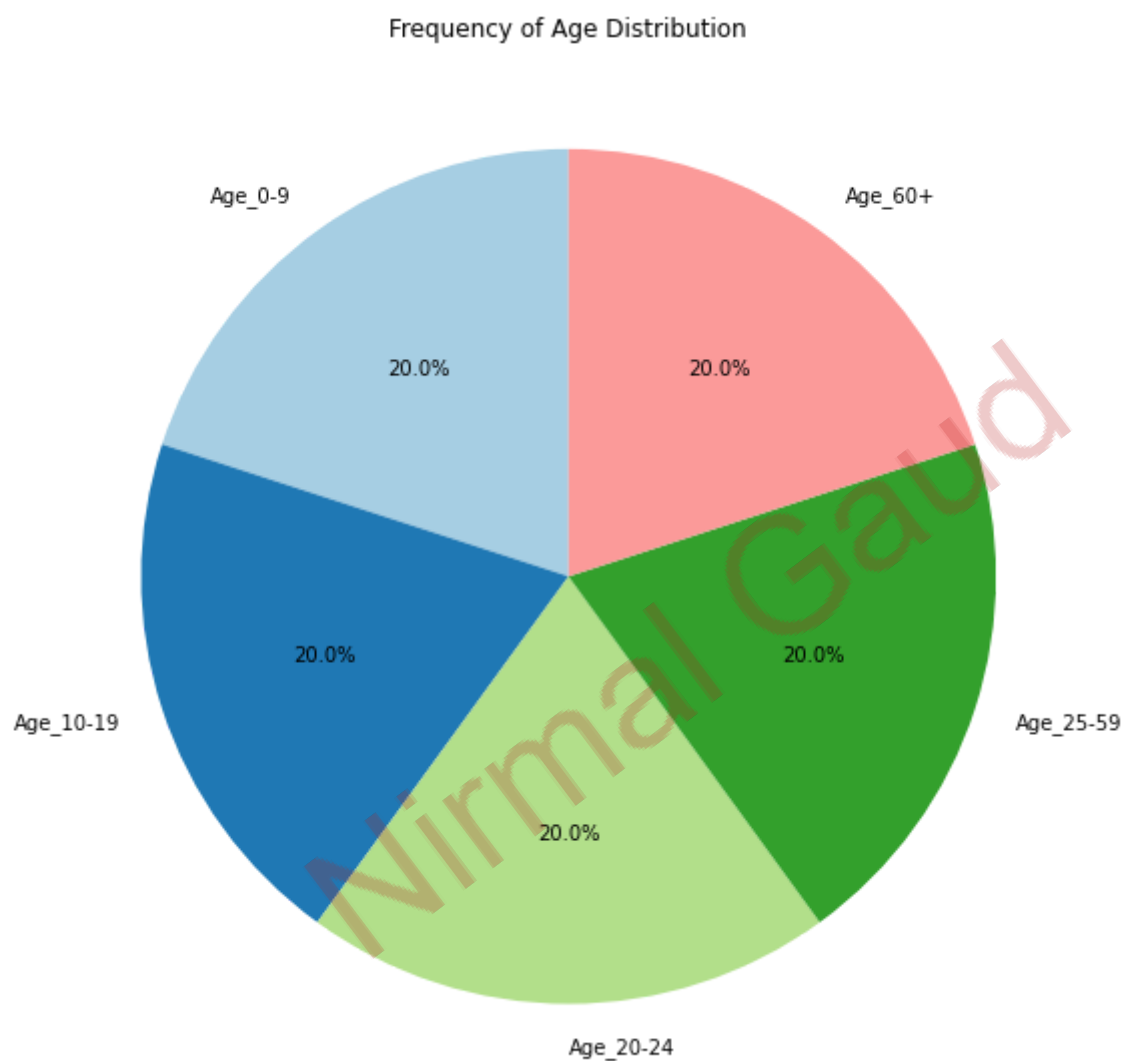
In [43]:

```
fig, ax = plt.subplots(figsize=(10, 6))  
age_distribution.sort_values().plot(kind='bar', ax=ax, color='skyblue',  
                                     edgecolor='black')  
ax.set_title('Frequency of Age Distribution')  
ax.set_xlabel('Count')  
ax.set_ylabel('Age')  
  
plt.tight_layout()  
plt.show()
```



In [44]:

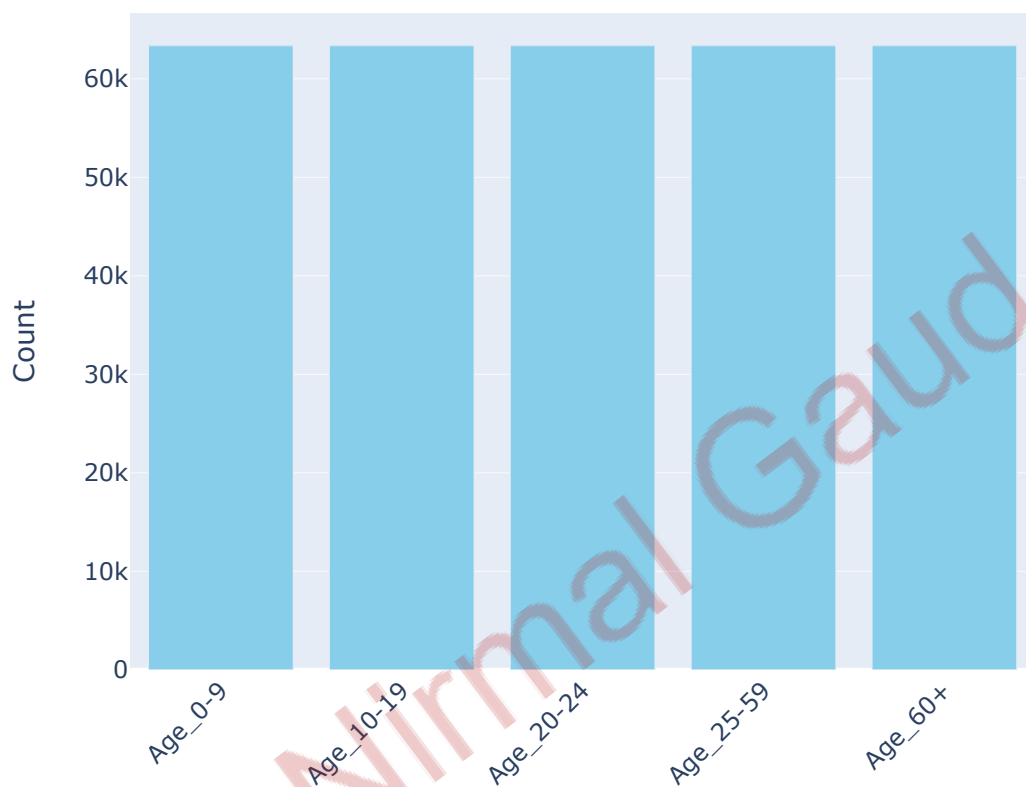
```
fig, ax = plt.subplots(figsize=(8, 8))
ax.pie(age_distribution, labels=age_distribution.index, autopct='%1.1f%%',
       startangle=90,
       colors=plt.cm.Paired(range(len(age_distribution))))
ax.set_title('Frequency of Age Distribution')
plt.axis('equal')
plt.tight_layout()
plt.show()
```



In [45]:

```
fig = px.bar(age_distribution, x=age_distribution.index, y=age_distribution.values,  
             labels={'x': 'Age Distribution', 'y': 'Count'},  
             title='Frequency of Age Distribution', color_discrete_sequence=['skyblue'])  
fig.update_layout(xaxis_tickangle=-45)  
fig.show()
```

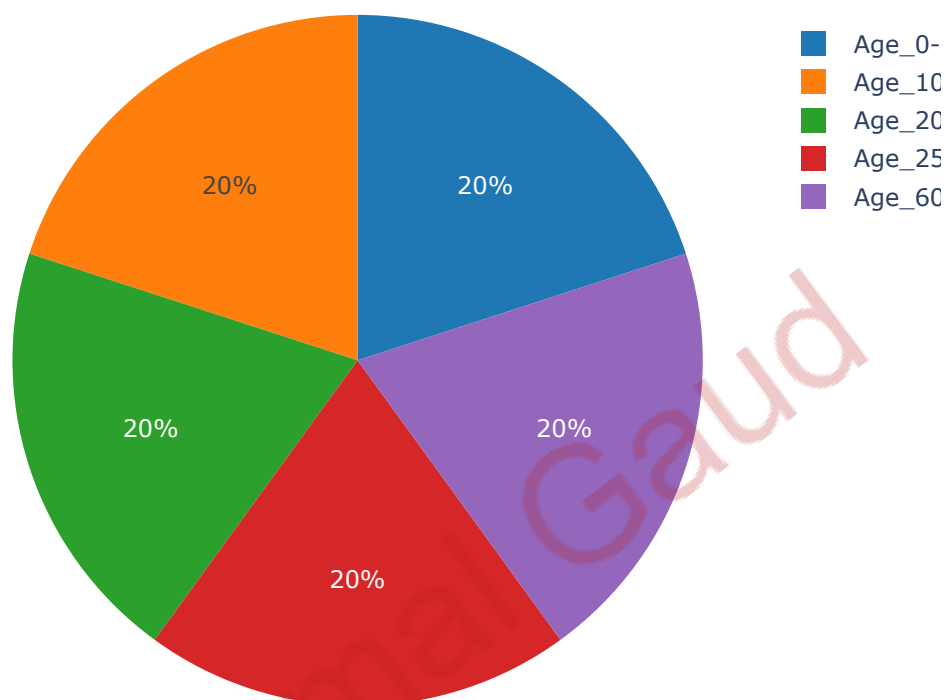
Frequency of Age Distribution



In [46]:

```
custom_colors = ['#1f77b4', '#ff7f0e', '#2ca02c', '#d62728', '#9467bd', '#8c564b', '#e377c2']  
fig = px.pie(age_distribution, values=age_distribution.values, names=age_distribution.index, title='Frequency of Age Distribution', color_discrete_sequence=custom_colors)  
fig.show()
```

Frequency of Age Distribution

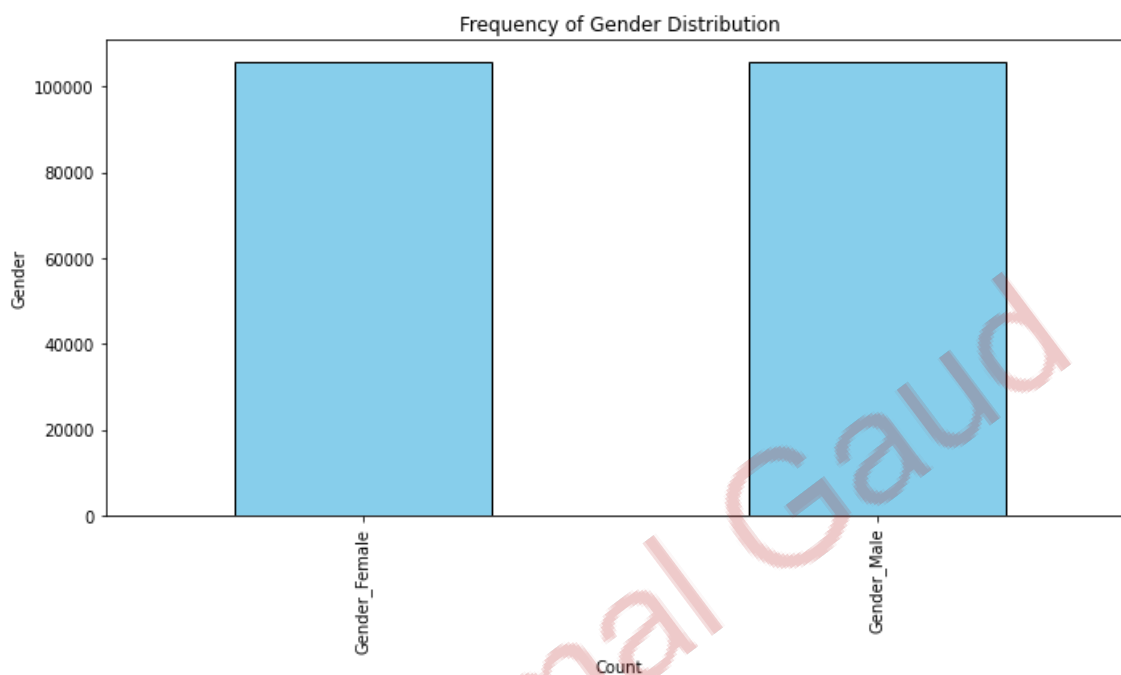


In [47]:

```
fig, ax = plt.subplots(figsize=(10, 6))

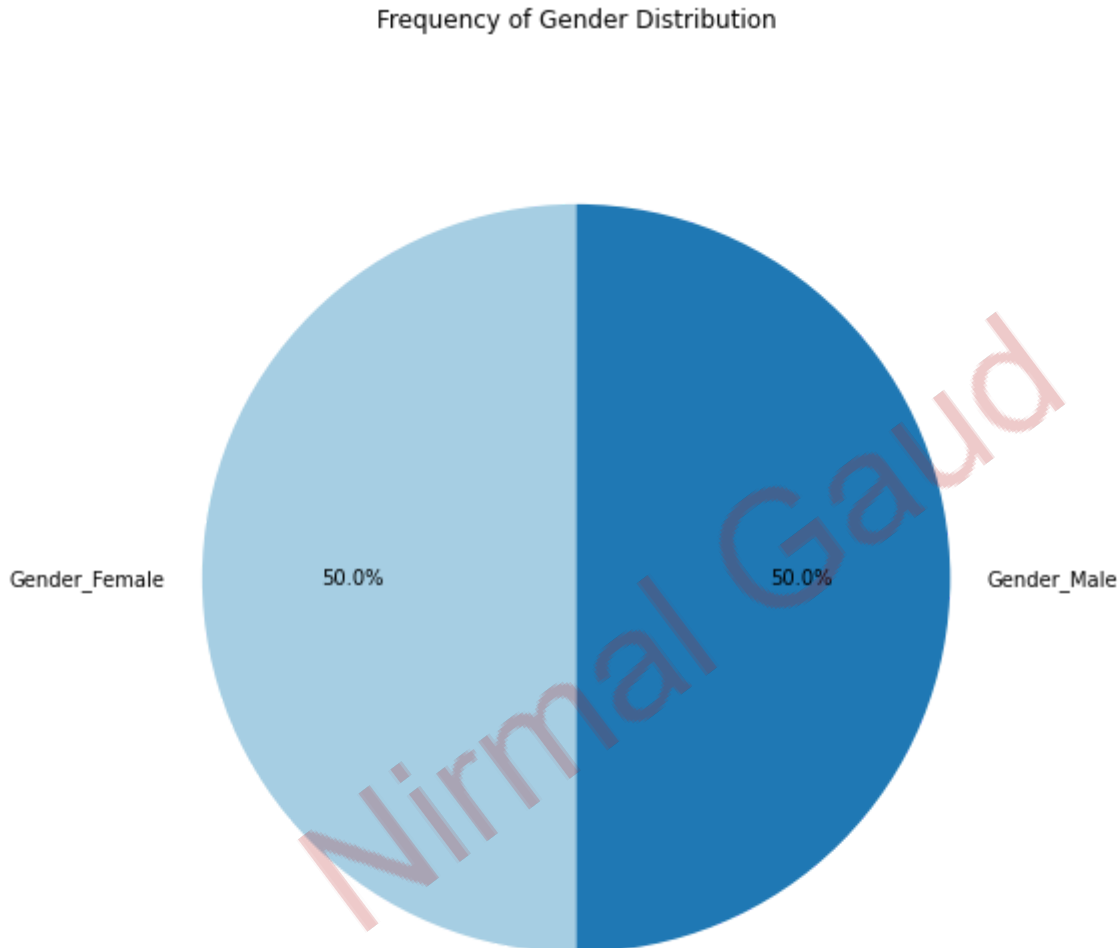
gender_distribution.sort_values().plot(kind='bar', ax=ax, color='skyblue',
                                       edgecolor='black')
ax.set_title('Frequency of Gender Distribution')
ax.set_xlabel('Count')
ax.set_ylabel('Gender')

plt.tight_layout()
plt.show()
```



In [48]:

```
fig, ax = plt.subplots(figsize=(8, 8))
ax.pie(gender_distribution, labels=gender_distribution.index, autopct='%1.1f%%',
      startangle=90,
      colors=plt.cm.Paired(range(len(age_distribution))))
ax.set_title('Frequency of Gender Distribution')
plt.axis('equal')
plt.tight_layout()
plt.show()
```

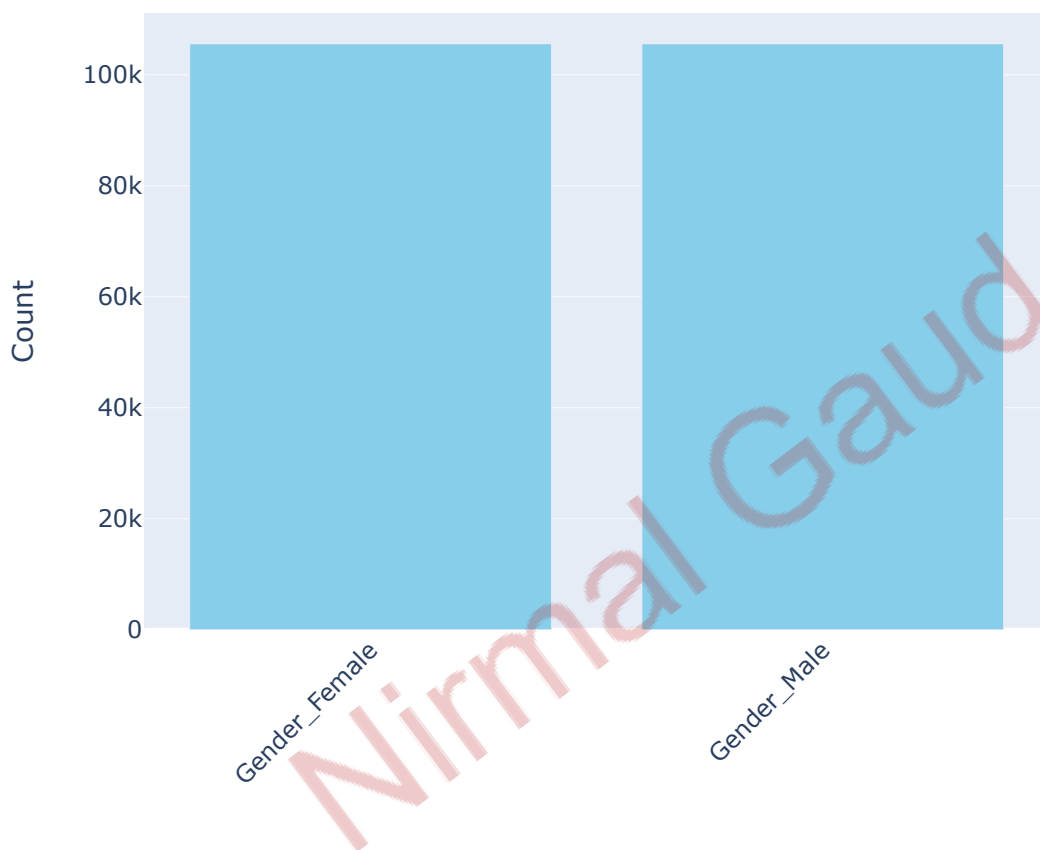




In [49]:

```
fig = px.bar(gender_distribution, x=gender_distribution.index, y=gender_distribution.val,
             labels={'x': 'Gender Distribution', 'y': 'Count'},
             title='Frequency of Gender Distribution', color_discrete_sequence=['skyblue'])
fig.update_layout(xaxis_tickangle=-45)
fig.show()
```

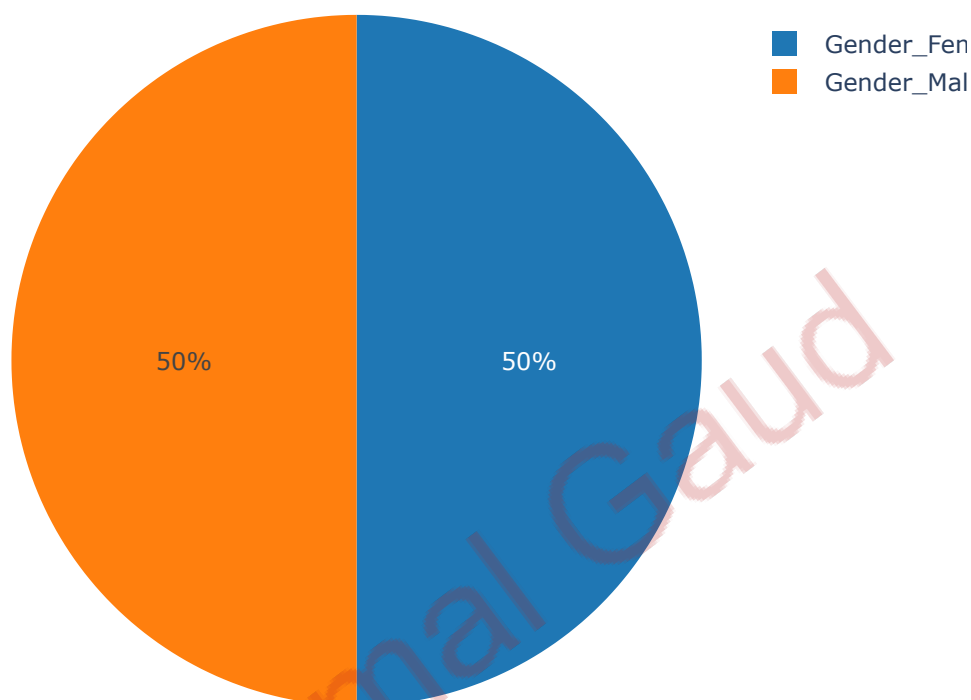
Frequency of Gender Distribution



In [50]:

```
custom_colors = ['#1f77b4', '#ff7f0e', '#2ca02c', '#d62728', '#9467bd', '#8c564b', '#e377c2']
fig = px.pie(gender_distribution, values=gender_distribution.values, names=gender_distribution.index,
             title='Frequency of Gender Distribution', color_discrete_sequence=custom_colors)
fig.show()
```

Frequency of Gender Distribution



In [51]:

```
corr = df[symptoms].corr()
```

In [52]:

```
corr
```

Out[52]:

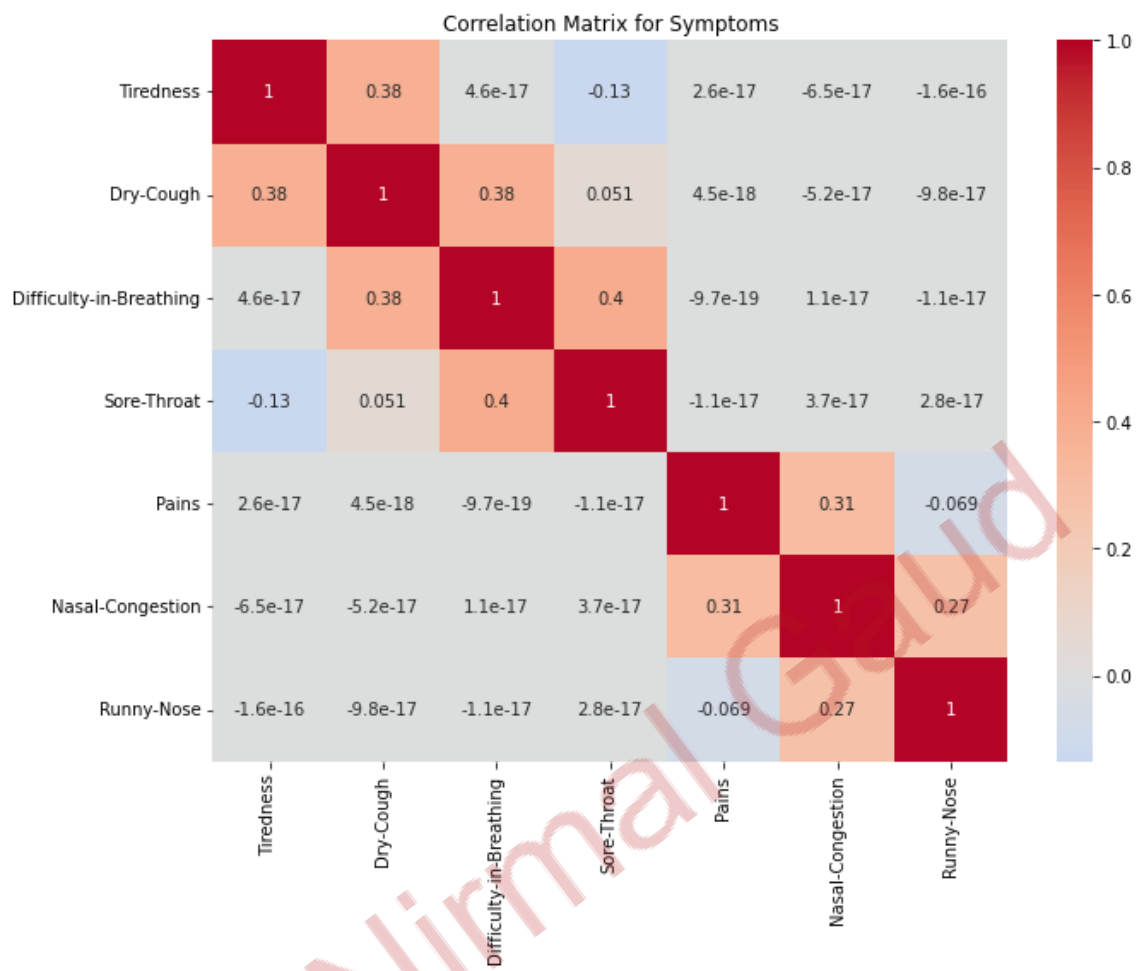
	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	Pains	N Conge
Tiredness	1.000000e+00	3.779645e-01	4.629153e-17	-1.348400e-01	2.612420e-17	-6.526
Dry-Cough	3.779645e-01	1.000000e+00	3.779645e-01	5.096472e-02	4.519459e-18	-5.180
Difficulty-in-Breathing	4.629153e-17	3.779645e-01	1.000000e+00	4.045199e-01	-9.663757e-19	1.07676
Sore-Throat	-1.348400e-01	5.096472e-02	4.045199e-01	1.000000e+00	-1.063529e-17	3.70577
Pains	2.612420e-17	4.519459e-18	-9.663757e-19	-1.063529e-17	1.000000e+00	3.10529
Nasal-Congestion	-6.526807e-17	-5.180861e-17	1.076763e-17	3.705775e-17	3.105295e-01	1.000000
Runny-Nose	-1.561936e-16	-9.767619e-17	-1.091121e-17	2.844633e-17	-6.900656e-02	2.66666



Nirmal Gaud

In [53]:

```
fig, ax = plt.subplots(figsize=(10, 8))
sns.heatmap(corr, annot=True, cmap='coolwarm', center=0, ax=ax)
ax.set_title('Correlation Matrix for Symptoms')
plt.tight_layout()
plt.show()
```



In [54]:

```
severity_symptom_means = df.groupby(by=[col for col in df.columns if "Asthma_Severity" i
```

In [55]:

```
severity_symptom_means
```

Out[55]:

	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	Pains	Nasal-Congestion	Runny-Nose
Asthma_Severity							
0	0.5	0.5625	0.5	0.3125	0.363636	0.545455	0.545455
1	0.5	0.5625	0.5	0.3125	0.363636	0.545455	0.545455

In [56]:

```
severity_symptom_means = severity_symptom_means.transpose()
```

In [57]:

```
severity_symptom_means
```

Out[57]:

<b>Asthma_Severity</b>	<b>0</b>	<b>1</b>
<b>Tiredness</b>	0.500000	0.500000
<b>Dry-Cough</b>	0.562500	0.562500
<b>Difficulty-in-Breathing</b>	0.500000	0.500000
<b>Sore-Throat</b>	0.312500	0.312500
<b>Pains</b>	0.363636	0.363636
<b>Nasal-Congestion</b>	0.545455	0.545455
<b>Runny-Nose</b>	0.545455	0.545455

In [58]:

```
data = {  
    'Asthma_Severity': [0, 1, 2],  
    'Tiredness': [0.500000, 0.500000, 0.500000],  
    'Dry-Cough': [0.562500, 0.562500, 0.562500],  
    'Difficulty-in-Breathing': [0.500000, 0.500000, 0.500000],  
    'Sore-Throat': [0.312500, 0.312500, 0.312500],  
    'Pains': [0.363636, 0.363636, 0.363636],  
    'Nasal-Congestion': [0.545455, 0.545455, 0.545455],  
    'Runny-Nose': [0.545455, 0.545455, 0.545455]  
}
```

In [59]:

```
df1 = pd.DataFrame(data)
```

In [60]:

```
df1.set_index('Asthma_Severity', inplace=True)
```

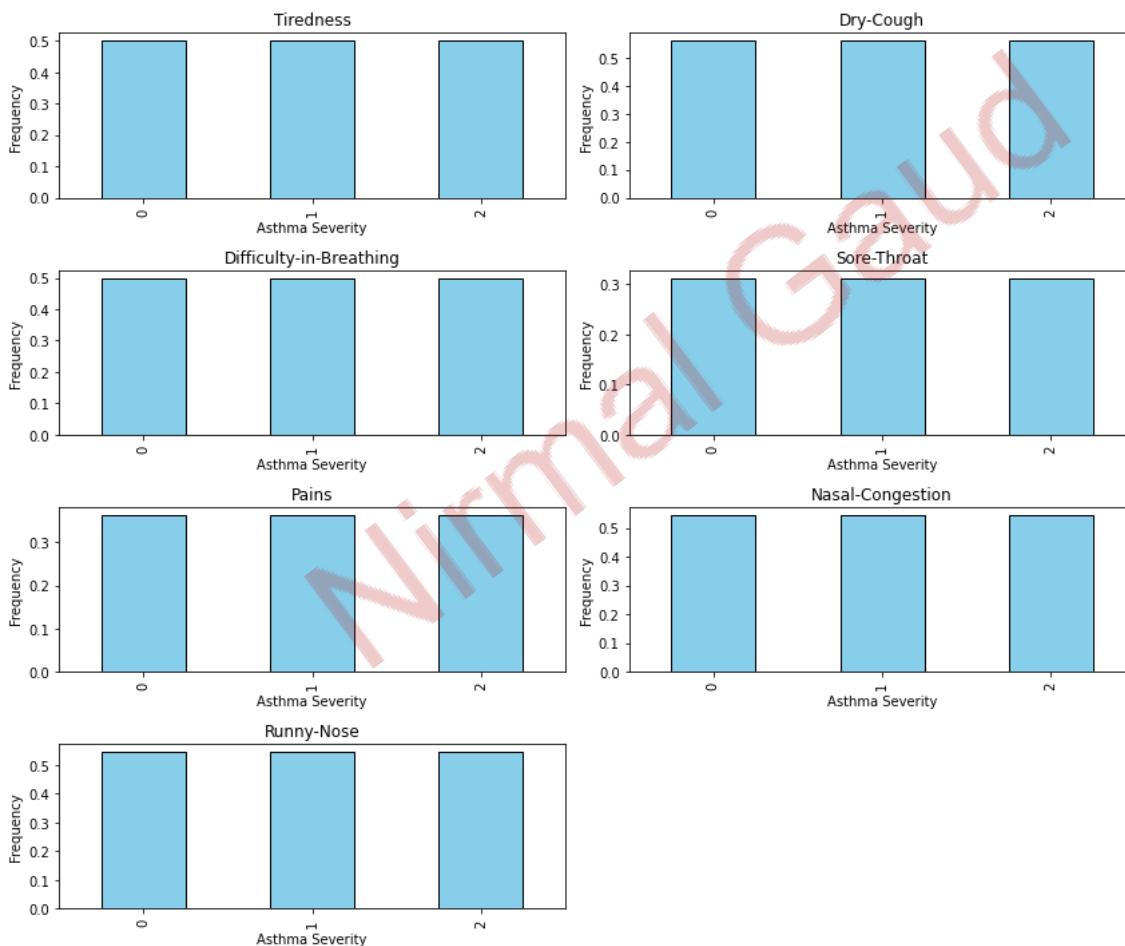
In [61]:

```
fig, axes = plt.subplots(nrows=4, ncols=2, figsize=(12, 10))
symptoms = ['Tiredness', 'Dry-Cough', 'Difficulty-in-Breathing', 'Sore-Throat', 'Pains',
            'Runny-Nose', 'Nasal-Congestion']

for i, symptom in enumerate(symptoms):
    row, col = divmod(i, 2)
    df1[symptom].plot(kind='bar', ax=axes[row, col], color='skyblue', edgecolor='black')
    axes[row, col].set_title(symptom)
    axes[row, col].set_xlabel('Asthma Severity')
    axes[row, col].set_ylabel('Frequency')

if len(symptoms) < len(axes.flat):
    for i in range(len(symptoms), len(axes.flat)):
        fig.delaxes(axes.flatten()[i])

plt.tight_layout()
plt.show()
```



In [62]:

```
df
```

Out[62]:

	Tiredness	Dry-Cough	Difficulty-in-Breathing	Sore-Throat	None_Sympton	Pains	Nasal-Congestion	Runny-Nose	N
0	1	1	1	1	0	1	1	1	
1	1	1	1	1	0	1	1	1	
2	1	1	1	1	0	1	1	1	
3	1	1	1	1	0	1	1	1	
4	1	1	1	1	0	1	1	1	
...	...	...	...	...	...	...	...	...	
316795	0	0	0	0	1	0	0	0	
316796	0	0	0	0	1	0	0	0	
316797	0	0	0	0	1	0	0	0	
316798	0	0	0	0	1	0	0	0	
316799	0	0	0	0	1	0	0	0	

316800 rows × 17 columns

In [63]:

```
X = df.drop(['Asthma_Severity'], axis = 1)
y = df['Asthma_Severity']
```

In [64]:

```
from sklearn.ensemble import RandomForestClassifier
```

In [65]:

```
clf = RandomForestClassifier(n_estimators=100, random_state=42)
```

In [66]:

```
clf.fit(X, y)
```

Out[66]:

```
RandomForestClassifier
RandomForestClassifier(random_state=42)
```

In [67]:

```
feature_importances = clf.feature_importances_
```

In [68]:

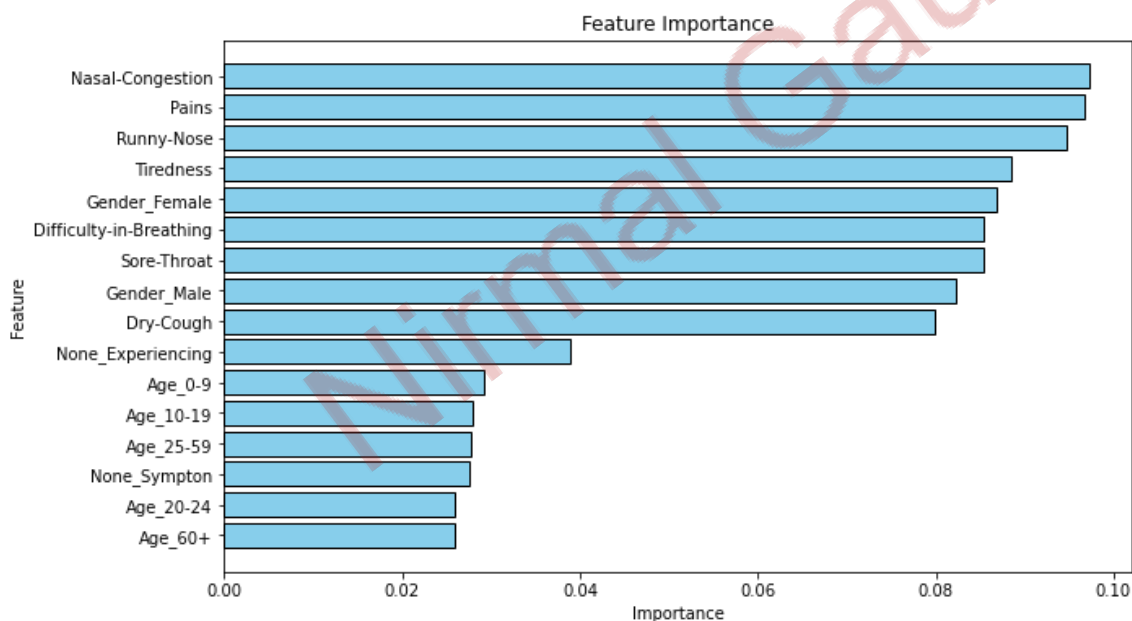
```
feature_importance_df = pd.DataFrame({'Feature': X.columns, 'Importance': feature_import
```

In [69]:

```
feature_importance_df = feature_importance_df.sort_values(by='Importance', ascending=False)
```

In [70]:

```
plt.figure(figsize=(10, 6))
plt.barh(feature_importance_df['Feature'], feature_importance_df['Importance'], color='s')
plt.xlabel('Importance')
plt.ylabel('Feature')
plt.title('Feature Importance')
plt.gca().invert_yaxis()
```



In [71]:

```
X = df[symptoms + gender_groups]
```

In [72]:

```
from sklearn.linear_model import LogisticRegression
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
```



In [73]:

```
X_train,X_test,y_train,y_test=train_test_split(X,
                                                y,
                                                test_size=0.2,
                                                stratify = y,
                                                random_state=42)
```

In [74]:

```
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report
```

In [75]:

```
lr = LogisticRegression(random_state=42)
```

In [76]:

```
lr.fit(X_train, y_train)
```

Out[76]:

▼	LogisticRegression
LogisticRegression(random_state=42)	

In [77]:

```
y_pred_lr = lr.predict(X_test)
```

In [78]:

```
accuracy = accuracy_score(y_test, y_pred_lr)
print(f'Accuracy: {accuracy:.2f}')
```

Accuracy: 0.75

In [79]:

```
dt = DecisionTreeClassifier()
```

In [80]:

```
model_dt = dt.fit(X_train, y_train)
```

In [81]:

```
y_pred_dt = model_dt.predict(X_test)
```

In [82]:

```
accuracy = accuracy_score(y_test, y_pred_dt)
print(f'Accuracy: {accuracy:.2f}')
```

Accuracy: 0.75

In [83]:

```
import xgboost as xgb
```

In [84]:

```
clf = xgb.XGBClassifier(
    objective='multi:softmax',
    num_class=len(pd.unique(y)),
    eval_metric='mlogloss',
    use_label_encoder=False
)
```

In [85]:

```
clf.fit(X_train, y_train)
```

Out[85]:

```
XGBClassifier
XGBClassifier(base_score=0.5, booster='gbtree', callbacks=None,
               colsample_bylevel=1, colsample_bynode=1, colsample_bytree=1,
               early_stopping_rounds=None, enable_categorical=False,
               eval_metric='mlogloss', gamma=0, gpu_id=-1,
               grow_policy='depthwise', importance_type=None,
               interaction_constraints='', learning_rate=0.300000012,
               max_bin=256, max_cat_to_onehot=4, max_delta_step=0, max_
depth=6,
               max_leaves=0, min_child_weight=1, missing=nan,
```

In [86]:

```
y_pred_xgb = clf.predict(X_test)
```

In [87]:

```
accuracy = accuracy_score(y_test, y_pred_xgb)
print(f'Accuracy: {accuracy:.2f}')
```

Accuracy: 0.75

# Thanks !!!