



Project Title	Drugs, Side Effects and Medical Condition arrow_drop_up
Tools	Python, ML, SQL, Excel
Technologies	Data Analyst & Data scientist
Project Difficulties level	intermediate

Dataset : Dataset is available in the given link. You can download it at your convenience.

[Click here to download data set](#)

## About Dataset

Data contains details of various drugs (used for conditions like Acne, Cancer, Heart Disease, etc. ) and their side effects

Drugs detail URLs were collected from following dataset

Major Column Descriptors:

generic\_name:

The chemical name of the drug (not brand name)

drug\_classes:

The drug belongs to which drug class, i.e a drug class is a set of medications and other compounds that have a similar chemical structure, the same mechanism of action (i.e. binding to the same biological target), a related mode of action, and/or are used to treat the same disease.

brand\_names:

brand names in which the drugs are being sold or available in the market.

activity:

Activity is based on recent site visitor activity relative to other medications in the list. Data was gathered from <https://www.drugs.com>

rx\_otc:

Rx-to-OTC switch is the transfer of proven prescription drugs to nonprescription, where

OTC (Over-the-counter) = Medication that can be purchased without a medical prescription

Rx = Prescription Needed

Rx/OTC = Prescription or Over-the-counter.

pregnancy\_category:

A = Adequate and well-controlled studies have failed to demonstrate a risk to the fetus in the first trimester of pregnancy (and there is no evidence of risk in later trimesters).

B = Animal reproduction studies have failed to demonstrate a risk to the fetus and there are no adequate and well-controlled studies in pregnant women.

C = Animal reproduction studies have shown an adverse effect on the fetus and there are no adequate and well-controlled studies in humans, but potential benefits may warrant use in pregnant women despite potential risks.

D = There is positive evidence of human fetal risk based on adverse reaction data from investigational or marketing experience or studies in humans, but potential benefits may warrant use in pregnant women despite potential risks.

X = Studies in animals or humans have demonstrated fetal abnormalities and/or there is positive evidence of human fetal risk based on adverse reaction data from investigational or marketing experience, and the risks involved in use in pregnant women clearly outweigh potential benefits.

N = FDA has not classified the drug.

csa:

Controlled Substances Act (CSA) Schedule

M = The drug has multiple schedules. The schedule may depend on the exact dosage form or strength of the medication.

U = CSA Schedule is unknown.

N = Is not subject to the Controlled Substances Act.

1 = Has a high potential for abuse. Has no currently accepted medical use in treatment in the United States. There is a lack of accepted safety for use under medical supervision.

2 = Has a high potential for abuse. Has a currently accepted medical use in treatment in the United States or a currently accepted medical use with severe restrictions. Abuse may lead to severe psychological or physical dependence.

3 = Has a potential for abuse less than those in schedules 1 and 2. Has a currently accepted medical use in treatment in the United States. Abuse may lead to moderate or low physical dependence or high psychological dependence.

4 = Has a low potential for abuse relative to those in schedule 3. It has a currently accepted medical use in treatment in the United States. Abuse may lead to limited physical dependence or psychological dependence relative to those in schedule 3.

5 = Has a low potential for abuse relative to those in schedule 4. Has a currently accepted medical use in treatment in the United States. Abuse may lead to limited physical dependence or psychological dependence relative to those in schedule 4.

alcohol:

X = Interacts with Alcohol.

rating:

For ratings, users were asked how effective they found the medicine while considering positive/adverse effects and ease of use (1 = not effective, 10 = most effective).

All other columns are self-explanatory.

#### NOTE :

1. this project is only for your guidance, not exactly the same you have to create. Here I am trying to show the way or idea of what steps you can follow and how your projects look. Some projects are very advanced (because it will be made with the help of flask, nlp, advance ai, advance DL and some advanced things ) which you can not understand .

2. You can make or analyze your project with yourself, with your idea, make it more creative from where we can get some information and understand about our business. make sure what overall things you have created all things you understand very well.

## **Example**

what steps you should have to follow

Here's a beginner-friendly guide to performing data analytics on a dataset involving Drugs, Side Effects, and Medical Conditions. The project will involve exploratory data analysis (EDA) using Python with the specified columns.

**Project Title:**

# Exploratory Data Analysis on Drugs, Side Effects, and Medical Conditions

## 1. Objective

The goal is to analyze the relationships between drugs, their side effects, and the medical conditions they treat, as well as to explore the ratings and reviews associated with these drugs.

## 2. Dataset Overview

The dataset contains the following columns:

- **drug\_name**: Name of the drug.
- **medical\_condition**: The condition the drug is used to treat.
- **side\_effects**: Common side effects of the drug.
- **generic\_name**: The generic name of the drug.
- **drug\_classes**: The class of the drug (e.g., antibiotic, antihistamine).
- **brand\_names**: Brand names under which the drug is sold.
- **activity**: The activity of the drug (e.g., active, inactive).
- **rx\_otc**: Indicates if the drug is prescription (Rx) or over-the-counter (OTC).
- **pregnancy\_category**: The drug's pregnancy risk category.
- **csa**: Controlled Substances Act schedule, if applicable.
- **alcohol**: Interactions with alcohol.
- **related\_drugs**: Other drugs related to the primary drug.
- **medical\_condition\_description**: A brief description of the medical condition.
- **rating**: Average user rating of the drug.
- **no\_of\_reviews**: Number of user reviews.
- **drug\_link**: URL link to more information about the drug.
- **medical\_condition\_url**: URL link to more information about the medical condition.

## 3. Tools Required

- **Python**: The primary programming language for data analysis.
- **Pandas**: For data manipulation and analysis.
- **Matplotlib/Seaborn**: For data visualization.
- **Jupyter Notebook**: To write and run Python code.

## 4. Step-by-Step Guide

Step 1: Import Libraries

```
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

### Step 2: Load the Dataset

```
# Load your dataset
df = pd.read_csv('path_to_your_dataset.csv')

# Display the first few rows of the dataset
df.head()
```

### Step 3: Data Cleaning

- Check for missing values:

```
# Check for missing values
df.isnull().sum()
```

- Handle missing values:
  - Drop or fill missing values depending on the context.

```
# Example: Drop rows with missing values
df_cleaned = df.dropna()
```

```
# Or fill missing values with a placeholder
df_filled = df.fillna('Unknown')
```

### Step 4: Basic Data Exploration

- Summary statistics:

```
# Summary statistics
df.describe()
```

- **Distribution of Ratings:**

```
# Distribution of drug ratings
plt.figure(figsize=(10, 6))
sns.histplot(df['rating'], bins=10, kde=True)
plt.title('Distribution of Drug Ratings')
plt.xlabel('Rating')
plt.ylabel('Frequency')
plt.show()
```

#### Step 5: Analyzing Relationships

- **Top Drugs by Condition:**

```
# Count the most common drugs for each medical condition
top_drugs =
df.groupby('medical_condition')['drug_name'].value_counts().nlargest(10)
print(top_drugs)
```

- **Side Effects Analysis:**

```
# Analyzing the most common side effects
side_effects = df['side_effects'].value_counts().head(10)
print(side_effects)
```

- **Drug Ratings by Class:**

```
# Boxplot of ratings by drug class
plt.figure(figsize=(12, 8))
sns.boxplot(x='drug_classes', y='rating', data=df)
plt.xticks(rotation=90)
plt.title('Drug Ratings by Class')
plt.show()
```

## Step 6: Conclusion

- **Summarize findings:**
  - Identify any trends or patterns in the data.
  - Discuss how certain drug classes or conditions are associated with specific side effects or ratings.

## 5. Next Steps

- **Advanced Analysis:** Perform more sophisticated statistical tests or machine learning techniques.
- **Reporting:** Create a report or presentation to share the findings.

## 6. Example Output

- **Distribution of Drug Ratings:**
  - A histogram showing how drug ratings are distributed, with peaks at certain rating values.
- **Top Drugs for a Condition:**
  - A list or bar chart showing the most commonly prescribed drugs for a particular condition.
- **Side Effects Analysis:**
  - A list of the most common side effects reported in the dataset.

By following this guide, a beginner can start exploring and analyzing the dataset effectively. Let me know if you need further assistance!

## Sample code

```
# Import dataset

import pandas as pd

import numpy as np
```

```
# Read the CSV file into a DataFrame
```

```
fpath =
```

```
'/kaggle/input/drugs-side-effects-and-medical-condition/drugs_side_effects_drugs_com  
.csv'
```

```
data = pd.read_csv(fpath)
```

```
# Display the columns quantity and names
```

```
print('The dataset has {} rows and {} columns'.format(data.shape[0], data.shape[1]))
```

```
print("column:")
```

```
print(data.columns)
```

```
The dataset has 2931 rows and 17 columns
```

```
column:
```

```
Index(['drug_name', 'medical_condition', 'side_effects', 'generic_name',  
      'drug_classes', 'brand_names', 'activity', 'rx_otc',  
      'pregnancy_category', 'csa', 'alcohol', 'related_drugs',  
      'medical_condition_description', 'rating', 'no_of_reviews', 'drug_link',  
      'medical_condition_url'],  
      dtype='object')
```



```
# Show the main information about dataset
```

```
data.info()
```

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

```
RangeIndex: 2931 entries, 0 to 2930
```

```
Data columns (total 17 columns):
```

#	Column	Non-Null Count	Dtype
---	-----	-----	-----
0	drug_name	2931 non-null	object
1	medical_condition	2931 non-null	object
2	side_effects	2807 non-null	object
3	generic_name	2888 non-null	object
4	drug_classes	2849 non-null	object
5	brand_names	1718 non-null	object
6	activity	2931 non-null	object
7	rx_otc	2930 non-null	object
8	pregnancy_category	2702 non-null	object
9	csa	2931 non-null	object

```
10  alcohol                1377 non-null  object
11  related_drugs          1462 non-null  object
12  medical_condition_description  2931 non-null  object
13  rating                 1586 non-null  float64
14  no_of_reviews          1586 non-null  float64
15  drug_link              2931 non-null  object
16  medical_condition_url   2931 non-null  object
```

```
dtypes: float64(2), object(15)
```

```
memory usage: 389.4+ KB
```

In [3]:

```
data.head()
```

Out[3]:

	drug_name	medical_condition	side_effects	generic_name	drug_classes	brand_names	activity	rx_category	pregnancy_category	csa	alcohol	related_drugs	medical_condition_description	rating	no_of_reviews	drug_link	medical_condition_url
0	doxycycline	Acne	(hives	doxycycline	Miscellaneous	Acetate	87	R	D	N	X	amoxicillin: <a href="https://www.drugs.com/">https://www.drugs.com/</a>	Acne Other names:	6	76	<a href="https://www.drugs.com/dox">https://www.drugs.com/dox</a>	<a href="https://www.drugs.com/condition/acne.h">https://www.drugs.com/condition/acne.h</a>

	ne	e	, difficult breathing, swelling in your ...	ne	eo us anti mal ari als, Tet rac ycli nes	, Ad ox a CK , Ad ox a Pa k, Ad ox a TT, Al od. ..	%	x				amoxicillin..	Acne Vulgaris; Blackhe ads; B...	8	0.0	ycycline.html	ml
1	spi ron ola ctone	Acne	hive s; difficulty breathing; swelling of your ...	spi ron ola ctone	Ald ost ero ne rec ept or ant ag oni sts, Pot ass ium -sp ...	Al da ctone, Ca ro Sp ir	82%	Rx	C	N	X	amlodipine: <a href="https://www.drugs.com/amlodipine.h...">https://www.drugs.com/amlodipine.h...</a>	Acne Other names: Acne Vulgaris; Blackhe ads; B...	7.2	449.0	<a href="https://www.drugs.com/spironolactone.html">https://www.drugs.com/spironolactone.html</a>	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>
2	mi no cyc line	Acne	skin rash, fever, s	mi no cyc line	Tet rac ycli nes	Dyna cin , Mi no cin , Mi	48%	Rx	D	N	N a N	amoxicillin: <a href="https://www.drugs.com/amoxicillin..">https://www.drugs.com/amoxicillin..</a>	Acne Other names: Acne Vulgaris; Blackhe ads; B...	5.7	482.0	<a href="https://www.drugs.com/minocycline.html">https://www.drugs.com/minocycline.html</a>	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>

			w oll e n gl a n ds , flu -li ke sy m ...			nol ira, So lod yn, Xi mi no, V...											
3	Ac cut an e	Acn e	pr o bl e m s wi th yo ur vi si o n or he ar in g; m us cl e o. ...	isot reti noi n (or al)	Mis cell an eo us anti ne opl asti cs, Mis cell an eo us u...	Na N	4 1 %	R x	X	N	X	doxycycline : https://www .drugs.com/ doxycycline ...	Acne Other names: Acne Vulgaris; Blackhe ads; B...	7 . 9	62 3.0	https://www.d rugs.com/acc utane.html	https://www.d rugs.com/co dition/acne.h ml
4	clin da my cin	Acn e	hi ve s; dif fic ult bre at	clin da my cin top ical	Top ical acn e ag ent s, Va gin	Cl eo cin T, Cli nd aci n ET	3 9 %	R x	B	N	N a N	doxycycline : https://www .drugs.com/ doxycycline ...	Acne Other names: Acne Vulgaris; Blackhe ads; B...	7 . 4	14 6.0	https://www.d rugs.com/mt m/clindamyci n-topical....	https://www.d rugs.com/co dition/acne.h ml

			hi n g; s w ell in g of yo ur ...		al anti -inf ecti ves	Z, Cli nd aci n P, Cli nd ag. ..											
--	--	--	--	--	-----------------------------------	---	--	--	--	--	--	--	--	--	--	--	--

In [4]:

*# Dropping the 'brand\_names' column and delete from dataset*

```
data.drop(columns=['brand_names'], inplace=True)
```

In [5]:

*# Find duplicate rows based on all columns*

```
duplicate_rows= data[data.duplicated()]
```

*#Count the duplicated rows*

```
duplicate_count = duplicate_rows.shape[0]
```

*# Print the count of duplicate rows*

```
print("Count of Duplicate Rows:", duplicate_count)
```

```
print(duplicate_rows) # Print the duplicate rows
```

Count of Duplicate Rows: 0

Empty DataFrame

```
Columns: [drug_name, medical_condition, side_effects, generic_name, drug_classes,
activity, rx_otc, pregnancy_category, csa, alcohol, related_drugs,
medical_condition_description, rating, no_of_reviews, drug_link,
medical_condition_url]
```

```
Index: []
```

In [6]:

```
# Convert 'rating' and 'no_of_reviews' attributes to numeric
```

```
data['rating'] = pd.to_numeric(data['rating'], errors='coerce')
```

```
# data['no_of_reviews'] = pd.to_numeric(data['no_of_reviews'], errors='coerce')
```

```
print(data.dtypes.value_counts())
```

```
object      14
```

```
float64      2
```

```
Name: count, dtype: int64
```

In [7]:

```
# Convert 'activity' to string, remove whitespace and '%' character, then convert to
float and divide by 100
```

```
data['activity'] = data['activity'].astype(str).str.replace(r'\s+', '',
```

```
regex=True).str.rstrip('%').astype('float')/100
```

```
# Display the updated 'activity' column
```

```
print(data['activity'].head())
```

```
0    0.87
```

```
1    0.82
```

```
2    0.48
```

```
3    0.41
```

```
4    0.39
```

```
Name: activity, dtype: float64
```

In [8]:

```
# Print the total number of missing values
```

```
print("There are {} missing values in this  
dataset".format(data.isnull().sum().sum()))
```

```
print('Number of instances = %d' % (data.shape[0]))
```

```
print('Number of attributes = %d' % (data.shape[1]))
```

```
print('Number of missing values:')
```

```
for col in data.columns:
```

```
print('\t%s: %d' % (col,data[col].isna().sum()))
```

There are 6192 missing values in this dataset

Number of instances = 2931

Number of attributes = 16

Number of missing values:

drug\_name: 0

medical\_condition: 0

side\_effects: 124

generic\_name: 43

drug\_classes: 82

activity: 0

rx\_otc: 1

pregnancy\_category: 229

csa: 0

alcohol: 1554

related\_drugs: 1469

medical\_condition\_description: 0

rating: 1345



```
no_of_reviews: 1345
```

```
drug_link: 0
```

```
medical_condition_url: 0
```

In [9]:

```
# In the alcohol column we have X and null(NaN) values, because the drug can interact with alcohol or not.
```

```
# Therefore, let's replace the values of alcohol column with boolean values.
```

```
# Let X will be 1 of interaction, NaN will be 0.
```

```
data['alcohol']=data['alcohol'].replace(np.NaN, '0')
```

```
data['alcohol']=data['alcohol'].replace({'X': 1})
```

In [10]:

```
# To avoid missing values let's fill them with some information
```

```
# In our case we will replace all them
```

```
# Fill the null values in 'side_effects' and 'related_drugs' with no
```

```
data["side_effects"] = data['side_effects'].fillna('Unknown')
```

```
data["related_drugs"] = data['related_drugs'].fillna('Unknown')
```

In [11]:

```
# Fill the null values with 0 as a base for 'rating' and 'no_of_reviews' columns
```

```
# It will show that there are no information about it
```

```
data["rating"] = data['rating'].fillna('0')
```

```
data["no_of_reviews"] = data['no_of_reviews'].fillna('0')
```

In [12]:

```
# Fill the null values with ?
```

```
data['generic_name']=data['generic_name'].replace(np.NaN, 'Unknown')
```

```
# Fill the null values with undefined for 'drug_classes'
```

```
data['drug_classes']=data['drug_classes'].replace(np.NaN, 'Unknown')
```

In [13]:

```
# For these two columns we already have some category values from dataset's description
```

```
# So, let's check the categorical values
```

```
# For Rx_OTC
```

```
data["rx_otc"].unique()
```

Out[13]:

```
array(['Rx', 'Rx/OTC', 'OTC', nan], dtype=object)
```

In [14]:

```
# For pregnancy categories
```

```
data["pregnancy_category"].unique()
```

Out[14]:

```
array(['D', 'C', 'X', 'B', 'N', nan, 'A'], dtype=object)
```

In [15]:

```
# Fill the null value with Unknown as a basic value
```

```
data['rx_otc']=data['rx_otc'].replace(np.NaN, 'Unknown')
```

```
# Fill the null value with Unknown as a basic value
```

```
data['pregnancy_category']=data['pregnancy_category'].replace(np.NaN, 'Unknown')
```

```
data['no_of_reviews'] = pd.to_numeric(data['no_of_reviews'], errors='coerce')
```

```
print(data.head())
```

```
dfs=data.copy()
```

drug\_name medical\_condition \

0	doxycycline	Acne
1	spironolactone	Acne
2	minocycline	Acne
3	Accutane	Acne
4	clindamycin	Acne

side\_effects generic\_name \

0	(hives, difficult breathing, swelling in your ...	doxycycline
1	hives ; difficulty breathing; swelling of your...	spironolactone
2	skin rash, fever, swollen glands, flu-like sym...	minocycline
3	problems with your vision or hearing; muscle o...	isotretinoin (oral)
4	hives ; difficult breathing; swelling of your ...	clindamycin topical

drug\_classes activity rx\_otc \

0	Miscellaneous antimalarials, Tetracyclines	0.87	Rx
1	Aldosterone receptor antagonists, Potassium-sp...	0.82	Rx
2	Tetracyclines	0.48	Rx

3	Miscellaneous antineoplastics, Miscellaneous u...	0.41	Rx
4	Topical acne agents, Vaginal anti-infectives	0.39	Rx

pregnancy\_category csa alcohol \

0	D	N	1
1	C	N	1
2	D	N	0
3	X	N	1
4	B	N	0

related\_drugs \

0	amoxicillin: <a href="https://www.drugs.com/amoxicillin...">https://www.drugs.com/amoxicillin...</a>
1	amlodipine: <a href="https://www.drugs.com/amlodipine.h...">https://www.drugs.com/amlodipine.h...</a>
2	amoxicillin: <a href="https://www.drugs.com/amoxicillin...">https://www.drugs.com/amoxicillin...</a>
3	doxycycline: <a href="https://www.drugs.com/doxycycline...">https://www.drugs.com/doxycycline...</a>
4	doxycycline: <a href="https://www.drugs.com/doxycycline...">https://www.drugs.com/doxycycline...</a>

medical\_condition\_description rating no\_of\_reviews \

0	Acne Other names: Acne Vulgaris; Blackheads; B...	6.8	760.0
---	---	-----	-------

1	Acne Other names: Acne Vulgaris; Blackheads; B...	7.2	449.0
2	Acne Other names: Acne Vulgaris; Blackheads; B...	5.7	482.0
3	Acne Other names: Acne Vulgaris; Blackheads; B...	7.9	623.0
4	Acne Other names: Acne Vulgaris; Blackheads; B...	7.4	146.0

drug\_link \

0	<a href="https://www.drugs.com/doxycycline.html">https://www.drugs.com/doxycycline.html</a>
1	<a href="https://www.drugs.com/spironolactone.html">https://www.drugs.com/spironolactone.html</a>
2	<a href="https://www.drugs.com/minocycline.html">https://www.drugs.com/minocycline.html</a>
3	<a href="https://www.drugs.com/accutane.html">https://www.drugs.com/accutane.html</a>
4	<a href="https://www.drugs.com/mtm/clindamycin-topical....">https://www.drugs.com/mtm/clindamycin-topical....</a>

medical\_condition\_url

0	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>
1	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>
2	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>
3	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>
4	<a href="https://www.drugs.com/condition/acne.html">https://www.drugs.com/condition/acne.html</a>

```
# Let's check is there any missing values left

print("There are {} missing values in this
dataset".format(data.isnull().sum().sum()))

print('Number of instances = %d' % (data.shape[0]))

print('Number of attributes = %d' % (data.shape[1]))

print('Number of missing values:')

for col in data.columns:

    print('\t%s: %d' % (col, data[col].isna().sum()))
```

There are 0 missing values in this dataset

Number of instances = 2931

Number of attributes = 16

Number of missing values:

drug\_name: 0

medical\_condition: 0

side\_effects: 0

generic\_name: 0

drug\_classes: 0

activity: 0

```
rx_otc: 0

pregnancy_category: 0

csa: 0

alcohol: 0

related_drugs: 0

medical_condition_description: 0

rating: 0

no_of_reviews: 0

drug_link: 0

medical_condition_url: 0
```

In [17]:

```
data_version2=data.copy()

print(data_version2.head())

# Print head of dataset to our check
```

```
drug_name medical_condition \
0    doxycycline           Acne
1  spironolactone           Acne
```



2	minocycline	Acne
3	Accutane	Acne
4	clindamycin	Acne

	side_effects	generic_name \
0	(hives, difficult breathing, swelling in your ...	doxycycline
1	hives ; difficulty breathing; swelling of your...	spironolactone
2	skin rash, fever, swollen glands, flu-like sym...	minocycline
3	problems with your vision or hearing; muscle o...	isotretinoin (oral)
4	hives ; difficult breathing; swelling of your ...	clindamycin topical

	drug_classes	activity	rx_otc \
0	Miscellaneous antimalarials, Tetracyclines	0.87	Rx
1	Aldosterone receptor antagonists, Potassium-sp...	0.82	Rx
2	Tetracyclines	0.48	Rx
3	Miscellaneous antineoplastics, Miscellaneous u...	0.41	Rx
4	Topical acne agents, Vaginal anti-infectives	0.39	Rx

pregnancy\_category csa alcohol \

0	D	N	1
1	C	N	1
2	D	N	0
3	X	N	1
4	B	N	0

related\_drugs \

0	amoxicillin: <a href="https://www.drugs.com/amoxicillin...">https://www.drugs.com/amoxicillin...</a>
1	amlodipine: <a href="https://www.drugs.com/amlodipine.h...">https://www.drugs.com/amlodipine.h...</a>
2	amoxicillin: <a href="https://www.drugs.com/amoxicillin...">https://www.drugs.com/amoxicillin...</a>
3	doxycycline: <a href="https://www.drugs.com/doxycycline...">https://www.drugs.com/doxycycline...</a>
4	doxycycline: <a href="https://www.drugs.com/doxycycline...">https://www.drugs.com/doxycycline...</a>

medical\_condition\_description rating no\_of\_reviews \

0	Acne Other names: Acne Vulgaris; Blackheads; B...	6.8	760.0
1	Acne Other names: Acne Vulgaris; Blackheads; B...	7.2	449.0
2	Acne Other names: Acne Vulgaris; Blackheads; B...	5.7	482.0
3	Acne Other names: Acne Vulgaris; Blackheads; B...	7.9	623.0
4	Acne Other names: Acne Vulgaris; Blackheads; B...	7.4	146.0

```
drug_link \
```

```
0      https://www.drugs.com/doxycycline.html
1      https://www.drugs.com/spironolactone.html
2      https://www.drugs.com/minocycline.html
3      https://www.drugs.com/accutane.html
4  https://www.drugs.com/mtm/clindamycin-topical....
```

```
medical_condition_url
```

```
0  https://www.drugs.com/condition/acne.html
1  https://www.drugs.com/condition/acne.html
2  https://www.drugs.com/condition/acne.html
3  https://www.drugs.com/condition/acne.html
4  https://www.drugs.com/condition/acne.html
```

In [18]:

```
# Save the data
```

```
data_version2.to_csv('drugs_side_effects_drugs_com_version2.csv', index=False)
```

In [19]:

```
# Read the new version dataset
```

```
data_ver3=pd.read_csv('drugs_side_effects_drugs_com_version2.csv')
```

```
data_ver3["pregnancy_category"].unique()
```

Out[19]:

```
array(['D', 'C', 'X', 'B', 'N', 'Unknown', 'A'], dtype=object)
```

In [20]:

```
data_ver3["csa"].unique()
```

Out[20]:

```
array(['N', '2', '4', 'U', 'M', '5', '3'], dtype=object)
```

In [21]:

```
data_ver3["rx_otc"].unique()
```

Out[21]:

```
array(['Rx', 'Rx/OTC', 'OTC', 'Unknown'], dtype=object)
```

In [22]:

```
data_ver3["generic_name"].unique()
```

Out[22]:

```
array(['doxycycline', 'spironolactone', 'minocycline', ...,  
      'fenfluramine', 'phendimetrazine tartrate', 'setmelanotide'],  
      dtype=object)
```

In [23]:

```
data_ver3["medical_condition"].unique()
```

Out[23]:

```
array(['Acne', 'ADHD', 'AIDS/HIV', 'Allergies', "Alzheimer's", 'Angina',  
      'Anxiety', 'Asthma', 'Bipolar Disorder', 'Bronchitis', 'Cancer',  
      'Cholesterol', 'Colds & Flu', 'Constipation', 'COPD', 'Covid 19',  
      'Depression', 'Diabetes (Type 1)', 'Diabetes (Type 2)', 'Diarrhea',  
      'Eczema', 'Erectile Dysfunction', 'Gastrointestinal',  
      'GERD (Heartburn)', 'Gout', 'Hair Loss', 'Hayfever', 'Herpes',  
      'Hypertension', 'Hypothyroidism', 'IBD (Bowel)', 'Incontinence',  
      'Insomnia', 'Menopause', 'Migraine', 'Osteoarthritis',  
      'Osteoporosis', 'Pain', 'Pneumonia', 'Psoriasis',  
      'Rheumatoid Arthritis', 'Schizophrenia', 'Seizures', 'Stroke',  
      'Swine Flu', 'UTI', 'Weight Loss'], dtype=object)
```

In [24]:

```
from sklearn.preprocessing import LabelEncoder

label_encoder = LabelEncoder()

data_ver3["csa"] = label_encoder.fit_transform(data_ver3["csa"])

data_ver3["rx_otc"] = label_encoder.fit_transform(data_ver3["rx_otc"])

data_ver3["generic_name"] = label_encoder.fit_transform(data_ver3["generic_name"])

data_ver3["medical_condition"] =
label_encoder.fit_transform(data_ver3["medical_condition"])

data_ver3["pregnancy_category"] =
label_encoder.fit_transform(data_ver3["pregnancy_category"])

data_ver3["side_effects"] = label_encoder.fit_transform(data_ver3["side_effects"])
```

In [25]:

```
data_ver3["generic_name"].unique()
```

Out[25]:

```
array([ 642, 1270, 1034, ..., 729, 1157, 1259])
```

In [26]:

```
data_ver3["rx_otc"].unique()
```

Out[26]:

```
array([1, 2, 0, 3])
```

In [27]:

```
data_ver3["csa"].unique()
```

Out[27]:

```
array([5, 0, 2, 6, 4, 3, 1])
```

In [28]:

```
data_ver3["side_effects"].unique()
```

Out[28]:

```
array([ 15, 1972, 2697, ..., 1647, 416, 1706])
```

In [29]:

```
data_ver3["medical_condition"].unique()
```

Out[29]:

```
array([ 2, 0, 1, 3, 4, 5, 6, 7, 8, 9, 11, 12, 13, 14, 10, 15, 16,  
       17, 18, 19, 20, 21, 23, 22, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33,  
       34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46])
```

In [30]:

```
data_ver3["pregnancy_category"].unique()
```

Out[30]:

```
array([3, 2, 6, 1, 4, 5, 0])
```

In [31]:

```
df=pd.DataFrame(data_ver3,columns=('generic_name', 'medical_condition',  
'no_of_reviews', 'side_effects', 'rating', 'csa', 'pregnancy_category', 'rx_otc',  
'alcohol'))
```

```
df.head(10)
```

Out[31]:

	generic_name	medical_condition	no_of_reviews	side_effects	rating	csa	pregnancy_category	rx_otc	alcohol
0	642	2	760.0	15	6.8	5	3	1	1
1	1270	2	449.0	1972	7.2	5	2	1	1
2	1034	2	482.0	2697	5.7	5	3	1	0
3	903	2	623.0	2570	7.9	5	6	1	1



4	505	2	146.0	1260	7.4	5	1	1	0
5	1270	2	8.0	1971	7.6	5	2	1	1
6	1335	2	439.0	1895	7.7	5	2	1	0
7	903	2	999.0	2577	8.0	5	6	1	1
8	1276	2	96.0	2702	8.5	5	3	1	1
9	162	2	86.0	2405	7.9	5	2	1	0

In [32]:

```
from sklearn.preprocessing import StandardScaler
```

```
scaler=StandardScaler()
```

```
scaler.fit(df)
```

```
scaled_data=scaler.transform(df)
```

```
print(scaled_data)
```

```
[[ -0.11111578 -1.43400434  5.10119829 ...  0.28892455 -0.17025661
```

```

1.06232778]

[ 1.50040103 -1.43400434  2.89586941 ... -0.43301735 -0.17025661

1.06232778]

[ 0.89479917 -1.43400434  3.12987537 ...  0.28892455 -0.17025661

-0.94132905]

...

[ 1.21043065  1.82918864 -0.28802985 ... -0.43301735 -0.17025661

1.06232778]

[ 1.47217383  1.82918864 -0.28802985 ...  1.73280834 -0.17025661

-0.94132905]

[ 1.47217383  1.82918864 -0.28802985 ...  1.73280834 -0.17025661

-0.94132905]]

```

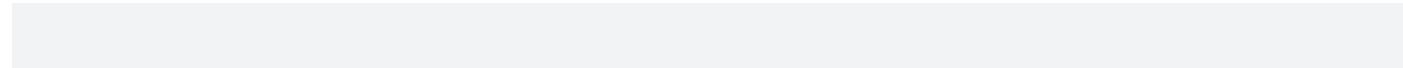
In [33]:

```

df_std = pd.DataFrame(scaler.fit_transform(df), columns=df.columns)

print(df_std)

```



	generic_name	medical_condition	no_of_reviews	side_effects	rating \
0	-0.111116	-1.434004	5.101198	-1.678954	0.819930

1	1.500401	-1.434004	2.895869	0.778579	0.925271
2	0.894799	-1.434004	3.129875	1.689009	0.530244
3	0.558639	-1.434004	4.129719	1.529527	1.109617
4	-0.462673	-1.434004	0.747269	-0.115526	0.977941
...	...	...	...	...	...
2926	-0.832193	1.829189	-0.167481	0.757231	1.004277
2927	0.112136	1.829189	-0.288030	0.370455	-0.970861
2928	1.210431	1.829189	-0.288030	-1.029724	-0.970861
2929	1.472174	1.829189	-0.288030	-1.175392	-0.970861
2930	1.472174	1.829189	-0.288030	0.444545	-0.970861
	csa	pregnancy_category	rx_otc	alcohol	
0	0.274178	0.288925	-0.170257	1.062328	
1	0.274178	-0.433017	-0.170257	1.062328	
2	0.274178	0.288925	-0.170257	-0.941329	
3	0.274178	2.454750	-0.170257	1.062328	
4	0.274178	-1.154959	-0.170257	-0.941329	
...	...	...	...	...	
2926	-3.424857	2.454750	-0.170257	1.062328	

```
2927 -2.500098          -0.433017 -0.170257  1.062328
2928 -3.424857          -0.433017 -0.170257  1.062328
2929  0.274178          1.732808 -0.170257 -0.941329
2930  0.274178          1.732808 -0.170257 -0.941329
```

```
[2931 rows x 9 columns]
```

In [34]:

```
import seaborn as sns

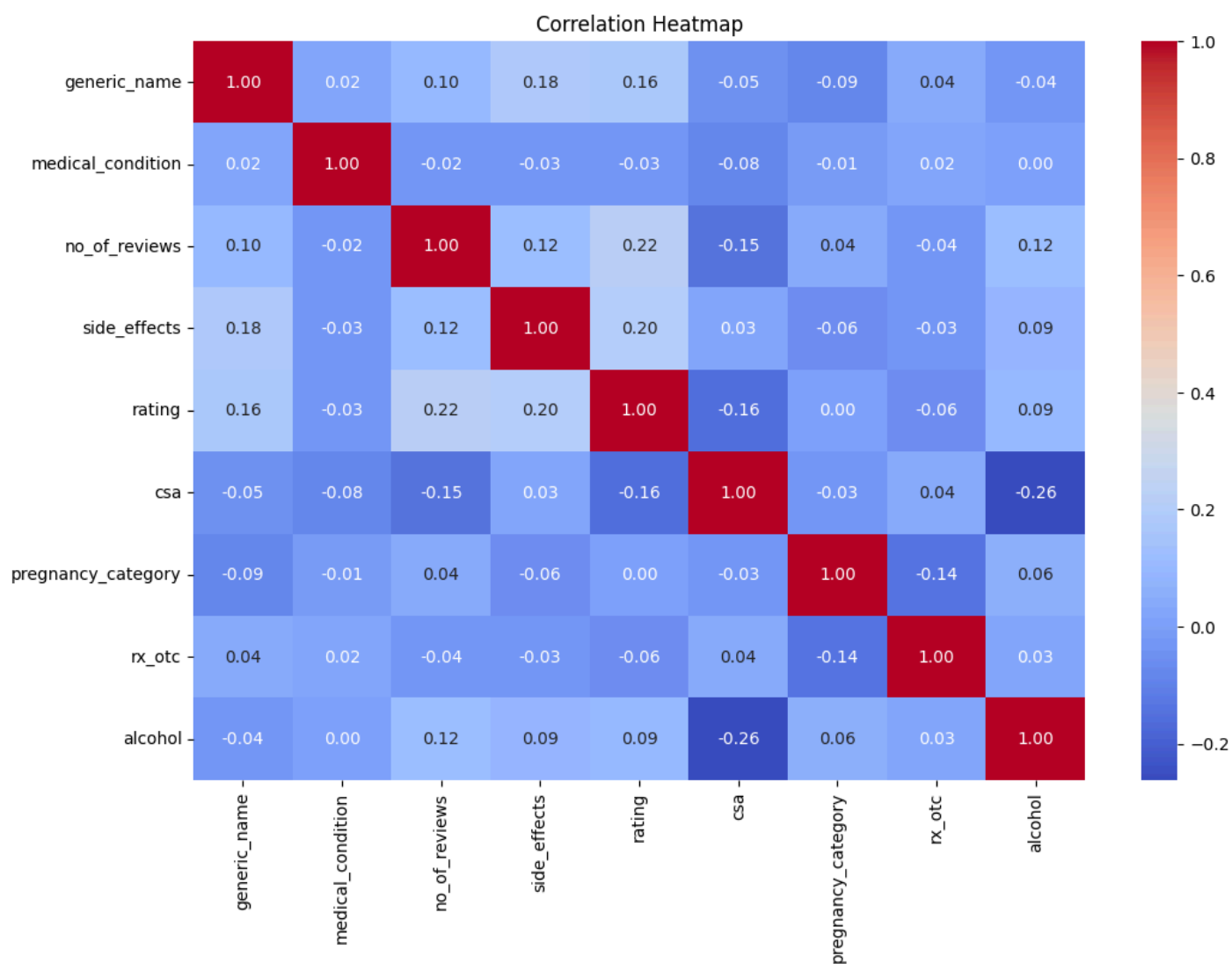
import matplotlib.pyplot as plt

plt.figure(figsize=(12, 8))

sns.heatmap(df.corr(), annot=True, cmap='coolwarm', fmt=".2f")

plt.title('Correlation Heatmap')

plt.show()
```



In [35]:

```
# Read the new version dataset
```

```
data_ver4 = pd.read_csv('drugs_side_effects_drugs_com_version2.csv')
```

```
# Importing necessary libraries
```

```
from mlxtend.frequent_patterns import apriori, association_rules
```

```
import matplotlib.pyplot as plt
```

```
import pandas as pd
```

```
# Check for occurrence and frequency of medical conditions, sorted from highest to lowest
```

```
medical_condition_counts =
```

```
data_ver4['medical_condition'].value_counts().sort_values(ascending=False)
```

```
print("\nMedical condition occurrence and frequency (sorted from highest to lowest):")
```

```
print(medical_condition_counts)
```

```
Medical condition occurrence and frequency (sorted from highest to lowest):
```

```
medical_condition
```

```
Pain                264
```

```
Colds & Flu         245
```

```
Acne                238
```

```
Hypertension        177
```

```
Osteoarthritis      129
```

```
Hayfever            124
```

```
Eczema              122
```

```
AIDS/HIV            109
```

Diabetes (Type 2)	104
Psoriasis	93
GERD (Heartburn)	77
Pneumonia	72
Angina	71
Bronchitis	71
Migraine	61
Insomnia	60
Constipation	60
Diabetes (Type 1)	57
Osteoporosis	56
ADHD	55
Depression	51
Seizures	50
Bipolar Disorder	47
UTI	46
Asthma	45
Anxiety	45
Cholesterol	45

Diarrhea	38
Covid 19	34
Rheumatoid Arthritis	33
Alzheimer's	27
Weight Loss	23
COPD	23
IBD (Bowel)	22
Schizophrenia	20
Cancer	20
Incontinence	19
Hypothyroidism	17
Allergies	14
Erectile Dysfunction	13
Hair Loss	11
Herpes	10
Gout	9
Menopause	7
Gastrointestinal	7
Stroke	5



Swine Flu

5

Name: count, dtype: int64

In [36]:

```
# Save the results to CSV files if needed
```

```
medical_condition_counts.to_csv('medical_condition_counts.csv')
```

In [37]:

```
# Importing necessary libraries for processing text
```

```
from collections import Counter
```

```
import re
```

```
# Function to extract side effects from text, split by semicolons
```

```
def extract_side_effects(text):
```

```
    # Split the text on semicolons then strip whitespace
```

```
    return [effect.strip() for effect in re.split(r'[;]', text)]
```

```
# Extract and count occurrences of side effects
```

```
side_effects =
```

```
data_ver4['side_effects'].dropna().apply(extract_side_effects).explode()
```

```
side_effect_counts = side_effects.value_counts().sort_values(ascending=False)

print("\nSide effects occurrence and frequency (sorted from highest to lowest):")

print(side_effect_counts)
```

Side effects occurrence and frequency (sorted from highest to lowest):

side\_effects

hives

1788

difficult breathing

1130

difficulty breathing

450

itching

275

a light-headed feeling, like you might pass out

272

...

swelling of your face, lips, tongue, or throat. Rizatriptan may cause serious side effects. Stop using rizatriptan and call your doctor at once if you have: sudden and severe stomach pain and bloody diarrhea

1

swelling of your face, lips, tongue, or throat. Report any new or worsening symptoms to your doctor, such as: mood or behavior changes, anxiety , panic attacks , trouble sleeping, or if you feel impulsive, irritable, agitated, hostile, aggressive, restless, hyperactive (mentally or physically), depressed, or have thoughts about suicide or hurting yourself. Zarontin may cause serious side effects. Call your doctor at once if you have: fever, chills, flu symptoms, sore throat , feeling very weak

1

or signs of a stroke--sudden numbness or weakness (especially on one side of the body), sudden severe headache, slurred speech, problems with vision or balance. Common side effects of rizatriptan may include: dizziness , drowsiness, feeling tired

1

Suddenly stopping or reducing the dose of Diastat AcuDial very quickly may precipitate acute withdrawal reactions, which can be life-threatening. In some cases, patients have developed withdrawal symptoms lasting weeks to more than 12 months, including but not limited to: anxiety difficulty thinking mental changes depression insomnia abnormal skin sensations muscle weakness tremors twitching ringing in your ears burning or prickling feeling in your hands, arms, or feet The most frequent side effect reported for Diastat AcuDial in clinical studies was somnolence (sleepiness or drowsiness). Other side effects included dizziness, headache, pain, abdominal pain, nervousness, vasodilation (increase in diameter of blood vessel), diarrhea, ataxia/incoordination (lack of coordination), euphoria (feeling of great happiness or well-being), asthma, rhinitis (irritation of the nose similar to an allergy or a cold), and rash. You are encouraged to report negative side effects of prescription drugs to the FDA. Visit [www.fda.gov/medwatch](http://www.fda.gov/medwatch), or call 1-800-FDA-1088. You may also contact Bausch Health Customer Service at

1-800-321-4576. Diastat AcuDial side effects 1

or nausea , vomiting , diarrhea , or stomach pain.

1

Name: count, Length: 8438, dtype: int64

In [38]:

*# Save the side effect counts to a CSV file*

```
side_effect_counts.to_csv('side_effect_counts.csv')
```

In [39]:

*# Function to extract drug classes from text, split by commas*

```
def extract_drug_classes(text):
```

```
    # Split the text on commas then strip whitespace
```

```
    return [effect.strip() for effect in re.split(r'[,]', text)]
```

*# Extract and count occurrences of drug classes*

```
drug_classes =
```

```
data_ver4['drug_classes'].dropna().apply(extract_drug_classes).explode()
```

```
drug_classes_counts = drug_classes.value_counts().sort_values(ascending=False)
```

```
print("\nDrug Classes occurrence and frequency (sorted from highest to lowest):")

print(drug_classes_counts)
```

Drug Classes occurrence and frequency (sorted from highest to lowest):

drug\_classes

Upper respiratory combinations	245
Topical acne agents	125
Topical steroids	94
Antihistamines	82
Unknown	82
...	
Immune globulins	1
Smoking cessation agents	1
Mouth and throat products	1
Skeletal muscle relaxant combinations	1
Anthelmintics	1

Name: count, Length: 244, dtype: int64

In [40]:

```
# Save the drug classes counts to a CSV file
```

```
drug_classes_counts.to_csv('drug_classes_counts.csv')
```

In [41]:

```
# Define functions to check for specific side effects and create new boolean columns
```

```
def has_hives(text):
```

```
    return 'hives' in text.lower()
```

```
data_ver4['Hives'] = data_ver4['side_effects'].apply(has_hives)
```

```
def has_difficult_breathing(text):
```

```
    return 'difficult breathing' in text.lower() or 'difficulty breathing' in  
text.lower()
```

```
data_ver4['Difficult Breathing'] =
```

```
data_ver4['side_effects'].apply(has_difficult_breathing)
```

```
def has_itching(text):
```

```
    return 'itching' in text.lower()
```

```
data_ver4['Itching'] = data_ver4['side_effects'].apply(has_itching)
```

In [42]:

```
# Define functions to check for specific drug classes and create new boolean columns
```

```
def is_usc(text):
```

```
    return 'Upper respiratory combinations' in text
```

```
data_ver4['Upper respiratory combinations'] =
```

```
data_ver4['drug_classes'].apply(is_usc)
```

```
def is_steroids(text):
```

```
    return 'Topical steroids' in text
```

```
data_ver4['Topical steroids'] = data_ver4['drug_classes'].apply(is_steroids)
```

```
def is_acne(text):
```

```
    return 'Topical acne agents' in text
```

```
data_ver4['Topical acne agents'] = data_ver4['drug_classes'].apply(is_acne)
```

In [43]:

```
# Define functions to check for specific medical conditions and create new boolean columns
```

```
def has_pain(text):
```

```
    return 'Pain' in text
```

```
data_ver4['Pain'] = data_ver4['medical_condition'].apply(has_pain)
```

```
def has_colds_and_flu(text):  
  
    return 'Colds & Flu' in text  
  
data_ver4['Colds & Flu'] = data_ver4['medical_condition'].apply(has_colds_and_flu)  
  
def has_acne(text):  
  
    return 'Acne' in text  
  
data_ver4['Acne'] = data_ver4['medical_condition'].apply(has_acne)
```

In [44]:

```
# Plot the count of occurrences for each side effect
```

```
import seaborn as sns
```

```
# Plot count of Hives
```

```
data_ver4['Hives'].value_counts().plot(kind='bar')
```

```
plt.title('Count of Hives')
```

```
plt.xlabel('Hives')
```

```
plt.ylabel('Count')
```

```
plt.xticks([0, 1], ['False', 'True'], rotation=0)
```



```
plt.show()

# Plot count of Difficult Breathing

data_ver4['Difficult Breathing'].value_counts().plot(kind='bar')

plt.title('Count of Difficult Breathing')

plt.xlabel('Difficult Breathing')

plt.ylabel('Count')

plt.xticks([0, 1], ['False', 'True'], rotation=0)

plt.show()

# Plot count of Itching

data_ver4['Itching'].value_counts().plot(kind='bar')

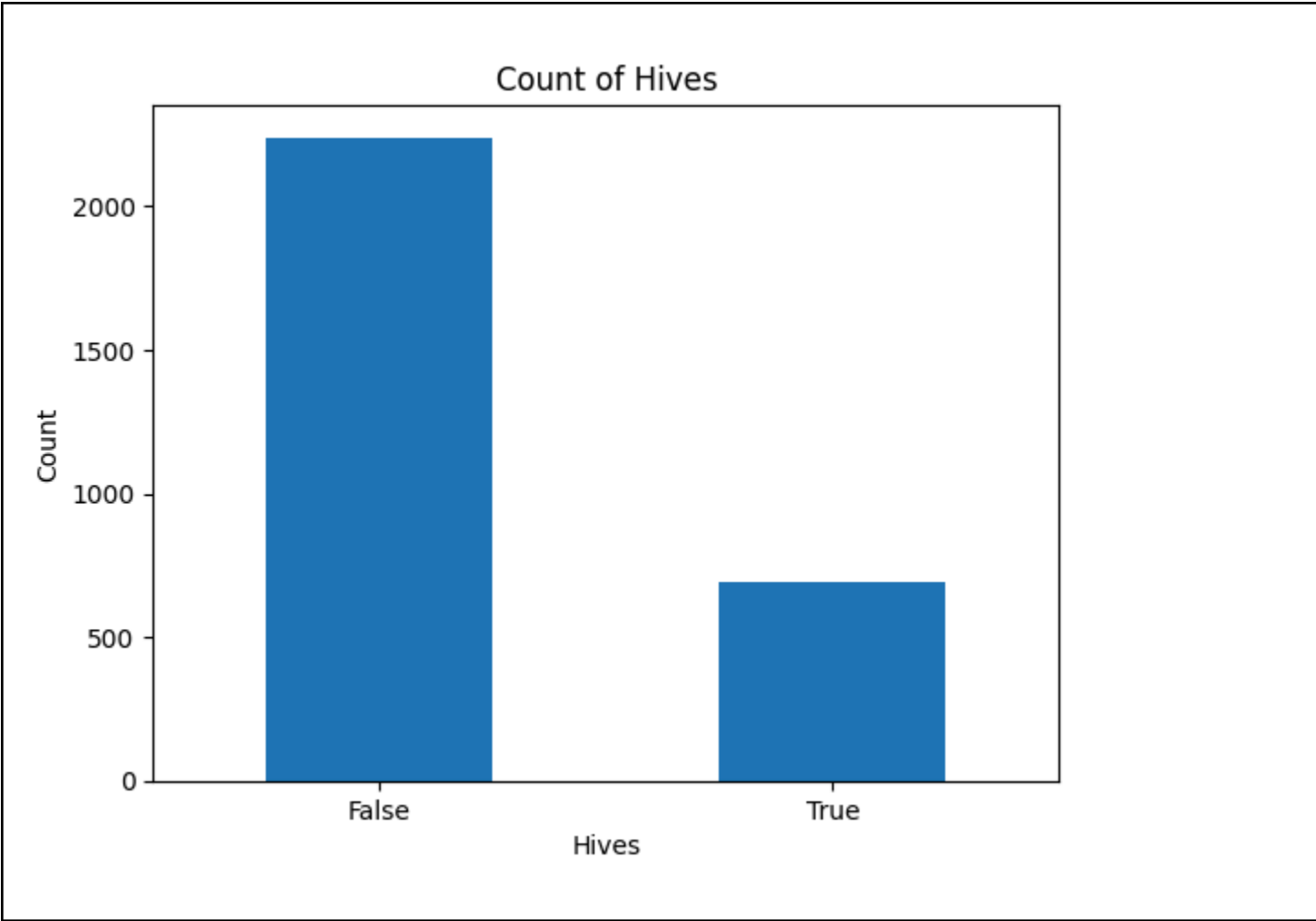
plt.title('Count of Itching')

plt.xlabel('Itching')

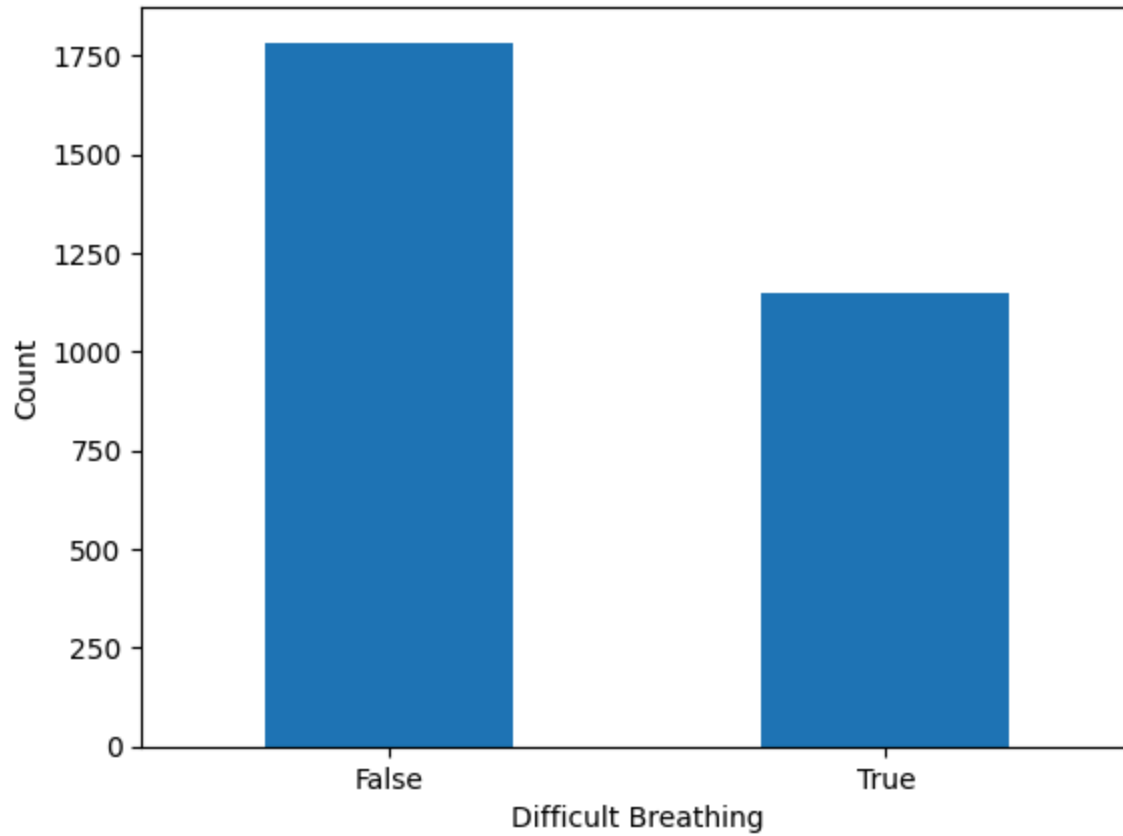
plt.ylabel('Count')

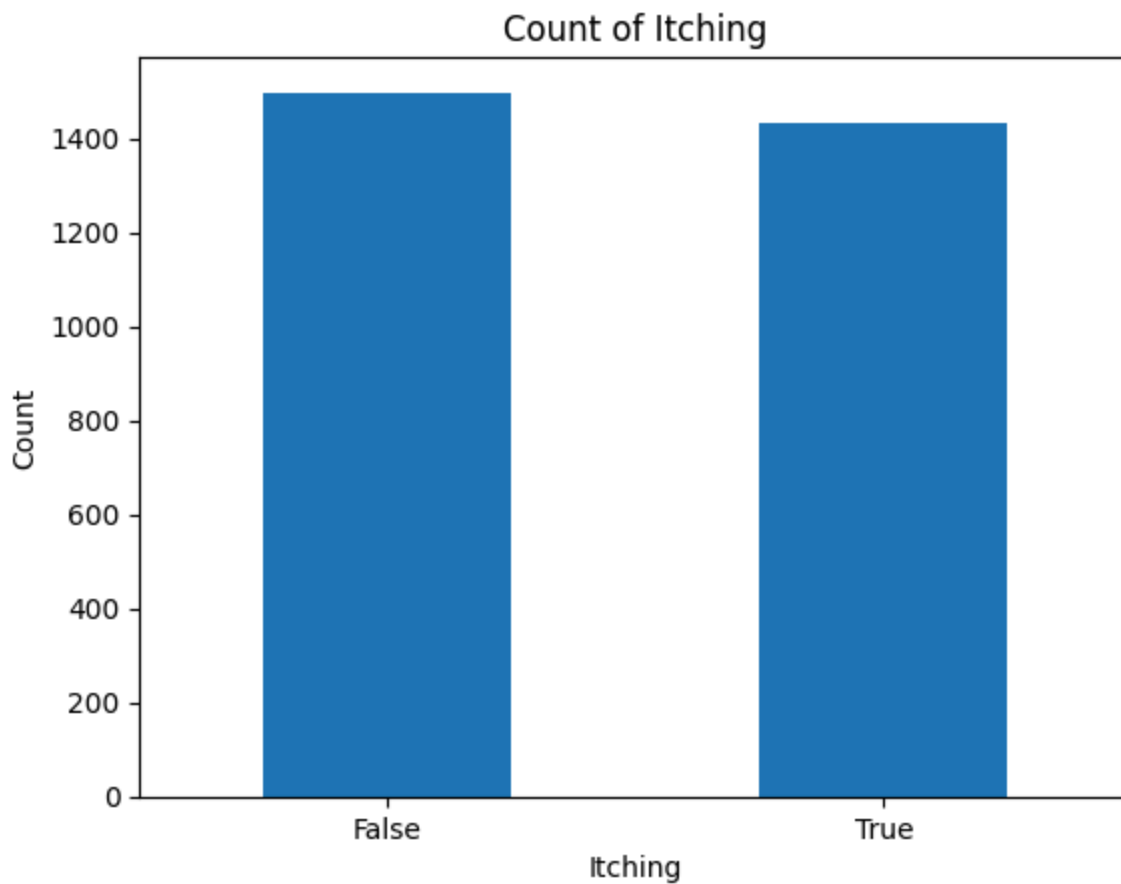
plt.xticks([0, 1], ['False', 'True'], rotation=0)

plt.show()
```



Count of Difficult Breathing





In [45]:

```
# Plot the count of occurrences for each drug class
```

```
# Plot count of Upper respiratory combinations
```

```
data_ver4['Upper respiratory combinations'].value_counts().plot(kind='bar')
```

```
plt.title('Count of Upper respiratory combinations')
```

```
plt.xlabel('Upper respiratory combinations')
```

```
plt.ylabel('Count')
```

```
plt.xticks([0, 1], ['False', 'True'], rotation=0)
```

```
plt.show()

# Plot count of Topical steroids

data_ver4['Topical steroids'].value_counts().plot(kind='bar')

plt.title('Count of Topical steroids')

plt.xlabel('Topical steroids')

plt.ylabel('Count')

plt.xticks([0, 1], ['False', 'True'], rotation=0)

plt.show()
```

1 [Reference link](#)

2 [Reference link](#) for ML project