

vaers

September 11, 2023

1 VAERS

1.1 <https://github.com/jgarza9788/vaers>

2 Disclaimer

THIS IS NOT MEDICAL ADVICE

if you want medical advice please go see your doctor.

please also read the disclaimer on the VAERS data on their website.

[VAERS website](#)

3 get started

3.1 1. get the data

download the data from VAERS website

[link to VAERS website](#)

make sure to download the 2020-2022 data 2022VAERSVAX.csv

2022VAERSSYMPTOMS.csv

2022VAERSDATA.csv

2021VAERSVAX.csv

2021VAERSSYMPTOMS.csv

2021VAERSDATA.csv

2020VAERSVAX.csv

2020VAERSSYMPTOMS.csv

2020VAERSDATA.csv

3.2 2. move and unzip

move the AllVAERSDataCSVs.zip to VAERS folder and unzip it

3.3 3. check the files

you should have at least 6 files

.../VAERS/AllVAERSDataCSVs/

2022VAERSVAX.csv

2022VAERSSYMPTOMS.csv

2022VAERSDATA.csv

```
...
2020VAERSVAX.csv
2020VAERSSYMPTOMS.csv
2020VAERSDATA.csv
```

4 THE CODE!

4.1 setup/imports

```
[1]: # Import needed libraries
import pandas as pd
import numpy as np
import os, sys, re, json
from collections import Counter

# used for graphs/charts
# import plotly.express as px # too large
import matplotlib.pyplot as plt
import seaborn as sns

from IPython.display import display, HTML # needed for IPYNB
```

```
[2]: # set pandas options
pd.options.display.float_format = '{:,.2f}'.format
pd.set_option('display.max_columns', None)
pd.set_option('display.max_rows', None)
pd.set_option('display.max_colwidth', None)
```

4.2 directory variables

DIR (String) : the current directory this file is in

DATAPATH (String) : the location where the data files are stored

```
[3]: # DIR = os.path.dirname(os.path.realpath(__file__)) #doesn't work due to IPYNB
      ↪file
DIR = '.' #this works in IPYNB
DATAPATH = os.path.join(DIR, 'AllVAERSDataCSVs')

print('DATAPATH: ', DATAPATH)
```

DATAPATH: .\AllVAERSDataCSVs

4.3 Saving and Loading JSON files

```
[4]: # LOADING AND SAVING JSON FILES

def load_json(path_to_file):
```

```

    """
    loads json from file
    """
    with open(path_to_file, 'r') as f:
        return json.load(f)

def save_json(path_to_file,data):
    """
    writes dict/json to file
    """
    with open(path_to_file, 'w') as f:
        json.dump(data, f,indent=4)

## test
# save_json(os.path.join(DATAPATH, 'test.json'),{"HELLO":0})
# print(load_json(os.path.join(DATAPATH, 'test.json'))

```

4.4 processing files

the functions flow will process the csv files into one all_data.json file

```

[5]: def has_covid(text):
    """
    returns 1 or 0 if text has \'COVID\' in it
    """
    if re.search('COVID',text.upper()):
        return 1
    else:
        return 0

def compile_files(directory,files):
    """
    compiles/adds/unions multiple csv files together and returns a dataframe
    """
    df = pd.DataFrame()
    for i,f in enumerate(files):
        df0 = pd.read_csv(os.path.
↪join(directory,f),encoding='cp1252',low_memory=False)
        df = pd.concat([df,df0])
    return df

def process_to_one_file():
    """
    processes VAERS data from 2020 - 2022, creates all_data.json, and returns a_
↪dataframe
    """

```

```

print('process to one file\n\tthis might take a while...go get a
↳drink \n')
df_data = compile_files(DATAPATH,['2020VAERSDATA.csv','2021VAERSDATA.
↳csv','2022VAERSDATA.csv'])
df_vax = compile_files(DATAPATH,['2020VAERSVAX.csv','2021VAERSVAX.
↳csv','2022VAERSVAX.csv'])
df_sym = compile_files(DATAPATH,['2020VAERSSYMPTOMS.csv','2021VAERSSYMPTOMS.
↳csv','2022VAERSSYMPTOMS.csv'])

# print("""
# symptoms are contained in columns (up to 5 symptoms per event)
# we must transform these symptoms into a single list for each event
# """)
print('dedup-ing Symptoms')
vid = list(df_sym['VAERS_ID'].unique())

symptom_columns = [
    'SYMPTOM1',
    'SYMPTOM2',
    'SYMPTOM3',
    'SYMPTOM4',
    'SYMPTOM5'
]

idf_sym = []
for index,v in enumerate(vid):
    if index%5000 == 0:
        print('{:.2f}'.format(index/len(vid)), end='\r')
    temp = df_sym[df_sym['VAERS_ID'] == v]
    temp = temp.to_dict(orient='records')
    syms = []
    for t in temp:
        for sc in symptom_columns:
            if isinstance(t[sc],str):
                syms.append(t[sc])
    idf_sym.append({'VAERS_ID':v,'SYMPTOMS':syms})
df_sym = pd.DataFrame(idf_sym)

print('merge data')
df = pd.merge(df_data,df_vax,how='outer',on='VAERS_ID')
df = df.drop_duplicates(ignore_index = True)
df = pd.merge(df,df_sym,how='outer',on='VAERS_ID')

df.reset_index()

# creating a new column depending if this is the covid vaccine or not
df['COVID_VAX'] = df['VAX_TYPE'].apply(has_covid)

```

```

df = df[df['COVID_VAX'] == 1]
# print(len(df))

#save json file
json_file = os.path.join(DATAPATH, 'all_data.json')
save_json(json_file, df.to_dict(orient='records'))
print('saved: ', json_file)

#save out csv file (not needed), but people might like a csv
csv_file = os.path.join(DATAPATH, 'all_data.csv')
df.to_csv(csv_file)
print('saved: ', csv_file)

return df

```

4.5 get data

the below will get data from the files or from all_data.json.

please note i am only using 25% of the data below, but you can see the .pdf for how this will look with all the data, or edit and run the notebook on your own.

df (DataFrame) : contains all the data from VAERS files listed below *

- .../VAERS/AllVAERSDataCSVs/
- * 2022VAERSVAX.csv
- * 2022VAERSSYMPTOMS.csv
- * 2022VAERSDATA.csv
- * ... * 2020VAERSVAX.csv
- * 2020VAERSSYMPTOMS.csv
- * 2020VAERSDATA.csv

```

[6]: def get_data():
    """
    gets the data and returns a dataframe
    """

    all_data = os.path.join(DATAPATH, 'all_data.json')

    if os.path.isfile(all_data):
        print('loading all_data.json (15sec-30sec)')
        df = load_json(all_data)
        df = pd.DataFrame(df)
    else:
        print('processing the 2020-202X files')
        print("""
        .../VAERS/AllVAERSDataCSVs/
        20??VAERSVAX.csv
        20??VAERSSYMPTOMS.csv

```

```

        20??VAERSDATA.csv
    """)
    df = process_to_one_file()

    return df

df = get_data() # this will get all the data
# df = get_data().sample(int(946527 * 0.25)) # only 25% of the data

# assuming null for these is No
df.loc[df['DIED'].isna(), 'DIED'] = 'N'
df.loc[df['L_THREAT'].isna(), 'L_THREAT'] = 'N'
df.loc[df['ER_VISIT'].isna(), 'ER_VISIT'] = 'N'
df.loc[df['HOSPITAL'].isna(), 'HOSPITAL'] = 'N'

print('\nloaded {:,} records/rows\n'.format(len(df)))
print('columns:\n', df.columns.to_list())
print('\n\ndf.head(10):\n')
display(df.head(2))

```

processing the 2020-202X files

```

.../VAERS/AllVAERSDataCSVS/
    20??VAERSVAX.csv
    20??VAERSSYMPTOMS.csv
    20??VAERSDATA.csv

```

process to one file

this might take a while...go get a drink

dedup-ing Symptoms

merge data

saved: .\AllVAERSDataCSVS\all_data.json

saved: .\AllVAERSDataCSVS\all_data.csv

loaded 983,305 records/rows

columns:

```

['VAERS_ID', 'RECVDATE', 'STATE', 'AGE_YRS', 'CAGE_YR', 'CAGE_MO', 'SEX',
'RPT_DATE', 'SYMPTOM_TEXT', 'DIED', 'DATEDIED', 'L_THREAT', 'ER_VISIT',
'HOSPITAL', 'HOSPDAYS', 'X_STAY', 'DISABLE', 'RECOVD', 'VAX_DATE', 'ONSET_DATE',
'NUMDAYS', 'LAB_DATA', 'V_ADMINBY', 'V_FUNDBY', 'OTHER_MEDS', 'CUR_ILL',
'HISTORY', 'PRIOR_VAX', 'SPLTTYPE', 'FORM_VERS', 'TODAYS_DATE', 'BIRTH_DEFECT',
'OFC_VISIT', 'ER_ED_VISIT', 'ALLERGIES', 'VAX_TYPE', 'VAX_MANU', 'VAX_LOT',
'VAX_DOSE_SERIES', 'VAX_ROUTE', 'VAX_SITE', 'VAX_NAME', 'SYMPTOMS', 'COVID_VAX']

```

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

	VAERS_ID	RECVDATE	STATE	AGE_YRS	CAGE_YR	CAGE_MO	SEX	RPT_DATE	\
46308	902418	12/15/2020	NJ	56.00	56.00	NaN	F	NaN	
46327	902440	12/15/2020	AZ	35.00	35.00	NaN	F	NaN	

	SYMPTOM_TEXT	\
46308	Patient experienced mild numbness traveling from injection site up and	
	down arm that subsided over 20 minutes.	

46327

C/O Headache

	DIED	DATEDIED	L_THREAT	ER_VISIT	HOSPITAL	HOSPDAYS	X_STAY	DISABLE	\
46308	N	NaN	N	N	N	NaN	NaN	NaN	
46327	N	NaN	N	N	N	NaN	NaN	NaN	

	RECOVD	VAX_DATE	ONSET_DATE	NUMDAYS	LAB_DATA	V_ADMINBY	V_FUNDBY	\
46308	Y	12/15/2020	12/15/2020	0.00	none	PVT	NaN	
46327	Y	12/15/2020	12/15/2020	0.00	none	PVT	NaN	

	OTHER_MEDS	CUR_ILL	HISTORY	PRIOR_VAX	SPLTTYPE	FORM_VERS	TODAYS_DATE	\
46308	latex	none	none	NaN	NaN	2	12/15/2020	
46327	NaN	NaN	NaN	NaN	NaN	2	12/15/2020	

	BIRTH_DEFECT	OFC_VISIT	ER_ED_VISIT	ALLERGIES	VAX_TYPE	VAX_MANU	\
46308	NaN	NaN	NaN	none	COVID19	PFIZER\BIONTECH	
46327	NaN	NaN	NaN	NaN	COVID19	PFIZER\BIONTECH	

	VAX_LOT	VAX_DOSE_SERIES	VAX_ROUTE	VAX_SITE	\
46308	EH9899	1	IM	LA	
46327	EH 9899	1	SYR	LA	

	VAX_NAME	\
46308	COVID19 (COVID19 (PFIZER-BIONTECH))	
46327	COVID19 (COVID19 (PFIZER-BIONTECH))	

	SYMPTOMS	COVID_VAX
46308	[Hypoaesthesia, Injection site hypoaesthesia]	1
46327	[Headache]	1

```
[7]: def print_row(items,column_lengths=[]):
    """
    comes in handy :)
    """
    row = ''
    for index,i in enumerate(items):
```

```

    try:
        cl = column_lengths[index]
    except IndexError:
        cl = 20
    row += str(i)[0:cl].ljust(cl)
print(row)

```

4.6 gets a list of symptoms

df_symptoms (DataFrame) : a list of all the symptoms and the counts of each all_symptoms (list):
a list of all the symptoms and the counts of each

note:

symptoms might be medical jargon or plain english
i.e. “RASH”, “ERYTHEMA”, and “ITCHY RED SKIN”
would be reported as different items (for now)

note:

the counts/percentages below are of the symptoms.
and one adverse reaction can have multiple symptoms.

```

[33]: def get_symptom_list(df, column='SYMPTOMS'):
        """
        returns a list of symptoms for the dataframe
        """
        s = df[column].to_list()
        l = []
        for i in s:
            try:
                for j in i:
                    if str(j) == 'nan':
                        pass
                    else:
                        l.append(str(j).upper())
            except:
                pass
        return l

all_symptoms = get_symptom_list(df, 'SYMPTOMS')
symptoms_count = len(all_symptoms)
all_symptoms = Counter(all_symptoms).most_common()

df_symptoms = pd.DataFrame(all_symptoms, columns=['SYMPTOM', 'COUNT'])
df_symptoms['PERCENT'] = (df_symptoms['COUNT']/symptoms_count)*100

file_name = os.path.join(DATAPATH, 'symptoms.csv')
df_symptoms.to_csv(file_name)
print('saved: ', file_name)

```



```

topX = 50
print('below are the top {} symptoms'.format(topX))
display(df_symptoms.head(topX))

```

saved: .\AllVAERSDataCSVs\symptoms.csv

below are the top 50 symptoms

	SYMPTOM	COUNT	PERCENT
0	HEADACHE	154222	3.64
1	PYREXIA	133598	3.15
2	FATIGUE	133167	3.14
3	PAIN	110322	2.60
4	CHILLS	108458	2.56
5	COVID-19	92983	2.19
6	NAUSEA	84493	1.99
7	PAIN IN EXTREMITY	82660	1.95
8	DIZZINESS	81874	1.93
9	SARS-COV-2 TEST POSITIVE	65186	1.54
10	MYALGIA	59229	1.40
11	DYSPOEA	56400	1.33
12	ARTHRALGIA	53648	1.26
13	INJECTION SITE PAIN	46943	1.11
14	RASH	44233	1.04
15	PRURITUS	39742	0.94
16	ASTHENIA	38566	0.91
17	VOMITING	37968	0.89
18	COUGH	37029	0.87
19	EXPIRED PRODUCT ADMINISTERED	32188	0.76
20	DIARRHOEA	30722	0.72
21	SARS-COV-2 TEST	30267	0.71
22	INJECTION SITE ERYTHEMA	30030	0.71
23	NO ADVERSE EVENT	29044	0.68
24	FEELING ABNORMAL	29000	0.68
25	MALaise	28990	0.68
26	PRODUCT STORAGE ERROR	28557	0.67
27	CHEST PAIN	28460	0.67
28	URTICARIA	26870	0.63
29	PARAESTHESIA	26726	0.63
30	LYMPHADENOPATHY	26171	0.62
31	ERYTHEMA	26053	0.61
32	HYPOAESTHESIA	25670	0.61
33	INJECTION SITE SWELLING	25562	0.60
34	HYPERHIDROSIS	25552	0.60
35	PERIPHERAL SWELLING	21227	0.50
36	BLOOD TEST	20948	0.49
37	CONDITION AGGRAVATED	20896	0.49
38	INJECTION SITE PRURITUS	20252	0.48

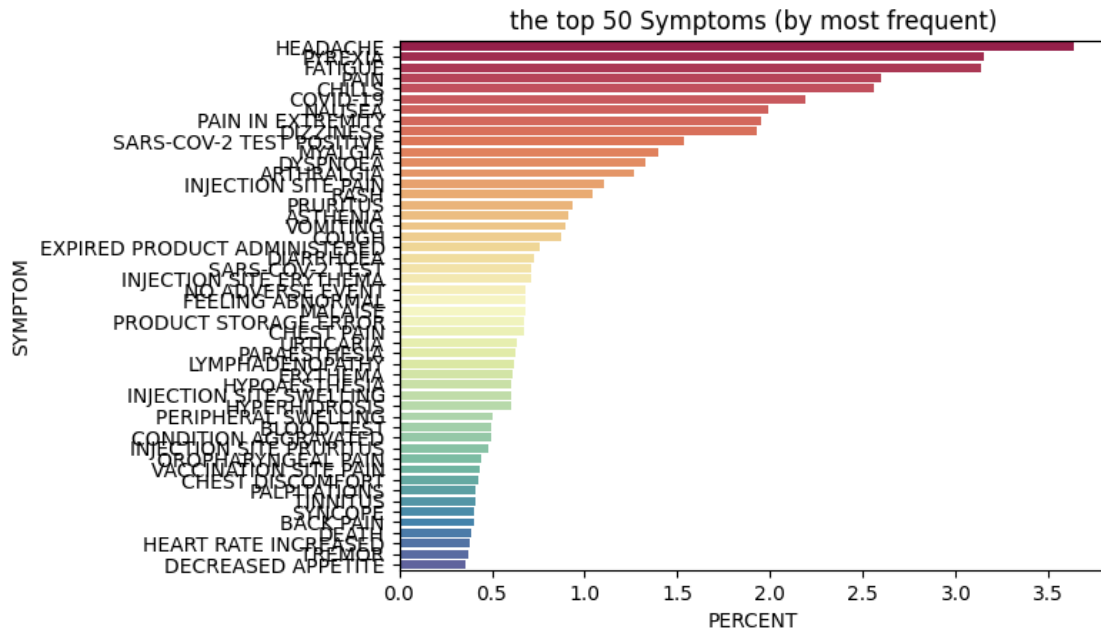
39	OROPHARYNGEAL PAIN	18683	0.44
40	VACCINATION SITE PAIN	18342	0.43
41	CHEST DISCOMFORT	18170	0.43
42	PALPITATIONS	17611	0.42
43	TINNITUS	17336	0.41
44	SYNCOPE	17070	0.40
45	BACK PAIN	16984	0.40
46	DEATH	16320	0.38
47	HEART RATE INCREASED	16234	0.38
48	TREMOR	15864	0.37
49	DECREASED APPETITE	15183	0.36

```
[35]: # lets put it in a histogram!

# fig = px.histogram(
#     df_symptoms.head(topX),
#     x="SYMPTOM",
#     y="PERCENT",
#     # marginal="rug", # can be `box`, `violin`
#     hover_data=df_symptoms.head(topX).columns
# )
# fig.show()

sns.barplot(
    data=df_symptoms.head(topX),
    x="PERCENT",
    y="SYMPTOM",
    # hue="PERCENT"
    palette = "Spectral"
).set(title=f'the top {topX} Symptoms (by most frequent)')
```

```
[35]: [Text(0.5, 1.0, 'the top 50 Symptoms (by most frequent)')]
```



4.7 break down functions...

these functions will help me breakdown the data

4.7.1 break_down_columns

```
[36]: def break_down_columns(idf,column):
    """
    shows what values there are for a given column (with counts and percent)
    """
    print('\nbreak down of {0}'.format(column))
    idf = pd.DataFrame(idf[column])
    idf = idf.fillna('nan')

    result = []

    l = list(idf[column].unique())
    for i in l:
        df0 = idf[idf[column]==i]
        result.append({'column':column,'value':i,'count':len(df0),'percent':
        ↪(len(df0)/len(idf))*100})

    result = pd.DataFrame(result)
    display(result)
    return result
```

```
# test
# break_down_columns(df, 'DIED')
```

4.7.2 break_down_buckets

```
[37]: def break_down_buckets(idf,column,buckets,message='',nan_value=-1):
    """
    breaks a column down into buckets/bins
    """
    idf = idf.fillna(nan_value)

    print('',message,'\ncolumn: ',column, '\nbuckets: ', buckets)

    idf = pd.DataFrame(idf[column])
    idf['bucket'] = pd.cut(idf[column], bins=buckets)
    idf = idf.groupby(by='bucket').count()
    idf['percent'] = (idf[column]/idf[column].sum())*100
    idf['percent'] = idf['percent'].round(2)
    display(idf)

    return idf

#test
# break_down_buckets(df, 'AGE_YRS', [-1,0,15,25,35,45,55,65,75,85,500])
```

4.8 Additional Numbers

these numbers are **not** part of the vaers data;
however they are important to analyzing the data

vaxx (int) : the number of vaccinated (1 or more shots) US citizens according to a quick google search (on 8/3/2021)

~~**google no longer shows partially vaxxed** so we are using the **Total doses given** number
“Total doses given” shows the number of vaccine doses given to people. Since some vaccines require more than 1 dose, the number of fully vaccinated people is likely lower. “People fully vaccinated” shows how many people have received the full amount of doses for the COVID-19 vaccine. ~~

~~This number was updated on 06/28/2022~~

full_vaxx (int) : the number of vaccinated (2 or more shots) US citizens according to a quick google search ~~(on 06/28/2022)~~

~~the numbers were taken from <https://usafacts.org/visualizations/covid-vaccine-tracker-states> on 11/18/2022~~

the numbers were taken from <https://usafacts.org/visualizations/covid-vaccine-tracker-states> on 9/10/2023

```
[79]: vaxx = 270_227_181 + 230_637_348
      full_vaxx = 230_637_348
```

```
[80]: #only death vaers
      df_death = df[df['DIED']=='Y']
```

4.9 Analyzing the df (all_data.json)

this is just a generic break down of the data

df_death (DataFrame) : adverse reactions that resulted in a death

4.9.1 DIED column

```
[81]: temp = break_down_columns(df, 'DIED')
      print('\n' + '-'*25 + '\n')

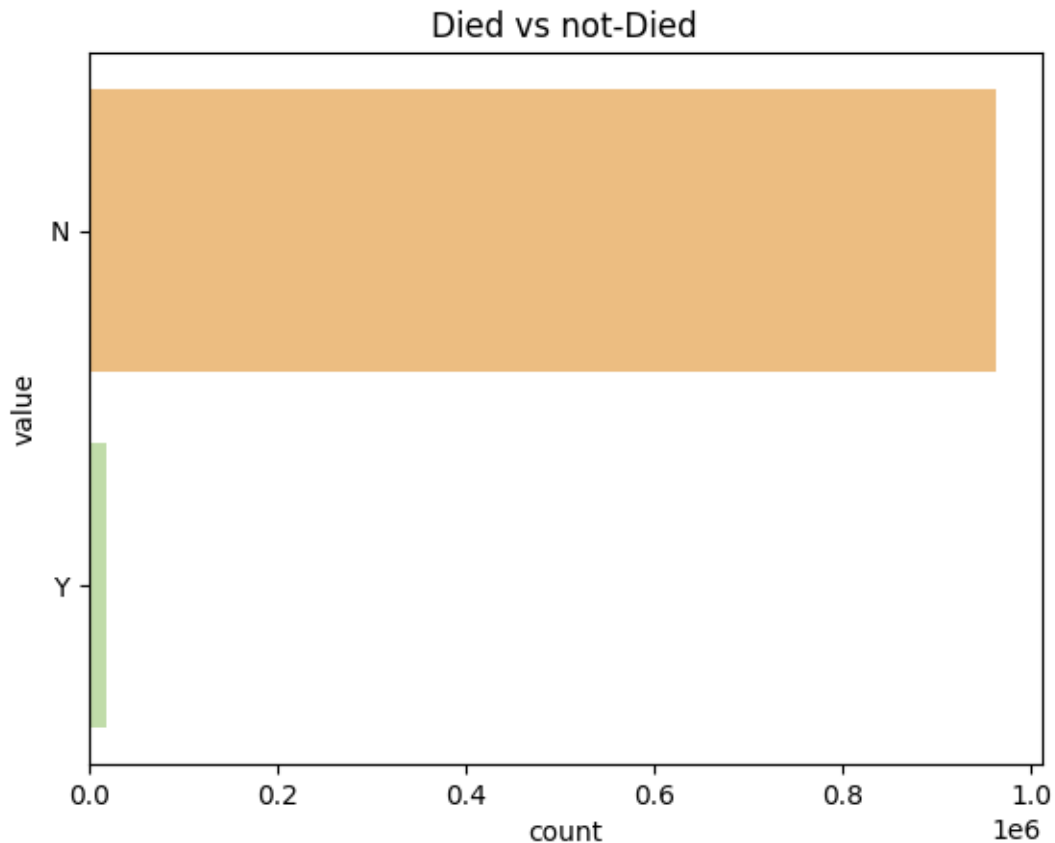
      # fig = px.pie(
      #     temp,
      #     values='count',
      #     names='value',
      #     title='Death as Symptoms from Vaccine',
      #     # hover_data = temp.columns
      # )
      # fig.show()

      sns.barplot(
          data=temp,
          x="count",
          y="value",
          # hue="PERCENT"
          palette = "Spectral"
      ).set(title='Died vs not-Died')
```

break down of DIED

	column	value	count	percent
0	DIED	N	963477	97.98
1	DIED	Y	19828	2.02

```
[81]: [Text(0.5, 1.0, 'Died vs not-Died')]
```



4.9.2 ER_VISIT column

```
[82]: print('did the adverse reaction result in an ER Visit')
temp = break_down_columns(df, 'ER_VISIT')
print('\n' + '-'*25 + '\n')

# temp.plot.
# bar(title='ER_VISIT',x='value',y='count',rot=0,figsize=(5,5),width=0.
# 25,fontsize=18)
# temp = temp.set_index('value')
# temp.plot.pie(title='ER_VISIT',y='percent',figsize=(5, 5),autopct='%.'
# 2f',fontsize=18)

# fig = px.bar(
#     temp,
#     x='count',
#     y='value',
#     title='Did the they need the ER?',
# )
```

```
# fig.show()

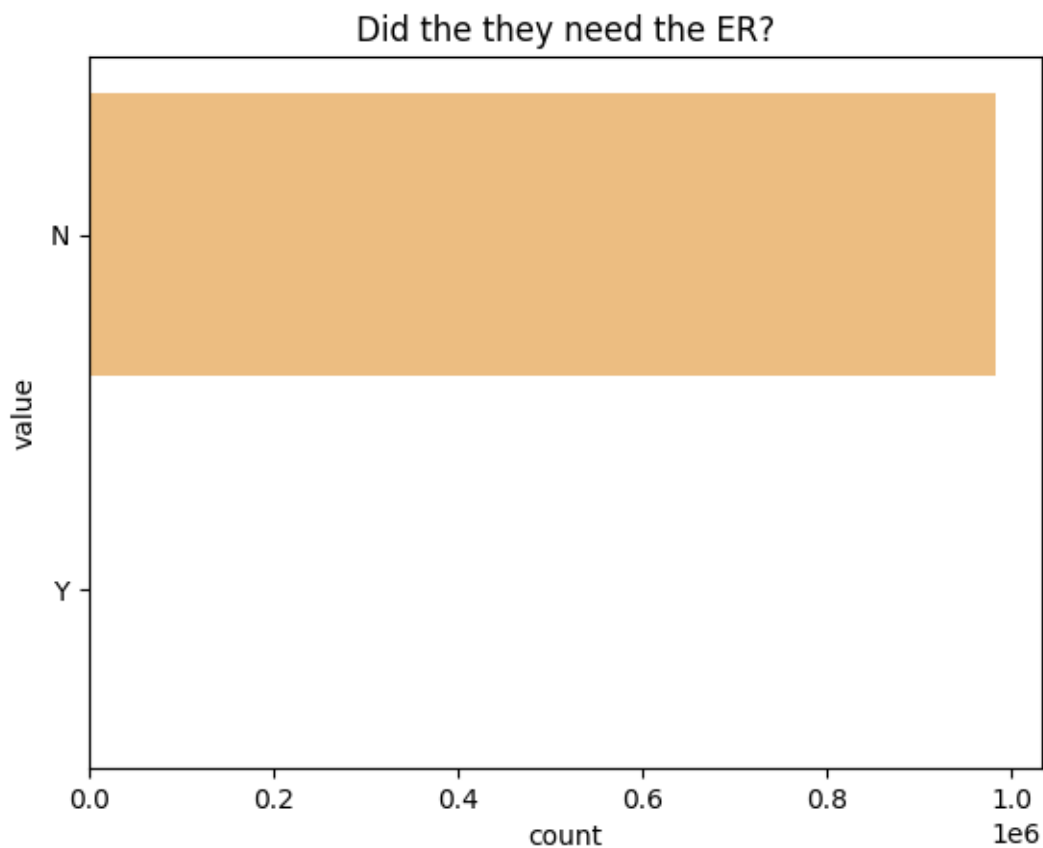
sns.barplot(
    data=temp,
    x="count",
    y="value",
    # hue="PERCENT"
    palette = "Spectral"
).set(title='Did the they need the ER?')
```

did the adverse reaction result in an ER Visit

break down of ER_VISIT

	column	value	count	percent
0	ER_VISIT	N	983165	99.99
1	ER_VISIT	Y	140	0.01

[82]: [Text(0.5, 1.0, 'Did the they need the ER?')]



4.9.3 L_THREAT column

```
[83]: print('Life Threatening/Leathal Threat')
temp = break_down_columns(df, 'L_THREAT')
print('\n' + '-'*25 + '\n')

# temp.plot.
↳ bar(title='L_THREAT',x='value',y='count',rot=0,figsize=(5,5),width=0.
↳ 25,fontsize=18)
# temp = temp.set_index('value')
# temp.plot.pie(title='L_THREAT',y='percent',figsize=(5, 5),autopct='%
↳ 2f',fontsize=18)

# fig = px.bar(
#     temp,
#     x='count',
#     y='value',
#     title="was this a lethal threat?"
# )
# fig.show()

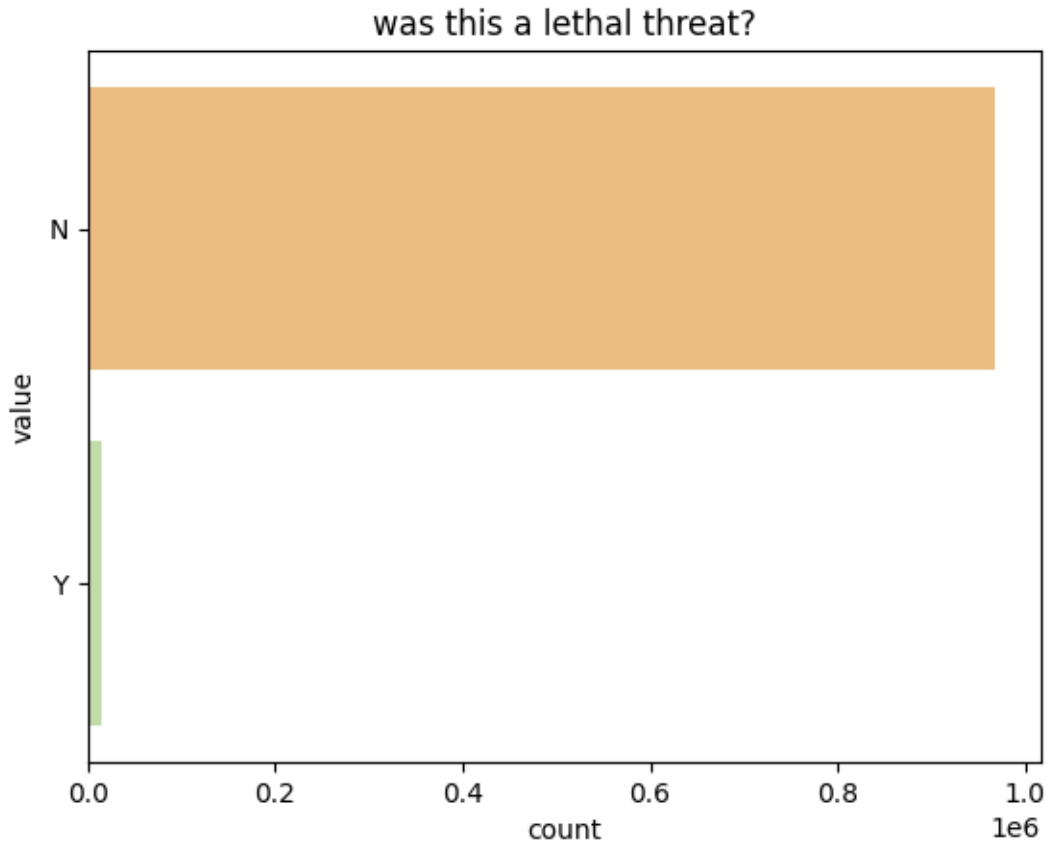
sns.barplot(
    data=temp,
    x="count",
    y="value",
    # hue="PERCENT"
    palette = "Spectral"
).set(title='was this a lethal threat?')
```

Life Threatening/Leathal Threat

break down of L_THREAT

	column	value	count	percent
0	L_THREAT	N	967762	98.42
1	L_THREAT	Y	15543	1.58

```
[83]: [Text(0.5, 1.0, 'was this a lethal threat?')]
```

4.9.4 RECOVD column

```
[84]: print('did the patient recover?')
temp = break_down_columns(df, 'RECOVD')

print('\n' + '-'*25 + '\n')

# temp.plot.bar(title='RECOVD',x='value',y='count',rot=0,figsize=(5,5),width=0.
# ↪25,fontsize=18)
# temp = temp.set_index('value')
temp.plot.pie(title='RECOVD',y='percent',figsize=(5, 5),autopct='%.'
# ↪2f',fontsize=18)

# fig = px.bar(
#     temp,
#     x='count',
#     y='value',
#     title="Did the patient recover? (Y=yes, N=no, U=unknown, nan=N/A )"
# )
```

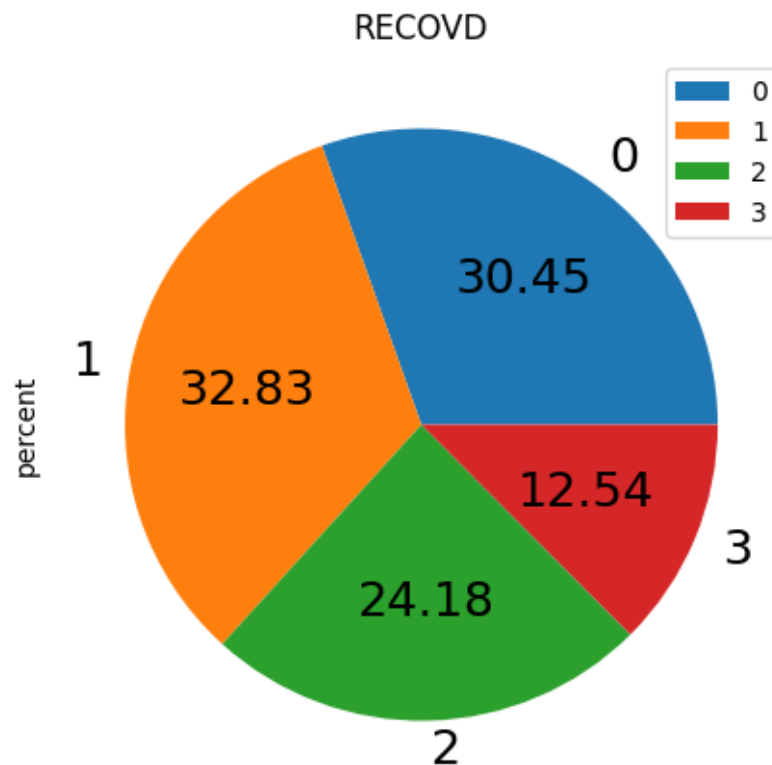
```
# fig.show()
```

did the patient recover?

break down of RECOVD

	column	value	count	percent
0	RECOVD	Y	299382	30.45
1	RECOVD	N	322851	32.83
2	RECOVD	U	237799	24.18
3	RECOVD	nan	123273	12.54

```
[84]: <Axes: title={'center': 'RECOVD'}, ylabel='percent'>
```



4.9.5 the Age of the patient

```
[85]: print('the Age of the patient')
temp = break_down_buckets(df, 'AGE_YRS', [0,15,25,35,45,55,65,75,85,500])
temp = temp.reset_index()
temp["bucket"] = temp["bucket"].astype(str)

# temp = temp.reset_index()
# temp.plot.
# bar(title='AGE_YRS',x='bucket',y='AGE_YRS',rot=90,figsize=(5,5),width=0.
# 25,fontsize=18)
# # temp.plot.
# bar(title='AGE_YRS',x='bucket',y='percent',rot=90,figsize=(5,5),width=0.
# 25,fontsize=18)

# fig = px.bar(
#     temp,
#     x='percent',
#     y='bucket',
#     title="The Age of the Patient"
# )
# fig.show()

sns.barplot(
    data=temp,
    x="percent",
    y="bucket",
    palette = "Spectral"
).set(title='The Age of the Patient')

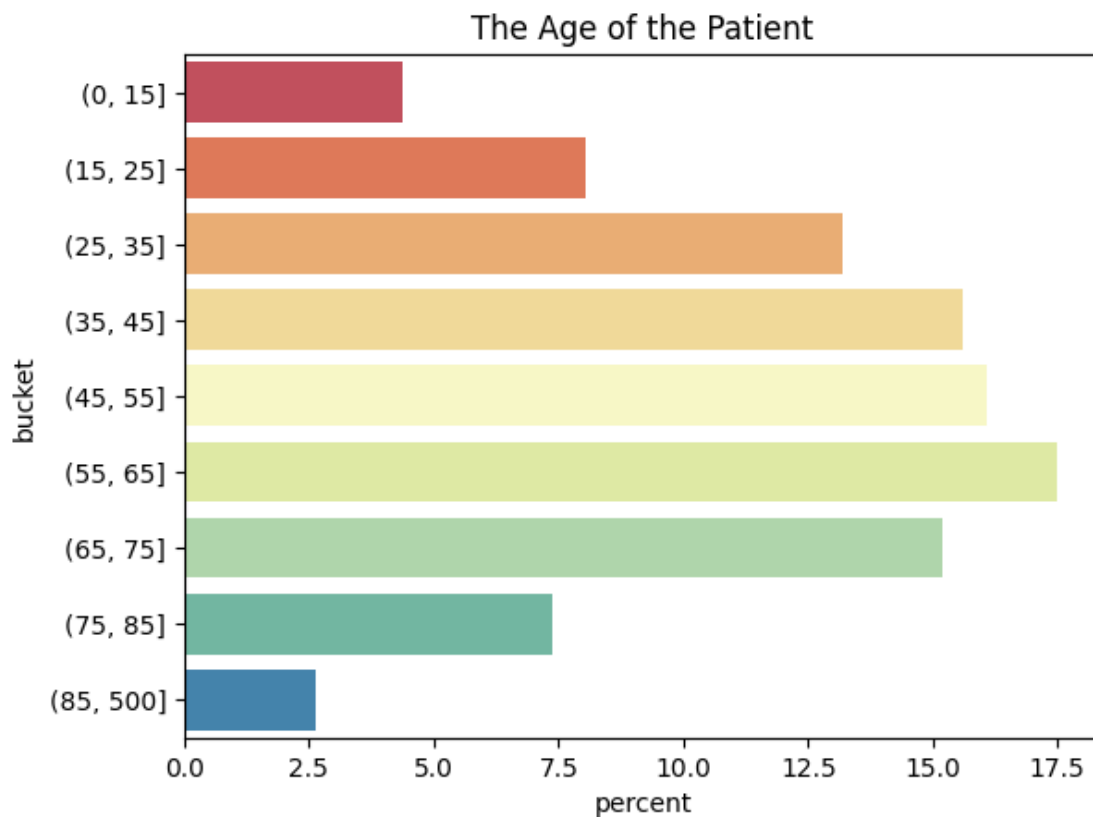
print('\n' + '-'*25 + '\n')
```

the Age of the patient

column: AGE_YRS

buckets: [0, 15, 25, 35, 45, 55, 65, 75, 85, 500]

	AGE_YRS	percent
bucket		
(0, 15]	38797	4.36
(15, 25]	71592	8.05
(25, 35]	117340	13.20
(35, 45]	138581	15.59
(45, 55]	142923	16.07
(55, 65]	155638	17.50
(65, 75]	135097	15.19
(75, 85]	65641	7.38
(85, 500]	23511	2.64



```
[86]: temp =
    ↳break_down_buckets(df_death, 'AGE_YRS', [0,15,25,35,45,55,65,75,85,500],message='***deaths_
    ↳only***')
temp = temp.reset_index()
temp["bucket"] = temp["bucket"].astype(str)

# temp = temp.reset_index()
# temp.plot.bar(title='AGE_YRS (death_
    ↳only)',x='bucket',y='AGE_YRS',rot=90,figsize=(5,5),width=0.25,fontsize=18)
# # temp.plot.bar(title='AGE_YRS (death_
    ↳only)',x='bucket',y='percent',rot=90,figsize=(5,5),width=0.25,fontsize=18)

# fig = px.bar(
#     temp,
#     x='percent',
#     y='bucket',
#     title="The Age of the Patient"
```

```

#         )
# fig.show()

sns.barplot(
    data=temp,
    x="percent",
    y="bucket",
    palette = "Spectral"
).set(title='The Age of the Patient (Deaths Only)')

print('\n' + '-'*25 + '\n')

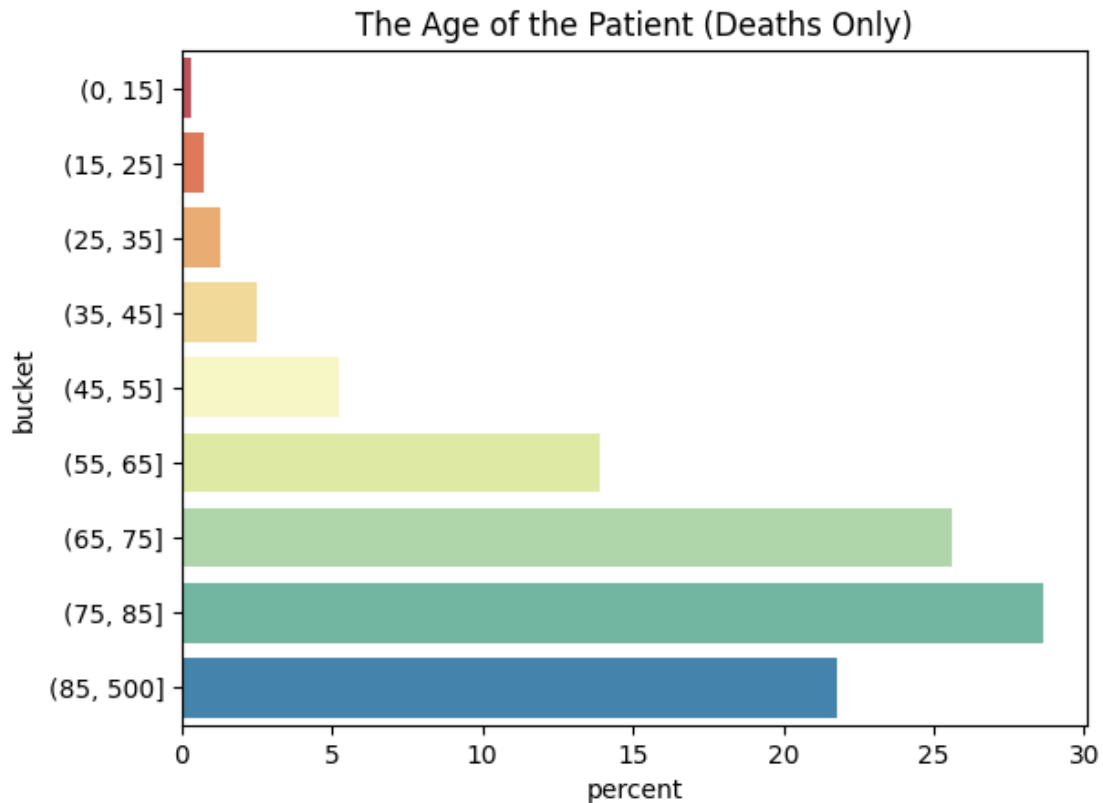
```

```

***deaths only***
column:  AGE_YRS
buckets: [0, 15, 25, 35, 45, 55, 65, 75, 85, 500]

```

bucket	AGE_YRS	percent
(0, 15]	54	0.29
(15, 25]	139	0.76
(25, 35]	233	1.27
(35, 45]	462	2.51
(45, 55]	963	5.23
(55, 65]	2558	13.90
(65, 75]	4710	25.60
(75, 85]	5272	28.66
(85, 500]	4007	21.78



4.9.6 the number of days between the vaccine and the adverse rection

```
[87]: print('the number of days between the vaccine and the adverse rection')
temp = break_down_buckets(df, 'NUMDAYS', [0,1,5,10,20,30,40,50,60])
temp = temp.reset_index()
temp["bucket"] = temp["bucket"].astype(str)

# temp.plot.
↳ barh(title='NUMDAYS',x='bucket',y='NUMDAYS',rot=0,figsize=(10,5),width=0.
↳ 25,fontsize=18)
# # temp.plot.
↳ bar(title='AGE_YRS',x='bucket',y='percent',rot=90,figsize=(5,5),width=0.
↳ 25,fontsize=18)

# fig = px.bar(
#     temp,
#     x='percent',
#     y='bucket',
#     title="the number of days between the vaccine and the adverse rection"
# )
# fig.show()
```

```

sns.barplot(
    data=temp,
    x="percent",
    y="bucket",
    palette = "Spectral"
).set(title='the number of days between the vaccine and the adverse_
reaction')

print('\n' + '-'*25 + '\n')

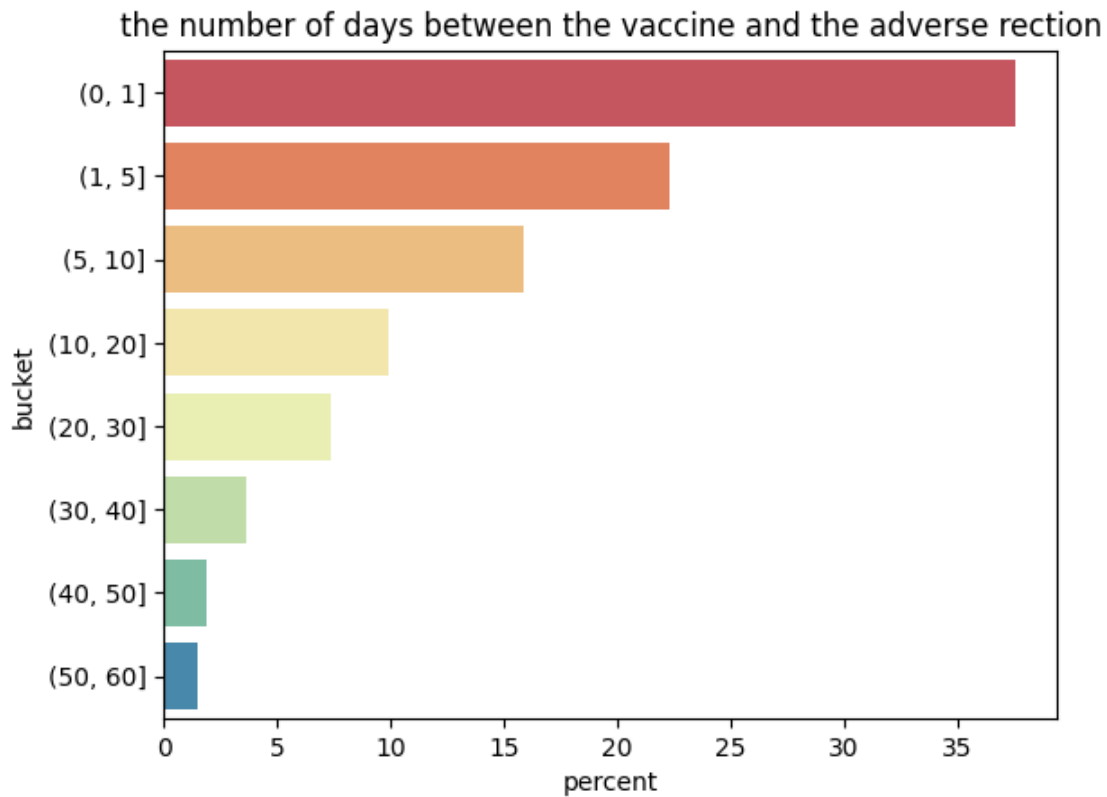
```

the number of days between the vaccine and the adverse rection

column: NUMDAYS

buckets: [0, 1, 5, 10, 20, 30, 40, 50, 60]

bucket	NUMDAYS	percent
(0, 1]	147363	37.54
(1, 5]	87665	22.33
(5, 10]	62496	15.92
(10, 20]	38932	9.92
(20, 30]	28835	7.35
(30, 40]	14159	3.61
(40, 50]	7293	1.86
(50, 60]	5796	1.48



```
[88]: temp =
    ↳break_down_buckets(df_death, 'NUMDAYS', [0,1,5,10,20,30,40,50,60], message='***deaths_
    ↳only***')
temp = temp.reset_index()
temp["bucket"] = temp["bucket"].astype(str)

# temp.plot.barh(title='NUMDAYS (deaths_
    ↳only)', x='bucket', y='NUMDAYS', rot=0, figsize=(10,5), width=0.25, fontsize=18)
# temp.plot.
    ↳bar(title='AGE_YRS', x='bucket', y='percent', rot=90, figsize=(5,5), width=0.
    ↳25, fontsize=18)

# fig = px.bar(
#     temp,
#     x='percent',
#     y='bucket',
#     title="the number of days between the vaccine and death"
# )
# fig.show()

sns.barplot(
```



```

data=temp,
x="percent",
y="bucket",
palette = "Spectral"
).set(title='the number of days between the vaccine and death')

print('\n' + '-'*25 + '\n')

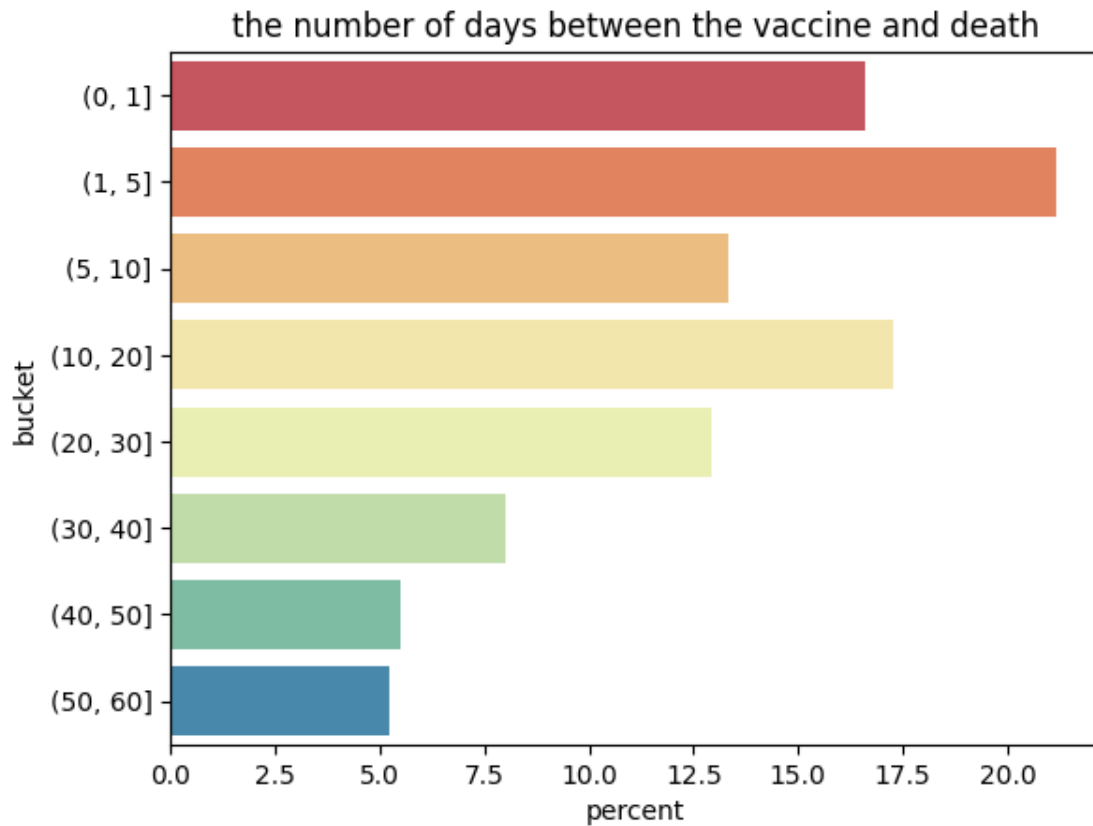
```

```

***deaths only***
column:  NUMDAYS
buckets: [0, 1, 5, 10, 20, 30, 40, 50, 60]

```

	NUMDAYS	percent
bucket		
(0, 1]	1261	16.59
(1, 5]	1608	21.15
(5, 10]	1015	13.35
(10, 20]	1312	17.26
(20, 30]	984	12.94
(30, 40]	610	8.02
(40, 50]	417	5.48
(50, 60]	396	5.21



```
[89]: temp = df[df['NUMDAYS'] <= 10]

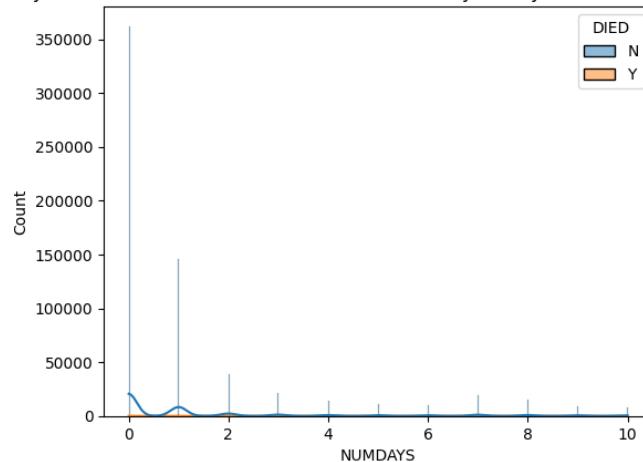
sns.histplot(
    data=temp,
    x="NUMDAYS",
    hue='DIED',
    kde=True
).set(title='showing number of days between vaccine and adverse reaction (
↳ only 10 days or less, broke out death into other color)')

# fig = px.histogram(
#     temp,
#     x="NUMDAYS",
#     color='DIED',
#     marginal="violin",
#     hover_data=temp.columns,
#     title="showing number of days between vaccine and adverse reaction ( only
↳ 25 days or less, 25% sample, broke out death into other color)"
# )
# fig.update_layout(barmode='overlay')
```

```
# fig.update_traces(opacity=0.75)
# fig.show()
```

[89]: [Text(0.5, 1.0, 'showing number of days between vaccine and adverse reaction (only 10 days or less, broke out death into other color)')]

showing number of days between vaccine and adverse reaction (only 10 days or less, broke out death into other color)



4.9.7 break down of the VAX_NAME column

```
[90]: print('break down of the VAX_NAME column')
df_VN = break_down_columns(df, 'VAX_NAME')

print('\n***deaths only***')
df_DVN = break_down_columns(df_death, 'VAX_NAME')

# df_CVN = df_VN.join(df_DVN[['value', 'count']], on='value')
df_DVN = df_DVN.rename(columns={"count": "death_count"})

print('calculating the death ratio ( death_count / count )')
df_CVN = pd.concat([df_VN, df_DVN], keys=['value'], join="inner", axis=1)

df_CVN = pd.merge(df_VN, df_DVN, on=['value'])
df_CVN = df_CVN.drop(columns=['percent_x', 'percent_y', 'column_y'])
df_CVN = df_CVN.rename(columns={"column_x": "column"})
df_CVN['death_ratio'] = df_CVN['death_count'] / df_CVN['count']
df_CVN['death_percent'] = df_CVN['death_ratio'] * 100

display(df_CVN)
```

```
print('\n' + '-'*25 + '\n')
```

break down of the VAX_NAME column

break down of VAX_NAME

	column	value	count	percent
0	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH))	447882	45.55
1	VAX_NAME	COVID19 (COVID19 (UNKNOWN))	4999	0.51
2	VAX_NAME	COVID19 (COVID19 (MODERNA))	443030	45.06
3	VAX_NAME	COVID19 (COVID19 (MODERNA BIVALENT))	6745	0.69
4	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH BIVALENT))	8890	0.90
5	VAX_NAME	COVID19 (COVID19 (JANSSEN))	71550	7.28
6	VAX_NAME	COVID19 (COVID19 (NOVAVAX))	209	0.02

deaths only

break down of VAX_NAME

	column	value	count	percent
0	VAX_NAME	COVID19 (COVID19 (MODERNA))	8666	43.71
1	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH))	9196	46.38
2	VAX_NAME	COVID19 (COVID19 (UNKNOWN))	66	0.33
3	VAX_NAME	COVID19 (COVID19 (JANSSEN))	1781	8.98
4	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH BIVALENT))	75	0.38
5	VAX_NAME	COVID19 (COVID19 (MODERNA BIVALENT))	44	0.22

calculating the death ratio (death_count / count)

	column	value	count \
0	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH))	447882
1	VAX_NAME	COVID19 (COVID19 (UNKNOWN))	4999
2	VAX_NAME	COVID19 (COVID19 (MODERNA))	443030
3	VAX_NAME	COVID19 (COVID19 (MODERNA BIVALENT))	6745
4	VAX_NAME	COVID19 (COVID19 (PFIZER-BIONTECH BIVALENT))	8890
5	VAX_NAME	COVID19 (COVID19 (JANSSEN))	71550

	death_count	death_ratio	death_percent
0	9196	0.02	2.05
1	66	0.01	1.32
2	8666	0.02	1.96
3	44	0.01	0.65
4	75	0.01	0.84
5	1781	0.02	2.49

4.10 Approximating actual adverse reaction numbers

VAERS only contains reported data and

'...fewer than 1% of vaccine adverse events are reported.'

Source: <https://digital.ahrq.gov/sites/default/files/docs/publication/r18hs017045-lazarus-final-report-2011.pdf> (page 6)

we will multiply the counts by 80 and 120,

in order to get an approximate min and max of what the numbers might actually be.

```
[91]: def print_percent(vmin,vmax,label0,vcount,label1):
    """
    vmin: vaers min
    vmax: vaers max
    label0: vaers label
    vcount: vaxxed count
    label1: vaxxed label
    """
    print(
        '({label0} / {label1}) * 100\n'.format(
            label0=label0,
            label1=label1
        ),
        'min: ({0:,} / {1:,}) * 100 \n'.format(vmin,vcount),
        'max: ({0:,} / {1:,}) * 100 \n'.format(vmax,vcount),
        '{:.2f} %'.format((vmin/vcount)*100),
        '-',
        '{:.2f} %'.format((vmax/vcount)*100),
        '\n'
    )

    print_percent(
        len(df)*80,
        len(df)*120,
        'approx adverse reactions',
        vaxx,
        'vaxxed [1 or more shots]'
    )

    print_percent(
        len(df_death)*80,
        len(df_death)*120,
        'approx adverse deaths',
        vaxx,
        'vaxxed [1 or more shots]'
    )

df_nrecovd = df[df['RECOVD']=='N']
```

```

print_percent(
    len(df_nrecovd)*80,
    len(df_nrecovd)*120,
    'approx no recovery',
    vaxx,
    'vaxxed [1 or more shots]'
)

df_urecovd = df[df['RECOVD']=='U']
print_percent(
    (len(df_nrecovd) + (len(df_urecovd)*0.5))*80,
    (len(df_nrecovd) + (len(df_urecovd)*0.5))*120,
    'approx no recovery + (50% of unknowns)',
    vaxx,
    'vaxxed [1 or more shots]'
)

```

```

( approx adverse reactions / vaxxed [1 or more shots] ) * 100
min: ( 78,664,400 / 500,864,529 ) * 100
max: ( 117,996,600 / 500,864,529 ) * 100
15.71 % - 23.56 %

```

```

( approx adverse deaths / vaxxed [1 or more shots] ) * 100
min: ( 1,586,240 / 500,864,529 ) * 100
max: ( 2,379,360 / 500,864,529 ) * 100
0.32 % - 0.48 %

```

```

( approx no recovery / vaxxed [1 or more shots] ) * 100
min: ( 25,828,080 / 500,864,529 ) * 100
max: ( 38,742,120 / 500,864,529 ) * 100
5.16 % - 7.74 %

```

```

( approx no recovery + (50% of unknowns) / vaxxed [1 or more shots] ) * 100
min: ( 35,340,040.0 / 500,864,529 ) * 100
max: ( 53,010,060.0 / 500,864,529 ) * 100
7.06 % - 10.58 %

```

4.11 Women's Reproductive Symptoms

Why? I have women in my life that were curious about this.

WRS_list (list) : a list of symptoms that effect or could cause effects to a women's reproductive system

df_WRS (DataFrame) : a dataframe that contains VAERS events that have at least 1 of the WRS symptoms

```
[92]: def symptom_filter_search(idf, search_list):
    """
    returns a dataframe pf vaers events where the patient
    has had 1 or more of the symptoms on the list
    """
    print('this could take between 20sec-60sec')

    data = idf.to_dict(orient='records')

    search_list = [i.upper() for i in search_list]

    results = []
    for index,d in enumerate(data):
        # if index%5000 == 0:
        #     print('{:.2f}'.format(index/len(data)))
        try:
            d['SYMPTOMS'] = [i.upper() for i in d['SYMPTOMS'] if isinstance(i,
↪str)]

            symptom_match = list(set(d['SYMPTOMS']) & set(search_list))
            d['SYMPTOMS_MATCH'] = symptom_match
            d['SYMPTOMS_MATCH_LENGTH'] = len(symptom_match)
            if len(symptom_match) > 0:
                results.append(d)
        except:
            pass

    return pd.DataFrame(results)
```

```
[93]: WRS_list = [
    'Intermenstrual bleeding',
    'Menopause',
    'Heavy menstrual bleeding',
    'dysmenorrhoea',
    'ABNORMAL UTERINE BLEEDING',
    'MATERNAL EXPOSURE BEFORE PREGNANCY',
    'MENSTRUATION IRREGULAR',
    'Oligomenorrhea',
    'OLIGOMENORRHOEA',
    'POLYMENORRHOEA',
    'MENSTRUAL DISORDER',
    'OLIGOMENORRHOEA',
    'ANOVULATORY CYCLE',
    'OVULATION DELAYED',
    'BACTERIAL VAGINOSIS',
    'GYNAECOLOGICAL EXAMINATION ABNORMAL',
    'OVARIAN CYST',
```

'BIOPSY UTERUS',
'UTERINE LEIOMYOMA',
'HOT FLUSH',
'BREAST TENDERNESS',
'BREAST SWELLING',
'BREAST PAIN',
'VAGINAL HAEMORRHAGE'

```
[94]: df_WRS = symptom_filter_search(df,WRS_list)

print('df_WRS.head(5)')
display(df_WRS.head(5))
```

```
this could take between 20sec-60sec
df_WRS.head(5)
```

	VAERS_ID	RECVDATE	STATE	AGE_YRS	CAGE_YR	CAGE_MO	SEX	RPT_DATE	\
0	902796	12/16/2020	TX	34.00	34.00	NaN	F	NaN	
1	903202	12/17/2020	NE	41.00	41.00	NaN	F	NaN	
2	903247	12/17/2020	AR	42.00	42.00	NaN	F	NaN	
3	903329	12/18/2020	PA	46.00	46.00	NaN	F	NaN	
4	903345	12/18/2020	VA	37.00	37.00	NaN	F	NaN	

0 I inserted my NuvaRing birth control on 12/14/2020. I have the Covid19 vaccine on 12/15/2020 at about 8:30am. The injection site was mildly sore and that continued into the next day. I woke up on 12/16/2020 feeling a little off and it progressed throughout the day. I felt a headache that I knew was turning into a migraine, threw up a few times (this is normal for me when I get migraines), and started having hot flashes. My temperature never went above 98. After sleeping a few hours and taking some Excedrin migraine I was feeling much better, though still a little sickly, by 4pm. I really think this is due to my migraines (which I get roughly once a month) even though it was a day later than normal.

1
↳
↳
↳
↳
↳
↳
↳
Severe dizzy spell about 5-10
↳following injection, helped to the floor, this lasted approximately 30-40
↳minutes after start of symptom. Hot flashes and visual disturbance lasting
↳into following day.

2
↳
↳
↳
↳
↳
↳
↳
↳
I had arm pain. Yesterday I had a hot flash with throwing up and
↳diarrhea.

3
↳
↳
↳
↳
↳
↳
↳
↳
↳
hot flushing feeling, light
↳headed, legs heavy gave patient a chair to sit and candy symptoms resolved by
↳11:36

4
↳
↳
↳
↳
↳
↳
↳
↳
↳
Right arm soreness, headache, hot
↳flashes.

	DIED	DATEDIED	L_THREAT	ER_VISIT	HOSPITAL	HOSPDAYS	X_STAY	DISABLE	RECOVD	\
0	N	NaN	N	N	N	NaN	NaN	NaN	Y	
1	N	NaN	N	N	N	NaN	NaN	NaN	Y	
2	N	NaN	N	N	N	NaN	NaN	NaN	N	
3	N	NaN	N	N	N	NaN	NaN	NaN	Y	
4	N	NaN	N	N	N	NaN	NaN	NaN	Y	

	VAX_DATE	ONSET_DATE	NUMDAYS	LAB_DATA	V_ADMINBY	V_FUNDBY	\
0	12/15/2020	12/16/2020	1.00	NaN	PVT	NaN	
1	12/16/2020	12/16/2020	0.00	None	OTH	NaN	
2	12/15/2020	12/16/2020	1.00	No	UNK	NaN	
3	12/17/2020	12/17/2020	0.00	NaN	PVT	NaN	
4	12/16/2020	12/17/2020	1.00	NaN	WRK	NaN	

↪

↪

↪

↪OTHER_MEDS \

0

↪

↪

↪Methylphenidate Citalopram Buspirone

↪Tylenol

1 Buspar, propranolol, escitalopram, crestor, amitriptyline, baby aspirin,
 ↪super B complex, black elderberry, calcium/mag/zinc supplement, CoQ10, vitamin
 ↪D3, biotin, antarctic Krill oil, women's multivitamin, tart cherry
 ↪supplement, ibupr

2

↪

↪I

↪take Estradiol, Metprolol, Lexapro, Multivitamin, Calcium, Ambien, Benadryl,
 ↪Melatonin

3

↪

↪

↪NaN

4

↪

↪

↪NaN

CUR_ILL \

0 None

1 None

2 No

3 NaN

4 None

↪

↪

↪

HISTORY \

0 I normally get a migraine a day or two after I insert my NuvaRing. I
 ↪inserted it on Monday, 12/14/2020 and I think all my symptoms are related to
 ↪that rather than the vaccination.

1
↳
↳depression, hyperlipidemia
2
↳
↳
↳ I have SVT
3
↳
↳
↳ NaN
4
↳
↳
↳ None

	PRIOR_VAX	SPLTTYPE	FORM_VERS	TODAYS_DATE	BIRTH_DEFECT	OFC_VISIT	\
0	NaN	NaN	2	12/16/2020	NaN	NaN	
1	NaN	NaN	2	12/17/2020	NaN	NaN	
2	NaN	vsafe	2	12/17/2020	NaN	NaN	
3	NaN	NaN	2	12/18/2020	NaN	NaN	
4	NaN	NaN	2	12/18/2020	NaN	NaN	

	ER_ED_VISIT		ALLERGIES	VAX_TYPE	VAX_MANU	VAX_LOT	\
0	NaN		None	COVID19	PFIZER\BIONTECH	EH9899	
1	NaN		Aimovig	COVID19	PFIZER\BIONTECH	EH9899	
2	NaN	I am allergic to	Sulphur	COVID19	PFIZER\BIONTECH	NaN	
3	NaN		NaN	COVID19	PFIZER\BIONTECH	EK5730	
4	NaN		None	COVID19	PFIZER\BIONTECH	NaN	

	VAX_DOSE_SERIES	VAX_ROUTE	VAX_SITE		VAX_NAME	\
0	1	SYR	LA	COVID19 (COVID19 (PFIZER-BIONTECH))		
1	1	IM	RA	COVID19 (COVID19 (PFIZER-BIONTECH))		
2	UNK	IM	LA	COVID19 (COVID19 (PFIZER-BIONTECH))		
3	1	IM	RA	COVID19 (COVID19 (PFIZER-BIONTECH))		
4	UNK	NaN	RA	COVID19 (COVID19 (PFIZER-BIONTECH))		

↳
SYMPTOMS \

0 [CONDITION AGGRAVATED, FEELING ABNORMAL, HOT FLUSH, INJECTION SITE PAIN, ↳
↳MALAISE, MIGRAINE, VOMITING]

1 [DIZZINESS, HOT ↳
↳FLUSH, VISUAL IMPAIRMENT]

2 [DIARRHOEA, HOT FLUSH, PAIN ↳
↳IN EXTREMITY, VOMITING]

3 [DIZZINESS, HOT ↳
↳FLUSH, LIMB DISCOMFORT]

4 [HEADACHE, HOT ↳
↳FLUSH, PAIN IN EXTREMITY]

	COVID_VAX	SYMPTOMS_MATCH	SYMPTOMS_MATCH_LENGTH
0	1	[HOT FLUSH]	1
1	1	[HOT FLUSH]	1
2	1	[HOT FLUSH]	1
3	1	[HOT FLUSH]	1
4	1	[HOT FLUSH]	1

4.11.1 What percent of women experienced WRS during their adverse reactions?

(women experiencing reproductive symptoms / number of women in VAERS data) * 100

WRS_ratio (float) : count WRS / count VAERS

```
[95]: #get count of women in VAERS data

w_df = df[df['SEX']=='F']
u_df = df[df['SEX']=='U']
w_count = len(w_df) + (len(u_df)/2) #half of unknown

WRS_ratio = (len(df_WRS)/w_count)

print( '{:.2f} %'.format(WRS_ratio*100))

file_name = os.path.join(DATAPATH, 'WRS.csv')
df_WRS.to_csv(file_name)

print('VAERS records of women experiencing reproductive symptoms have been_
↪ saved.')
print('saved: ',file_name)
```

3.35 %

VAERS records of women experiencing reproductive symptoms have been saved.
saved: .\AllVAERSDataCSVs\WRS.csv

4.11.2 Approximate the number of WRS in reality

since '~63% of the people who are vaccinated are women' source:
<https://www.statista.com/statistics/1212103/share-of-persons-initiating-covid-vaccinations-by-gender-us-first-month/>

we will be multiplying the total number of people vaccinated (1 or more shots) by 0.63 to get the count of women vaccinated.

women_vaxx (float) : an approximate number of women who have had 1 or more vaccine shot.

note: 0.63 is an estimate, and the actual could be somewhere between 0.50 and 0.70

```
[96]: women_vaxx = vaxx * 0.63
print(women_vaxx)
```

315544653.27

4.11.3 approximate real WRS

if we assume that the VAERS data is a random sample (or close to it)
then the ratio of WRS systems should be the same...

and thus we can get an approximate number of women that would be experiencing reproductive symptoms by multiplying the number of vaxxed women by the ratio

WRS (float) : the approximate number of actual women experiencing reproductive symptoms
min_WRS (float) : $WRS * 0.80$ max_WRS (float) : $WRS * 1.20$

```
[97]: WRS = women_vaxx * WRS_ratio
      min_WRS = WRS*0.80
      max_WRS = WRS*1.20
```

```
[98]: cl = [35,5,15]
      print_row(['total vaxxed (1 or more)', '---', '{:,.2f}'.
        ↪format(vaxx)],column_lengths=cl)
      print_row(['women vaxxed ~0.63%', '---', '{:,.2f}'.
        ↪format(women_vaxx)],column_lengths=cl)
      print_row(['repro sympt / women count', '---', '{:,.4f}'.
        ↪format(WRS_ratio)],column_lengths=cl)
      print_row(['(repro sympt / women count) * 100', '---', '{:,.2f} %'.
        ↪format(WRS_ratio*100)],column_lengths=cl)
      print_row(['women w/ repro symptoms', '---', '{:,.2f}'.
        ↪format(WRS)],column_lengths=cl)
      print_row(['min women w/ repro symptoms', '---', '{:,.2f}'.
        ↪format(min_WRS)],column_lengths=cl)
      print_row(['max women w/ repro symptoms', '---', '{:,.2f}'.
        ↪format(max_WRS)],column_lengths=cl)
```

total vaxxed (1 or more)	---	500,864,529.00
women vaxxed ~0.63%	---	315,544,653.27
repro sympt / women count	---	0.0335
(repro sympt / women count) * 100	---	3.35 %
women w/ repro symptoms	---	10,575,525.70
min women w/ repro symptoms	---	8,460,420.56
max women w/ repro symptoms	---	12,690,630.84

4.11.4 Where do the WRS (Women reproductive symptoms) rank against with the other symptoms?

```
[99]: temp = []

      for index,i in enumerate(all_symptoms):
          if i[0].upper() in WRS_list:
              temp.append(
                  {
                      'index':index,
```

```

        'symptoms':i[0],
        'count': '{:,.2f}'.format(i[1]),
        'percent': '{:,.2f}'.format((i[1]/len(all_symptoms))*100)
    }
)

temp = pd.DataFrame(temp)
display(temp)

```

	index	symptoms	count	percent
0	133	HEAVY MENSTRUAL BLEEDING	5,702.00	46.80
1	172	MENSTRUATION IRREGULAR	4,346.00	35.67
2	173	HOT FLUSH	4,333.00	35.56
3	198	MENSTRUAL DISORDER	3,285.00	26.96
4	248	DYSMENORRHOEA	2,399.00	19.69
5	251	BREAST PAIN	2,335.00	19.16
6	279	VAGINAL HAEMORRHAGE	1,996.00	16.38
7	310	INTERMENSTRUAL BLEEDING	1,688.00	13.85
8	480	BREAST SWELLING	910.00	7.47
9	497	POLYMENORRHOEA	855.00	7.02
10	539	OLIGOMENORRHOEA	769.00	6.31
11	598	BREAST TENDERNESS	648.00	5.32
12	782	MATERNAL EXPOSURE BEFORE PREGNANCY	428.00	3.51
13	1272	OVARIAN CYST	194.00	1.59
14	1286	MENOPAUSE	192.00	1.58
15	1461	UTERINE LEIOMYOMA	152.00	1.25
16	1674	ABNORMAL UTERINE BLEEDING	118.00	0.97
17	2987	BACTERIAL VAGINOSIS	36.00	0.30
18	3149	ANOVULATORY CYCLE	32.00	0.26
19	3374	BIOPSY UTERUS	27.00	0.22
20	3667	OVULATION DELAYED	22.00	0.18
21	5243	GYNAECOLOGICAL EXAMINATION ABNORMAL	9.00	0.07

[]:

[]: