

Ingesting Data

We are going to Ingest Data directly from Kaggle

```
# IMPORTANT: RUN THIS CELL IN ORDER TO IMPORT YOUR KAGGLE DATA SOURCES
# TO THE CORRECT LOCATION (/kaggle/input) IN YOUR NOTEBOOK,
# THEN FEEL FREE TO DELETE THIS CELL.
# NOTE: THIS NOTEBOOK ENVIRONMENT DIFFERS FROM KAGGLE'S PYTHON
# ENVIRONMENT SO THERE MAY BE MISSING LIBRARIES USED BY YOUR
# NOTFBOOK.
import os
import sys
from tempfile import NamedTemporaryFile
from urllib.request import urlopen
from urllib.parse import unquote, urlparse
from urllib.error import HTTPError
from zipfile import ZipFile
import tarfile
import shutil
CHUNK STZF = 40960
\textbf{DATA\_SOURCE\_MAPPING} = \text{'widsdatathon2024-challenge2:https%3A%2F%2Fstorage.googleapis.com%2Fkaggle-competitions-data%2Fkaggle-v2%2F73478%2F812'} \\
KAGGLE_INPUT_PATH='/kaggle/input'
KAGGLE_WORKING_PATH='/kaggle/working'
KAGGLE_SYMLINK='kaggle'
!umount /kaggle/input/ 2> /dev/null
shutil.rmtree('/kaggle/input', ignore_errors=True)
os.makedirs(KAGGLE_INPUT_PATH, 0o777, exist_ok=True)
os.makedirs(KAGGLE_WORKING_PATH, 0o777, exist_ok=True)
try:
  os.symlink(KAGGLE_INPUT_PATH, os.path.join("..", 'input'), target_is_directory=True)
except FileExistsError:
 pass
try:
 os.symlink(KAGGLE_WORKING_PATH, os.path.join("..", 'working'), target_is_directory=True)
except FileExistsError:
 pass
for data_source_mapping in DATA_SOURCE_MAPPING.split(','):
    directory, download_url_encoded = data_source_mapping.split(':')
    download_url = unquote(download_url_encoded)
    filename = urlparse(download_url).path
    destination_path = os.path.join(KAGGLE_INPUT_PATH, directory)
    try:
        with \ urlopen(download\_url) \ as \ fileres, \ NamedTemporaryFile() \ as \ tfile:
            total_length = fileres.headers['content-length']
            print(f'Downloading {directory}, {total_length} bytes compressed')
            dl = 0
            data = fileres.read(CHUNK_SIZE)
            while len(data) > 0:
                dl += len(data)
                tfile.write(data)
                done = int(50 * dl / int(total_length))
                sys.stdout.write(f"\r[{'=' * done}{{' ' * (50-done)}}] {dl} bytes downloaded")
                sys.stdout.flush()
                data = fileres.read(CHUNK_SIZE)
            if filename.endswith('.zip'):
              with ZipFile(tfile) as zfile:
                zfile.extractall(destination_path)
            else:
              with tarfile.open(tfile.name) as tarfile:
                tarfile.extractall(destination_path)
            print(f'\nDownloaded and uncompressed: {directory}')
    except HTTPError as e:
        print(f'Failed to load (likely expired) {download_url} to path {destination_path}')
        continue
    except OSError as e:
        print(f'Failed to load {download_url} to path {destination_path}')
        continue
print('Data source import complete.')
Downloading widsdatathon2024-challenge2, 5985935 bytes compressed
[======] 5985935 bytes downloaded
Downloaded and uncompressed: widsdatathon2024-challenge2
Data source import complete.
```

```
# This Python 3 environment comes with many helpful analytics libraries installed
# It is defined by the kaggle/python Docker image: https://github.com/kaggle/docker-python
# For example, here's several helpful packages to load
import numpy as np # linear algebra
{\color{red} \textbf{import pandas as pd}} \ \textit{\# data processing, CSV file I/O (e.g. pd.read\_csv)}
# Input data files are available in the read-only "../input/" directory
# For example, running this (by clicking run or pressing Shift+Enter) will list all files under the input directory
import os
for dirname, _, filenames in os.walk('/kaggle/input'):
    for filename in filenames:
        print(os.path.join(dirname, filename))
# You can write up to 20GB to the current directory (/kaggle/working/) that gets preserved as output when you create a version using "Save &
# You can also write temporary files to /kaggle/temp/, but they won't be saved outside of the current session
/kaggle/input/widsdatathon2024-challenge2/solution_template.csv
/kaggle/input/widsdatathon2024-challenge2/test.csv
/kaggle/input/widsdatathon2024-challenge2/train.csv
```

```
train_df=pd.read_csv('/kaggle/input/widsdatathon2024-challenge2/train.csv')
test_df=pd.read_csv('/kaggle/input/widsdatathon2024-challenge2/test.csv')
```

Exploring Data

```
#printing rows and columns
train_df.shape

(13173, 152)
```

	patient_id int64	patient_race object	payer_type object	patient_state object	patient_zip3 int64	Region object	Division object	patient_age int64	pati
0	268700	nan	COMMERCIAL	AR	724	South	West South Central	39	F
1	484983	White	nan	IL	629	Midwest	East North Central	55	F
2	277055	nan	COMMERCIAL	CA	925	West	Pacific	59	F
3	320055	Hispanic	MEDICAID	CA	900	West	Pacific	59	F
4	190386	nan	COMMERCIAL	CA	934	West	Pacific	71	F
4	190386	nan	COMMERCIAL	CA	934	West	Pacific		71

```
### No of patients
train_df['patient_id'].nunique()

13173
```

Asi Black His Oth Wh 5 rows, show	etastatic_diagno 96.6986711 106.7469636 104.404416 90.4664723 93.38932147 96.74847561	rpage
Asi Black His Oth Wh 5 rows, show ## crea train_d: me Eas Eas Mid Mo Pac	98.19571046 98.01035375 82.42379182 98.4869281 91.68415147 wing 10	ple for average diagnosis period for each Division
His Oth Wh 5 rows, show ## crea train_d me Eas Eas Mid Mo Pac	82.42379182 98.4869281 91.68415147 wing 10	ple for average diagnosis period for each Division
Oth Wh 5 rows, show ## crea train_d: me Eas Eas Mid Mo Pac	98.4869281 91.68415147 wing 10 pe ating pivot tab If .pivot_table(etastatic_diagno 96.6986711 106.7469636 104.404416 90.4664723 93.38932147 96.74847561	ple for average diagnosis period for each Division
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## creatrain_d: me Eas Eas Mid Mo Pac	ating pivot tall If.pivot_table(etastatic_diagno 96.6986711 106.7469636 104.404416 90.4664723 93.38932147 96.74847561	ple for average diagnosis period for each Division
## crea train_d: me Eas Eas Mid Mo Pac	ating pivot tal If.pivot_table(etastatic_diagno 96.6986711 106.7469636 104.404416 90.4664723 93.38932147 96.74847561	ple for average diagnosis period for each Division
train_d me Eas Eas Mid Mo Pac	Hf.pivot_table(etastatic_diagno 96.6986711 106.7469636 104.404416 90.4664723 93.38932147 96.74847561	
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Eas Mid Mo Pac	106.7469636 104.404416 90.4664723 93.38932147 96.74847561	
Mid Mo Pac	104.404416 90.4664723 93.38932147 96.74847561	
Mo Pac	90.4664723 93.38932147 96.74847561	
Pac	93.38932147 96.74847561	
	96.74847561	
So		
We	93.72089947	
We	94.59546061	
		ole for average diagnosis period for each Region index=['Region','Division'],values='metastatic_diagnosis_period',aggfunc='mean')
me	etastatic_diagno	
('Mi	96.6986711	
('Mi	93.72089947	
('N	104.404416	
('S	106.7469636	
('S	96.74847561	
('S	94.59546061	
('W	90.4664723	
('W	93.38932147	
8 rows, show	wing 10 v pe	r page
train_d		ple for average diagnosis period for each type of payer [index='payer_type',values='metastatic_diagnosis_period',aggfunc='mean')
CO	102.5312053	
ME	93.6278442	
ME	98.63621922	
3 rows, show	wing 10 v pe	er page

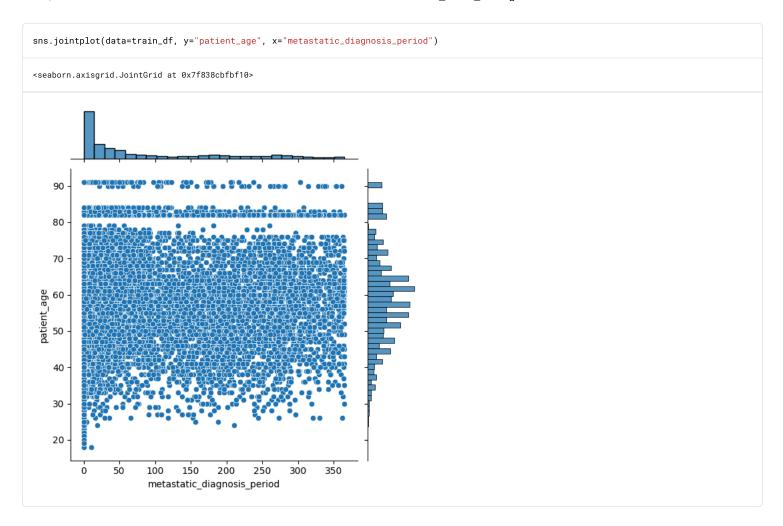


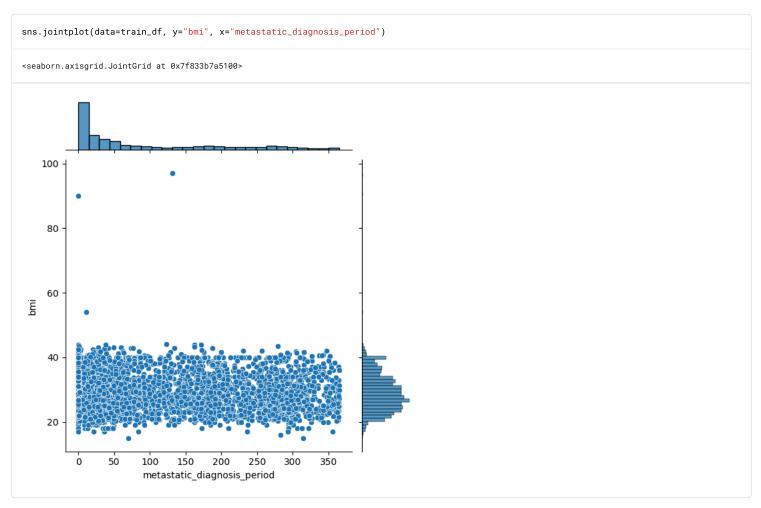
Prelimnary insights :-

- Hispanic race patients have lowest diagnosis time
- Patients in northeast and southeastern states have higher diagnosis time that western states.
- Patients whose payment type is medicaid or medicare advantage have less diagnosis time.

Generating plots for numerical columns like age and BMI of patients

```
import matplotlib.pyplot as plt
import seaborn as sns
fig, axes = plt.subplots(1, 2, figsize=(15, 5))
sns.scatterplot(data=train\_df, \ y="patient\_age", \ x="metastatic\_diagnosis\_period", \ ax=axes[\emptyset])
sns.scatterplot(data=train_df, y="bmi", x="metastatic_diagnosis_period", ax=axes[1])
plt.show()
                                                                                   100
   90
   80
                                                                                    80
   70
patient_age
% 9
                                                                                    60
                                                                                 bmi
    40
                                                                                    40
   30
                                                                                    20
   20
                 50
                         100
                                  150
                                          200
                                                  250
                                                           300
                                                                   350
                                                                                                  50
                                                                                                                  150
                                                                                                                          200
                                                                                                                                   250
                                                                                                                                            300
                                                                                                                                                    350
                            metastatic_diagnosis_period
                                                                                                             metastatic_diagnosis_period
```





Creating New Features

New features can be created i.e

- Age group
- BMI group
- Patient age X Bmi
- removes letter and keeps numbers from 'breast_cancer_diagnosis_code'

```
train_df['patient_age'].isna().sum()
0
```

```
##define age group
def age_group(x):
    if x<=20 and x>0:
        tag='Age 0-20'
    elif x>20 and x<=40:
        tag= 'Age 20-40'
    elif x>40 and x<=60:
        tag= 'Age 40-60'
    elif x>60 and x<=80:
        tag='Age 60-80'
    else:
        tag= 'Age >80'
    return tag
```

```
train_df['Age_group']=train_df['patient_age'].apply(age_group)
```

97.89440338

102.4603922

90.1641563 89.6738437

5 rows, showing 10 v per page

Ag...

Ag...

Ag...

```
train_df['Age_group'].value_counts()
Age_group
             6375
Age 40-60
Age 60-80
             4581
Age >80
             1254
Age 20-40
              947
Age 0-20
              16
Name: count, dtype: int64
train_df.pivot_table(index='Age_group', values='metastatic_diagnosis_period', aggfunc='mean')
      metastatic_diagno...
Ag...
                0.6875
```

train_df['bmi'].isna().sum()
9071

of 1 > >>

Page 1

```
from sklearn.impute import SimpleImputer
imputer=SimpleImputer(strategy='mean')
train_df['bmi_new']=imputer.fit_transform(train_df[['bmi']])
```

	n_df.head() patient_id int64	patient_race object	payer_type object	patient_state object	patient_zip3 int64	Region object	Division object	patient_age int64	patie
	patient_id into4	patient_race object	payer_type object	patient_state object	patient_zip3 into4	Region object	Division object	patient_age into4	patie
0	268700	nan	COMMERCIAL	AR	724	South	West South Central	39	F
1	484983	White	nan	IL	629	Midwest	East North Central	55	F
2	277055	nan	COMMERCIAL	CA	925	West	Pacific	59	F
3	320055	Hispanic	MEDICAID	CA	900	West	Pacific	59	F
4	190386	nan	COMMERCIAL	CA	934	West	Pacific	71	F
←									•
5 rows, showing 10 v per page « < Page 1 of 1 > >>							+		

```
##define bmi group

def bmi_group(x):
    if x<=20:
        tag='BMI 0-20'
    elif x>20 and x<=30:
        tag= 'BMI 20-30'
    elif x>30 and x<=40:
        tag= 'BMI 30-40'
    else:
        tag='BMI >40'
    return tag
```

```
train_df['Bmi_group']=train_df['bmi_new'].apply(bmi_group)
```

 $\underline{\downarrow}$

```
fig, axes = plt.subplots(1, 2, figsize=(15, 5))
sns.barplot(data=train\_df, \ x="Age\_group", \ y="metastatic\_diagnosis\_period", \ ax=axes[\theta])
sns.barplot(data=train\_df, \ x="Bmi\_group", \ y="metastatic\_diagnosis\_period", \ ax=axes[1])
plt.show()
                                                                                              140
    100
                                                                                              120
 metastatic diagnosis period
     80
                                                                                           metastatic_diagnosis_period
                                                                                              100
                                                                                               80
     60
                                                                                               60
     40
                                                                                               40
     20
                                                                                               20
       0
                                                                                                0
           Age 20-40
                           Age 40-60
                                         Age 60-80
                                                                        Age 0-20
                                                                                                       BMI 20-30
                                                         Age >80
                                                                                                                          BMI 30-40
                                                                                                                                             BMI 0-20
                                                                                                                                                                BMI >40
                                         Age_group
                                                                                                                                   Bmi_group
```



```
train_df.drop(['patient_age','bmi'],axis=1,inplace=True)
```

Handling Missing values

```
train_df.shape
(13173, 153)
```

```
train_df.iloc[:,0:8].info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 13173 entries, 0 to 13172
Data columns (total 8 columns):
# Column
                  Non-Null Count Dtype
                   -----
0
   patient_id
                  13173 non-null int64
    patient_race
                  6516 non-null object
1
                   11408 non-null object
    payer_type
3
    patient_state 13173 non-null object
                  13173 non-null int64
4
    patient_zip3
                   13173 non-null object
5
    Region
6
    Division
                  13173 non-null object
    patient_gender 13173 non-null object
dtypes: int64(2), object(6)
memory usage: 823.4+ KB
```

```
pd.concat([train_df.iloc[:,0:8],train_df.iloc[:,79:149]],axis=1).isna().sum()
patient_id
                       0
                    6657
patient_race
payer_type
                    1765
patient_state
                       0
                       0
patient_zip3
Average of Aug-18
                      16
Average of Sep-18
Average of Oct-18
                      7
Average of Nov-18
                      12
Average of Dec-18
                      33
Length: 78, dtype: int64
```

```
train_df_cat=train_df.iloc[:,[1,2]]
```

```
from sklearn.impute import SimpleImputer
imputer=SimpleImputer(strategy='most_frequent')
train_df_cat=pd.DataFrame(imputer.fit_transform(train_df_cat),columns=['Patient_race_new','Payer_type_new'])
```

```
train_df.drop(['patient_race','payer_type'],axis=1,inplace=True)
```

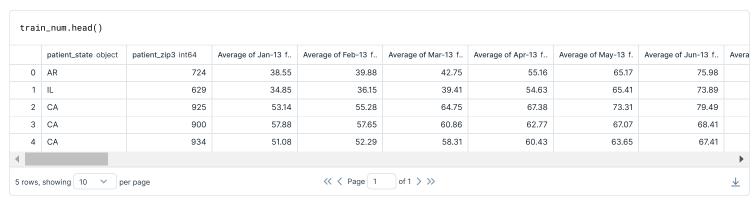
```
train_df.drop(['bmi_new','patient_gender'],axis=1,inplace=True)
```

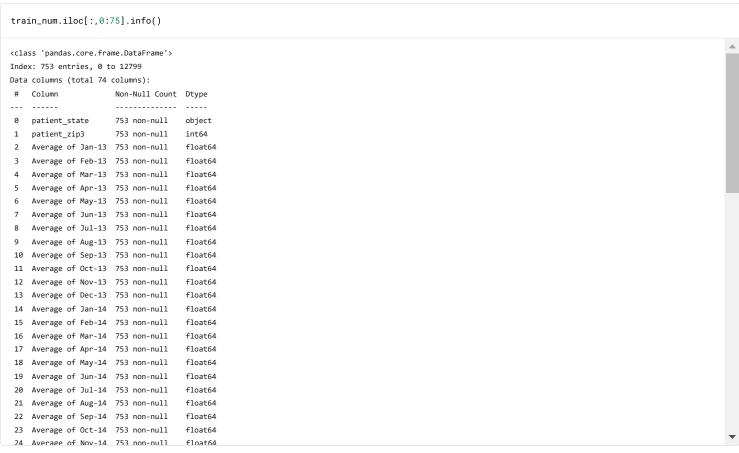
```
train_df=pd.concat([train_df,train_df_cat],axis=1)
```

```
train_df.iloc[:,0:20].head()
                                                                                                 Division object
        patient_id int64
                              patient_state object
                                                    patient_zip3 int64
                                                                           Region object
                                                                                                                       breast_cancer_dia...
                                                                                                                                             breast_cancer_dia...
                                                                                                                                                                   metastatic_cancer...
                                                                                                                                                                                          metas
                                                                                                 West South Central
                                                                                                                       C50912
   0
                    268700
                              AR
                                                                    724
                                                                          South
                                                                                                                                              Malignant neopla...
                                                                                                                                                                   C773
                                                                                                                                                                                          nan
    1
                   484983
                              \mathsf{IL}
                                                                          Midwest
                                                                                                 East North Central
                                                                                                                       C50412
                                                                                                                                              Malig neoplasm o...
                                                                                                                                                                   C773
                                                                    629
                                                                                                                                                                                          nan
   2
                                                                                                 Pacific
                                                                                                                       1749
                    277055
                              CA
                                                                    925
                                                                          West
                                                                                                                                              Malignant neopla...
                                                                                                                                                                   C773
                                                                                                                                                                                          nan
   3
                                                                                                 Pacific
                                                                                                                       C50911
                    320055
                              CA
                                                                    900
                                                                          West
                                                                                                                                              Malignant neopla...
                                                                                                                                                                   C773
                                                                                                                                                                                          nan
   4
                    190386
                              CA
                                                                                                 Pacific
                                                                                                                       1748
                                                                                                                                                                   C7951
                                                                    934
                                                                          West
                                                                                                                                             Malignant neopla...
                                                                                                                                                                                          nan
5 rows, showing 10
                                                                                << < Page 1</pre>
                                                                                                   of 1 > >>
                                                                                                                                                                                          \underline{\downarrow}
```

	patient_id int64	patient_state object	patient_zip3 int64	Region object	Division object	breast_cancer_dia	breast_cancer_dia	metastatic_cancer	meta
0	268700	AR	724	South	West South Central	C50912	Malignant neopla	C773	nan
1	484983	IL	629	Midwest	East North Central	C50412	Malig neoplasm o	C773	nan
2	277055	CA	925	West	Pacific	1749	Malignant neopla	C773	nan
3	320055	CA	900	West	Pacific	C50911	Malignant neopla	C773	nan
4	190386	CA	934	West	Pacific	1748	Malignant neopla	C7951	nan
									•

train_df_new=train_df[['patient_id','patient_state','patient_zip3','Region','Division','Age_group','Bmi_group','Patient_race_new','Payer_typ train_df_new.shape (13173, 10) # prompt: export to csv file #train_df_new.to_csv('/content/patient_data.csv', index=False) train_num=train_df.iloc[:,74:146] imputer=SimpleImputer(strategy='mean') train_num=pd.DataFrame(imputer.fit_transform(train_num),columns=train_num.columns.to_list()) train_num.head() Average of Jan-13 f.. Average of Feb-13 f.. Average of Mar-13 f.. Average of Apr-13 f.. Average of May-13 f. Average of Jun-13 f.. Average of Jul-13 f... Average of Aug-13 f... Avera 0 38.55 39.88 75.98 42 75 5516 65 17 76.75 76 45 1 34.85 36.15 39.41 54.63 65.41 73.89 74.07 74.37 2 53.14 55.28 64.75 67.38 73.31 79.49 84.01 83.28 3 57.88 57.65 60.86 62.77 67.07 68.41 70.69 71.19 4 51.08 52.29 58.31 60.43 63.65 67.41 67.95 << < Page 1</p> of 1 > >> $\underline{\downarrow}$ 5 rows, showing 10 per page train_num=pd.concat([train_df.iloc[:,[1,2]],train_num],axis=1) train_num.head() patient state object patient zip3 int64 Average of Jan-13 f... Average of Feb-13 f... Average of Mar-13 f... Average of Apr-13 f... Average of May-13 f. Average of Jun-13 f... Avera 0 AR 724 38.55 39.88 42.75 55.16 65.17 75.98 1 IL 629 34.85 36.15 39.41 54.63 65.41 73.89 2 CA 925 53.14 55.28 64.75 67.38 73.31 79.49 3 CA 57.88 57.65 60.86 900 62.77 67.07 68.41 4 CA 51.08 58.31 60.43 63.65 67.41 52.29 << < Page 1</pre> of 1 > >> $\underline{\downarrow}$ 5 rows, showing 10 per page train_num.drop_duplicates(inplace=True)





```
min_value=train_num.iloc[:,2:75].min(axis=1)

max_value=train_num.iloc[:,2:75].max(axis=1)

avg_value=train_num.iloc[:,2:75].mean(axis=1)

train_num['min_temp']=min_value
```

```
train_num['avg_temp']=avg_value
```

train_num['max_temp']=max_value

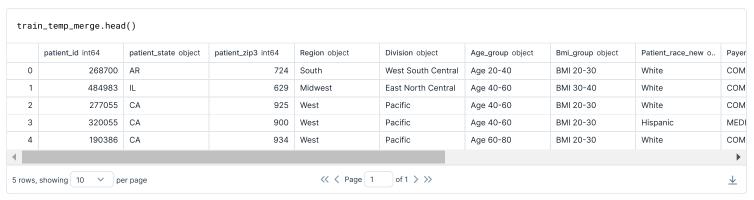




train_num=train_num.iloc[:,[0,1,74,75,76,77]]



train_temp_merge=pd.merge(train_df_new,train_num,how='left',on=['patient_state','patient_zip3'])



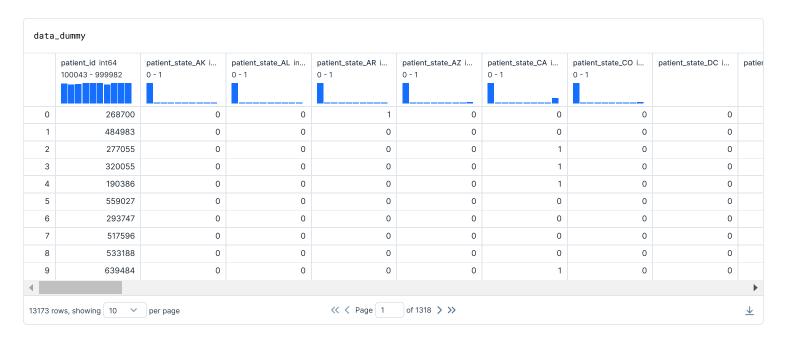
```
train_temp_merge.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 13173 entries, 0 to 13172
Data columns (total 14 columns):
    Column
#
                             Non-Null Count Dtype
                             -----
                           13173 non-null int64
0
    patient_id
                           13173 non-null object
1
    patient_state
                          13173 non-null int64
2
   patient_zip3
                           13173 non-null object
3 Region
4 Division
                           13173 non-null object
5 Age_group
                           13173 non-null object
6 Bmi_group
                           13173 non-null object
7 Patient_race_new
                           13173 non-null object
                            13173 non-null object
8 Payer_type_new
9
    metastatic_diagnosis_period 13173 non-null int64
                             13173 non-null float64
10 min_temp
11 max_temp
                             13173 non-null float64
12 avg_temp
                            13173 non-null float64
13 temp_diff
                            13173 non-null float64
dtypes: float64(4), int64(3), object(7)
memory usage: 1.4+ MB
```

```
train_temp_merge.drop('patient_zip3',axis=1,inplace=True)
```

Creating Dummy variables

```
data_dummy=train_temp_merge.iloc[:,[0,1,2,3,4,5,6,7]]
```

```
data_dummy=pd.get_dummies(data_dummy,dtype=int)
```



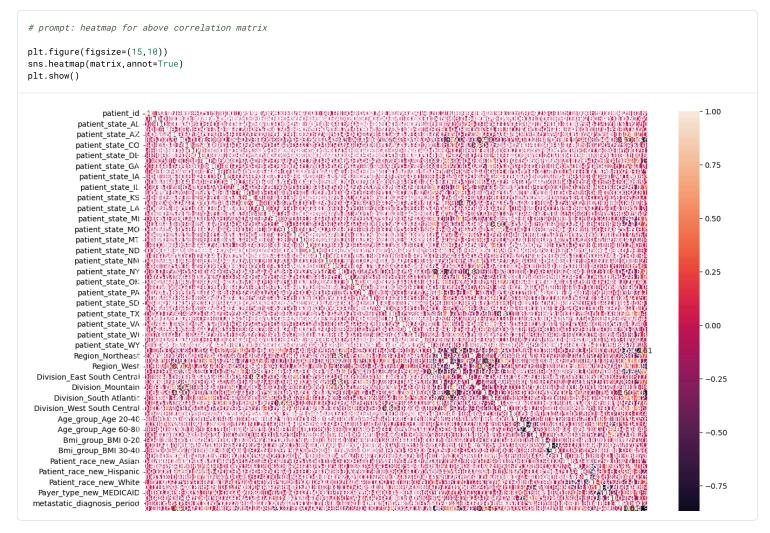
train_temp_merge=pd.concat([data_dummy,train_temp_merge.iloc[:,[8,9,10,11,12]]],axis=1)

```
train_temp_merge.info()
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 13173 entries, 0 to 13172
Data columns (total 79 columns):
    Column
                                     Non-Null Count Dtype
                                      -----
0
    patient_id
                                     13173 non-null int64
1
    patient_state_AK
                                     13173 non-null int64
2
    patient_state_AL
                                     13173 non-null int64
3
    patient_state_AR
                                     13173 non-null int64
4
    patient_state_AZ
                                     13173 non-null int64
5
    patient_state_CA
                                     13173 non-null int64
6
    patient_state_CO
                                     13173 non-null int64
    patient_state_DC
7
                                     13173 non-null int64
    patient_state_DE
                                     13173 non-null int64
8
    patient_state_FL
                                     13173 non-null int64
10 patient_state_GA
                                     13173 non-null int64
11 patient_state_HI
                                     13173 non-null int64
12 patient_state_IA
                                     13173 non-null int64
13 patient state ID
                                     13173 non-null int64
14 patient_state_IL
                                     13173 non-null
                                                    int64
                                      13173 non-null
15
    patient_state_IN
    patient_state_KS
                                     13173 non-null
17 patient_state_KY
                                     13173 non-null int64
18 patient state LA
                                     13173 non-null int64
19 patient_state_MD
                                     13173 non-null int64
20 patient_state_MI
                                     13173 non-null int64
21 patient_state_MN
                                      13173 non-null int64
22
    patient_state_MO
                                     13173 non-null int64
                                     13173 non-null int64
23 patient_state_MS
24 natient state MT
                                      13173 non-null int64
```

Correlation matrix

```
matrix=train_temp_merge.corr()

m=matrix['metastatic_diagnosis_period']
```



Selecting the features based on correlation values

```
# prompt: sort the series in decending
sorted_m = m.sort_values(ascending=False)
print(sorted_m[1:20])
Age_group_Age 40-60
                                     0.052835
patient_state_NY
                                     0.039525
                                     0.028067
Region_Northeast
Division_Middle Atlantic
                                     0.028067
patient_state_KY
                                     0.027646
temp_diff
                                     0.025150
                                     0.024021
patient state ID
Division_East South Central
                                     0.018534
Bmi_group_BMI 30-40
                                     0.018420
patient_state_NM
                                     0.015971
patient_state_GA
                                     0.014711
                                     0.013616
max_temp
                                     0.013042
Patient_race_new_White
                                     0.012243
patient_state_AZ
patient_state_MN
                                     0.011989
patient_state_NV
                                     0.010530
patient_state_IL
                                     0.009754
Bmi_group_BMI 0-20
                                     0.009609
Payer_type_new_MEDICARE ADVANTAGE
                                     0.009462
Name: metastatic_diagnosis_period, dtype: float64
```

```
X=train_temp_merge.set_index('patient_id')
Y=X['metastatic_diagnosis_period']
```

```
columns=sorted_m[1:20].index.to_list()
```

```
X=X[columns]
```

```
Y.head()

patient_id
268700    191
484983    33
277055    157
320055    146
190386    286
Name: metastatic_diagnosis_period, dtype: int64
```

Splitting the values in test and train datasets

```
# Split into training and test set
from sklearn.model_selection import train_test_split
X_train, X_val, y_train, y_val = train_test_split(X, Y, test_size=0.2)
```

```
print(X_train.shape)
print(X_val.shape)
print(y_train.shape)
print(y_val.shape)

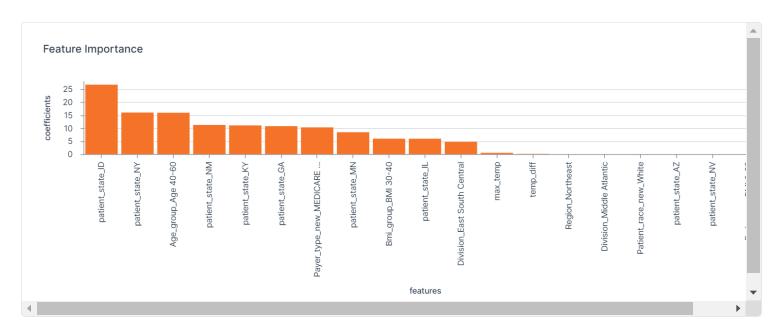
(10538, 19)
(2635, 19)
(10538,)
(2635,)
```

Model Training with linear regression

```
from sklearn.linear_model import Lasso
from sklearn.metrics import mean_squared_error
alpha = 0.1 # Regularization strength
lasso_reg = Lasso(alpha=alpha)
lasso_reg.fit(X_train, y_train)
print(X_train.columns.to_list())
print(lasso_reg.coef_)
y_pred = lasso_reg.predict(X_val)
# Calculate Mean Squared Error
mse = mean_squared_error(y_val, y_pred)
print("Mean Squared Error:", mse)
print()
['Age_group_Age 40-60', 'patient_state_NY', 'Region_Northeast', 'Division_Middle Atlantic', 'patient_state_KY', 'temp_diff', 'patient_state_ID', 'Division_East Sout
[15.91443646 15.98695251 -0. -0.
                                         11.03873633 0.10962569
26.7035212 4.76151777 5.94147095 11.22007212 10.75830615 0.50365612
 0.
                    8.42428329 0.
                                          5.92902273 0.
10.28892926]
Mean Squared Error: 11659.600993144202
```

```
cols=(X_train.columns.to_list())
coefficients=lasso_reg.coef_
key_value_pairs = pd.DataFrame({'features':cols,'coefficients':coefficients})
```

```
print(key_value_pairs)
                          features coefficients
0
                Age_group_Age 40-60
                                    15.914436
1
                   patient_state_NY
                                     15.986953
                                    -0.000000
2
                   Region Northeast
3
           Division Middle Atlantic
                                   -0.000000
                  patient_state_KY 11.038736
5
                         temp_diff 0.109626
                  patient_state_ID 26.703521
7
                                    4.761518
        Division_East South Central
8
               Bmi_group_BMI 30-40
                                     5.941471
9
                  patient state NM
                                     11.220072
10
                                     10.758306
                  patient_state_GA
11
                                       0.503656
                         max_temp
12
             Patient_race_new_White
                                       0.000000
13
                  patient_state_AZ
                                       9.999999
                                       8.424283
14
                  patient state MN
15
                  patient_state_NV
                                       0.000000
16
                  patient_state_IL
                                       5.929023
17
                 Bmi_group_BMI 0-20
                                       0.000000
18 Payer_type_new_MEDICARE ADVANTAGE
                                      10.288929
```



print('Root Mean Squared Error',np.sqrt(mse))

Root Mean Squared Error 107.97963230695038