

```
In [ ]: import pandas as pd
import numpy as np
import datetime as dt
import matplotlib.pyplot as plt
import seaborn as sns
import missingno as msno
pd.options.display.max_columns = None
```

LA GRAN PANDEMIA

contenido del data set COVID

The dataset was provided by the Mexican government (link). This dataset contains an enormous number of anonymized patient-related information including pre-conditions. The raw dataset consists of 21 unique features and 1,048,576 unique patients. In the Boolean features, 1 means "yes" and 2 means "no". values as 97 and 99 are missing data.

- sex: 1 for female and 2 for male.
- age: of the patient.
- classification: covid test findings. Values 1-3 mean that the patient was diagnosed with covid in different
- degrees. 4 or higher means that the patient is not a carrier of covid or that the test is inconclusive.
- patient type: type of care the patient received in the unit. 1 for returned home and 2 for hospitalization.
- pneumonia: whether the patient already have air sacs inflammation or not.
- pregnancy: whether the patient is pregnant or not.
- diabetes: whether the patient has diabetes or not.
- copd: Indicates whether the patient has Chronic obstructive pulmonary disease or not.
- asthma: whether the patient has asthma or not.
- inmsupr: whether the patient is immunosuppressed or not.
- hypertension: whether the patient has hypertension or not.
- cardiovascular: whether the patient has heart or blood vessels related disease.
- renal chronic: whether the patient has chronic renal disease or not.
- other disease: whether the patient has other disease or not.
- obesity: whether the patient is obese or not.
- tobacco: whether the patient is a tobacco user.
- usmr: Indicates whether the patient treated medical units of the first, second or third level.
- medical unit: type of institution of the National Health System that provided the care.
- intubed: whether the patient was connected to the ventilator.
- icu: Indicates whether the patient had been admitted to an Intensive Care Unit.
- date died: If the patient died indicate the date of death, and 9999-99-99 otherwise.

Limpieza de datos

Realizaremos una exploracion de los datos para buscar datos faltantes, duplicados y en su caso extremos.

```
In [ ]: df_covid = pd.read_csv('../Covid Data.csv')
In [ ]: df_covid.head()
```

Out[]:

	USMER	MEDICAL_UNIT	SEX	PATIENT_TYPE	DATE_DIED	INTUBED	PNEUMONIA	AGE	PREGNANT	DIABETES	COPD	ASTHMA	INMSUPR	HIPERTENSION	OTHER_DISEASE
0	2		1	1	1 03/05/2020	97	1	65	2	2	2	2	2	1	2
1	2		1	2	1 03/06/2020	97	1	72	97	2	2	2	2	1	2
2	2		1	2	2 09/06/2020	1	2	55	97	1	2	2	2	2	2
3	2		1	1	1 12/06/2020	97	2	53	2	2	2	2	2	2	2
4	2		1	2	1 21/06/2020	97	2	68	97	1	2	2	2	1	2

```
In [ ]: df_covid.shape
Out [ ]: (1048575, 21)
```

```
In [ ]: df_covid.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1048575 entries, 0 to 1048574
Data columns (total 21 columns):
#   Column                Non-Null Count  Dtype
---  -
0   USMER                  1048575 non-null  int64
1   MEDICAL_UNIT           1048575 non-null  int64
2   SEX                    1048575 non-null  int64
3   PATIENT_TYPE           1048575 non-null  int64
4   DATE_DIED              1048575 non-null  object
5   INTUBED                1048575 non-null  int64
6   PNEUMONIA              1048575 non-null  int64
7   AGE                    1048575 non-null  int64
8   PREGNANT               1048575 non-null  int64
9   DIABETES               1048575 non-null  int64
10  COPD                   1048575 non-null  int64
11  ASTHMA                 1048575 non-null  int64
12  INMSUPR                1048575 non-null  int64
13  HIPERTENSION           1048575 non-null  int64
14  OTHER_DISEASE          1048575 non-null  int64
15  CARDIOVASCULAR         1048575 non-null  int64
16  OBESITY                 1048575 non-null  int64
17  RENAL_CHRONIC          1048575 non-null  int64
18  TOBACCO                1048575 non-null  int64
19  CLASIFICATION_FINAL    1048575 non-null  int64
20  ICU                    1048575 non-null  int64
dtypes: int64(20), object(1)
memory usage: 168.0+ MB
```

Buscamos datos nulos

En nuestro dataset los valores 97 y 99 se consideran como data faltante. Por lo tanto, los reemplazaremos por NaN y luego se quitaran.

```
In [ ]: # reemplazamos todos Los 97 y 99 por NaN

df_covid.replace(97, np.nan, inplace=True)
df_covid.replace(98, np.nan, inplace=True)
df_covid.replace(99, np.nan, inplace=True)

df_covid.isnull().sum()
```

```
Out[ ]: USMER                0
MEDICAL_UNIT            0
SEX                     0
PATIENT_TYPE            0
DATE_DIED               0
INTUBED                855869
PNEUMONIA              16003
AGE                    345
PREGNANT               527265
DIABETES               3338
COPD                   3003
ASTHMA                 2979
INMSUPR               3404
HIPERTENSION          3104
OTHER_DISEASE         5045
CARDIOVASCULAR        3076
OBESITY               3032
RENAL_CHRONIC         3006
TOBACCO               3220
CLASIFFICATION_FINAL   0
ICU                   856032
dtype: int64
```

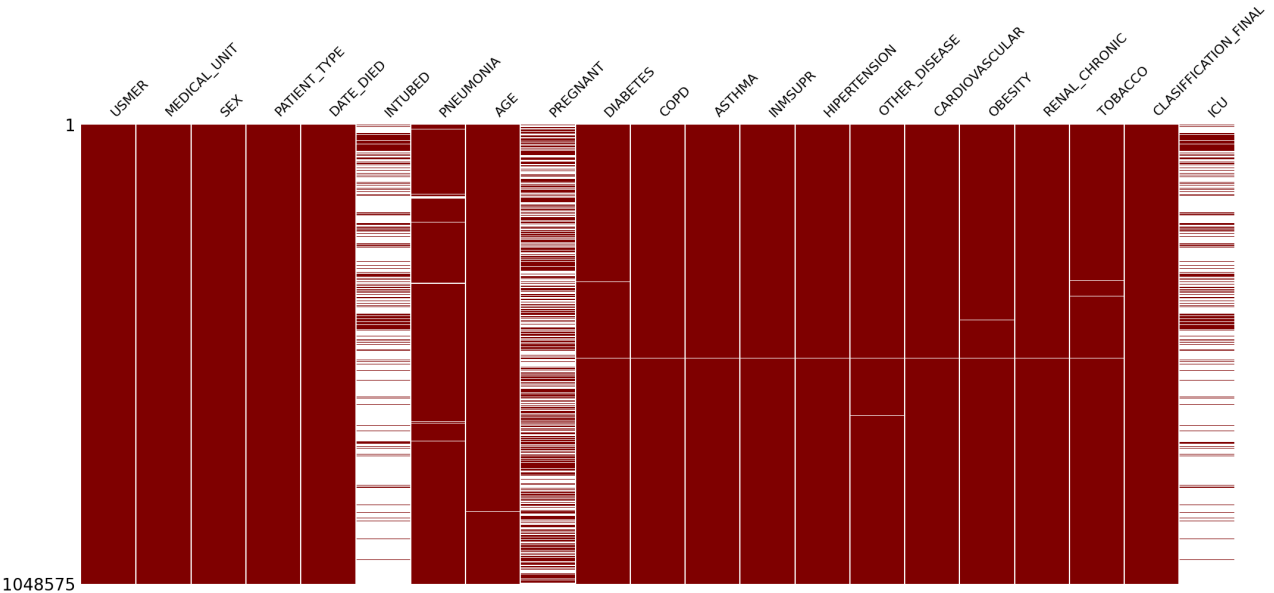
```
In [ ]: #porcentaje de datos faltantes por columna

df_covid.isnull().sum()/df_covid.shape[0]
```

```
Out[ ]: USMER                0.000000
MEDICAL_UNIT            0.000000
SEX                     0.000000
PATIENT_TYPE            0.000000
DATE_DIED               0.000000
INTUBED                0.816221
PNEUMONIA              0.015262
AGE                    0.000329
PREGNANT               0.502840
DIABETES               0.003183
COPD                   0.002864
ASTHMA                 0.002841
INMSUPR               0.003246
HIPERTENSION          0.002960
OTHER_DISEASE         0.004811
CARDIOVASCULAR        0.002934
OBESITY               0.002892
RENAL_CHRONIC         0.002867
TOBACCO               0.003071
CLASIFFICATION_FINAL  0.000000
ICU                   0.816377
dtype: float64
```

```
In [ ]: msno.matrix(df_covid, color=(0.5, 0, 0), sparkline=False)
```

Out[]: <Axes: >



```
In [ ]: # ver datos unicos

df_covid.nunique()
```

```
Out[ ]: USMER                2
        MEDICAL_UNIT       13
        SEX                 2
        PATIENT_TYPE        2
        DATE_DIED           400
        INTUBED              2
        PNEUMONIA           2
        AGE                 117
        PREGNANT             2
        DIABETES             2
        COPD                 2
        ASTHMA               2
        INMSUPR              2
        HIPERTENSION         2
        OTHER_DISEASE        2
        CARDIOVASCULAR       2
        OBESITY              2
        RENAL_CHRONIC        2
        TOBACCO              2
        CLASIFFICATION_FINAL  7
        ICU                  2
        fallecidos           2
        mes                  12
        dtype: int64
```

Podemos observar que tenemos 5 campos con datos faltantes en el dataset. El dataset cuenta con un total de 1048575 de registros. Como vemos en la tabla, los campos con datos faltantes son: Intubados, neumonia, edad, embarazo y terapita_intensiva. De estos campos, Intubados y terapia_intensiva tienen un 81% de datos faltantes y embarazos 49%. Si no fueran variables con un significativo peso dentro de la evolucion de la enfermedad se podría eliminar del dataset pero se decidio mantenerlas para su analisis.

```
In [ ]: df_covid.shape
```

```
Out[ ]: (1048575, 21)
```

Buscamos datos duplicados

```
In [ ]: df_covid.duplicated().sum()
```

```
Out[ ]: 812074
```

Tenemos 814264 registros duplicados. Pero en este caso es normal por que son registros de personas contagiadas y es posible que compartan las características con otras personas.

Creamos el campo fallecido

El campo Date died indica mediante la fecha si el paciente murio y si no murio pone 9999-99-99. Creamos un campo de fallecido que sera 1 si murio y 2 si no murio. Esto permitira una mejor lectura y un mejor uso de los datos en el momento de realizar el modelo de prediccion.

```
In [ ]: # creamos el campo fallecidos a partir del campo date died. Si hay fecha el paciente fallecio, si la fecha es 9999-99-99 el paciente no fallecio.

df_covid['fallecidos'] = df_covid['DATE_DIED'].apply(lambda x: 2 if x == '9999-99-99' else 1)

df_covid.head()
```

	USMER	MEDICAL_UNIT	SEX	PATIENT_TYPE	DATE_DIED	INTUBED	PNEUMONIA	AGE	PREGNANT	DIABETES	COPD	ASTHMA	INMSUPR	HIPERTENSION	OTHER_DISEASE
0	2	1	1	1	03/05/2020	NaN	1.0	65.0	2.0	2.0	2.0	2.0	2.0	1.0	2.0
1	2	1	2	1	03/06/2020	NaN	1.0	72.0	NaN	2.0	2.0	2.0	2.0	1.0	2.0
2	2	1	2	2	09/06/2020	1.0	2.0	55.0	NaN	1.0	2.0	2.0	2.0	2.0	2.0
3	2	1	1	1	12/06/2020	NaN	2.0	53.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0
4	2	1	2	1	21/06/2020	NaN	2.0	68.0	NaN	1.0	2.0	2.0	2.0	1.0	2.0

```
In [ ]: # paso a datetime la columna date_died

df_covid['DATE_DIED'] = df_covid['DATE_DIED'].replace('9999-99-99', np.nan)

df_covid['DATE_DIED'] = pd.to_datetime(df_covid['DATE_DIED'], format='%d/%m/%Y')

# agrego una columna mes para poder hacer un analisis por mes

df_covid['mes'] = pd.to_datetime(df_covid['DATE_DIED']).dt.month

df_covid['mes_nombre'] = df_covid['DATE_DIED'].dt.strftime('%B')
```

```
In [ ]: df_covid.head()
```

	USMER	MEDICAL_UNIT	SEX	PATIENT_TYPE	DATE_DIED	INTUBED	PNEUMONIA	AGE	PREGNANT	DIABETES	COPD	ASTHMA	INMSUPR	HIPERTENSION	OTHER_DISEASE
0	2	1	1	1	2020-05-03	NaN	1.0	65.0	2.0	2.0	2.0	2.0	2.0	1.0	2.0
1	2	1	2	1	2020-06-03	NaN	1.0	72.0	NaN	2.0	2.0	2.0	2.0	1.0	2.0
2	2	1	2	2	2020-06-09	1.0	2.0	55.0	NaN	1.0	2.0	2.0	2.0	2.0	2.0
3	2	1	1	1	2020-06-12	NaN	2.0	53.0	2.0	2.0	2.0	2.0	2.0	2.0	2.0
4	2	1	2	1	2020-06-21	NaN	2.0	68.0	NaN	1.0	2.0	2.0	2.0	1.0	2.0

```
In [ ]: # # paso columna mes de float a int

# df_covid['mes'].fillna(0, inplace=True)

# df_covid['mes'] = df_covid['mes'].astype(int)

# df_covid.dtypes

In [ ]: # reemplazo los valores 2 por 0 en todo el dataset pero menos en CLASIFICACION_FINAL, USMER y mes

# Lista de columnas en las que se reemplazarán los valores 2 por 0
columnas_reemplazo = [col for col in df_covid.columns if col not in ["CLASIFICACION_FINAL", "USMER", "mes"]]
```

```
# Reemplazo de valores

df_covid[columnas_reemplazo] = df_covid[columnas_reemplazo].replace(2, 0)

df_covid.head()
```

Out[]:

	USMER	MEDICAL_UNIT	SEX	PATIENT_TYPE	DATE_DIED	INTUBED	PNEUMONIA	AGE	PREGNANT	DIABETES	COPD	ASTHMA	INMSUPR	HIPERTENSION	OTHER_DISEASE
0	2		1	1	1	2020-05-03	NaN	1.0	65.0	0.0	0.0	0.0	0.0	1.0	0.0
1	2		1	0	1	2020-06-03	NaN	1.0	72.0	NaN	0.0	0.0	0.0	1.0	0.0
2	2		1	0	0	2020-06-09	1.0	0.0	55.0	NaN	1.0	0.0	0.0	0.0	0.0
3	2		1	1	1	2020-06-12	NaN	0.0	53.0	0.0	0.0	0.0	0.0	0.0	0.0
4	2		1	0	1	2020-06-21	NaN	0.0	68.0	NaN	1.0	0.0	0.0	1.0	0.0

Guardamos el dataset limpio

```
In [ ]: df_covid.to_csv('./covid_clean_total.csv', index=False)
```