

```
%matplotlib inline
import matplotlib.pyplot as plt
import numpy
import pandas
import seaborn
import statistics as stats
```

```
# Let's install Tensorflow 2.0:
!pip install tensorflow==2.0.0
```

```
# And verify that it is now in its latest version:
import tensorflow as tf
print(tf.__version__)
```

```
Looking in indexes: https://pypi.org/simple, https://us-python.pkg.dev/colab-wheels/
Requirement already satisfied: tensorflow==2.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: absl-py>=0.7.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: opt-einsum>=2.3.2 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: termcolor>=1.1.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: wheel>=0.26 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: keras-preprocessing>=1.0.5 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: tensorboard<2.1.0,>=2.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: wrapt>=1.11.1 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: numpy<2.0,>=1.16.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: protobuf>=3.6.1 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: grpcio>=1.8.6 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: astor>=0.6.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: six>=1.10.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: google-pasta>=0.1.6 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: keras-applications>=1.0.8 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: gast==0.2.2 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: tensorflow-estimator<2.1.0,>=2.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: h5py in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: werkzeug>=0.11.15 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: google-auth-oauthlib<0.5,>=0.4.1 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: markdown>=2.6.8 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: requests<3,>=2.21.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: google-auth<2,>=1.6.3 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: setuptools>=41.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: cachetools<5.0,>=2.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: pyasn1-modules>=0.2.1 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: rsa<5,>=3.1.4 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: requests-oauthlib>=0.7.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: importlib-metadata>=4.4 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: zipp>=0.5 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: typing-extensions>=3.6.4 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: pyasn1<0.5.0,>=0.4.6 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: chardet<4,>=3.0.2 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: certifi>=2017.4.17 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: idna<3,>=2.5 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: urllib3!=1.25.0,!1.25.1,<1.26,>=1.21.1 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: oauthlib>=3.0.0 in /usr/local/lib/python3.7/dist-packages
Requirement already satisfied: cached-property in /usr/local/lib/python3.7/dist-packages
2.0.0
```

```

from google.colab import drive
drive.mount('/content/drive')
import os
os.chdir("/content/drive/MyDrive/pdatos")
!ls

```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.
dis2.csv dis_b.csv disb.csv

```

import numpy as np
import pandas as pd

```

```

#fetal = pd.read_table("/content/drive/MyDrive/pdatos/dis_b.data", header=None)
#print(fetal.head)
data = pd.read_csv("/content/drive/MyDrive/pdatos/data_t.csv")
X=data[0:8]
y=data['class']
#print(y)
print(X)
from sklearn.model_selection import train_test_split, ShuffleSplit
from sklearn.neural_network import MLPClassifier
#model = MLPClassifier(hidden_layer_sizes=(2126,) , max_iter=1000)
#model.fit(X, y)

```

	age	sex	on thyroxine	query on thyroxine	on antithyroid medication	sick	\
0	41	F	f	f	f	f	f
1	23	F	f	f	f	f	f
2	46	M	f	f	f	f	f
3	70	F	t	f	f	f	f
4	70	F	f	f	f	f	f
5	18	F	t	f	f	f	f
6	59	F	f	f	f	f	f
7	80	F	f	f	f	f	f

	pregnant	thyroid surgery	I131 treatment	query hypothyroid	...	TT4 measured	\
0	f	f	f	f	...	t	
1	f	f	f	f	...	t	
2	f	f	f	f	...	t	
3	f	f	f	f	...	t	
4	f	f	f	f	...	t	
5	f	f	f	f	...	t	
6	f	f	f	f	...	t	
7	f	f	f	f	...	t	

	TT4	T4U measured	T4U	FTI measured	FTI	TBG measured	TBG	referral	source	\
0	125	t	1.14	t	109	f	?	SVHC		
1	102	f	?	f	?	f	?	other		
2	109	t	0.91	t	120	f	?	other		
3	175	f	?	f	?	f	?	other		
4	61	t	0.87	t	70	f	?	SVI		
5	183	t	1.3	t	141	f	?	other		
6	72	t	0.92	t	78	f	?	other		
7	80	t	0.7	t	115	f	?	SVI		

class

```

0 negative.
1 negative.
2 negative.
3 negative.
4 negative.
5 negative.
6 negative.
7 negative.

```

```
[8 rows x 30 columns]
```

```

#Informacion del dataset
data.info()

```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 2800 entries, 0 to 2799
Data columns (total 30 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   age                                   2800 non-null   int64
1   sex                                   2800 non-null   object
2   on thyroxine                         2800 non-null   object
3   query on thyroxine                   2800 non-null   object
4   on antithyroid medication            2800 non-null   object
5   sick                                  2800 non-null   object
6   pregnant                             2800 non-null   object
7   thyroid surgery                      2800 non-null   object
8   I131 treatment                      2800 non-null   object
9   query hypothyroid                   2800 non-null   object
10  query hyperthyroid                   2800 non-null   object
11  lithium                              2800 non-null   object
12  goitre                               2800 non-null   object
13  tumor                                2800 non-null   object
14  hypopituitary                       2800 non-null   object
15  psych                                2800 non-null   object
16  TSH measured                         2800 non-null   object
17  TSH                                   2800 non-null   object
18  T3 measured                          2800 non-null   object
19  T3                                    2800 non-null   object
20  TT4 measured                         2800 non-null   object
21  TT4                                   2800 non-null   object
22  T4U measured                        2800 non-null   object
23  T4U                                   2800 non-null   object
24  FTI measured                        2800 non-null   object
25  FTI                                   2800 non-null   object
26  TBG measured                        2800 non-null   object
27  TBG                                   2800 non-null   object
28  referral source                     2800 non-null   object
29  class                               2800 non-null   object
dtypes: int64(1), object(29)
memory usage: 656.4+ KB

```

```
# Calcular media mediana y moda
```

```
data['age'].describe()
```

```
count    2800.000000
```

```
mean      51.686786
std       18.994013
min        1.000000
25%       36.000000
50%       54.000000
75%       67.000000
max       94.000000
Name: age, dtype: float64
```

```
data['referral source'].describe()
data['age'].dtype

dtype('int64')
```

```
data.groupby(by=['class','referral source','age']).describe()
```

			sex				on thyroxine				query thyro:
			count	unique	top	freq	count	unique	top	freq	count
class	referral source	age									
discordant.	STMW	2	1	1	F	1	1	1	f	1	1
		18	1	1	F	1	1	1	f	1	1
		45	1	1	?	1	1	1	t	1	1
	SVHC	30	1	1	F	1	1	1	f	1	1
		36	1	1	M	1	1	1	f	1	1
...
negative.	other	88	1	1	F	1	1	1	f	1	1
		89	2	1	F	2	2	1	f	2	2
		90	2	2	F	1	2	1	f	2	2
		92	1	1	F	1	1	1	t	1	1
		93	1	1	F	1	1	1	f	1	1

322 rows × 108 columns



```
data.groupby(by=['class','referral source','age','tumor']).describe()
```

	class	referral source	age	tumor	sex				on thyroxine			
					count	unique	top	freq	count	unique	top	freq
discordant.	STMW		2	f	1	1	F	1	1	1	f	1
			18	f	1	1	F	1	1	1	f	1
			45	f	1	1	?	1	1	1	t	1
	SVHC		30	f	1	1	F	1	1	1	f	1
			36	f	1	1	M	1	1	1	f	1

negative.	other		88	f	1	1	F	1	1	1	f	1
			89	f	2	1	F	2	2	1	f	2
			90	f	2	2	F	1	2	1	f	2
			92	t	1	1	F	1	1	1	t	1
			93	f	1	1	F	1	1	1	f	1

```
data.groupby(['age']).min()
```

```

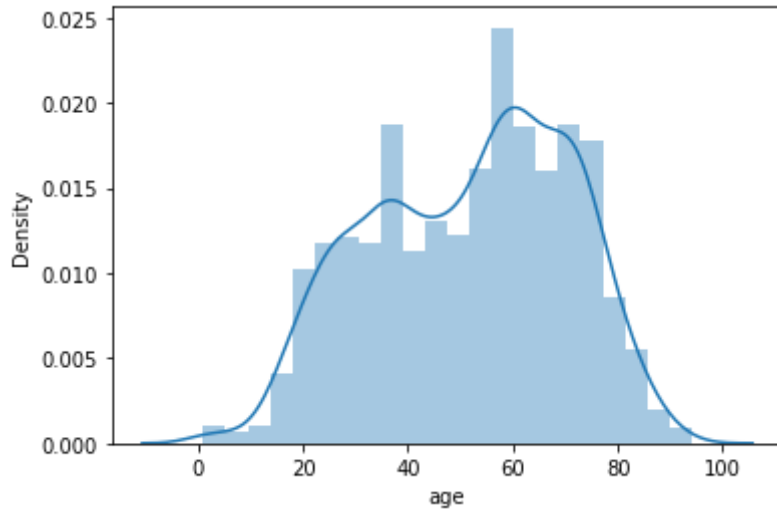
sex      on query on      on
thyroxine thyroxine antithyroid sick pregnant thyroid I131
surgery treatment hyp
seaborn.distplot(data.age.dropna(axis=0, how='all'))

```

```

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarning:
  warnings.warn(msg, FutureWarning)
<matplotlib.axes._subplots.AxesSubplot at 0x7f5f91053cd0>

```



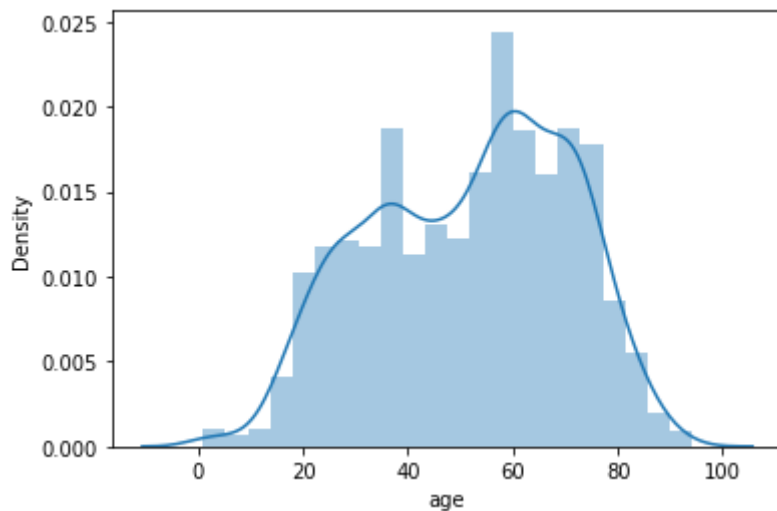
En la figura se puede apreciar que a las personas al rededor de los 21 años son los mas afectados por la Diabetes

```
seaborn.distplot(data.age.dropna(axis=0, how='all'))
```

```

/usr/local/lib/python3.7/dist-packages/seaborn/distributions.py:2619: FutureWarning:
  warnings.warn(msg, FutureWarning)
<matplotlib.axes._subplots.AxesSubplot at 0x7f5f90f0f650>

```



Es la dispersion de datos en la que se centran mas o en la que se agrtupan mas datos

```

# calculo de la media de Años
print(stats.mean(data['age']))

```

```
51.68678571428571
```

```
#calculo de la mediana de Años  
print(stats.median(data['age']))
```

```
54.0
```

```
#calculo de la moda de Años  
print(stats.mode(data['age']))
```

```
59
```

Moda, Mediana y media no coinciden en la variable Longitud, con lo cual para esta variable hay asimetria positiva.

```
data['age'].describe()
```

```
count    2800.000000  
mean      51.686786  
std       18.994013  
min        1.000000  
25%       36.000000  
50%       54.000000  
75%       67.000000  
max       94.000000  
Name: age, dtype: float64
```

La forma mas facil de saber todo esto es con la funcion .describe()

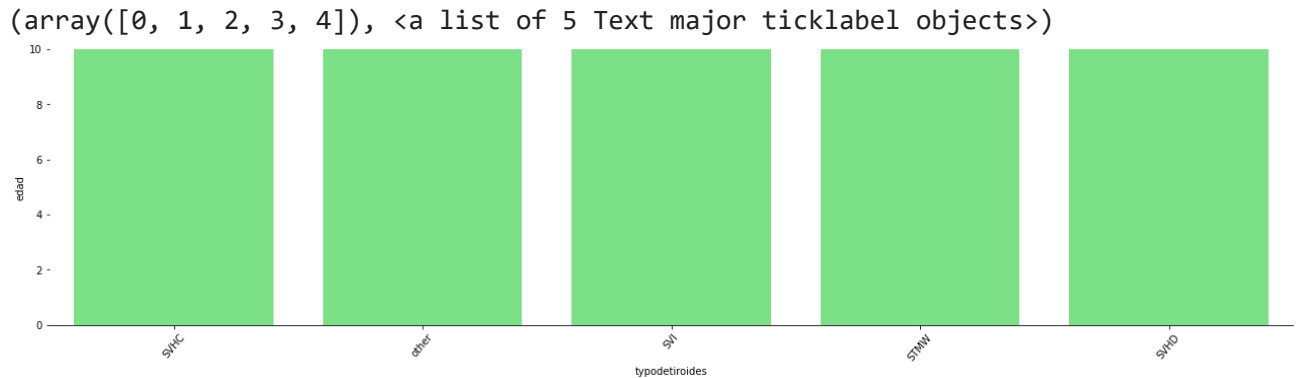
```
BLUE = '#35A7FF'  
RED = '#FF5964'  
GREEN = '#6BF178'  
YELLOW = '#FFE74C'
```

```
data.describe()
```

age



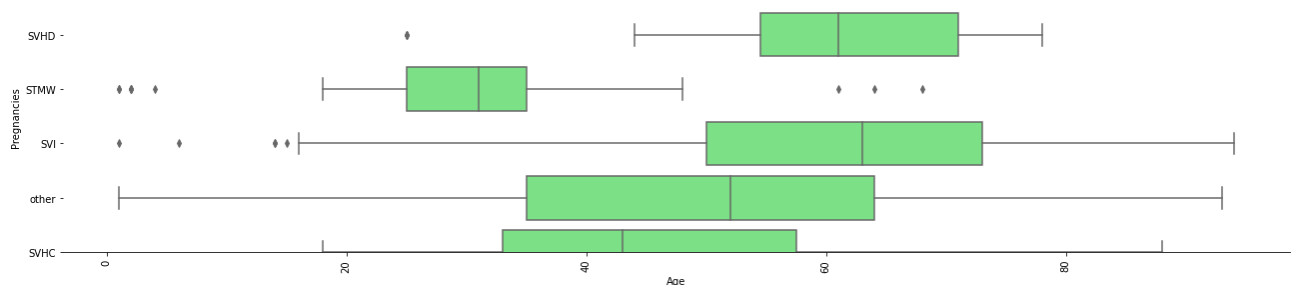
```
plt.figure(figsize=(22,5))
seaborn.barplot(data=data, x='referral source', y='age',color=GREEN)
plt.ylabel('edad')
plt.xlabel('tipodetiroides')
plt.ylim(0, 10)
seaborn.despine(left=True)
plt.xticks(rotation=50)
```



Aqui podemos observar la tabla de tipos y edades

```
plt.figure(figsize=(22,5))
seaborn.boxplot(data=data, x='age', y='referral source', color=GREEN)
plt.ylabel('Pregnancies')
plt.xlabel('Age')
plt.ylim(0, 5)
seaborn.despine(left=True)
plt.xticks(rotation=90)
```

(array([-20., 0., 20., 40., 60., 80., 100.]),
<a list of 7 Text major ticklabel objects>)



```
data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 2800 entries, 0 to 2799
Data columns (total 30 columns):
#   Column                Non-Null Count  Dtype
---  -
0   age                   2800 non-null  int64
1   sex                   2800 non-null  object
2   on thyroxine          2800 non-null  object
```

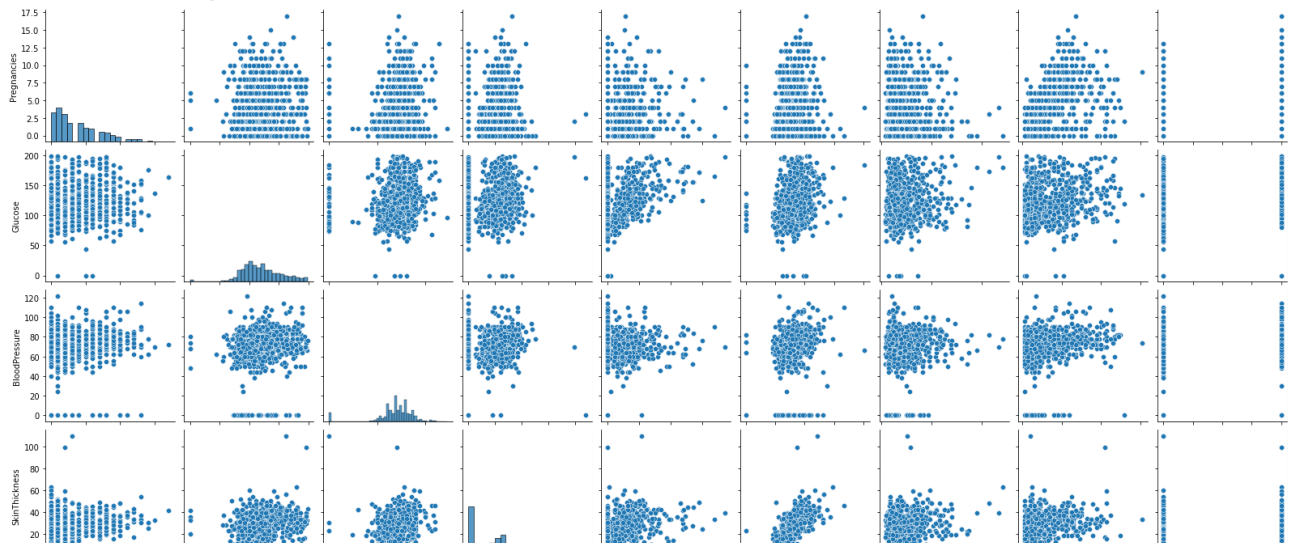

3	query on thyroxine	2800	non-null	object
4	on antithyroid medication	2800	non-null	object
5	sick	2800	non-null	object
6	pregnant	2800	non-null	object
7	thyroid surgery	2800	non-null	object
8	I131 treatment	2800	non-null	object
9	query hypothyroid	2800	non-null	object
10	query hyperthyroid	2800	non-null	object
11	lithium	2800	non-null	object
12	goitre	2800	non-null	object
13	tumor	2800	non-null	object
14	hypopituitary	2800	non-null	object
15	psych	2800	non-null	object
16	TSH measured	2800	non-null	object
17	TSH	2800	non-null	object
18	T3 measured	2800	non-null	object
19	T3	2800	non-null	object
20	TT4 measured	2800	non-null	object
21	TT4	2800	non-null	object
22	T4U measured	2800	non-null	object
23	T4U	2800	non-null	object
24	FTI measured	2800	non-null	object
25	FTI	2800	non-null	object
26	TBG measured	2800	non-null	object
27	TBG	2800	non-null	object
28	referral source	2800	non-null	object
29	class	2800	non-null	object

dtypes: int64(1), object(29)
memory usage: 656.4+ KB

```
important_cols =data[['class','age','sex','referral source']]
imp_cols2=data[['sex','T3','TT4','T4U','referral source','class']]
```

```
seaborn.pairplot(
    data=important_cols,
    markers='o')
```

```
<seaborn.axisgrid.PairGrid at 0x7f557bb0a750>
```



```
#a nivel numerico computar las correlaciones
```

```
import pandas as pd
import numpy as np
```

```
def get_corrs(important_cols):
    col_correlations = important_cols.corr()
    col_correlations.loc[:, :] = np.tril(col_correlations, k=-1)
    cor_pairs = col_correlations.stack()
    return cor_pairs.to_dict()
```

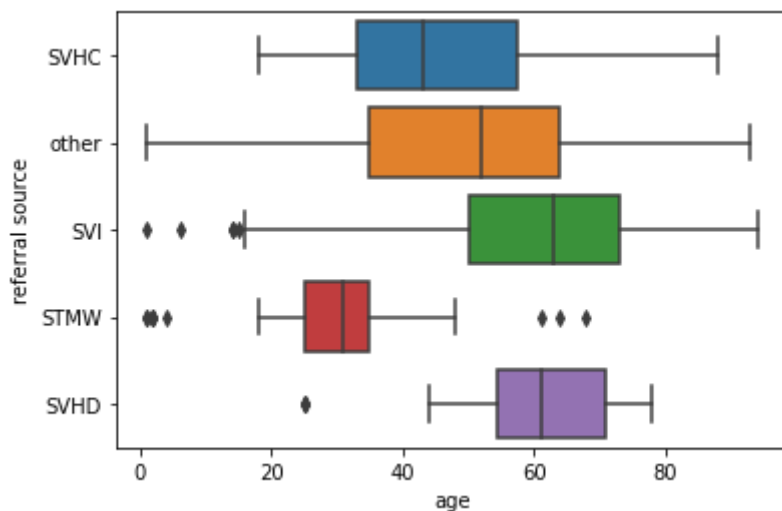
```
my_corrs = get_corrs(important_cols)
print(my_corrs)
```

```
{('age', 'age'): 0.0}
```

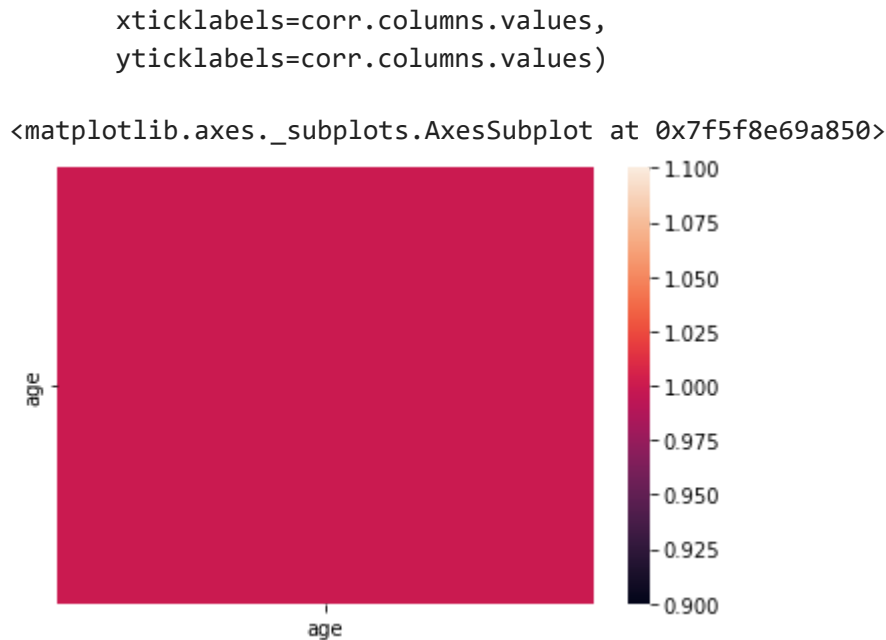
```
0.0 | | | | | | | | | |
```

```
import seaborn as sns
sns.boxplot( x=data["age"], y=data["referral source"] )
```

```
<matplotlib.axes._subplots.AxesSubplot at 0x7f5f8e7a0390>
```



```
import seaborn as sns
corr = important_cols.corr()
sns.heatmap(corr,
```



Correlaciones vistas en este caso con los valores de r

Correlaciones que se marcan en el heatmap: principalmente entre Age - Pregnacies y Glucose

```
## calculamos spearman's correlaciones
from scipy import stats
coef, p = stats.spearmanr(important_cols['age'].dropna(), important_cols['referra
print('Spearman's correlacion coeficiente al: %.3f' % coef)
# Interpretando el significado
alpha = 0.05
if p > alpha:
    print('Las muestras no están correlacionadas (fallo para rechazar H0) p=%.3f' %
else:
    print('Las muestras están correlacionadas (rechazar H0) p=%.3f' % p)

Spearman's correlacion coeficiente al: -0.046
Las muestras están correlacionadas (rechazar H0) p=0.016
```

```
## calculamos kendall's correlacion
from scipy import stats
coef, p = stats.kendalltau(important_cols['age'].dropna(), important_cols['referr
print('kendall correlacion coeficiente al: %.3f' % coef)
# interpret the significance
alpha = 0.05
if p > alpha:
    print('Las muestras no están correlacionadas (fallo para rechazar H0) p=%.3f' %
else:
    print('Las muestras están correlacionadas (rechazar H0) p=%.3f' % p)
```

↗ kendall correlacion coeficiente al: -0.036
Las muestras están correlacionadas (rechazar H_0) $p=0.015$

```
import seaborn as sns
seaborn.regplot(data=important_cols,
                 x='age', y='referral source',
                 marker='o')
seaborn.despine()
```

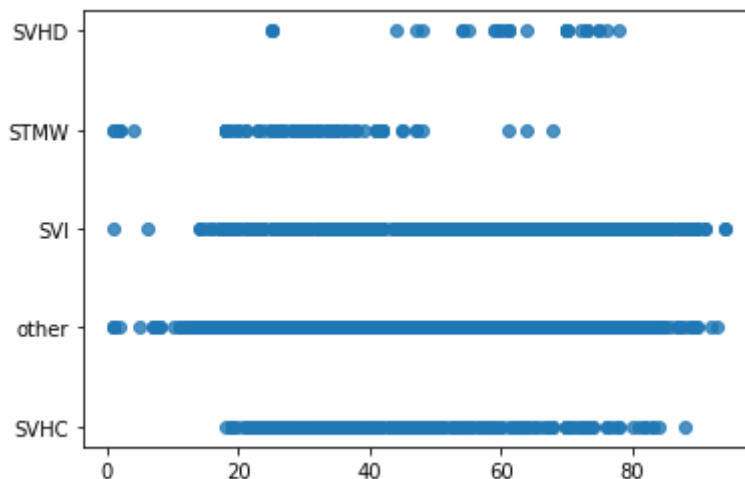
```
-----
TypeError                                Traceback (most recent call last)
<ipython-input-206-a049d3de3573> in <module>()
      2 seaborn.regplot(data=important_cols,
      3                 x='age', y='referral source',
----> 4                 marker='o')
      5 seaborn.despine()
```

⏏ 6 frames

```
/usr/local/lib/python3.7/dist-packages/seaborn/regression.py in reg_func(_x, _y)
    232     """Low-level regression and prediction using linear algebra."""
    233     def reg_func(_x, _y):
--> 234         return np.linalg.pinv(_x).dot(_y)
    235
    236     X, y = np.c_[np.ones(len(self.x)), self.x], self.y
```

TypeError: can't multiply sequence by non-int of type 'float'

SEARCH STACK OVERFLOW



Z1 sigue la dirección en la que las observaciones varían más (línea azul). La proyección de cada observación sobre esa dirección equivale al valor de la primera componente para dicha observación

 0 s completado a las 15:31

