

Uso de framework o biblioteca de aprendizaje máquina para la implementación de una solución

September 9, 2022

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[Enlace al repositorio en GitHub](#)

1 Librerías:

```
[701]: import pandas as pd
import numpy as np
import seaborn as sns
```

Importamos las librerías necesarias para algunos procesos a realizar antes de la implementación del modelo.

2 Importación de la base de datos:

```
[702]: url = "https://raw.githubusercontent.com/G4LF0/MLAlgorithmFramework/main/
→breast-cancer.csv"
```

```
[703]: df = pd.read_csv(url)
df.head(2)
```

```
[703]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
0	842302	M	17.99	10.38	122.8	1001.0	
1	842517	M	20.57	17.77	132.9	1326.0	

	smoothness_mean	compactness_mean	concavity_mean	concave	points_mean	\
0	0.11840	0.27760	0.3001		0.14710	
1	0.08474	0.07864	0.0869		0.07017	

	...	radius_worst	texture_worst	perimeter_worst	area_worst	\
0	...	25.38	17.33	184.6	2019.0	
1	...	24.99	23.41	158.8	1956.0	

	smoothness_worst	compactness_worst	concavity_worst	concave	points_worst	\
--	------------------	-------------------	-----------------	---------	--------------	---

0	0.1622	0.6656	0.7119	0.2654
1	0.1238	0.1866	0.2416	0.1860

	symmetry_worst	fractal_dimension_worst
0	0.4601	0.11890
1	0.2750	0.08902

[2 rows x 32 columns]

Declaramos el dataframe con ayuda de la libreria de pandas, la base de datos se encuentra alojada en nuestro repositorio en GitHub.

3 Analisis exploratorio de los datos:

```
[704]: df.isnull().sum()
```

```
[704]: id          0
diagnosis        0
radius_mean      0
texture_mean     0
perimeter_mean   0
area_mean        0
smoothness_mean  0
compactness_mean 0
concavity_mean   0
concave points_mean 0
symmetry_mean    0
fractal_dimension_mean 0
radius_se        0
texture_se       0
perimeter_se     0
area_se          0
smoothness_se    0
compactness_se   0
concavity_se     0
concave points_se 0
symmetry_se      0
fractal_dimension_se 0
radius_worst     0
texture_worst    0
perimeter_worst  0
area_worst       0
smoothness_worst 0
compactness_worst 0
concavity_worst  0
concave points_worst 0
symmetry_worst   0
```

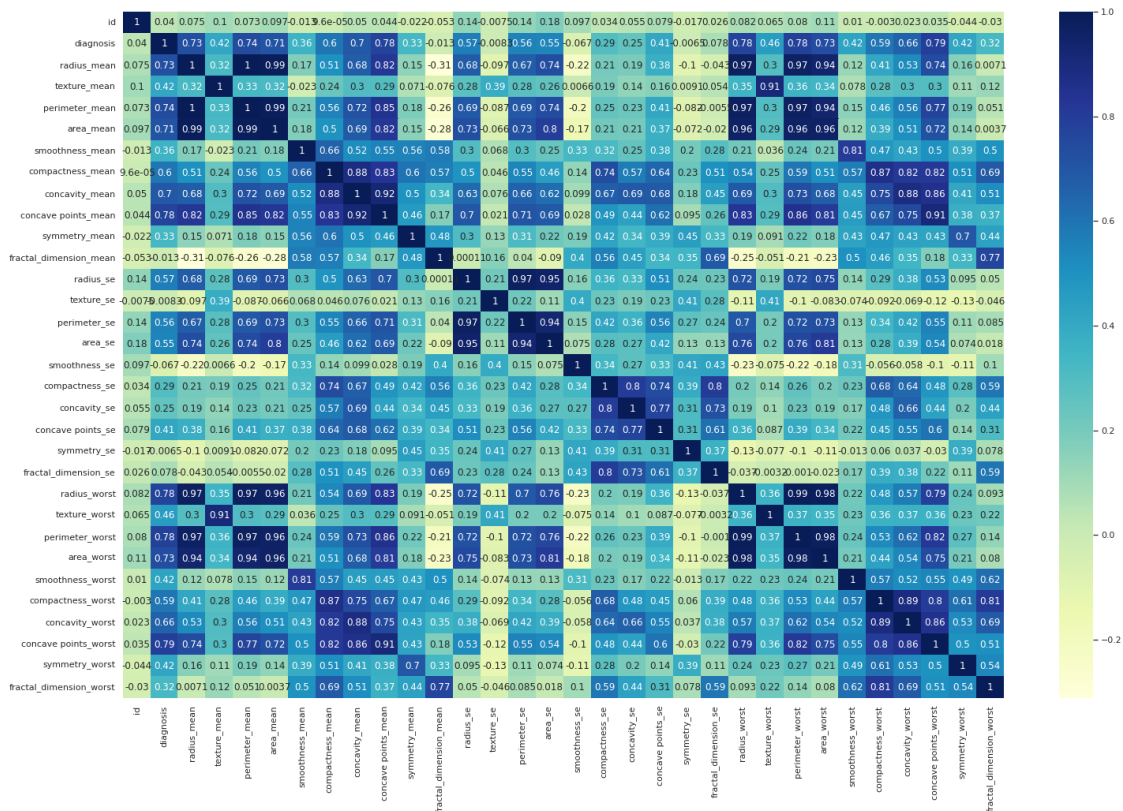
```
fractal_dimension_worst    0
dtype: int64
```

```
[705]: df.diagnosis = df.diagnosis.map({"M":1, "B":0})
df.shape
```

```
[705]: (569, 32)
```

```
[706]: sns.set(rc = {'figure.figsize':(25,16)})
sns.heatmap(df.corr(), annot=True, cmap= 'YlGnBu')
```

```
[706]: <matplotlib.axes._subplots.AxesSubplot at 0x7f7ebbc6e150>
```



```
[707]: df = df.drop(["id",
    →"fractal_dimension_se", "symmetry_se", "smoothness_se", "texture_se",
    →"fractal_dimension_mean"], axis = 1)
df.head(1)
```

```
[707]:   diagnosis  radius_mean  texture_mean  perimeter_mean  area_mean \
0         1         17.99         10.38         122.8         1001.0
```

```

smoothness_mean compactness_mean concavity_mean concave points_mean \
0 0.1184 0.2776 0.3001 0.1471

```

```

symmetry_mean ... radius_worst texture_worst perimeter_worst \
0 0.2419 ... 25.38 17.33 184.6

```

```

area_worst smoothness_worst compactness_worst concavity_worst \
0 2019.0 0.1622 0.6656 0.7119

```

```

concave points_worst symmetry_worst fractal_dimension_worst
0 0.2654 0.4601 0.1189

```

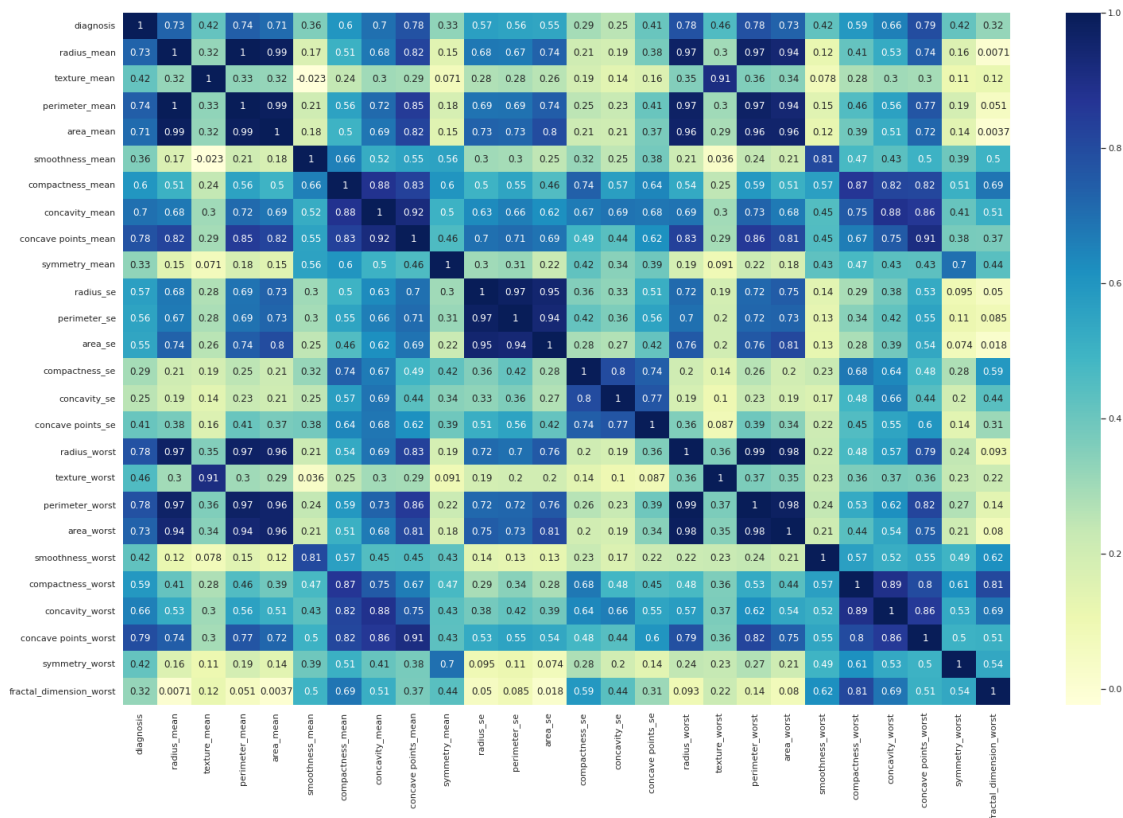
[1 rows x 26 columns]

```

[708]: sns.set(rc = {'figure.figsize':(25,16)})
sns.heatmap(df.corr(), annot=True, cmap= 'YlGnBu')

```

[708]: <matplotlib.axes._subplots.AxesSubplot at 0x7f7eb6f81110>



En esta parte lo que hicimos fue analizar columnas que no tuvieran relacion con la variable a predecir, asi como tambien cambiamos el valor de algunos variables para convertirlas en binarias.

4 Division de los datos en trainig, validating and testing:

```
[709]: x = df.drop(["diagnosis"], axis = 1)
      y = df.diagnosis
```

```
[710]: from sklearn.model_selection import train_test_split

      x_train, x_test, y_train, y_test = train_test_split(x.values, y.values,
      →train_size=0.9, random_state=16)
```

```
[711]: x_train, x_val, y_train, y_val = train_test_split(x_train, y_train, train_size=0.
      →7, random_state=16)
```

```
[712]: datos_n = [x_train.shape[0], x_val.shape[0], x_test.shape[0]]
```

```
[713]: df_mcd = pd.DataFrame({"Datos": ["Training", "Validating", "Testing"],
      "Registros": datos_n})
      df_mcd
```

```
[713]:      Datos:  Registros
0   Training      358
1 Validating      154
2   Testing       57
```

En esta parte lo que hacemos es dividir el data set, al principio lo dividimos en un 90% para training y un 10% para test, después lo que hacemos es que del 30% del 90% de training lo convertimos en validating.

5 Implementacion del modelo:

```
[714]: from sklearn.linear_model import LogisticRegression
      from sklearn.metrics import accuracy_score

      logreg = LogisticRegression(max_iter= 10000)
      logreg.fit(x_train, y_train)
      y_pred = logreg.predict(x_train)
```

Implementamos el algoritmo de regresión logística con un número de iteraciones máxima de 10,000.

Accuracy for training:

```
[715]: score = accuracy_score(y_train, y_pred)
      accuracy_training = score
      print("Accuracy for the training set is: ", round(score*100, 4))
```

Accuracy for the training set is: 95.8101

Accuracy for validating:

```
[716]: y_pred = logreg.predict(x_val)
score =accuracy_score(y_val,y_pred)
accuracy_validating = score
print("Accuracy for the training set is: ", round(score*100, 4))
```

Accuracy for the training set is: 96.1039

Accuracy for testing

```
[717]: y_pred = logreg.predict(x_test)
score =accuracy_score(y_test,y_pred)
accuracy_testing = score
print("Accuracy for the training set is: ", round(score*100, 4))
```

Accuracy for the training set is: 94.7368

6 Metrics:

Metrics del training:

```
[718]: from sklearn import metrics

y_pred = logreg.predict(x_train)
confusion_matrix = metrics.confusion_matrix(y_train, y_pred)
confusion_matrix
```

```
[718]: array([[220,  6],
        [ 9, 123]])
```

```
[719]: sns.heatmap(confusion_matrix, annot=True)
```

```
[719]: <matplotlib.axes._subplots.AxesSubplot at 0x7f7eb6a08350>
```



```
[720]: from sklearn.metrics import classification_report

target_names = ['Sin tumor', 'Con tumor']
print(classification_report(y_train, y_pred, target_names=target_names))
```

	precision	recall	f1-score	support
Sin tumor	0.96	0.97	0.97	226
Con tumor	0.95	0.93	0.94	132
accuracy			0.96	358
macro avg	0.96	0.95	0.95	358
weighted avg	0.96	0.96	0.96	358

Metricas del validating:

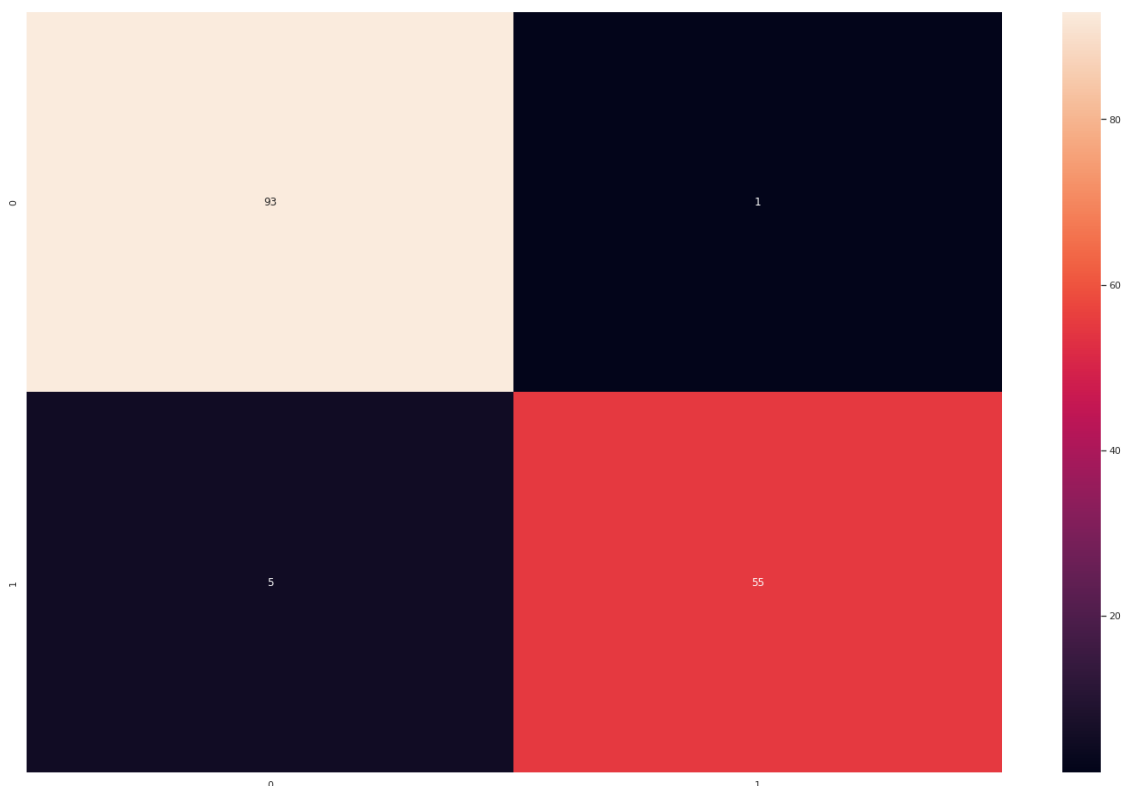
```
[721]: from sklearn import metrics

y_pred = logreg.predict(x_val)
confusion_matrix = metrics.confusion_matrix(y_val, y_pred)
confusion_matrix
```

```
[721]: array([[93,  1],
              [ 5, 55]])
```

```
[722]: sns.heatmap(confusion_matrix, annot=True)
```

```
[722]: <matplotlib.axes._subplots.AxesSubplot at 0x7f7eb69f40d0>
```



```
[723]: from sklearn.metrics import classification_report

target_names = ['Sin tumor', 'Con tumor']
print(classification_report(y_val, y_pred, target_names=target_names))
```

	precision	recall	f1-score	support
Sin tumor	0.95	0.99	0.97	94
Con tumor	0.98	0.92	0.95	60
accuracy			0.96	154
macro avg	0.97	0.95	0.96	154
weighted avg	0.96	0.96	0.96	154

Metricas del testing:

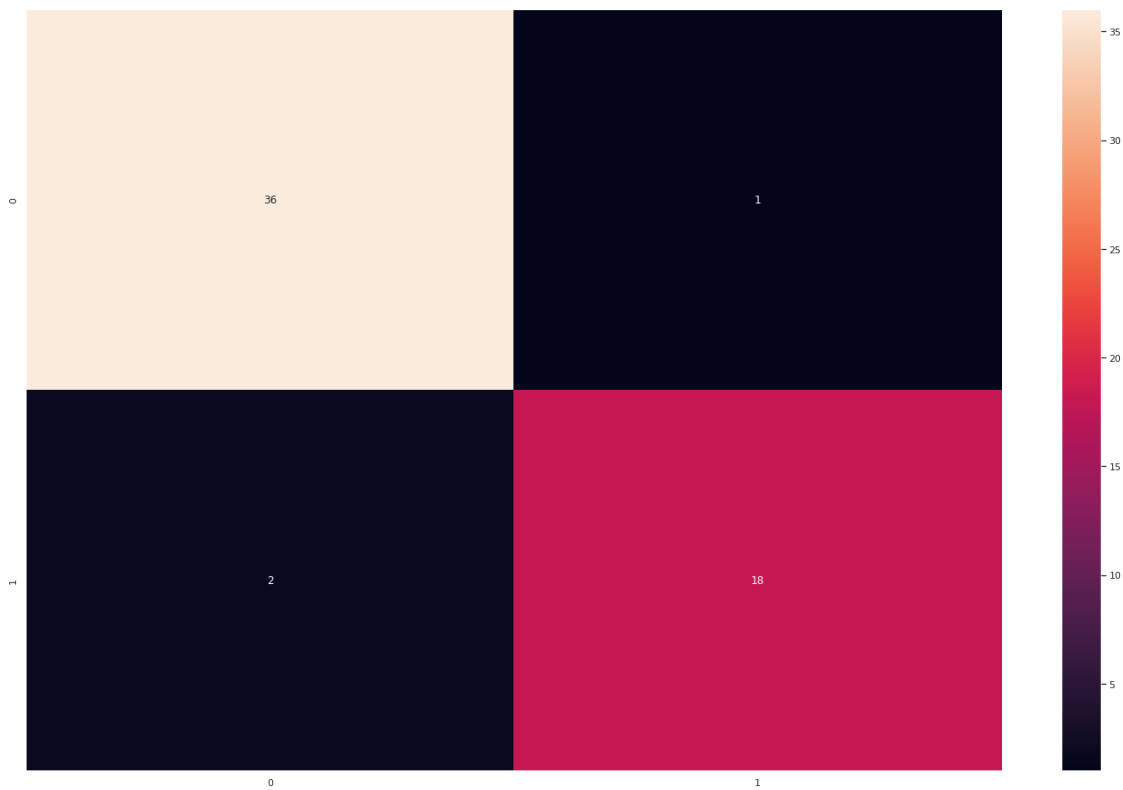

```
[724]: from sklearn import metrics
```

```
y_pred = logreg.predict(x_test)
confusion_matrix = metrics.confusion_matrix(y_test, y_pred)
confusion_matrix
```

```
[724]: array([[36,  1],
           [ 2, 18]])
```

```
[725]: sns.heatmap(confusion_matrix, annot=True)
```

```
[725]: <matplotlib.axes._subplots.AxesSubplot at 0x7f7eb68cd190>
```



```
[726]: from sklearn.metrics import classification_report
```

```
target_names = ['Sin tumor', 'Con tumor']
print(classification_report(y_test, y_pred, target_names=target_names))
```

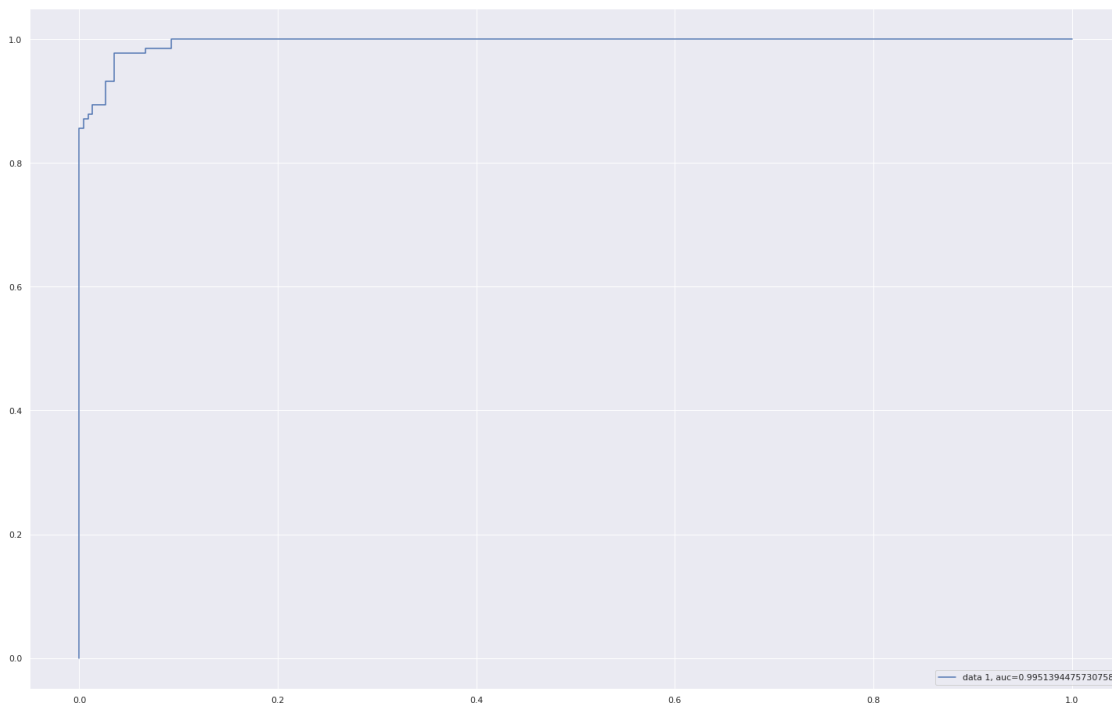
	precision	recall	f1-score	support
Sin tumor	0.95	0.97	0.96	37
Con tumor	0.95	0.90	0.92	20

accuracy			0.95	57
macro avg	0.95	0.94	0.94	57
weighted avg	0.95	0.95	0.95	57

7 Curva ROC

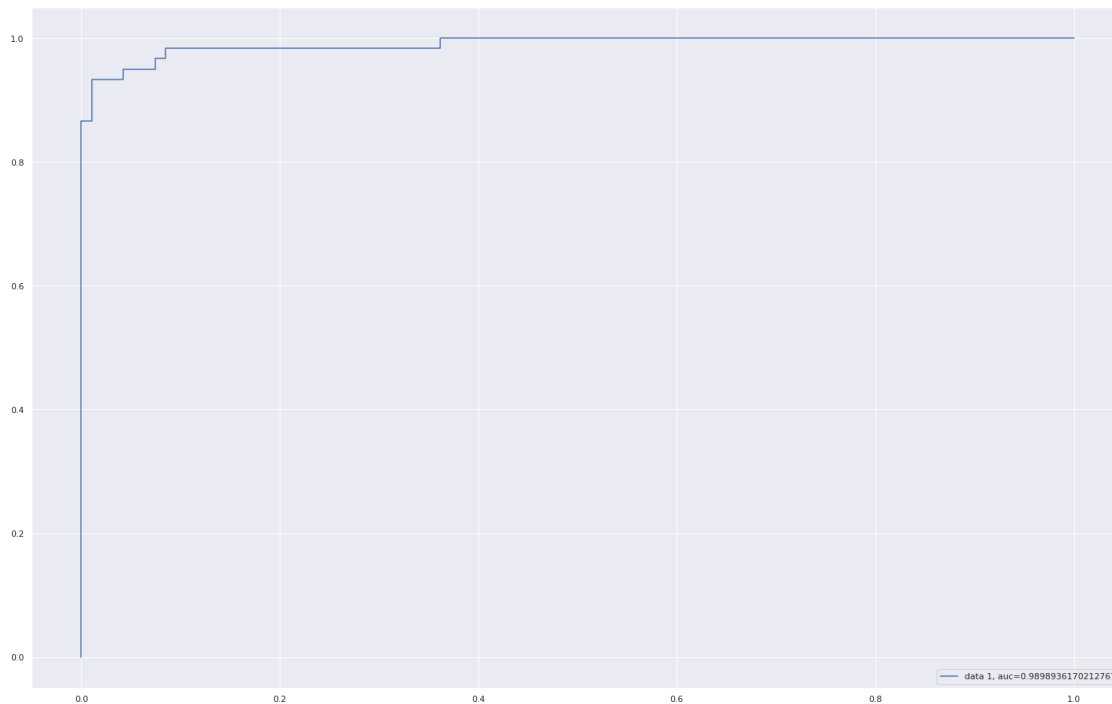
Training:

```
[727]: y_pred_proba = logreg.predict_proba(x_train)[:,:1]
fpr, tpr, _ = metrics.roc_curve(y_train, y_pred_proba)
auc = metrics.roc_auc_score(y_train, y_pred_proba)
plt.plot(fpr,tpr,label="data 1, auc="+str(auc))
plt.legend(loc=4)
plt.show()
```



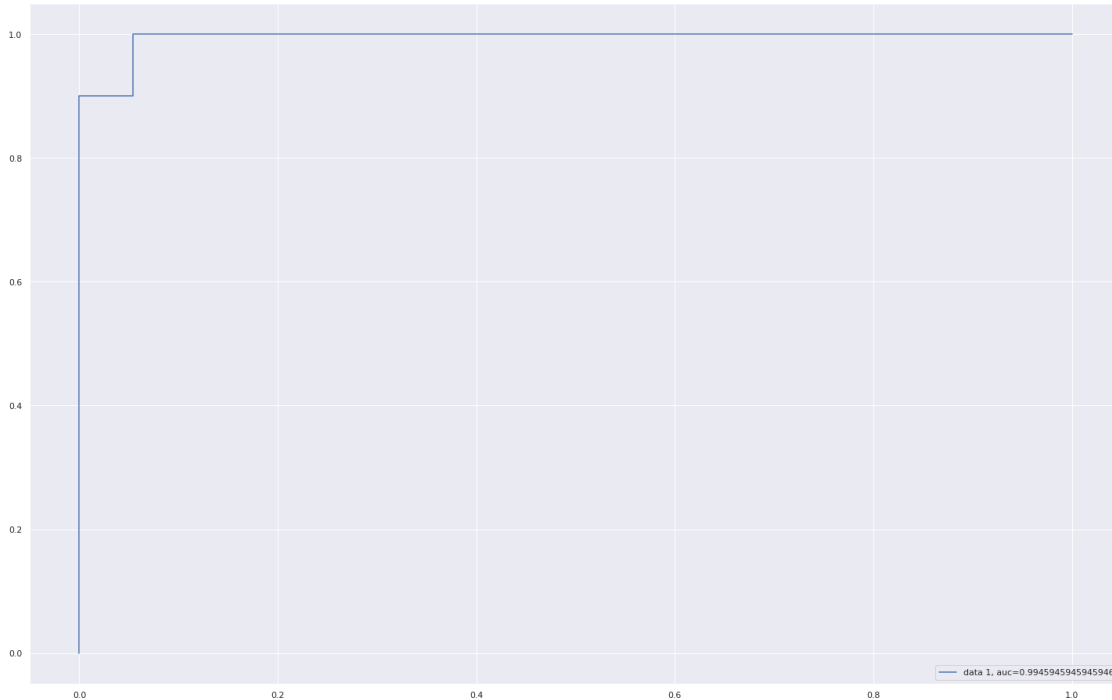
Validating:

```
[728]: y_pred_proba = logreg.predict_proba(x_val)[:,:1]
fpr, tpr, _ = metrics.roc_curve(y_val, y_pred_proba)
auc = metrics.roc_auc_score(y_val, y_pred_proba)
plt.plot(fpr,tpr,label="data 1, auc="+str(auc))
plt.legend(loc=4)
plt.show()
```



Testing:

```
[729]: y_pred_proba = logreg.predict_proba(x_test)[:,:1]
fpr, tpr, _ = metrics.roc_curve(y_test, y_pred_proba)
auc = metrics.roc_auc_score(y_test, y_pred_proba)
plt.plot(fpr,tpr,label="data 1, auc="+str(auc))
plt.legend(loc=4)
plt.show()
```



Tenemos que decir que la curva ROC es un grafico en el que se ponen los valores positivos y falsos en cada eje, y como podemos ver dieron un muy buen accuracy en los tres datasets.

8 Tabla de accuracys:

```
[730]: df_accuracys = pd.DataFrame({"Datos": ["Training", "Validating", "Testing"],
                                   "Registros": [accuracy_training,
                                   accuracy_validating, accuracy_testing]})
df_accuracys
```

```
[730]:   Datos  Registros
0  Training    0.958101
1 Validating    0.961039
2   Testing    0.947368
```

Como podemos ver nuestro modelo dispone de muy poco error en el dataset de training y mucho menos error en el dataset de validacion, por lo que decimos que se encuentra en **balance**.

9 Predicciones:

```
[731]: df.head(2)
```

```
[731]:
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
0	1	17.99	10.38	122.8	1001.0	
1	1	20.57	17.77	132.9	1326.0	

	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	\
0	0.11840	0.27760	0.3001	0.14710	
1	0.08474	0.07864	0.0869	0.07017	

	symmetry_mean	...	radius_worst	texture_worst	perimeter_worst	\
0	0.2419	...	25.38	17.33	184.6	
1	0.1812	...	24.99	23.41	158.8	

	area_worst	smoothness_worst	compactness_worst	concavity_worst	\
0	2019.0	0.1622	0.6656	0.7119	
1	1956.0	0.1238	0.1866	0.2416	

	concave points_worst	symmetry_worst	fractal_dimension_worst	
0	0.2654	0.4601	0.11890	
1	0.1860	0.2750	0.08902	

[2 rows x 26 columns]

```
[732]: df.tail(2)
```

```
[732]:
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
567	1	20.60	29.33	140.10	1265.0	
568	0	7.76	24.54	47.92	181.0	

	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	\
567	0.11780	0.27700	0.3514	0.152	
568	0.05263	0.04362	0.0000	0.000	

	symmetry_mean	...	radius_worst	texture_worst	perimeter_worst	\
567	0.2397	...	25.740	39.42	184.60	
568	0.1587	...	9.456	30.37	59.16	

	area_worst	smoothness_worst	compactness_worst	concavity_worst	\
567	1821.0	0.16500	0.86810	0.9387	
568	268.6	0.08996	0.06444	0.0000	

	concave points_worst	symmetry_worst	fractal_dimension_worst	
567	0.265	0.4087	0.12400	
568	0.000	0.2871	0.07039	

[2 rows x 26 columns]

Necesito retro de esto, entiendo que es generando un dataframe de prueba, pero como le especifico a la funcion el rango de valor minimo o maximo que puede tomar cada feature????

```
[734]: from sklearn.datasets import make_blobs
x_pred, _ = make_blobs(n_samples=5, centers=2, n_features=25, random_state=1)
y_pred = logreg.predict(x_pred)
for i in range(len(x_pred)):
    print("X=%s, Prediccion=%s" % (x_pred[i], y_pred[i]))
```

```
X=[-0.72845782  4.69207719 -9.11257134 -4.70774649 -5.81201403 -7.64029828
-6.57288861 -2.60026731 -2.14022223  1.90796407 -0.0962929  5.88996541
-7.30745134  6.11823492 -9.956714   3.56938727 -0.77773503  1.48943152
-9.21446244 -6.34417423  6.84286602  9.59532625 -2.96950526  3.62412417
 7.32702498], Prediccion=0
X=[ 6.66207513e+00 -7.74857828e+00 -8.42609747e+00 -7.22692234e+00
 8.08342641e+00 -9.17740471e+00 -7.75986468e-01  9.20435790e+00
 4.76735928e-01  3.73579641e+00 -2.82080122e+00  4.48043019e+00
 7.22197876e+00 -9.49653324e+00  5.08070743e+00  1.03956020e+01
 5.19580765e+00 -3.70856875e+00  5.47546980e+00 -1.03703176e+01
-3.30487462e-03  1.03588897e+01 -3.68635259e+00 -4.34464846e+00
-7.53587330e+00], Prediccion=0
X=[-1.47299851  4.81654152 -9.79941278 -3.8343399  -7.73554447 -7.77566432
-6.1529745  -1.95930155 -0.86573264  0.9614911  -1.99139466  3.0656596
-5.48746065  7.6396888  -9.79610181  3.45294706 -2.2739048  1.8718286
-7.63938979 -4.81346251  6.41838302  9.95881004 -4.82642828  4.01583475
 8.2683395 ], Prediccion=0
X=[ -2.34673261  3.56128423 -10.66895863  -3.96601315  -8.18219253
-7.91881241  -4.6149936  -2.3467413  -2.25648607  -0.11129428
-2.36326801  5.39684461  -5.86014725  6.92535308  -9.26133265
 5.50960534  -1.533745   1.79099968  -6.89209091  -6.39022006
 4.87237318  9.01588879  -3.94041067  4.4330755   8.36676646], Prediccion=0
X=[ 6.93843267 -8.56533428 -9.18628979 -7.97650893  7.87800946 -7.18690268
-2.43736344  9.50833658 -0.64897771  3.79884677 -5.30545973  4.85143626
 7.10141398 -9.65885141  4.22772468 11.05097771  6.93041484 -6.24910202
 7.0217506  -6.30782912 -0.70411778  6.97264203 -3.26437171 -4.42541353
-8.00334919], Prediccion=0
```

```
[735]: l = np.random.randint(low=0, high=52)
x_pred = x_test[l:l+5]
```

```
[736]: y_pred = logreg.predict(x_pred)
```

```
[737]: y_pred
```

```
[737]: array([1, 0, 0, 0, 1])
```

```
[739]: pd.DataFrame({"Numero de prediccion":["1", "2", "3", "4", "5"],
                    "Prediccion":y_pred})
```

```
[739]: Numero de prediccion  Prediccion
0                          1          1
```

1	2	0
2	3	0
3	4	0
4	5	1

10 Conclusion:

Si bien el dataset que le pasamos al algoritmo implementado era un dataset muy limpio al cual no le tuvimos que hacer una gran cantidad de modificaciones o transformacion, al implementar el algoritmo mediante un framework nos dimos cuenta que esto es mas facil que hacerlo a mano, de hecho, en la entrega anterior yo implemente este algoritmo a mano, [enlace](#). En esta entrega el hecho de implementarlo con un framewor facilita el estarle jugando a algunos parametros, por lo que hace esto mucho mas facil.