Setup

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import sklearn
import seaborn as sns

from sklearn.model_selection import train_test_split
from sklearn.metrics import mean_squared_error, accuracy_score, confusion_matrix, precision_score, recall_score, ConfusionMatrixDisplay
from math import sqrt

import keras
from keras.models import Sequential
from keras.layers import Dense
from keras.utils import to_categorical
```

Set Data

rawData = 'https://raw.githubusercontent.com/Rwyld/Data-Science-Models/main/RedesNeuronales/diabetes%20CSV.csv'

data = pd.read_csv(rawData)
data.head(3)

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedi
0	6	148	72	35	0	33.6	
1	1	85	66	29	0	26.6	
2	8	183	64	0	0	23.3	

Analisis Exploratorio

data.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):

#	Column	Non-Null Count	Dtype
0	Pregnancies	768 non-null	int64
1	Glucose	768 non-null	int64
2	BloodPressure	768 non-null	int64
3	SkinThickness	768 non-null	int64
4	Insulin	768 non-null	int64
5	BMI	768 non-null	float64
6	DiabetesPedigreeFunction	768 non-null	float64
7	Age	768 non-null	int64
8	Outcome	768 non-null	int64

dtypes: float64(2), int64(7)
memory usage: 54.1 KB

fig.tight_layout()

plt.subplots_adjust(top = 0.9)

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BM
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.00000
mean	3.845052	120.894531	69.105469	20.536458	79.799479	31.99257
std	3.369578	31.972618	19.355807	15.952218	115.244002	7.88416
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.00000
25%	1.000000	99.000000	62.000000	0.000000	0.000000	27.30000
50%	3.000000	117.000000	72.000000	23.000000	30.500000	32.00000
75%	6.000000	140.250000	80.000000	32.000000	127.250000	36.60000
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.10000

```
fig, axes = plt.subplots(nrows=2, ncols=4, figsize=(10,5))
axes = axes.flat
#data = data.select_dtypes(include=['float64', 'int']).columns
graphData = data.drop('Outcome', axis=1)
for i, colum in enumerate(graphData):
   sns.histplot(
       data
               = graphData,
               = colum,
       X
       stat
               = "count",
       kde
               = True,
       color = (list(plt.rcParams['axes.prop_cycle'])*2)[i]["color"],
       line_kws= {'linewidth': 2},
       alpha = 0.3,
               = axes[i]
   axes[i].set_title(colum, fontsize = 10, fontweight = "bold")
   axes[i].tick_params(labelsize = 8)
   axes[i].set_xlabel("")
```

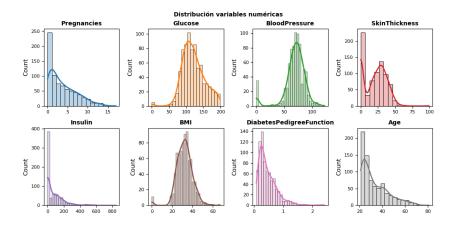


fig.suptitle('Distribución variables numéricas', fontsize = 10, fontweight = "bold");

Resumen: La data cargada no presenta datos faltantes o nulos. En cuanto a los datos visualizados en histogramas, no se observan posibles datos anomalos y las variables podemos observar una distribucion normal en Glucosa, BMI y BlooPressure.

Modelando Keras

Split test/train data

```
X = data.drop('Outcome', axis = 1).values
y = data['Outcome'].values

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.1, random_state=2023)

y_train = to_categorical(y_train)
y_test = to_categorical(y_test)

Definiendo la red

model = Sequential()
model.add(Dense(100, activation='relu', input_dim=8))
model.add(Dense(50, activation='relu'))
model.add(Dense(2, activation='rsigmoid'))

Compilando
```

Ajustando

loss='categorical_crossentropy',

metrics=['accuracy'])

Predicciones y conclusion

model.compile(optimizer='adam',

Conclusión: La red obtuvo un 89% de precision en los datos de entrenamiento y un 81% en los datos de testeo, a pesar de ser una precision buena, el modelo puede mejorar su precision si se configura con otros hiperparametros y con una mayor cantidad de datos.