```
In [1]: import pandas as pd
        import pylab as pl
        import numpy as np
        from sklearn.linear_model import LogisticRegression
        from sklearn.model_selection import train_test_split
        from sklearn import metrics
        import matplotlib.pyplot as plt
        import warnings
        from sklearn.metrics import confusion matrix
        from sklearn.metrics import classification_report
        warnings.filterwarnings('ignore')
        import os
        os.chdir('/Users/Lenovo/Desktop/EBAC')
In [3]: data = pd.read csv('drugs1.csv')
        data
Out[3]:
             Age Sex
                           BP Cholesterol Na_to_K Drug
                    F
                         HIGH
                                     HIGH
              23
                                            25.355 drugY
              47
                   M
                          LOW
                                     HIGH
                                            13.093 drugC
              47
                          LOW
                                     HIGH
                                            10.114 drugC
          3
              28
                    F NORMAL
                                     HIGH
                                             7.798 drugX
          4
                    F
                                     HIGH
                                            18.043 drugY
              61
                          LOW
        195
              56
                    F
                          LOW
                                     HIGH
                                            11.567 drugC
                          LOW
        196
              16
                                     HIGH
                                            12.006 drugC
              52
                   M NORMAL
                                     HIGH
                                             9.894 drugX
        197
        198
              23
                      NORMAL
                                  NORMAL
                                            14.020 drugX
                                  NORMAL
        199
              40
                          LOW
                                            11.349 drugX
        200 rows × 6 columns
In [5]: features_cols = ['Age', 'Sex', 'BP', 'Cholesterol', 'Na_to_K']
        X = data[features_cols].values
        y = data.Drug
In [7]: from sklearn import preprocessing
        Cod_Sex = preprocessing.LabelEncoder()
        Cod_Sex.fit(['F', 'M'])
        X[:,1] = Cod_Sex.transform(X[:,1])
        Cod_BP = preprocessing.LabelEncoder()
        Cod BP.fit(['HIGH', 'LOW', 'NORMAL'])
        X[:,2] = Cod_BP.transform(X[:,2])
        Cod_Cholesterol = preprocessing.LabelEncoder()
        Cod_Cholesterol.fit(['HIGH', 'LOW', 'NORMAL'])
        X[:,3] = Cod Cholesterol.transform(X[:,3])
In [9]: # Creacion de grupos de entrenamiento y prueba
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.3, random_state = 1)
```

## Regresion logistica con Solucionador (Solver): sag

```
# Reporte de clasificacion
 print(classification report(y test, y pred))
Intercepto (Beta0): [ 0.25207028 -0.08295724  0.09391024  0.21916712 -0.48219041]
Pesos de cada variable (Beta1, Beta2, ..., Beta7): [[-0.02543513 0.15505068 -0.63735024 0.12174788 0.1007136
 [ 0.07208856 -0.06031722 -0.51824822 -0.11326672 -0.29048744]
 [-0.00437994  0.01393438  0.14886401  -0.39560134  0.00341319]
 [ 0.01153286 -0.08992082 1.17809626 0.53268686 -0.12734421]
[-0.05380635 -0.01874702 -0.17136182 -0.14556668 0.31370484]]
precision recall f1-score support
                      0.25
0.67
                                 0.29
      drugA
               0.33
      drugB
               0.57
                                 0.62
                                             6
                       0.00
              0.00
      drugC
                                0.00
                                             4
              0.85
      drugX
                       0.89
                              0.87
                                            19
               0.83
                        0.93
                                 0.88
                                           27
      drugY
                                 0.78
                                           60
   accuracv
              0.52 0.55
  macro avg
                              0.53
                                           60
weighted avg
              0.72
                       0.78
                                 0.75
                                            60
```

#### Resgresion logistica con Solucionador (Solver): newton-cg

```
In [26]: model = LogisticRegression(solver = 'newton-cg')
        clf = model.fit(X_train, y_train)
        # Prediccion de etiquetas de clase sobre datos de prueba
        y pred = model.predict(X test)
        # Impresion de coeficientes de la regresion de puntajes
        print('Intercepto (Beta0): ', clf.intercept )
        print('Pesos de cada variable (Beta1, Beta2, ..., Beta7): ', clf.coef_)
        # Evaluacion de la precision del modelo
        score = model.score(X_test, y_test)
        print('Score de precision: ', score)
        print('-----
        # Reporte de clasificacion
        print(classification_report(y_test, y_pred))
       Intercepto (Beta0): [ 14.48790715 -0.08512375 10.50580619 5.75638881 -30.6649784 ]
       Pesos de cada variable (Beta1, Beta2, ..., Beta7): [[-8.96856681e-02 6.77268617e-02 -2.33960214e+00 -2.3674471
       8e-01
         -6.33653119e-01]
        [ 1.42467196e-01 -1.52212913e-01 -1.59788400e+00 -1.32018905e-01
         -4.42440147e-01]
        [-3.38662547e-02 2.54380976e-04 4.21069578e-01 -1.14094176e+00
         -5.85695514e-01]
        [-7.31005987e-03 -3.94719449e-01 3.09381357e+00 1.26917979e+00
         -5.67169846e-01]
        [-1.16052125e-02 4.78951120e-01 4.22602991e-01 2.40525596e-01
          2.22895863e+00]]
       Score de precision: 0.95
                    precision recall f1-score support
                       0.80 1.00
                                          0.89
                                0.83
0.50
                        0.71
                                           0.77
                                                       6
              druaB
              drugC
                        1.00
                                           0.67
                                                       4
                       1.00
                                1.00
                                          1.00
             druaX
                                                      19
              {\tt drugY}
                       1.00
                                1.00
                                          1.00
                                                     27
           accuracy
                                           0.95
                                                      60
                      0.90 0.87
          macro avq
                                           0.86
                                                      60
                       0.96
                                0.95
                                           0.95
       weighted avg
                                                      60
```

## Resgresion logistica con Solucionador (Solver): liblinear

```
In [18]: model = LogisticRegression(solver = 'liblinear')
        clf = model.fit(X train, y train)
        # Prediccion de etiquetas de clase sobre datos de prueba
        y_pred = model.predict(X_test)
        # Impresion de coeficientes de la regresion de puntajes
        print('Intercepto (Beta0): ', clf.intercept )
        print('Pesos de cada variable (Beta1, Beta2, ..., Beta7): ', clf.coef_)
        # Evaluacion de la precision del modelo
        score = model.score(X_test, y_test)
        print('Score de precision: ', score)
        # Reporte de clasificacion
        print(classification_report(y_test, y_pred))
       Intercepto (Beta0): [ 2.31089578 -0.65080868 0.6825165 0.16513608 -4.27580351]
       Pesos de cada variable (Beta1, Beta2, ..., Beta7): [[-0.02428697 0.42805745 -2.29889241 -0.14361067 -0.1657302
        [ 0.11187746 -0.52110997 -2.33619921 -0.54425809 -0.45569985]
        [-0.00360594  0.0611196  0.01560035 -1.39441786 -0.17622994]
        [ 0.01055564 -0.71011324 3.12578883 1.45357325 -0.47400147]
        Score de precision: 0.85
                   precision recall f1-score support
                       0.60 0.75
                                         0.67
             drugA
                                                       4
                                0.67
0.25
              drugB
                        0.57
                                           0.62
                      0.57
1.00
                                         0.40
             drugC
                                                      4
                      0.86 1.00 0.93
0.96 0.89 0.92
             drugX
                                                     19
                                                     27
             drugY
                                          0.85
                                                     60
           accuracv
                   0.80 0.71 0.71
0.87 0.85 0.84
          macro avg
                                                      60
       weighted avg
```

#### Resgresion logistica con Solucionador (Solver): saga

```
Intercepto (Beta0): [ 0.1265274 -0.0522943 0.04611565 0.13371509 -0.25406383]
Pesos de cada variable (Beta1, Beta2, ..., Beta7): [[-0.02337324 0.08785222 -0.37363311 0.08269469 0.0892573
21
[ 0.06354411 -0.03265927 -0.3150302 -0.06978496 -0.27327782]
[-0.00356606 0.00812699 0.07605 -0.24470664 -0.00516823]
[ 0.01628445 -0.04410388  0.74523258  0.35058191 -0.09611223]
[-0.05288925 -0.01921606 -0.13261927 -0.118785 0.28530096]]
______
Score de precision: 0.75
           precision recall f1-score support
                      0.00
0.67
     drugA
               0.00
                                0.00
              0.57
                               0.62
     druaB
                                           6
              0.00 0.00
     drugC
                               0.00
                      0.79
              0.75
                                0.77
                                          19
     drugX
     drugY
               0.79
                       0.96
                                0.87
                                          27
                               0.75
   accuracy
           0.42 0.48
0.65 0.75
                             0.45
  macro avg
                                          60
                                0.70
weighted avg
                                          60
```

#### Resgresion logistica con Solucionador (Solver): Ibfgs

```
In [24]: model = LogisticRegression(solver = 'lbfgs')
        clf = model.fit(X train, y train)
        # Prediccion de etiquetas de clase sobre datos de prueba
        y pred = model.predict(X test)
        # Impresion de coeficientes de la regresion de puntajes
        print('Intercepto (Beta0): ', clf.intercept_)
        print('Pesos de cada variable (Beta1, Beta2, ..., Beta7): ', clf.coef_)
        # Evaluacion de la precision del modelo
        score = model.score(X test, y test)
        print('Score de precision: ', score)
        # Reporte de clasificacion
        print(classification_report(y_test, y_pred))
       Intercepto (Beta0): [ 1.16939214 -0.31971945 0.54355859 0.7087925 -2.10202379]
       Pesos de cada variable (Beta1, Beta2, ..., Beta7): [[-0.03833878 0.47809653 -2.01089716 -0.08237991 0.1626778
        [ 0.09852575 -0.23784054 -1.80071268 -0.39446806 -0.28193157]
        [-0.01344921 -0.43843358 3.43060162 1.59338483 -0.30371246]
        [-0.05675346  0.12082599  -0.11028058  0.02044065  0.47145735]]
       Score de precision: 0.83333333333333334
       -----
                   precision recall f1-score support
             druaA
                       0.60
                                0.75
                                          0.67
                      0.57
                                0.67
             drugB
                                         0.62
                                                     6
                      0.00 0.00
0.83 1.00
0.96 0.89
             drugC
                                          0.00
                                                       4
                                                     19
             drugX
                                          0.90
             drugY
                                         0.92
                                                    27
                                          0.83
                                                     60
           accuracy
                    0.59 0.66 0.62
0.79 0.83 0.81
          macro avg
                                                     60
       weighted avg
                                                    60
```

#### Conclusion

Despues de haber evaluado todos los modelos, el mejor modelo de acuerdo al accuracy en F1 Score, fue el de Newton-cg, ya que alcanzo un 95% de acertividad en su pronostico.

# Eficacia del modelo Newton-cg

```
In [44]: classes = np.unique(y_test)
         n classes = len(classes)
         from sklearn.preprocessing import label binarize
         from sklearn.metrics import roc_curve, auc, roc_auc_score
         y_test_bin = label_binarize(y_test, classes=classes)
In [50]: y_score = model.predict_proba(X_test)
In [58]: for i in range(n_classes):
             fpr, tpr, _ = roc_curve(y_test_bin[:, i], y_score[:, i])
             roc_auc = auc(fpr, tpr)
             plt.plot(fpr, tpr, label=f'Clase {classes[i]} (AUC = {roc auc:.2f})')
         plt.plot([0, 1], [0, 1], 'k--')
         plt.xlabel('False Positive Rate')
         plt.ylabel('True Positive Rate')
         plt.title('Curvas ROC por clase (One-vs-Rest)')
         plt.legend(loc='lower right')
         plt.show()
```

#### Curvas ROC por clase (One-vs-Rest) 1.0 0.8 True Positive Rate 0.6 Clase drugA (AUC = 1.00) 0.2 Clase drugB (AUC = 0.98) Clase drugC (AUC = 0.98) Clase drugX (AUC = 1.00) Clase drugY (AUC = 1.00) 0.0 0.2 0.0 0.4 0.6 0.8 1.0 False Positive Rate

En la grafica ROC podemos observar que cada una de las clases esta muy pegadas a los bordes, por lo que podemos confirmar que la eficiencia en el modelo es muy alta.

In [ ]:

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