

intro_aprendizaje_

June 28, 2019

1 INTRODUCCIÓN AL APRENDIZAJE AUTOMÁTICO

1.1 LABORATORIO 1

1.1.1 FERRARO, MARÍA EUGENIA

```
In [4]: import matplotlib.pyplot as plt
import numpy as np
import matplotlib as mp

from matplotlib.colors import ListedColormap
from sklearn.datasets import load_boston, load_breast_cancer, load_iris
from sklearn.linear_model import LinearRegression, LogisticRegression, Perceptron, Ridge
from sklearn.metrics import accuracy_score, confusion_matrix, mean_squared_error
from sklearn.neighbors import KNeighborsClassifier
from sklearn.preprocessing import PolynomialFeatures

from ml.visualization import plot_confusion_matrix, classifier_boundary

np.random.seed(1234) # Setup seed to be more deterministic\n",

import pandas as pd
import seaborn as sb

%matplotlib inline
import warnings
warnings.filterwarnings('ignore')
```

2 REGRESION

```
In [29]: def regression_lineal(xtf,yt,xvf,yv,regularizacion,alpha,feature):
    if regularizacion==False:
        model = LinearRegression()
        model.fit(xtf, yt)
    else:
```

```

        model = Ridge(alpha=alpha)
        model.fit(xtf, yt)

    err_train = mean_squared_error(yt, model.predict(xtf))
    err_val = mean_squared_error(yv, model.predict(xvf))

    return [feature,regularizacion,alpha,err_train,err_val,model.coef_,model.intercept_]

def regresion_polinomial(xtf,yt,xvf,yv,grado_polinomio,regularizacion,alpha,feature):
    poly_features = PolynomialFeatures(grado_polinomio)
    poly_features.fit(xtf)
    X_poly_train = poly_features.transform(xtf)
    X_poly_val = poly_features.transform(xvf)

    if regularizacion==False:
        model = LinearRegression()
        model.fit(X_poly_train, y_train)

    else:
        model = Ridge(alpha=alpha)
        model.fit(X_poly_train, y_train)

    err_train = mean_squared_error(y_train, model.predict(X_poly_train))
    err_val = mean_squared_error(y_val, model.predict(X_poly_val))

    return [feature,grado_polinomio,regularizacion,alpha,err_train,err_val,model.coef_]

def modelo(x,coef,x0):
    ymodel = np.zeros(len(x))
    for i in range(len(x)):
        if len(coef)==1:
            ymodel[i] = coef[0]*x[i]
        else:
            for c in range(len(coef)):
                ymodel[i] = ymodel[i] + coef[c]*x[i]**c
    return ymodel+x0

def plot_fit(feature_resultados,tip,c,lw):
    X_range_start = np.min(np.r_[X_train_feature, X_val_feature])
    X_range_stop = np.max(np.r_[X_train_feature, X_val_feature])
    y_range_start = np.min(np.r_[y_train, y_val])
    y_range_stop = np.max(np.r_[y_train, y_val])

    X_linspace = np.linspace(X_range_start, X_range_stop, 200)

```

```

# Conjunto de entrenamiento
plt.scatter(X_train_feature, y_train, facecolor="lightgray", edgecolor="gray", label="Entrenamiento")
if tipo=='L':
    for j in range(len(feature_resultados)):
        coef = feature_resultados.Coefficients.values[j]
        x0 = feature_resultados.Intercept.values[j]
        plt.plot(X_linspace, modelo(X_linspace,coef,x0), color=c, label="modelo",)
elif tipo=='P':
    grados_ = feature_resultados.Degree.unique()
    cc=0
    for g in grados_:
        sub_df = feature_resultados[feature_resultados.Degree==g]
        for j in range(len(sub_df)):
            coef = sub_df.Coefficients.values[j]
            x0 = sub_df.Intercept.values[j]
            plt.plot(X_linspace, modelo(X_linspace,coef,x0),color=c[cc],label="modelo",)
            cc+=1

plt.ylim(y_range_start, y_range_stop)

# Conjunto de validación
# plt.subplot(1, 2, 2)
# plt.scatter(X_val_feature, y_val, facecolor="dodgerblue", edgecolor="black", label="Validación")
# plt.plot(X_linspace, model.predict(X_linspace_poly), color="tomato", label="Modelo")
# plt.ylim(y_range_start, y_range_stop)
# plt.title("Conjunto de Validación")

def plot_err(feature_resultados,Err,tipo,c):
    if tipo=='P':
        grados_ = feature_resultados.Degree.unique()
        cc=0
        for g in grados_:
            sub_df = feature_resultados[feature_resultados.Degree==g]
            if Err=='T':
                plt.plot(sub_df.Alpha,sub_df.TrainMSE,color=c[cc],label='Grado: '+str(g))
            elif Err=='V':
                plt.plot(sub_df.Alpha,sub_df.ValMSE,color=c[cc],label='Grado: '+str(g))
            cc+=1
        plt.legend(loc='lower right',frameon=False,ncol=3)
    elif tipo=='L':
        if Err=='T':
            plt.plot(feature_resultados.Alpha,feature_resultados.TrainMSE,color=c)
        elif Err=='V':
            plt.plot(feature_resultados.Alpha,feature_resultados.ValMSE,color=c)
    plt.xlabel('alpha')
    plt.ylabel('ECM')

def plot_err_g(feature_resultados,Err,c):

```

```

    if Err=='T':
        plt.plot(feature_resultados.Degree,feature_resultados.TrainMSE,color=c,label=
    elif Err=='V':
        plt.plot(feature_resultados.Degree,feature_resultados.ValMSE,color=c,label='V
plt.xlabel('Degree')
plt.ylabel('ECM')
plt.legend(frameon=False,loc='upper left')

```

```
In [26]: boston_data = load_boston()
```

```
In [27]: regresion_lineal_ECM = pd.DataFrame(columns=['Feature','Regularization','Alpha','Train
regresion_polynomial_ECM = pd.DataFrame(columns=['Feature','Degree','Regularization',
```

```

shuff_data = np.random.permutation(506)
shuff_train = shuff_data[:400]
shuff_val = shuff_data[400:]

X_train = boston_data['data'][shuff_train]
X_val = boston_data['data'][shuff_val]

y_train = boston_data['target'][shuff_train]
y_val = boston_data['target'][shuff_val]

feature_map = {feature: idx for idx, feature in enumerate(boston_data['feature_names'])

features = boston_data['feature_names']
categorical_features = ['CHAS','RAD','MEDV']
non_categorical_features = set(features)-set(categorical_features)

#alpha = np.arange(0.1, 1.1, 0.1)
grado = np.arange(1,10,1)
alpha = np.arange(1e-5,1e3,10)

cmap = mp.cm.get_cmap('Spectral')
color = []
for i in range(len(grado)):
    color.append(cmap(0.1*i))

for feature in non_categorical_features:
    feature_col = feature_map[feature]
    X_train_feature = X_train[:, feature_col].reshape(-1, 1)
    X_val_feature = X_val[:, feature_col].reshape(-1, 1)

    # regresion lineal
    ## sin regularizacion
    row = regresion_lineal(X_train_feature,y_train,X_val_feature,y_val,False,0,feature

```

```

regresion_lineal_ECM = regresion_lineal_ECM.append(pd.Series(row, index=regresion.
## con regularizacion
for i in alpha:
    row = regresion_lineal(X_train_feature,y_train,X_val_feature,y_val,True,i,fea
    regresion_lineal_ECM = regresion_lineal_ECM.append(pd.Series(row, index=regre

# regresion polinomial
## sin regularizacion
for g in grado:
    row = regresion_polinomial(X_train_feature,y_train,X_val_feature,y_val,g,False)
    regresion_polinomial_ECM = regresion_polinomial_ECM.append(pd.Series(row, ind
## con regularizacion
    for i in alpha:
        row = regresion_polinomial(X_train_feature,y_train,X_val_feature,y_val,g,
        regresion_polinomial_ECM = regresion_polinomial_ECM.append(pd.Series(row,

```

A continuación se exponen gráficos de regresión lineal con y sin regularización, de polinomios de distintos grados, para cada una de las variables no categóricas del dataset. A su vez se muestra la evolución del error cuadrático medio del conjunto de entrenamiento y del de validación, en función del grado del polinomio. Por otro lado se muestra, en color verde agua, que los distintos métodos, ya sea polynomial feature o la regresión directa sin pasar por el método anterior, llevan al mismo resultado cuando se trata de ajustar una recta. El punto verde agua representa el error cuando se hace la regresión directa, para un polinomio de grado 1, y la recta verde agua ancha, representa el resultado de dicho método. El resto de las curvas se corresponden con resultados de pasar primero por polynomial features y luego por la regresión.

```

In [31]: diferencias=[]
for feature in non_categorical_features:
    print(feature)
    feature_col = feature_map[feature]
    X_train_feature = X_train[:, feature_col].reshape(-1, 1)
    X_val_feature = X_val[:, feature_col].reshape(-1, 1)

    plt.figure(figsize=(20,10))

    feature_resultadosl = regresion_lineal_ECM[(regresion_lineal_ECM.Feature==feature,
                                                (regresion_lineal_ECM.Regularization==1)]
    punto_err_metodo_lineal = [1,feature_resultadosl.TrainMSE.values[0]]

    #plt.subplot(421)

    feature_resultados = regresion_polinomial_ECM[(regresion_polinomial_ECM.Feature==feature,
                                                    (regresion_polinomial_ECM.Regularization==1)]

    plt.subplot(221)
    plot_fit(feature_resultadosl,'L',color[-1],lw=10)
    plot_fit(feature_resultados,'P',color,lw=1)

```

```

plt.xlabel(feature)
plt.title('REGRESION POLINOMIAL SIN REGULAIZACION')

feature_resultados = regresion_polinomial_ECM[(regresion_polinomial_ECM.Alpha==0)
                                                (regresion_polinomial_ECM.Feature==feature)]

diferencias.append(punto_err_metodo_lineal[1]-feature_resultados.TrainMSE.values[0])

plt.subplot(222)
plot_err_g(feature_resultados, 'T', color[0])
plot_err_g(feature_resultados, 'V', color[2])
plt.scatter(punto_err_metodo_lineal[0], punto_err_metodo_lineal[1], color=color[-1])
plt.legend(frameon=False, loc='upper right')
plt.xlabel('GRADO')

feature_resultados = regresion_lineal_ECM[(regresion_lineal_ECM.Feature==feature)
                                           (regresion_lineal_ECM.Regularization==True)]

#plt.subplot(323)
#plot_fit(feature_resultados, 'L', color[0], lw=1)
#plt.subplot(324)
#plot_err(feature_resultados, 'L', color[0])

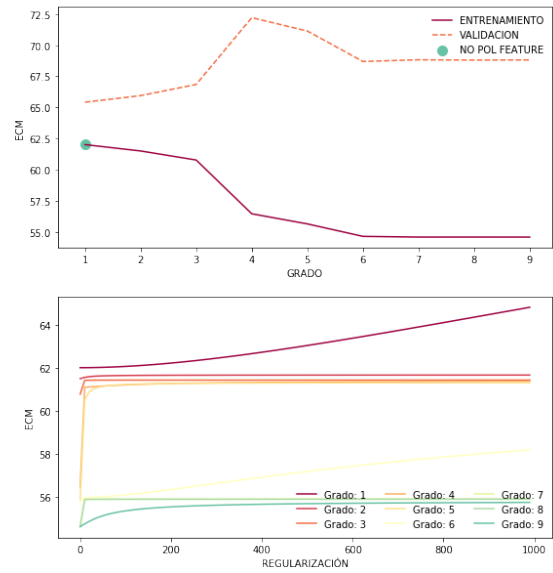
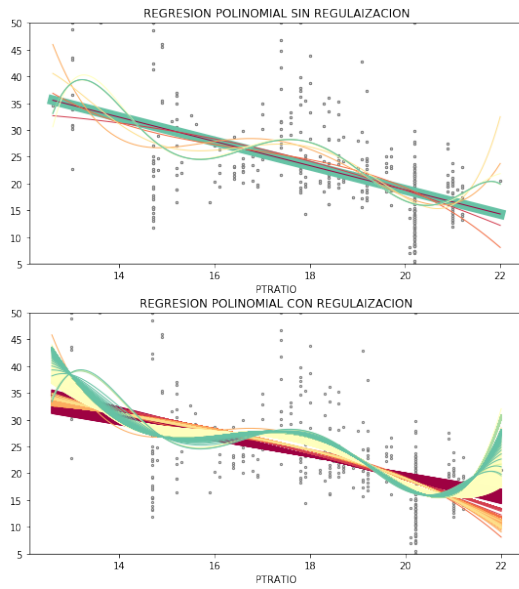
feature_resultados = regresion_polinomial_ECM[(regresion_polinomial_ECM.Feature==feature)
                                                (regresion_polinomial_ECM.Regularization==True)]

plt.subplot(223)
plot_fit(feature_resultados, 'P', color, lw=1)
plt.xlabel(feature)
plt.title('REGRESION POLINOMIAL CON REGULAIZACION')
plt.subplot(224)
plot_err(feature_resultados, 'T', 'P', color)
plt.xlabel('REGULARIZACIÓN')
#plot_err(feature_resultados, 'V', 'P', color)

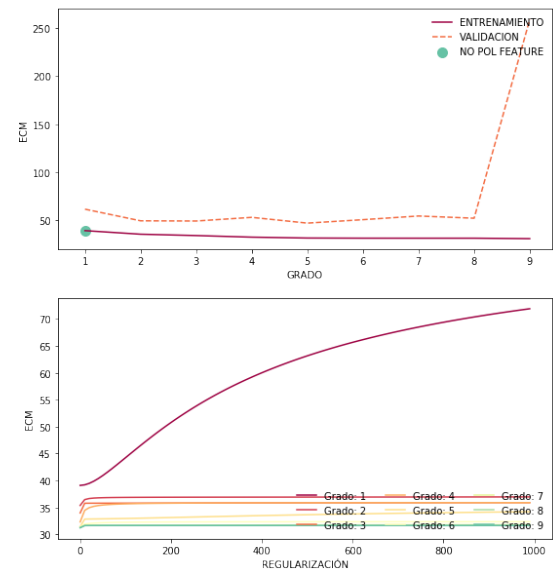
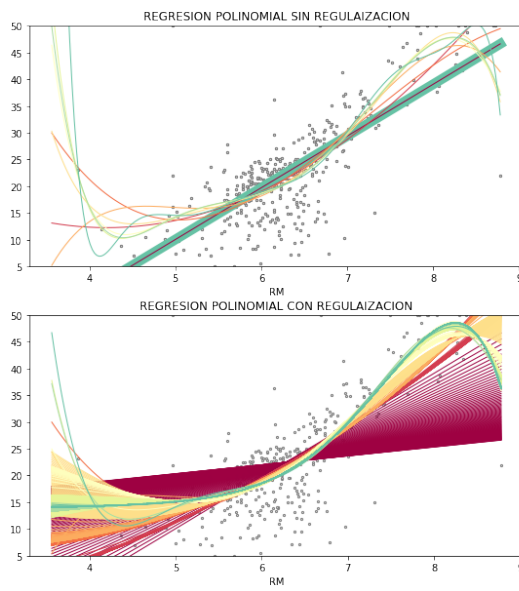
plt.show()

```

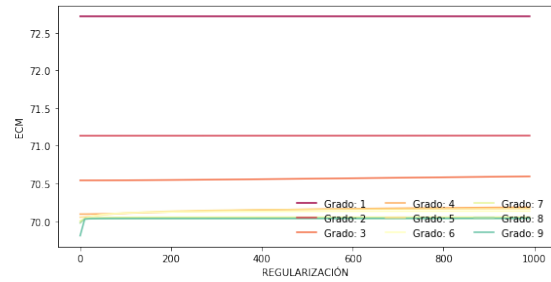
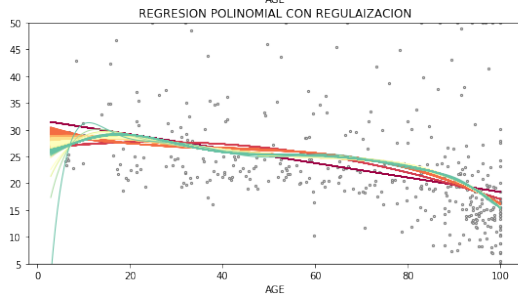
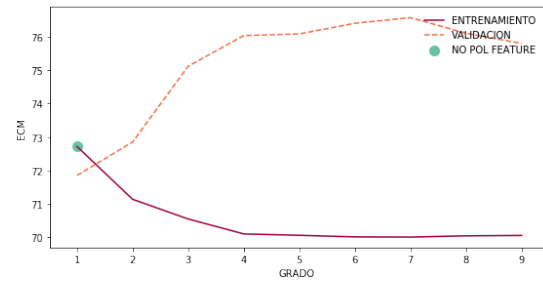
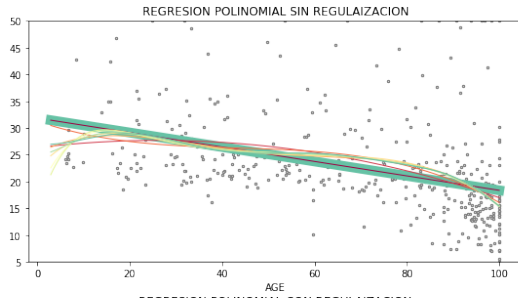
PTRATIO



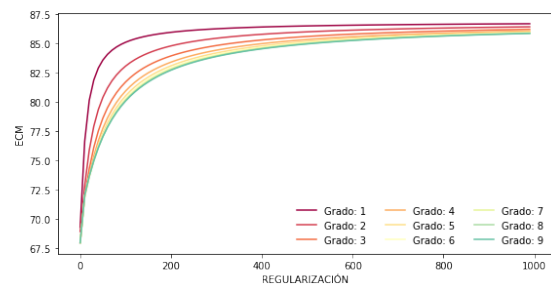
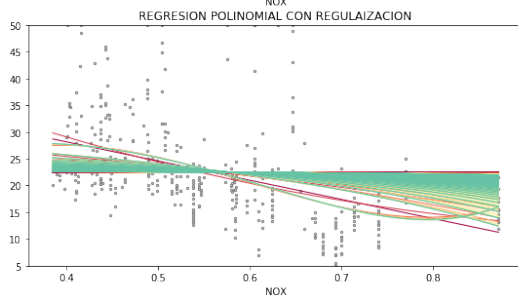
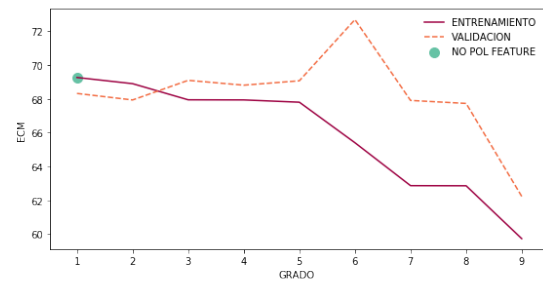
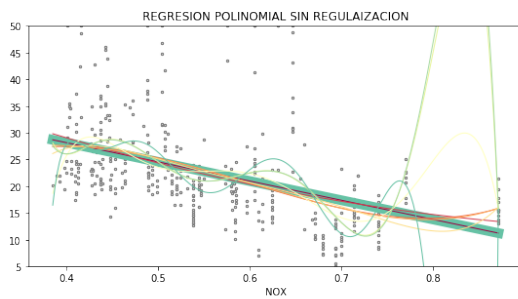
RM



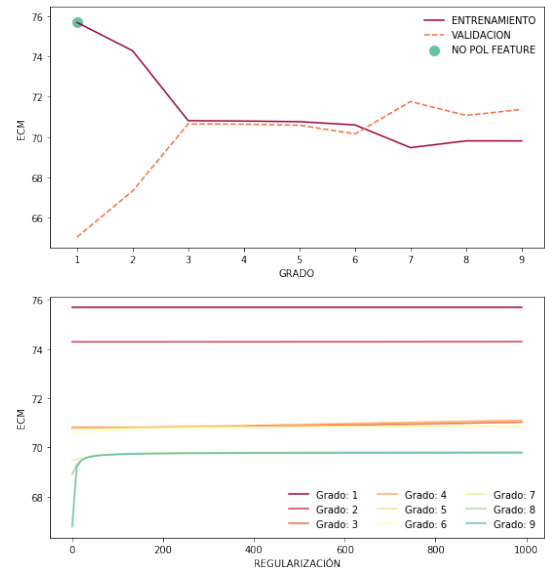
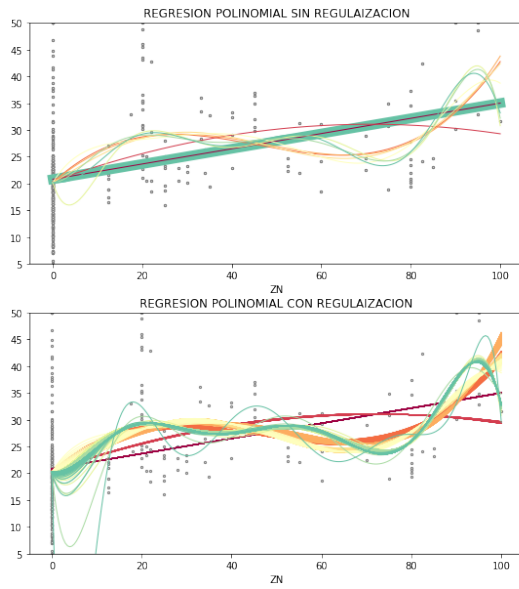
AGE



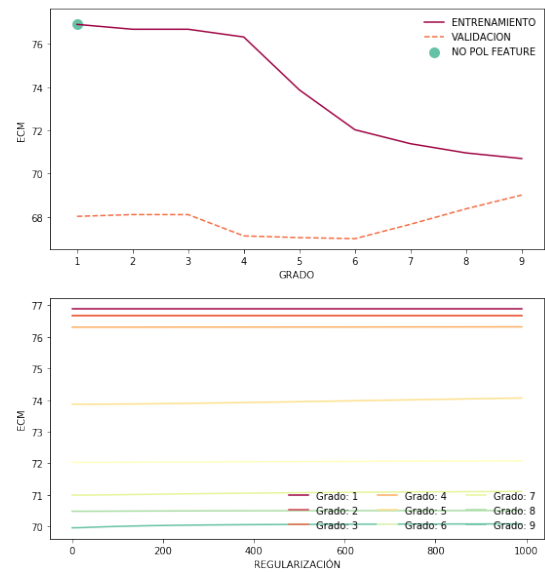
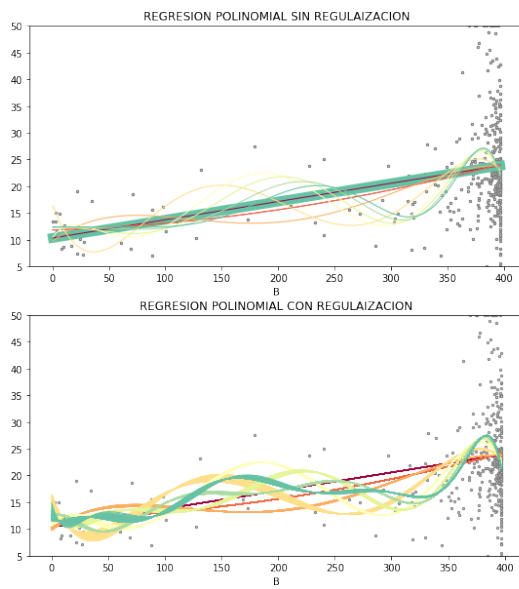
NOX



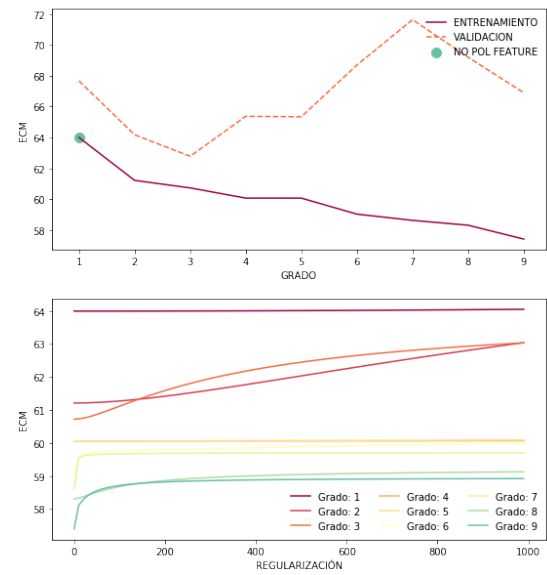
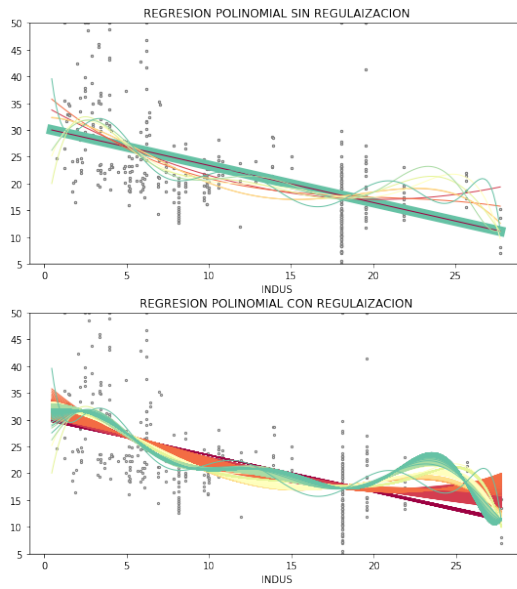
ZN



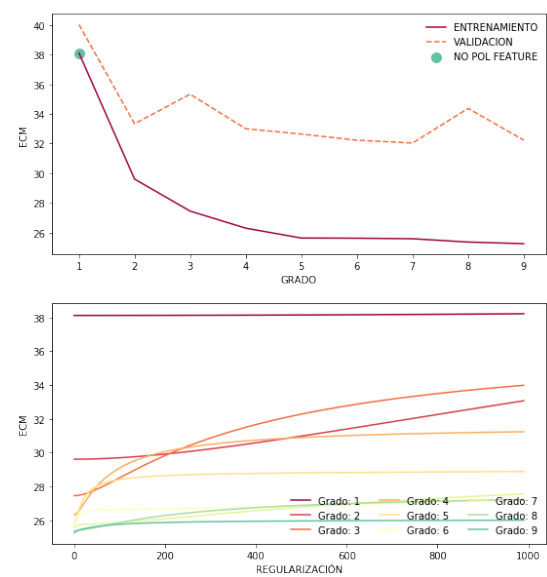
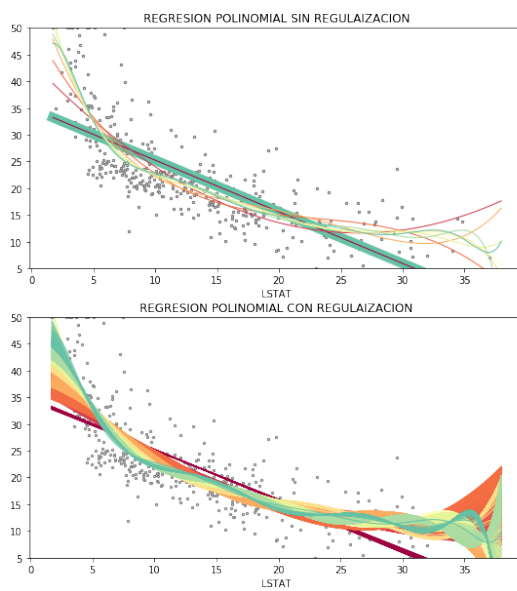
B



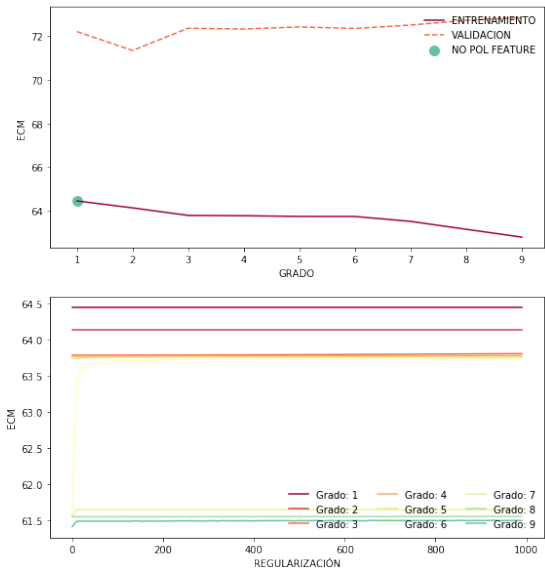
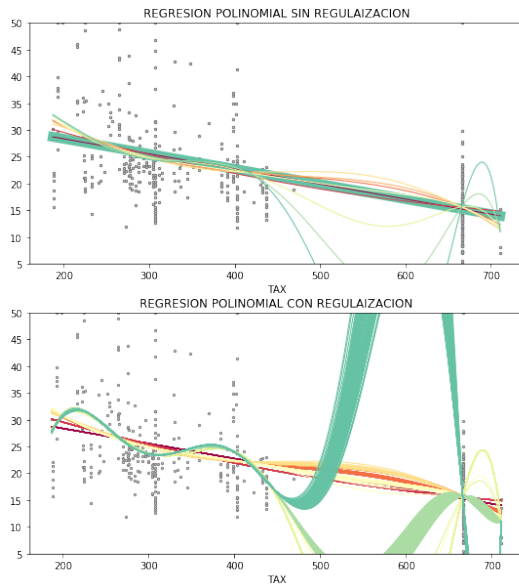
INDUS



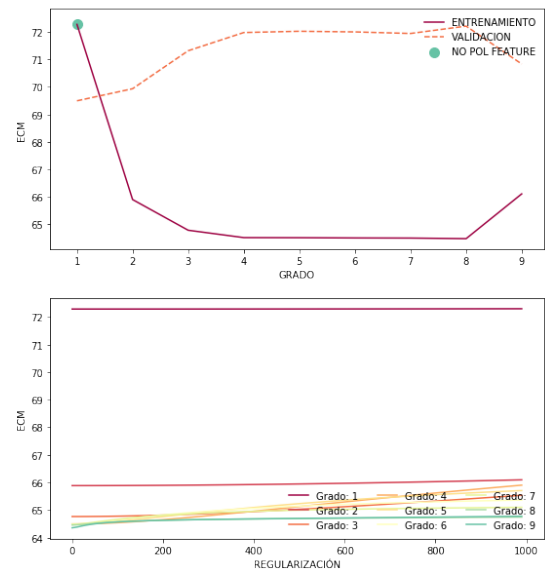
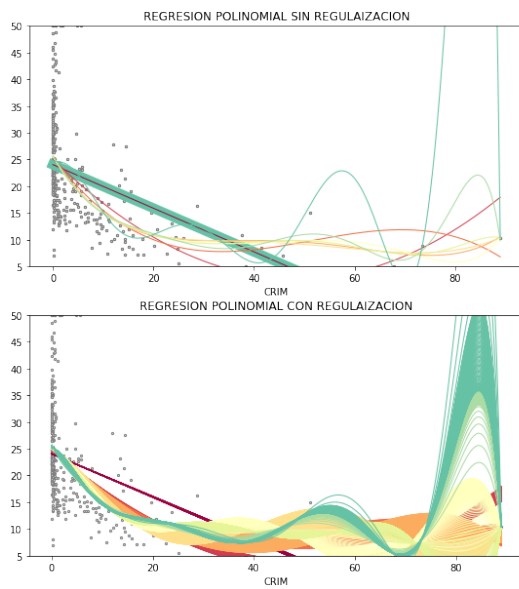
LSTAT



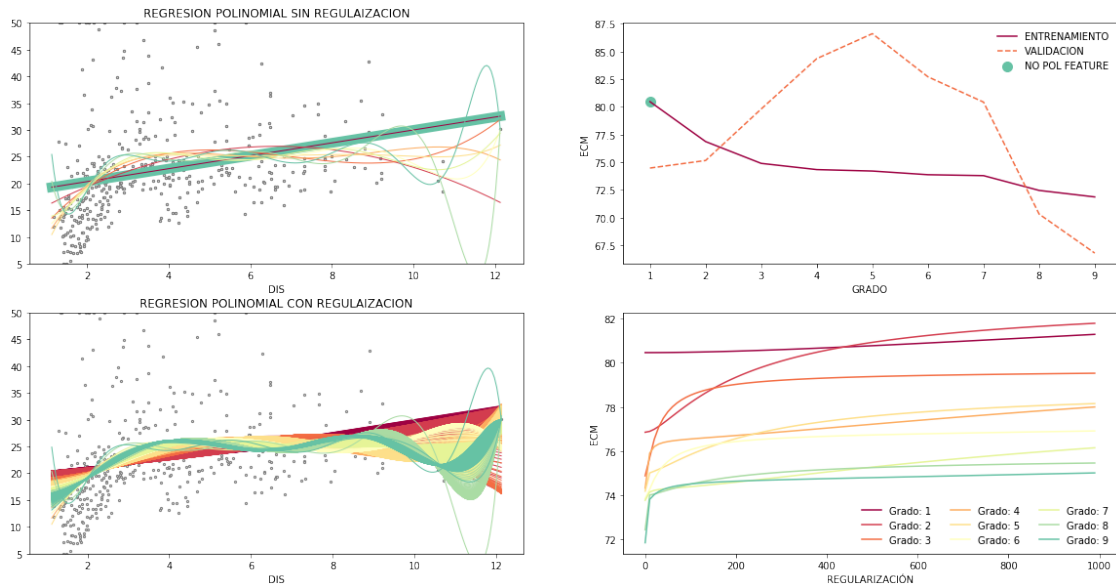
TAX



CRIM



DIS



A grandes rasgos se observa: 1. el error por lo general disminuye a medida que aumenta el grado 2. a medida que el grado aumenta, se observa como el overfitting genera grandes amplitudes 3. para grados grandes la regularización no es buena 4. el error en función de la regularización disminuye mientras el grado crece y crece mientras el valor de alpha (lambda en la teoría) crece

3 CLASIFICACION BINARIA

```
In [25]: breast_cancer_data = load_breast_cancer()
```

```
In [26]: # Utilizamos aproximadamente 80% de los datos para entrenamiento y 20% para validación
shuff_data = np.random.permutation(569)
shuff_train = shuff_data[:400]
shuff_val = shuff_data[400:]
```

```
X_train = breast_cancer_data['data'][shuff_train]
X_val = breast_cancer_data['data'][shuff_val]
```

```
y_train = breast_cancer_data['target'][shuff_train]
y_val = breast_cancer_data['target'][shuff_val]
```

```
data = pd.DataFrame(breast_cancer_data['data'], columns=breast_cancer_data['feature_names'])
data['cancer_label'] = breast_cancer_data['target']
data.describe()
```

```
Out[26]:
```

	mean radius	mean texture	mean perimeter	mean area \
count	569.000000	569.000000	569.000000	569.000000

mean	14.127292	19.289649	91.969033	654.889104
std	3.524049	4.301036	24.298981	351.914129
min	6.981000	9.710000	43.790000	143.500000
25%	11.700000	16.170000	75.170000	420.300000
50%	13.370000	18.840000	86.240000	551.100000
75%	15.780000	21.800000	104.100000	782.700000
max	28.110000	39.280000	188.500000	2501.000000

	mean smoothness	mean compactness	mean concavity	mean concave points \
count	569.000000	569.000000	569.000000	569.000000
mean	0.096360	0.104341	0.088799	0.048919
std	0.014064	0.052813	0.079720	0.038803
min	0.052630	0.019380	0.000000	0.000000
25%	0.086370	0.064920	0.029560	0.020310
50%	0.095870	0.092630	0.061540	0.033500
75%	0.105300	0.130400	0.130700	0.074000
max	0.163400	0.345400	0.426800	0.201200

	mean symmetry	mean fractal dimension	... worst texture \
count	569.000000	569.000000	... 569.000000
mean	0.181162	0.062798	... 25.677223
std	0.027414	0.007060	... 6.146258
min	0.106000	0.049960	... 12.020000
25%	0.161900	0.057700	... 21.080000
50%	0.179200	0.061540	... 25.410000
75%	0.195700	0.066120	... 29.720000
max	0.304000	0.097440	... 49.540000

	worst perimeter	worst area	worst smoothness	worst compactness \
count	569.000000	569.000000	569.000000	569.000000
mean	107.261213	880.583128	0.132369	0.254265
std	33.602542	569.356993	0.022832	0.157336
min	50.410000	185.200000	0.071170	0.027290
25%	84.110000	515.300000	0.116600	0.147200
50%	97.660000	686.500000	0.131300	0.211900
75%	125.400000	1084.000000	0.146000	0.339100
max	251.200000	4254.000000	0.222600	1.058000

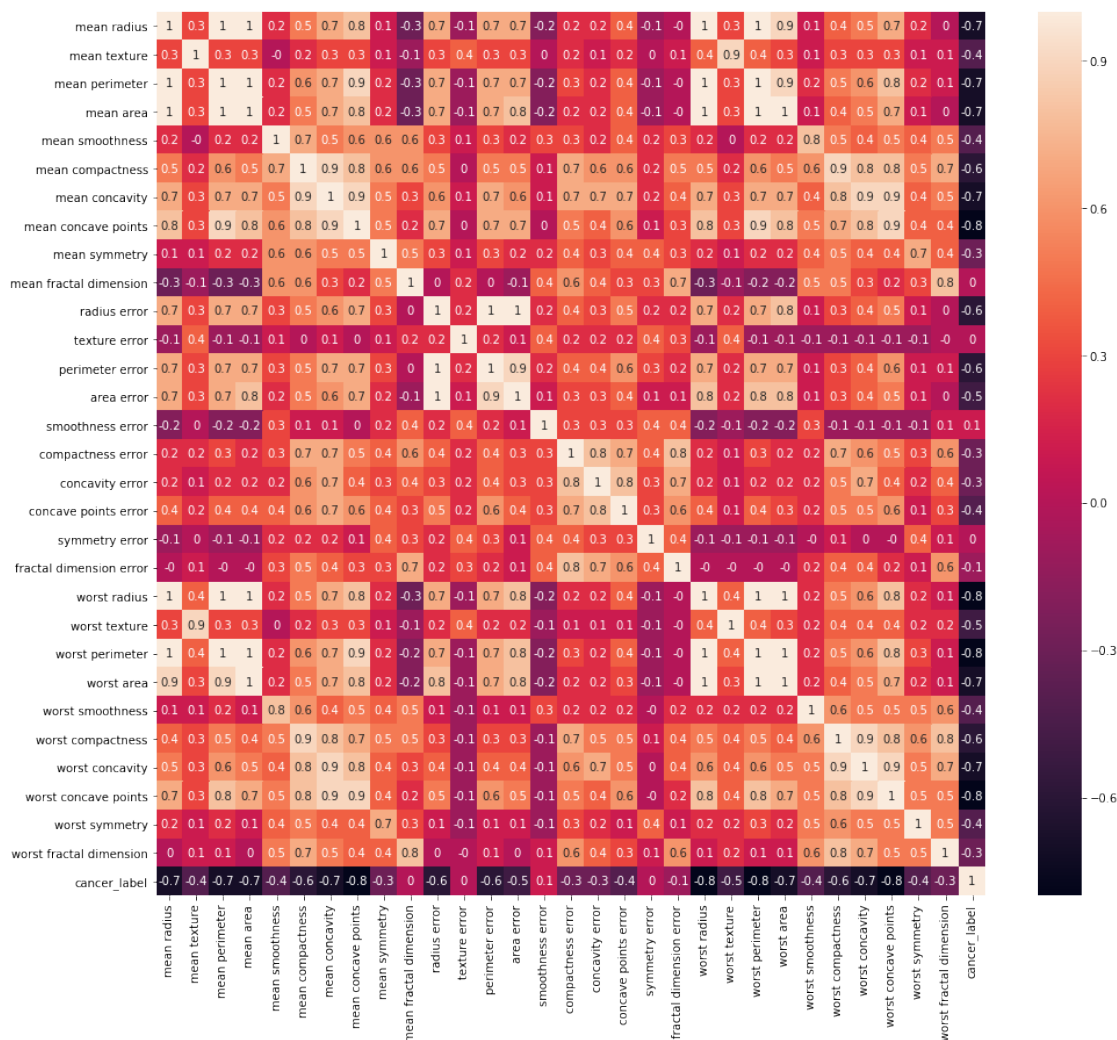
	worst concavity	worst concave points	worst symmetry \
count	569.000000	569.000000	569.000000
mean	0.272188	0.114606	0.290076
std	0.208624	0.065732	0.061867
min	0.000000	0.000000	0.156500
25%	0.114500	0.064930	0.250400
50%	0.226700	0.099930	0.282200
75%	0.382900	0.161400	0.317900
max	1.252000	0.291000	0.663800

	worst fractal dimension	cancer_label
count	569.000000	569.000000
mean	0.083946	0.627417
std	0.018061	0.483918
min	0.055040	0.000000
25%	0.071460	0.000000
50%	0.080040	1.000000
75%	0.092080	1.000000
max	0.207500	1.000000

[8 rows x 31 columns]

```
In [27]: correlation_matrix = data.corr().round(1)
fig, ax = plt.subplots(figsize=(16,14))
sb.heatmap(data=correlation_matrix, annot = True)
```

Out[27]: <matplotlib.axes._subplots.AxesSubplot at 0x7feb426b54e0>



```

In [28]: features_original = breast_cancer_data['feature_names']

In [34]: features = []
         for f in features_original:
             if 'error' in f: continue
             features.append(f)

In [35]: features

Out[35]: ['mean radius',
          'mean texture',
          'mean perimeter',
          'mean area',
          'mean smoothness',
          'mean compactness',
          'mean concavity',
          'mean concave points',
          'mean symmetry',
          'mean fractal dimension',
          'worst radius',
          'worst texture',
          'worst perimeter',
          'worst area',
          'worst smoothness',
          'worst compactness',
          'worst concavity',
          'worst concave points',
          'worst symmetry',
          'worst fractal dimension']

In [45]: feature_map = {feature: idx for idx, feature in enumerate(features)}
         e4 = 'verdaderos_positivos'
         e3 = 'falsos_positivos'
         e2 = 'falsos_negativos'
         e1 = 'verdaderos_negativos'

         penalties = ['None', 'l1', 'l2', 'elasticnet']

         iterations = [10, 100, 1000]
         alphas = [1e-5, 1e-3, 1e-1, 1e1]
         columnas = ['fx', 'fy', 'penalty', 'alpha', 'iter', 'ee', 'ev', e1, e2, e3, e4]
         dtCB = pd.DataFrame(columns=columnas)

         for fx in features:
             x_feature = fx

```

```

print('\n\n\n\n\n***** ',x_feature)
for fy in list(set(features)-set([fx])):
    print('-',fy)
    y_feature = fy

    x_feature_col = feature_map[x_feature]
    y_feature_col = feature_map[y_feature]
    X_train_feature = X_train[:, [x_feature_col, y_feature_col]]
    X_val_feature = X_val[:, [x_feature_col, y_feature_col]]

    for p in penalties:
        penalty = p
        for a in alphas:
            alpha = a
            for it in iterations:
                max_iter = it

                model = Perceptron(penalty=penalty, alpha=alpha, max_iter=max_iter)
                model.fit(X_train_feature, y_train)

                ee=accuracy_score(y_train, model.predict(X_train_feature))
                ev=accuracy_score(y_val, model.predict(X_val_feature))
                #print('Exactitud para entrenamiento: %.2f' % ee)
                #print('Exactitud para validación: %.2f' % ev)

                matriz = confusion_matrix(y_train, model.predict(X_train_feature))

                dtCB = dtCB.append({'fx':fx,'fy':fy,'penalty':penalty,'alpha':alpha,
                                   'ee':ee,'ev':ev,e1:matriz[0][0],e2:matriz[0][1],
                                   e3:matriz[1][0],e4:matriz[1][1]},ignore_index=True)

```

```

***** mean radius
- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- worst radius
- worst compactness
- worst concavity
- worst perimeter

```


- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean texture

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean perimeter

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

- ***** mean area
- mean smoothness
 - worst symmetry
 - mean compactness
 - worst texture
 - mean perimeter
 - worst concave points
 - mean concavity
 - mean fractal dimension
 - mean radius
 - worst radius
 - worst compactness
 - worst concavity
 - worst perimeter
 - mean symmetry
 - mean texture
 - worst smoothness
 - worst area
 - worst fractal dimension
 - mean concave points

- ***** mean smoothness
- worst symmetry
 - mean compactness
 - worst texture
 - mean perimeter
 - worst concave points
 - mean concavity
 - mean fractal dimension
 - mean radius
 - worst radius
 - worst compactness
 - worst concavity
 - worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean compactness

- mean smoothness
- worst symmetry
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean concavity

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean concave points

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean area

***** mean symmetry

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity

- worst perimeter
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** mean fractal dimension

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst radius

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst compactness
- worst concavity
- worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst texture

- mean smoothness
- worst symmetry
- mean compactness
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst perimeter

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst area

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst fractal dimension
- mean concave points
- mean area

***** worst smoothness

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity

- worst perimeter
- mean symmetry
- mean texture
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst compactness

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst concavity

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst concave points

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst symmetry

- mean smoothness
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter

- mean symmetry
- mean texture
- worst smoothness
- worst area
- worst fractal dimension
- mean concave points
- mean area

***** worst fractal dimension

- mean smoothness
- worst symmetry
- mean compactness
- worst texture
- mean perimeter
- worst concave points
- mean concavity
- mean fractal dimension
- mean radius
- worst radius
- worst compactness
- worst concavity
- worst perimeter
- mean symmetry
- mean texture
- worst smoothness
- worst area
- mean concave points
- mean area

In [46]: dtCB

Out[46]:

	fx	fy	penalty	alpha	iter	\
0	mean radius	mean smoothness	None	0.00001	10	
1	mean radius	mean smoothness	None	0.00001	100	
2	mean radius	mean smoothness	None	0.00001	1000	
3	mean radius	mean smoothness	None	0.00100	10	
4	mean radius	mean smoothness	None	0.00100	100	
5	mean radius	mean smoothness	None	0.00100	1000	
6	mean radius	mean smoothness	None	0.10000	10	
7	mean radius	mean smoothness	None	0.10000	100	
8	mean radius	mean smoothness	None	0.10000	1000	
9	mean radius	mean smoothness	None	10.00000	10	
10	mean radius	mean smoothness	None	10.00000	100	
11	mean radius	mean smoothness	None	10.00000	1000	

12		mean radius	mean smoothness	11	0.00001	10
13		mean radius	mean smoothness	11	0.00001	100
14		mean radius	mean smoothness	11	0.00001	1000
15		mean radius	mean smoothness	11	0.00100	10
16		mean radius	mean smoothness	11	0.00100	100
17		mean radius	mean smoothness	11	0.00100	1000
18		mean radius	mean smoothness	11	0.10000	10
19		mean radius	mean smoothness	11	0.10000	100
20		mean radius	mean smoothness	11	0.10000	1000
21		mean radius	mean smoothness	11	10.00000	10
22		mean radius	mean smoothness	11	10.00000	100
23		mean radius	mean smoothness	11	10.00000	1000
24		mean radius	mean smoothness	12	0.00001	10
25		mean radius	mean smoothness	12	0.00001	100
26		mean radius	mean smoothness	12	0.00001	1000
27		mean radius	mean smoothness	12	0.00100	10
28		mean radius	mean smoothness	12	0.00100	100
29		mean radius	mean smoothness	12	0.00100	1000
...	
18210	worst fractal dimension	mean area		11	0.10000	10
18211	worst fractal dimension	mean area		11	0.10000	100
18212	worst fractal dimension	mean area		11	0.10000	1000
18213	worst fractal dimension	mean area		11	10.00000	10
18214	worst fractal dimension	mean area		11	10.00000	100
18215	worst fractal dimension	mean area		11	10.00000	1000
18216	worst fractal dimension	mean area		12	0.00001	10
18217	worst fractal dimension	mean area		12	0.00001	100
18218	worst fractal dimension	mean area		12	0.00001	1000
18219	worst fractal dimension	mean area		12	0.00100	10
18220	worst fractal dimension	mean area		12	0.00100	100
18221	worst fractal dimension	mean area		12	0.00100	1000
18222	worst fractal dimension	mean area		12	0.10000	10
18223	worst fractal dimension	mean area		12	0.10000	100
18224	worst fractal dimension	mean area		12	0.10000	1000
18225	worst fractal dimension	mean area		12	10.00000	10
18226	worst fractal dimension	mean area		12	10.00000	100
18227	worst fractal dimension	mean area		12	10.00000	1000
18228	worst fractal dimension	mean area	elasticnet	0.00001	10	
18229	worst fractal dimension	mean area	elasticnet	0.00001	100	
18230	worst fractal dimension	mean area	elasticnet	0.00001	1000	
18231	worst fractal dimension	mean area	elasticnet	0.00100	10	
18232	worst fractal dimension	mean area	elasticnet	0.00100	100	
18233	worst fractal dimension	mean area	elasticnet	0.00100	1000	
18234	worst fractal dimension	mean area	elasticnet	0.10000	10	
18235	worst fractal dimension	mean area	elasticnet	0.10000	100	
18236	worst fractal dimension	mean area	elasticnet	0.10000	1000	
18237	worst fractal dimension	mean area	elasticnet	10.00000	10	
18238	worst fractal dimension	mean area	elasticnet	10.00000	100	

18239 worst fractal dimension mean area elasticnet 10.00000 1000

	ee	ev	verdaderos_negativos	falsos_negativos	\
0	0.3675	0.390533	146	0	
1	0.7750	0.781065	138	8	
2	0.8350	0.863905	130	16	
3	0.3675	0.390533	146	0	
4	0.7750	0.781065	138	8	
5	0.8350	0.863905	130	16	
6	0.3675	0.390533	146	0	
7	0.7750	0.781065	138	8	
8	0.8350	0.863905	130	16	
9	0.3675	0.390533	146	0	
10	0.7750	0.781065	138	8	
11	0.8350	0.863905	130	16	
12	0.3675	0.390533	146	0	
13	0.7750	0.781065	138	8	
14	0.8350	0.863905	130	16	
15	0.3675	0.390533	146	0	
16	0.7775	0.781065	138	8	
17	0.8775	0.905325	112	34	
18	0.3675	0.390533	146	0	
19	0.7075	0.704142	142	4	
20	0.6975	0.674556	143	3	
21	0.3650	0.390533	146	0	
22	0.3650	0.390533	146	0	
23	0.3650	0.390533	146	0	
24	0.3675	0.390533	146	0	
25	0.7775	0.781065	138	8	
26	0.8550	0.881657	96	50	
27	0.3675	0.390533	146	0	
28	0.7150	0.727811	140	6	
29	0.7125	0.727811	140	6	
...	
18210	0.3650	0.390533	146	0	
18211	0.3650	0.390533	146	0	
18212	0.3650	0.390533	146	0	
18213	0.3650	0.390533	146	0	
18214	0.6375	0.615385	1	145	
18215	0.3650	0.390533	146	0	
18216	0.3650	0.390533	146	0	
18217	0.3650	0.390533	146	0	
18218	0.8525	0.893491	120	26	
18219	0.3650	0.390533	146	0	
18220	0.3650	0.390533	146	0	
18221	0.3650	0.390533	146	0	
18222	0.3650	0.390533	146	0	
18223	0.3650	0.390533	146	0	

18224	0.3650	0.390533	146	0
18225	0.3650	0.390533	146	0
18226	0.3650	0.390533	146	0
18227	0.3650	0.390533	146	0
18228	0.3650	0.390533	146	0
18229	0.3650	0.390533	146	0
18230	0.8525	0.893491	120	26
18231	0.3650	0.390533	146	0
18232	0.3650	0.390533	146	0
18233	0.3650	0.390533	146	0
18234	0.3650	0.390533	146	0
18235	0.3650	0.390533	146	0
18236	0.3650	0.390533	146	0
18237	0.3650	0.390533	146	0
18238	0.3650	0.390533	146	0
18239	0.3650	0.390533	146	0

	falsos_positivos	verdaderos_positivos
0	253	1
1	82	172
2	50	204
3	253	1
4	82	172
5	50	204
6	253	1
7	82	172
8	50	204
9	253	1
10	82	172
11	50	204
12	253	1
13	82	172
14	50	204
15	253	1
16	81	173
17	15	239
18	253	1
19	113	141
20	118	136
21	254	0
22	254	0
23	254	0
24	253	1
25	81	173
26	8	246
27	253	1
28	108	146
29	109	145

...
18210	254	0
18211	254	0
18212	254	0
18213	254	0
18214	0	254
18215	254	0
18216	254	0
18217	254	0
18218	33	221
18219	254	0
18220	254	0
18221	254	0
18222	254	0
18223	254	0
18224	254	0
18225	254	0
18226	254	0
18227	254	0
18228	254	0
18229	254	0
18230	33	221
18231	254	0
18232	254	0
18233	254	0
18234	254	0
18235	254	0
18236	254	0
18237	254	0
18238	254	0
18239	254	0

[18240 rows x 11 columns]

```
In [47]: dtCB.to_csv(r'clasificacion_cancer.csv', index=None, header=True, sep='|')
```

```
In [184]: fig=plt.figure(figsize=(15,4))
```

```
fx = 'mean radius'
```

```
fy = 'mean concave points'
```

```
fig.suptitle('FEATURES: '+fx+ ' - '+fy , fontsize=20)
```

```
j=1
```

```
ls=['--', ':', '-.', '-']
```

```
for p in penalties:
```

```
    plt.subplot(1,4,j)
```

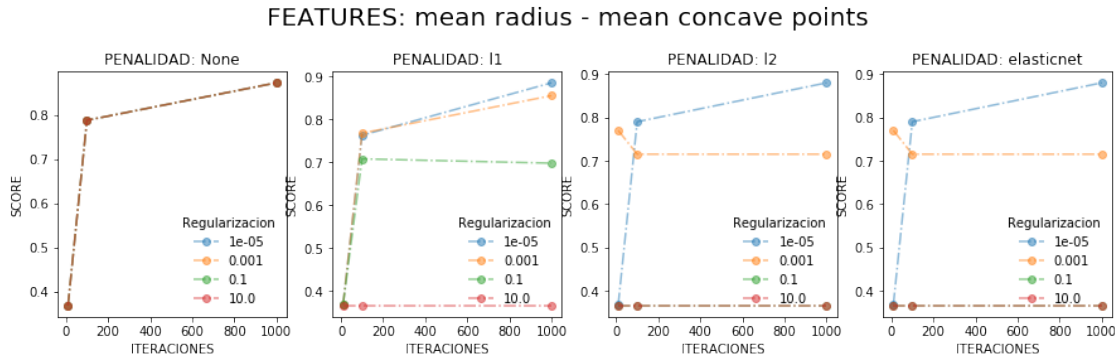
```
    sub=dtCB[(dtCB.penalty==p)&(dtCB.fx==fx)&(dtCB.fy==fy)]
```

```

for a in alphas:
    plt.plot(sub[sub.alpha==a].iter.values,sub[sub.alpha==a].ee.values,label=str(a))
plt.legend(title='Regularizacion',frameon=False,loc='lower right')
plt.title('PENALIDAD: ' + p)
plt.xlabel('ITERACIONES')
plt.ylabel('SCORE')
j+=1

plt.subplots_adjust(wspace=0.2, top=0.8)

```

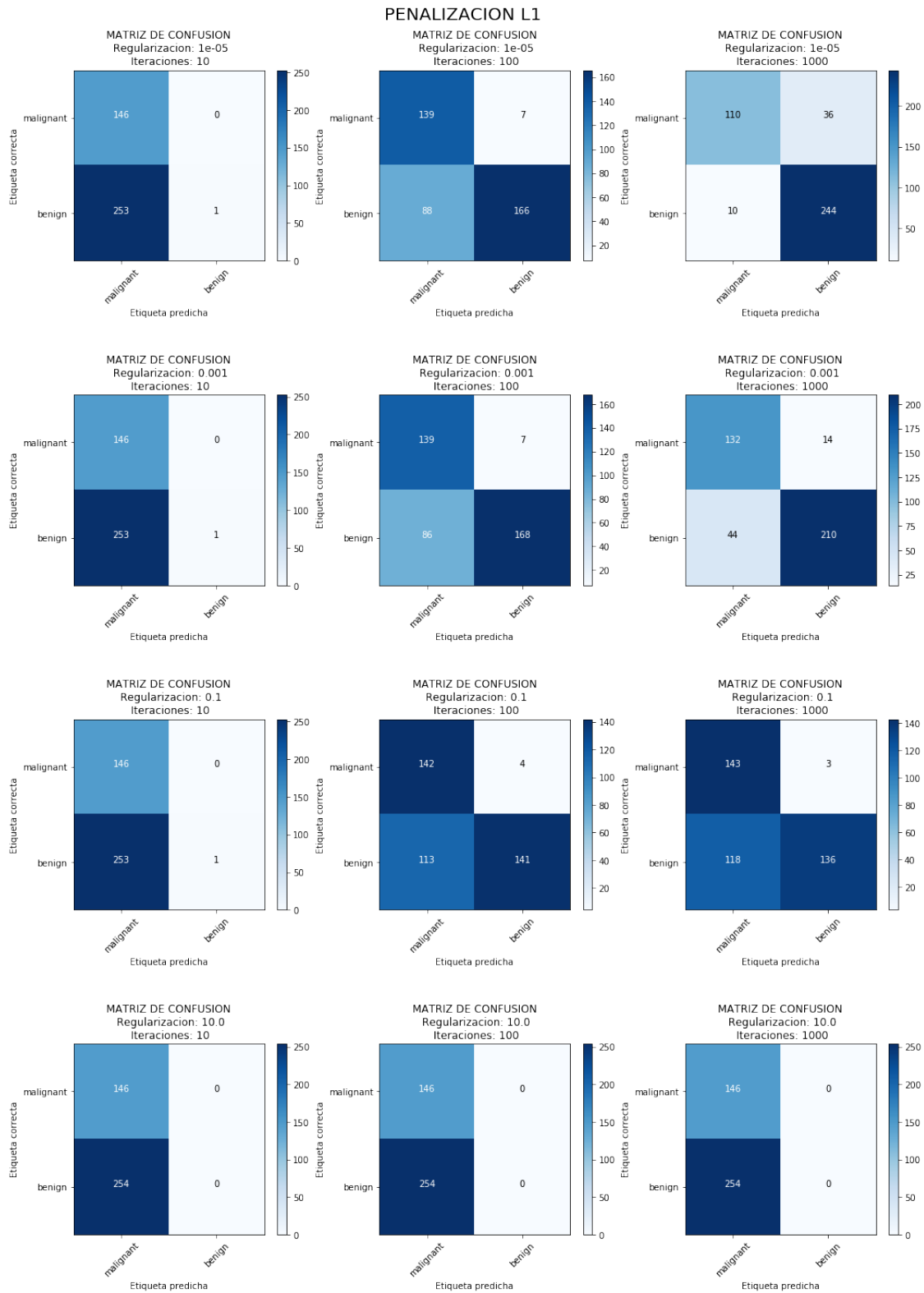


Se puede observar que la penalidad l1, es la única que permite diferenciar los distintos valores que toma el hiperparámetro de regularización para las tres números de iteraciones en estudio., por otro lado se observa que mientras menor es alpha y mayor es el número de iteraciones, la exactitud aumenta, tanto al no considerar penalidad como al tomar la penalidad l1, lo mismo no sucede para la norma al cuadrado o la combinación lineal de ambos tipos (l1,l2)

```

In [185]: sub=dtCB[(dtCB.penalty=='l1')&(dtCB.fx==fx)&(dtCB.fy==fy)]
fig = plt.figure(figsize=(15,20))
fig.suptitle('PENALIZACION L1', fontsize=20)
for i in range(len(sub)):
    plt.subplot(4,3,i+1)
    matriz=np.array([[sub.verdaderos_negativos.values[i],sub.falsos_negativos.values[i],
                      sub.falsos_positivos.values[i],sub.verdaderos_positivos.values[i]]])
    plot_confusion_matrix(matriz,classes=breast_cancer_data.target_names,
                          title='MATRIZ DE CONFUSION\n Regularizacion: '+str(sub.alpha.values[i])+'\n Iteraciones: '+str(sub.iter.values[i]))
plt.subplots_adjust( wspace=0.2, top=0.93, hspace=0.7)

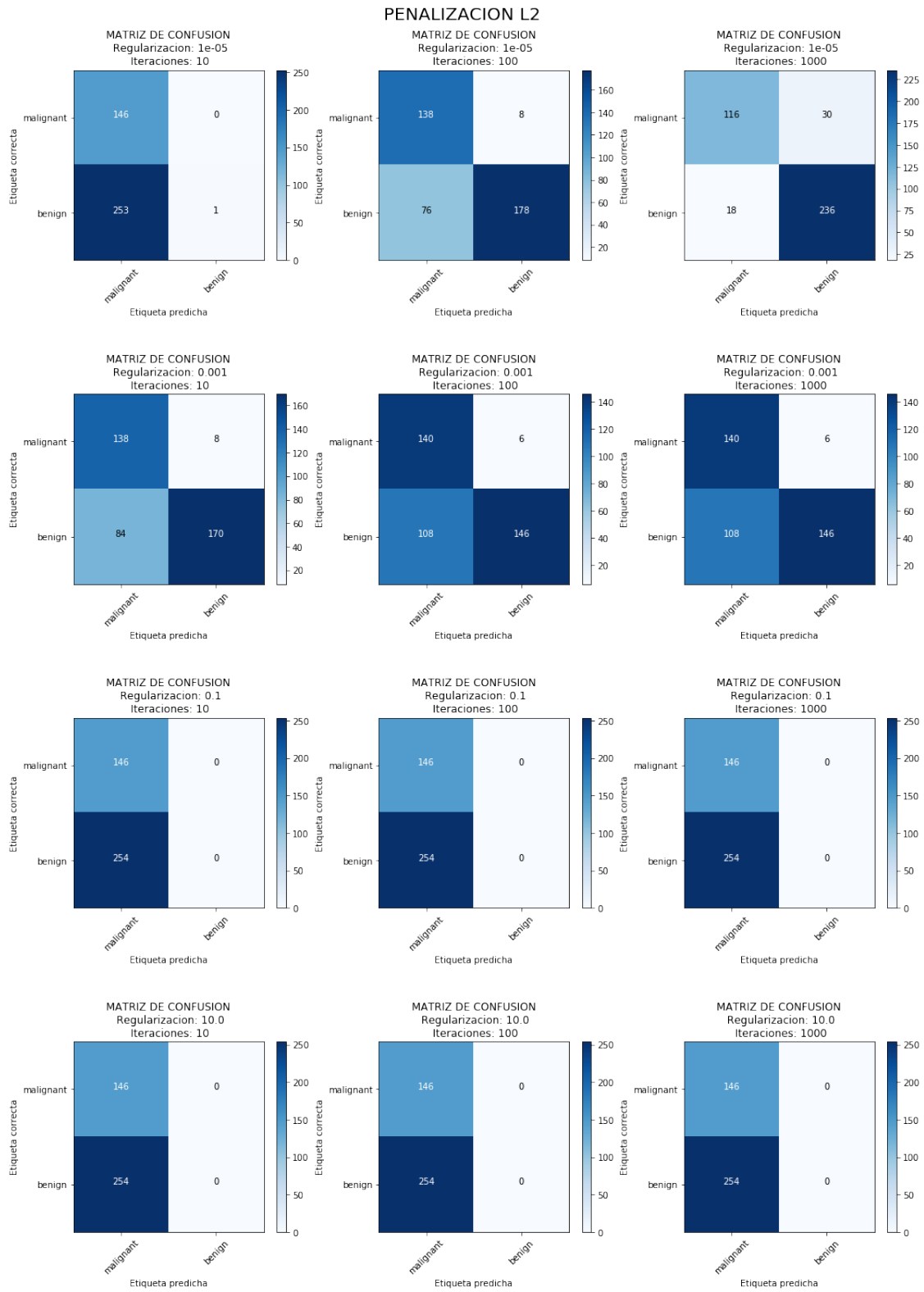
```



Lo óptimo es minimizar los casos: etiqueta true diagnóstico false, es decir, cancer positivo,

diagnóstico o predicción negativa. En el caso de penalidad L1, se puede ver que cuando alpha es muy chico, si bien se observó que la exactitud aumenta, también lo hacen los casos de falsos negativos cuando crece el número de iteraciones, lo cual no es bueno. Lo que se quiere decir, es que el análisis no es trivial y requiere de mucho estudio y de optar por la solución de compromiso más óptima. Tampoco se puede aceptar que el caso de nulidad del número de falsos negativos, pues como se puede ver en la matriz de confusión, tales predicciones conllevan una gran inexactitud, ya que no se predice ningún tumor benigno.

```
In [186]: sub=dtCB[(dtCB.penalty=='l2')&(dtCB.fx==fx)&(dtCB.fy==fy)]
fig = plt.figure(figsize=(15,20))
fig.suptitle('PENALIZACION L2', fontsize=20)
for i in range(len(sub)):
    plt.subplot(4,3,i+1)
    matriz=np.array([[sub.verdaderos_negativos.values[i],sub.falsos_negativos.values[i],
                      sub.falsos_positivos.values[i],sub.verdaderos_positivos.values[i]]])
    plot_confusion_matrix(matriz,classes=breast_cancer_data.target_names,
                          title='MATRIZ DE CONFUSION\n Regularizacion: '+str(sub.alpha.values[i])+'\n Iteraciones: '+str(sub.iterations.values[i]))
    plt.subplots_adjust( wspace=0.2, top=0.93, hspace=0.7)
```



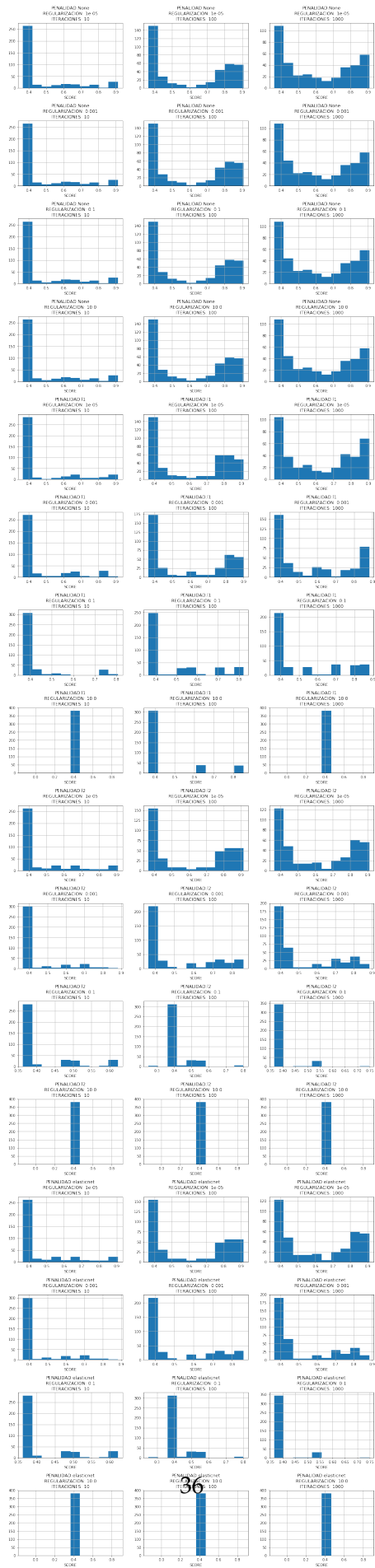
In []:

```

In [187]: fig = plt.figure(figsize=(15,20))
          j=1
          for p in penalties:
              for a in alphas:
                  for i in iterations:
                      plt.subplot(16,3,j)
                      dtCB[(dtCB.iter==i)&(dtCB.penalty==p)&(dtCB.alpha==a)].ee.hist()
                      #plt.title('MATRIZ DE CONFUSION\n Regularizacion: '+str(sub.alpha.values)
                      plt.title('PENALIDAD: '+str(p)+'\nREGULARIZACION: '+str(a)+'\nITERACIONES
                      plt.xlabel('SCORE')
                      j+=1

          plt.subplots_adjust(left=0.01, wspace=0.2, top=3, hspace=0.5)

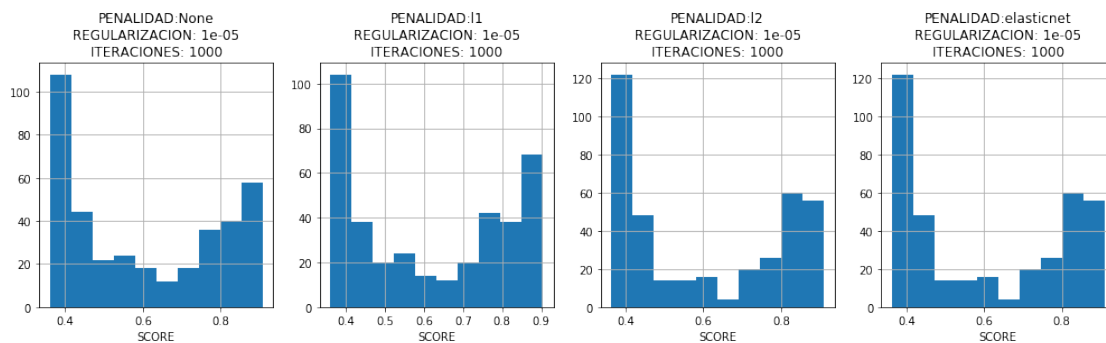
```



Los histogramas de exactitud en función de alpha, tipo de penalidad y número de iteraciones muestran: 1. la exactitud tiende a mejorar a partir de 100 iteraciones 2. la penalidad de tipo L1 parece mejor que el resto de las penalidades 3. regularizaciones más chicas implican mayor exactitud

```
In [188]: fig = plt.figure(figsize=(15,4))
          j=1
          for p in penalties:
              plt.subplot(1,4,j)
              dtCB[(dtCB.iter==1000)&(dtCB.penalty==p)&(dtCB.alpha==1e-5)].ee.hist()
              #plt.title('MATRIZ DE CONFUSION\n Regularizacion: '+str(sub.alpha.values[i])+'\n')
              plt.title('PENALIDAD: '+str(p)+'\nREGULARIZACION: '+str(1e-5)+'\nITERACIONES: '+str(1000))
              plt.xlabel('SCORE')
              j+=1
```

```
plt.subplots_adjust(left=0.01, wspace=0.2)
```



```
In [ ]:
```

```
In [191]: x_feature = fx
          y_feature = fy

          x_feature_col = feature_map[x_feature]
          y_feature_col = feature_map[y_feature]
          X_train_feature = X_train[:, [x_feature_col, y_feature_col]]
          X_val_feature = X_val[:, [x_feature_col, y_feature_col]]

          penalty = 'l1'
          alpha = 1e-5
          j=1

          fig=plt.figure(figsize=(14, 4), dpi=80, facecolor='w', edgecolor='k')
          fig.suptitle('PENALIZACION L1', fontsize=20)
```

```

for i in iterations:
    max_iter = i
    model = Perceptron(penalty=penalty, alpha=alpha, max_iter=max_iter)
    model.fit(X_train_feature, y_train)

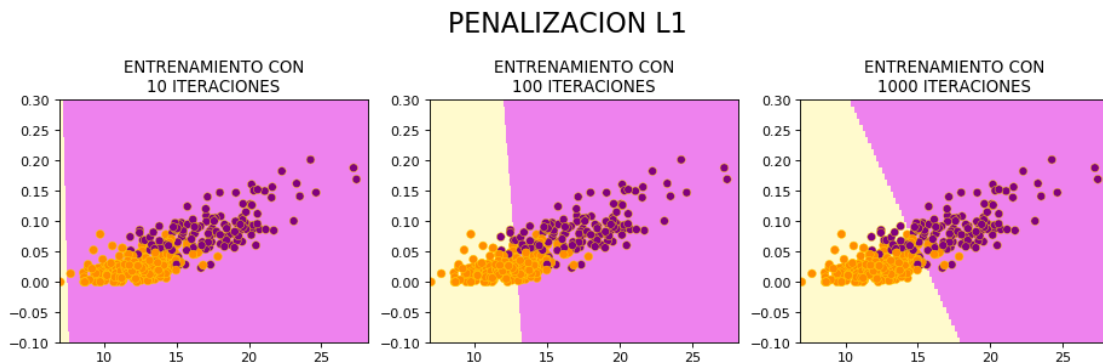
    xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature], model)

    cmap_dots = ListedColormap(['purple', 'darkorange'])
    cmap_edge = ListedColormap(['darkviolet', 'gold'])
    cmap_back = ListedColormap(['violet', 'lemonchiffon'])

    # Conjunto de entrenamiento
    plt.subplot(1, 3, j)
    plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
    plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, cmap=cmap_dots)
    plt.xlim(xx.min(), xx.max())
    plt.ylim(yy.min(), yy.max())
    plt.title('ENTRENAMIENTO CON\n'+str(i)+' ITERACIONES')

    j+=1
plt.subplots_adjust( top=0.75)

```



Se observa una gran mejora de la caracterización de la frontera a medida que aumenta el número de iteraciones, al menos, dentro de los tres valores, del número iteraciones, considerados

```

In [192]: x_feature = fx
          y_feature = fy

          x_feature_col = feature_map[x_feature]
          y_feature_col = feature_map[y_feature]
          X_train_feature = X_train[:, [x_feature_col, y_feature_col]]
          X_val_feature = X_val[:, [x_feature_col, y_feature_col]]

          penalty = 'l2'

```

```

alpha = 1e-5
j=1

fig=plt.figure(figsize=(14, 4), dpi=80, facecolor='w', edgecolor='k')
fig.suptitle('PENALIZACION L2', fontsize=20)
for i in iterations:
    max_iter = i
    model = Perceptron(penalty=penalty, alpha=alpha, max_iter=max_iter)
    model.fit(X_train_feature, y_train)

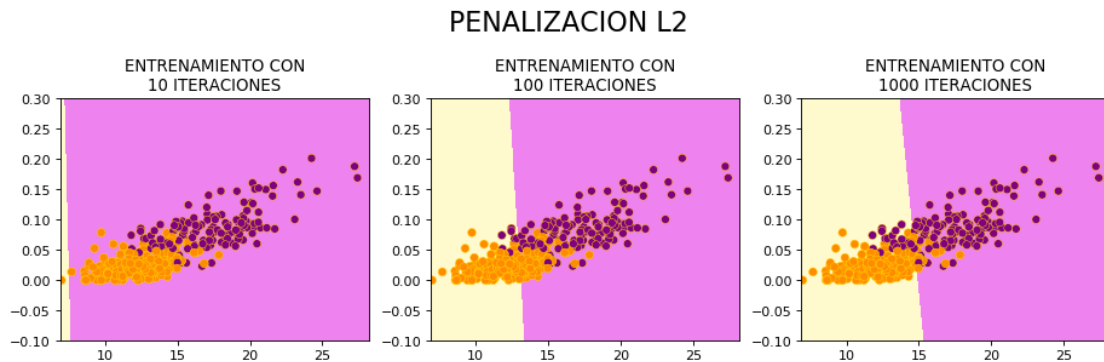
    xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature], model)

    cmap_dots = ListedColormap(['purple', 'darkorange'])
    cmap_edge = ListedColormap(['darkviolet', 'gold'])
    cmap_back = ListedColormap(['violet', 'lemonchiffon'])

    # Conjunto de entrenamiento
    plt.subplot(1, 3, j)
    plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
    plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, cmap=cmap_dots)
    plt.xlim(xx.min(), xx.max())
    plt.ylim(yy.min(), yy.max())
    plt.title('ENTRENAMIENTO CON\n'+str(i)+' ITERACIONES')

    j+=1
plt.subplots_adjust( top=0.75)

```



4 VECINOS MAS CERCANOS

```

In [209]: neighbors = [3,5,10]
          metricas = ['cosine', 'euclidean', 'manhattan']

plt.figure(figsize=(14,15), dpi= 80, facecolor='w', edgecolor='k')

```

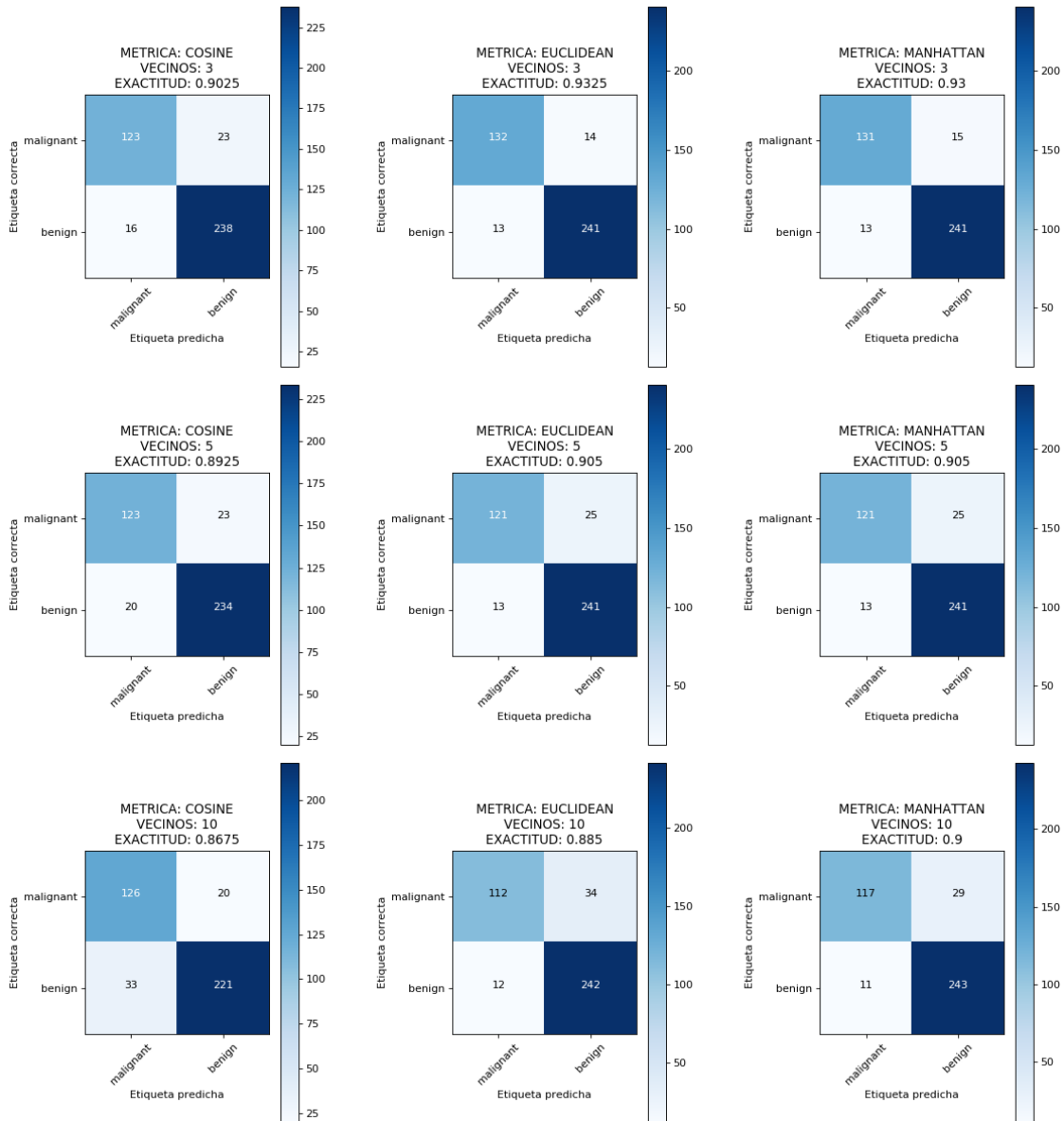
```

j=1
for n in neighbors:
    for m in metricas:
        n_neighbors=n
        metric = m
        model = KNeighborsClassifier(n_neighbors=n_neighbors, metric=metric)
        model.fit(X_train_feature, y_train)

        exactitud = accuracy_score(y_train, model.predict(X_train_feature))

        plt.subplot(3, 3, j)
        plot_confusion_matrix(confusion_matrix(y_train, model.predict(X_train_feature)),
                               classes=breast_cancer_data.target_names,
                               title='METRICA: '+m.upper()+'\nVECINOS: '+str(n)+'\nEXACTITUD: '+str(exactitud))
        j+=1
plt.subplots_adjust( wspace=0.6)

```

Se observa que para el menor número de vecinos considerado, la métrica euclídea muetsra la mejor exactitud y la menor cantidad de casos falsos negativos, sin embargo, para 10 vecinos es la métrica manhattan la que otorga los mejores predicciones

```
In [215]: plt.figure(figsize=(14,4), dpi= 80, facecolor='w', edgecolor='k')
          j=1
          for n in neighbors:
              for m in metricas:
                  if m!='euclidean':continue
                  n_neighbors=n
                  metric = m
                  model = KNeighborsClassifier(n_neighbors=n_neighbors, metric=metric)
```

```

model.fit(X_train_feature, y_train)

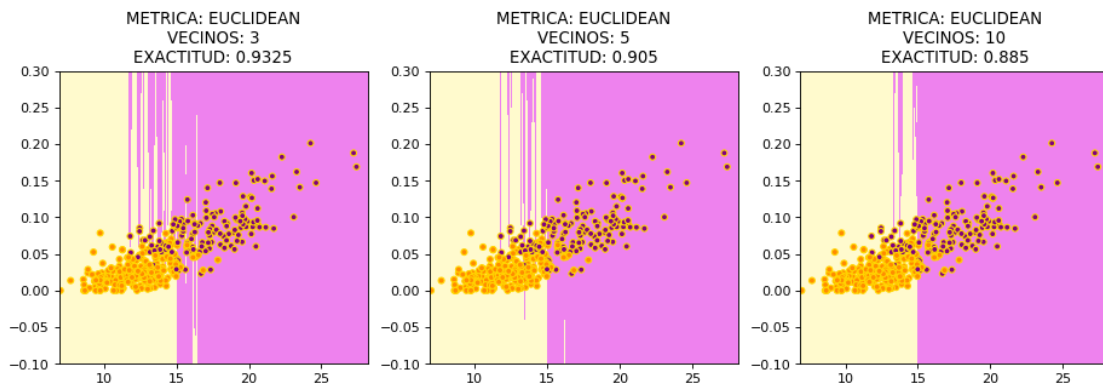
exactitud = accuracy_score(y_train, model.predict(X_train_feature))

plt.subplot(1, 3, j)

xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature], model)
plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, cmap=cm)
plt.xlim(xx.min(), xx.max())
plt.ylim(yy.min(), yy.max())
plt.title('METRICA: '+m.upper()+'\nVECINOS: '+str(n)+'\nEXACTITUD: '+str(exa

j+=1

```



5 CLASIFICACION MULTICLASE

```

In [2]: iris_data = load_iris()

shuff_data = np.random.permutation(150)
shuff_train = shuff_data[:120]
shuff_val = shuff_data[120:]

X_train = iris_data['data'][shuff_train]
X_val = iris_data['data'][shuff_val]

y_train = iris_data['target'][shuff_train]
y_val = iris_data['target'][shuff_val]

feature_map = {feature: idx for idx, feature in enumerate(iris_data['feature_names'])}

print(iris_data['DESCR'])

```

```
.. _iris_dataset:
```

```
Iris plants dataset
```

```
-----
```

```
**Data Set Characteristics:**
```

```
:Number of Instances: 150 (50 in each of three classes)
```

```
:Number of Attributes: 4 numeric, predictive attributes and the class
```

```
:Attribute Information:
```

- sepal length in cm
- sepal width in cm
- petal length in cm
- petal width in cm
- class:
 - Iris-Setosa
 - Iris-Versicolour
 - Iris-Virginica

```
:Summary Statistics:
```

```
=====  ====  ====  =====  =====  =====
                        Min  Max   Mean    SD    Class Correlation
=====  ====  ====  =====  =====  =====
sepal length:    4.3  7.9   5.84   0.83    0.7826
sepal width:     2.0  4.4   3.05   0.43   -0.4194
petal length:    1.0  6.9   3.76   1.76    0.9490 (high!)
petal width:     0.1  2.5   1.20   0.76    0.9565 (high!)
=====  ====  ====  =====  =====  =====
```

```
:Missing Attribute Values: None
```

```
:Class Distribution: 33.3% for each of 3 classes.
```

```
:Creator: R.A. Fisher
```

```
:Donor: Michael Marshall (MARSHALL%PLU@io.arc.nasa.gov)
```

```
:Date: July, 1988
```

The famous Iris database, first used by Sir R.A. Fisher. The dataset is taken from Fisher's paper. Note that it's the same as in R, but not as in the UCI Machine Learning Repository, which has two wrong data points.

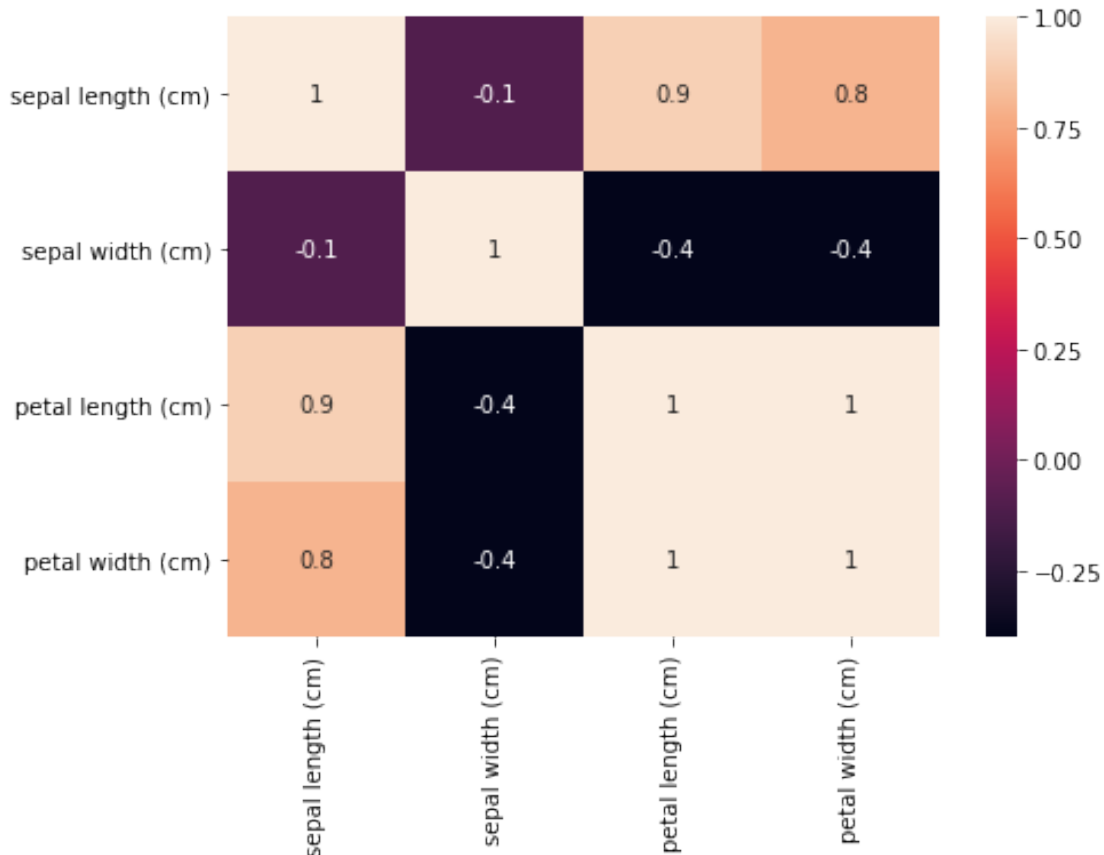
This is perhaps the best known database to be found in the pattern recognition literature. Fisher's paper is a classic in the field and is referenced frequently to this day. (See Duda & Hart, for example.) The data set contains 3 classes of 50 instances each, where each class refers to a type of iris plant. One class is linearly separable from the other 2; the latter are NOT linearly separable from each other.

```
.. topic:: References
```

- Fisher, R.A. "The use of multiple measurements in taxonomic problems" Annual Eugenics, 7, Part II, 179-188 (1936); also in "Contributions to Mathematical Statistics" (John Wiley, NY, 1950).
- Duda, R.O., & Hart, P.E. (1973) Pattern Classification and Scene Analysis. (Q327.D83) John Wiley & Sons. ISBN 0-471-22361-1. See page 218.
- Dasarathy, B.V. (1980) "Nosing Around the Neighborhood: A New System Structure and Classification Rule for Recognition in Partially Exposed Environments". IEEE Transactions on Pattern Analysis and Machine Intelligence, Vol. PAMI-2, No. 1, 67-71.
- Gates, G.W. (1972) "The Reduced Nearest Neighbor Rule". IEEE Transactions on Information Theory, May 1972, 431-433.
- See also: 1988 MLC Proceedings, 54-64. Cheeseman et al's AUTOCLASS II conceptual clustering system finds 3 classes in the data.
- Many, many more ...

```
In [6]: data = pd.DataFrame(iris_data['data'],columns=iris_data['feature_names'])
iris_data['iris_label'] = iris_data['target']
correlation_matrix = data.corr().round(1)
fig, ax = plt.subplots(figsize=(7,5))
sb.heatmap(data=correlation_matrix, annot = True)
```

```
Out[6]: <matplotlib.axes._subplots.AxesSubplot at 0x7f5c74dbfb70>
```



```

In [17]: x_feature = 'petal length (cm)'
         y_features = ['sepal width (cm)', 'petal width (cm)']
         penalties=['l1', 'l2']
         alphas=[1e-5, 1e1]

         cmap_dots = ListedColormap(['tomato', 'dodgerblue', 'goldenrod'])
         cmap_back = ListedColormap(['lightcoral', 'skyblue', 'palegoldenrod'])

plt.figure(figsize=(20, 8), dpi= 80, facecolor='w', edgecolor='k')
j=0
for yf in y_features:
    y_feature = yf

    x_feature_col = feature_map[x_feature]
    y_feature_col = feature_map[y_feature]
    X_train_feature = X_train[:, [x_feature_col, y_feature_col]]
    X_val_feature = X_val[:, [x_feature_col, y_feature_col]]

    for p in penalties:
        if p!=penalties[0]:continue
        penalty = p
        for a in alphas:
            if a!= alphas[0]:continue
            alpha = a
            model = LogisticRegression(penalty=penalty, C=1./alpha, multi_class='ovr')
            model.fit(X_train_feature, y_train)
            exactitud = accuracy_score(y_train, model.predict(X_train_feature))

            plt.subplot(2, 4, 1+j*4)
            plot_confusion_matrix(confusion_matrix(y_train, model.predict(X_train_feature)),
                                classes=iris_data.target_names,
                                title='Matriz de confusión')

            xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature], model)

            plt.subplot(2, 4, 2+j*4)
            plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
            plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, cmap=cmap_dots)
            plt.xlim(xx.min(), xx.max())
            plt.ylim(yy.min(), yy.max())
            plt.title("Conjunto de Entrenamiento")

```

```
n_neighbors = 3 # TODO: Cantidad de vecinos a tener en cuenta
metric = 'euclidean' # TODO: Medida de distancia. Algunas opciones: cosin
```

```
model = KNeighborsClassifier(n_neighbors=n_neighbors, metric=metric)
model.fit(X_train_feature, y_train)
```

```
exactitud =accuracy_score(y_train, model.predict(X_train_feature))
```

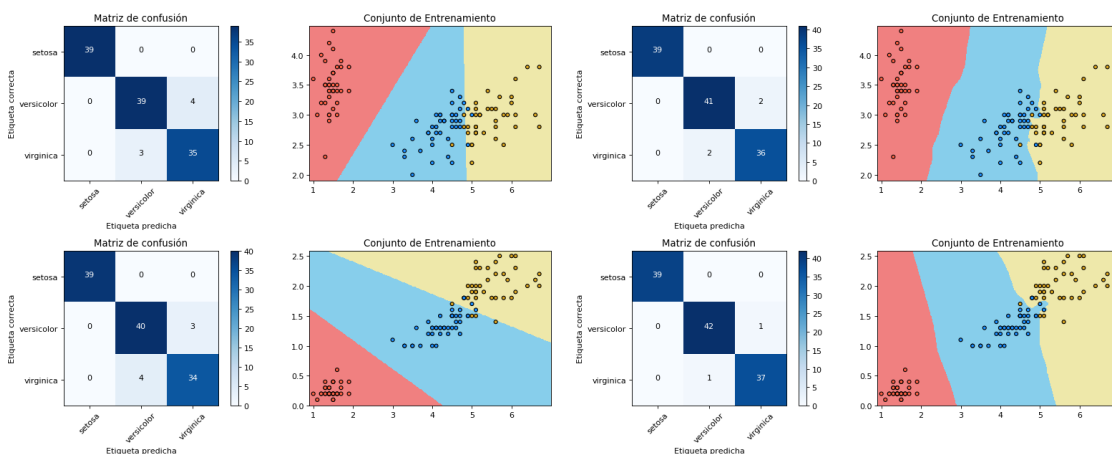
```
plt.subplot(2, 4, 3+j*4)
plot_confusion_matrix(confusion_matrix(y_train, model.predict(X_train_fea
                        classes=iris_data.target_names,
                        title='Matriz de confusión')
```

```
xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature], mo
```

```
cmap_dots = ListedColormap(['tomato', 'dodgerblue', 'goldenrod'])
cmap_back = ListedColormap(['lightcoral', 'skyblue', 'palegoldenrod'])
```

```
plt.subplot(2, 4, 4+j*4)
plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, cmap=
plt.xlim(xx.min(), xx.max())
plt.ylim(yy.min(), yy.max())
plt.title("Conjunto de Entrenamiento")
```

j+=1



```
In [ ]:
```

```
In [24]: x_feature = 'petal length (cm)'
y_features = ['sepal width (cm)', 'petal width (cm)']
penalties=['l1', 'l2']
alphas=[1e-5, 1e1]

cmap_dots = ListedColormap(['tomato', 'dodgerblue', 'goldenrod'])
cmap_back = ListedColormap(['lightcoral', 'skyblue', 'palegoldenrod'])

plt.figure(figsize=(20, 30), dpi= 80, facecolor='w', edgecolor='k')
fig.suptitle('x_feature: '+x_feature.upper(), fontsize=20)
j=0
for yf in y_features:

    y_feature = yf

    x_feature_col = feature_map[x_feature]
    y_feature_col = feature_map[y_feature]
    X_train_feature = X_train[:, [x_feature_col, y_feature_col]]
    X_val_feature = X_val[:, [x_feature_col, y_feature_col]]

    for p in penalties:
        if p!=penalties[0]:continue
        penalty = p
        for a in alphas:
            if a!= alphas[0]:continue
            alpha = a
            model = LogisticRegression(penalty=penalty, C=1./alpha, multi_class='ovr')
            model.fit(X_train_feature, y_train)

            i=1
            for n in neighbors:
                n_neighbors = n
                metric = 'euclidean'

                model = KNeighborsClassifier(n_neighbors=n_neighbors, metric=metric)
                model.fit(X_train_feature, y_train)

                exactitud = round(accuracy_score(y_train, model.predict(X_train_feature)), 2)

            plt.subplot(6, 2, i+j*6)
            plot_confusion_matrix(confusion_matrix(y_train, model.predict(X_train_feature)),
                                classes=iris_data.target_names,
                                title=y_feature.upper()+ ' MATRIZ DE CONFUSION\ñVECIOS')
```

```

xx, yy, Z = classifier_boundary(np.r_[X_train_feature, X_val_feature])

cmap_dots = ListedColormap(['tomato', 'dodgerblue', 'goldenrod'])
cmap_back = ListedColormap(['lightcoral', 'skyblue', 'palegoldenrod'])

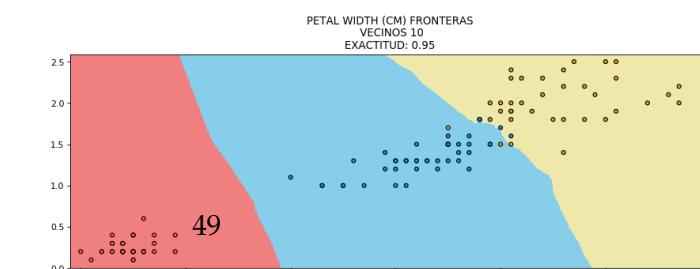
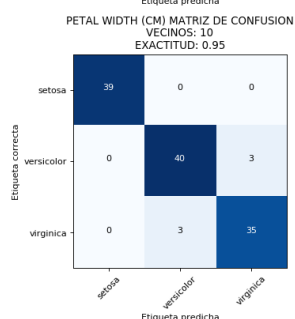
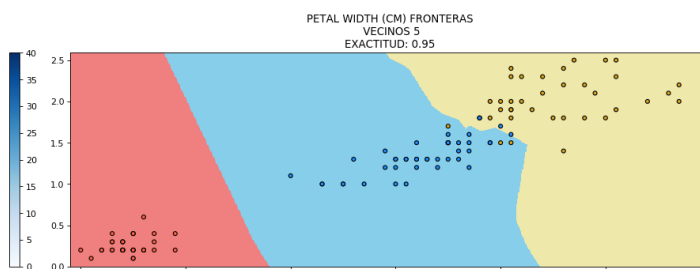
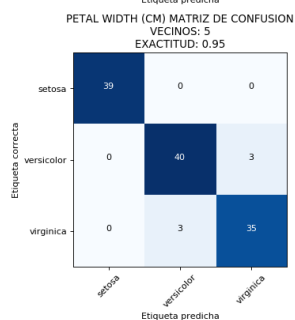
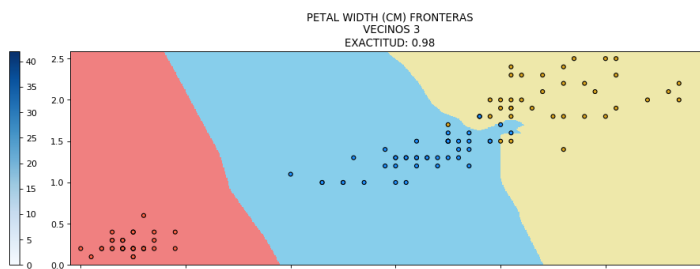
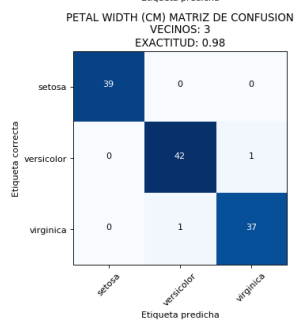
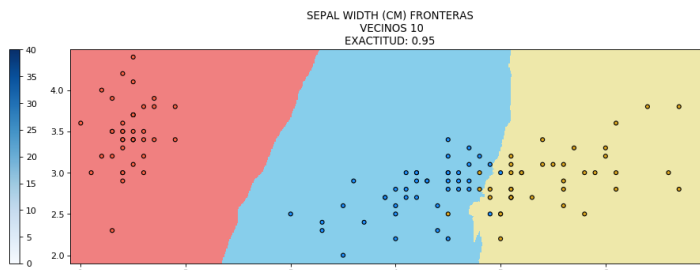
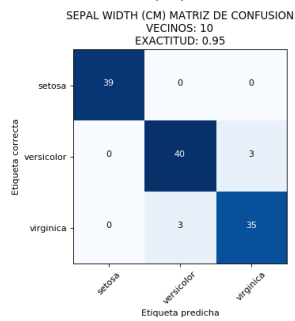
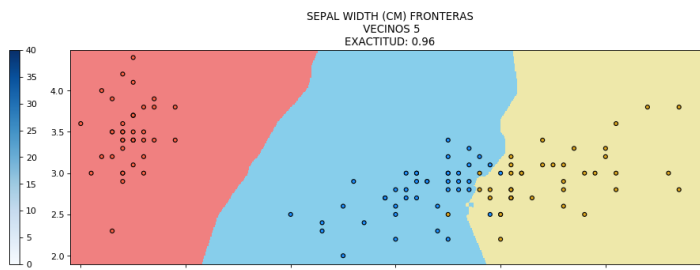
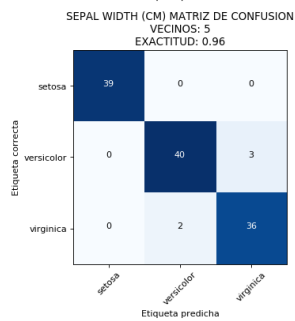
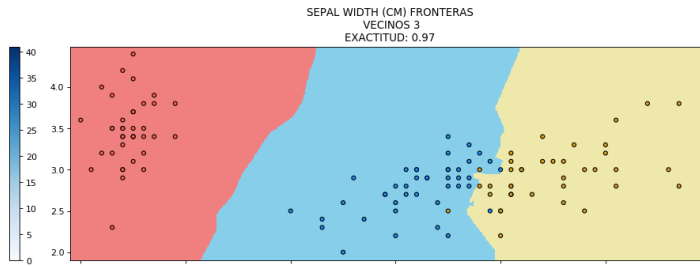
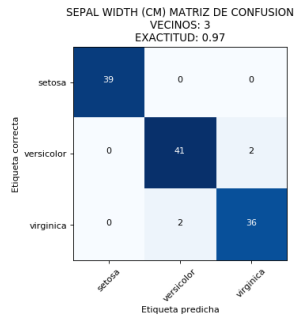
i+=1

plt.subplot(6, 2, i+j*6)
plt.pcolormesh(xx, yy, Z, cmap=cmap_back)
plt.scatter(X_train_feature[:, 0], X_train_feature[:, 1], c=y_train, c=cmap_dots)
plt.xlim(xx.min(), xx.max())
plt.ylim(yy.min(), yy.max())
plt.title(y_feature.upper()+' FRONTERAS\n VECINOS '+str(n)+'\nEXACTITUDE')

i+=1

j+=1

```

Si bien al considerar menos vecinos, la predicción es mejor, puede verse que la predicción también depende de los features utilizados. En este caso se escogieron pares de features fuertemente correlacionados y se ve que al variar `y_feature` entre dos valores, cuando se tienen 3 vecinos, la exactitud varía en 0.01.

In []: