hw4

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1 HW 4:

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1.1 Question 1:

```
[49]: import numpy as np
  import warnings
  import matplotlib . pyplot as plt
  warnings.simplefilter('ignore')
  from sklearn . metrics import mean_squared_error
  def data_generator ( n_samples ):
        x = np . random . uniform ( -10 , 10 , n_samples )
        y = np . cos (0.5 + np . exp ( - x )) + 1/(1 + np . exp ( - x ))
        noise = np . random . normal (0 , 0.01 , n_samples )
        y += noise
        return x , y
        complete_X , complete_Y = data_generator (5000)
        train_X , train_Y = complete_X [:100] , complete_Y [:100]
        large_X , large_Y = complete_X [100:] , complete_Y [100:]
        loss_func = mean_squared_error
```

1.1.1 Part A

```
for i in range((2,d+1)):
           rv = np.column_stack((rv,x ** i))
    else:
        return x
    return rv
def proyection(Y,x, cons=True):
    if cons:
        X = np.column_stack((x,np.ones([len(x),1])))
    else:
        X=x.copy()
    y_hat = np.dot(X,beta_est(Y,x,cons))
    return y_hat
def error_poly(n_poly, X, y):
    errors = []
    for i in range(n_poly):
        if i == 0:
            x2 = np.ones([len(X),1])
        else:
            x2 = np.column_stack((np.ones([len(X),1]),polynomial(X,i)))
        coef = ((np.linalg.inv(np.transpose(x2)@x2)@np.transpose(x2)@y))
        errors.append(((x2@coef-y)**2).sum()/len(x2))
    return errors
def error_poly_cross_val(n_poly, X, y,Xl,yl):
    errors = []
    for i in range(n_poly):
        if i == 0:
            x2 = np.ones([len(X),1])
            x21 = np.ones([len(X1),1])
        else:
            x2 = np.column_stack((np.ones([len(X),1]),polynomial(X,i)))
            x21 = np.column stack((np.ones([len(X1),1]),polynomial(X1,i)))
        coef = ((np.linalg.inv(np.transpose(x2)@x2)@np.transpose(x2)@y))
        errors.append(((x21@coef-y1)**2).sum()/len(x21))
    return errors
```

```
x2 = np.ones([len(train_X),1])
else:
    x2 = np.column_stack((np.ones([len(train_X),1]),polynomial(train_X,i)))
coef = ((np.linalg.inv(np.transpose(x2)@x2)@np.transpose(x2)@train_Y))
error_sq.append(((x2@coef- train_Y)**2).sum()/len(x2))
'''
error_sq = error_poly(31,train_X, train_Y)
```

[4]: print('Mininmal Square Error:',(np.array(error_sq)).min())

Mininmal Square Error: 0.20932454287986946

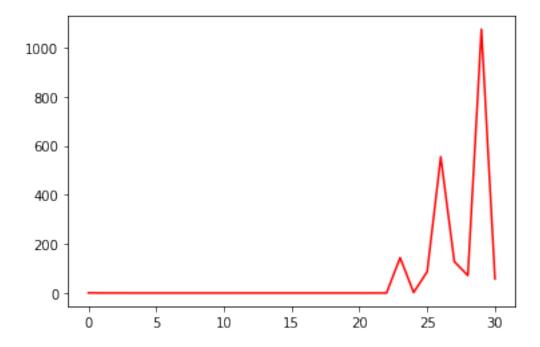
[5]: print('Degree of polynomail that minimize risk',np.argmin(np.array(error_sq)))

Degree of polynomail that minimize risk 20

1.1.2 Part B

```
[6]: import matplotlib.pyplot as plt
plt.plot(error_sq,color='r', label='error square')
```

[6]: [<matplotlib.lines.Line2D at 0x1020003790>]



1.1.3 Part C

1.1.4 R^*

Since we know the true process behind $y = f(x) + \epsilon$, and we know that $\epsilon \sim N(0, 0.01)$ we can argue that R^* on average is the variance of ϵ . Since the MSE of would be:

$$MSE = \frac{1}{n} \sum_{i=1}^{n} (y - f(x))^{2}$$

$$MSE = \frac{1}{n} \sum_{i=1}^{n} (f(x) + \epsilon - f(x))^{2}$$

$$MSE = \frac{1}{n} \sum_{i=1}^{n} (\epsilon)^{2}$$

$$MSE = \sigma_{\epsilon}$$

We present the calculation below

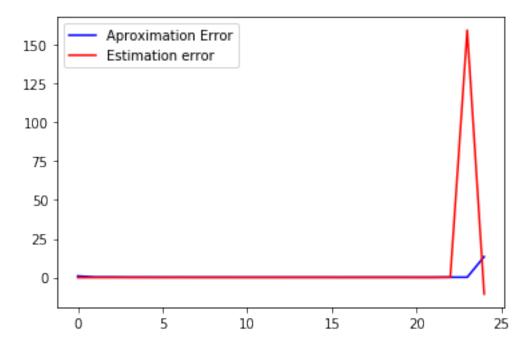
```
[35]: y_non_noice_train =train_Y-np . random . normal (0 , 0.01 , 100 )
y_non_noice_large =large_Y-np . random . normal (0 , 0.01 , 4900 )
r_star= ((large_Y-y_non_noice_large)**2).mean()
print("R^* is",r_star)
```

R^* is 0.00010019491220344034

```
[39]: error_sq_cv =error_poly_cross_val(25,train_X, train_Y,large_X,large_Y)
```

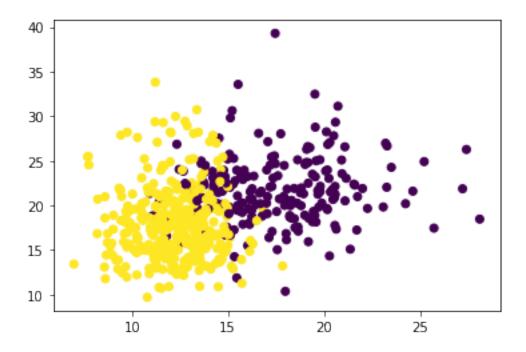
```
plt.plot(estimation_error,color='r', label='Estimation error')
plt.legend()
```

[59]: <matplotlib.legend.Legend at 0x1a2a1615d0>



1.2 Question 2

```
[44]: from sklearn.model_selection import train_test_split
    from sklearn . datasets import load_breast_cancer
    from sklearn.metrics import accuracy_score
    import matplotlib . pyplot as plt
    X , y = load_breast_cancer ( return_X_y = True )
    X = X [: , :2]
    plt . clf ()
    plt . scatter ( X [: , 0] , X [: , 1] , c = y )
    plt . show ()
```



1.2.1 Part A

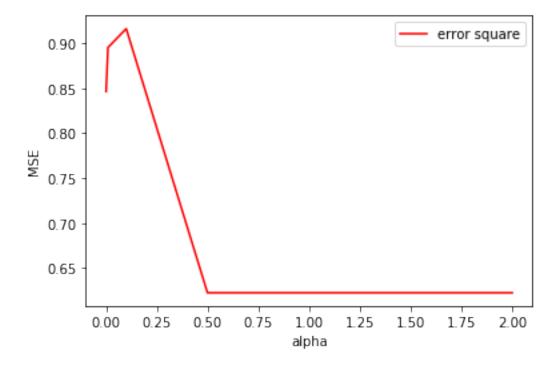
In sample accuracy: 1.0
In out of sample accuracy: 0.8461538461538461

1.2.2 Part B

```
[47]: alpha_list = [1/1000,1/100,1/10,1/2,3/4,1,2]
acc_test = []
acc_train = []
for ccp in alpha_list:
```

```
clf = tree.DecisionTreeClassifier(ccp_alpha=ccp)
  clf = clf.fit(X_train, y_train)
  y_pred_train = clf.predict(X_train)
  y_pred_test = clf.predict(X_test)
  acc_train.append(sklearn.metrics.accuracy_score(y_train, y_pred_train))
  acc_test.append(sklearn.metrics.accuracy_score(y_test, y_pred_test))
plt.plot(alpha_list,acc_test,color='r', label='error square')
plt.legend()
plt.xlabel('alpha')
plt.ylabel('MSE')
```

[47]: Text(0, 0.5, 'MSE')

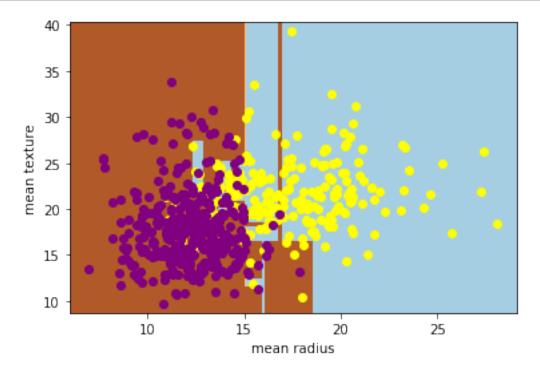


1.2.3 Part C

```
[13]: # Python
def plot_regions ( tree ):
    plot_colors = ['yellow', 'purple']
    plot_step = 0.02
    breast_cancer = load_breast_cancer ()
    X = breast_cancer.data
    y = breast_cancer.target
    plt.clf ()
```

```
idx = np . arange ( X . shape [0])
  np . random . shuffle ( idx )
  X = X [idx]
  y = y [idx]
  x_{min}, x_{max} = X [: , 0]. min () - 1 , X [: , 0]. max () + 1
  y_{min}, y_{max} = X [: , 1]. min () - 1 , X [: , 1]. max () + 1
  xx , yy = np.meshgrid ( np . arange ( x_min , x_max , plot_step ), np .__
→arange ( y_min , y_max , plot_step ))
  Z = tree.predict ( np.c_ [ xx.ravel() , yy.ravel()])
  Z = Z.reshape (xx.shape)
  cs = plt.contourf ( xx , yy , Z , cmap = plt . cm . Paired )
  plt.xlabel( breast_cancer . feature_names [0])
  plt.ylabel ( breast_cancer . feature_names [1])
  plt.axis ('tight')
  for i , color in zip( range (2) , plot_colors ):
       idx = np . where ( y == i )
      plt . scatter ( X [idx , 0] , X [idx , 1] , c = color , label = 
→breast_cancer.target_names [ i ] ,cmap = plt.cm.Paired )
```

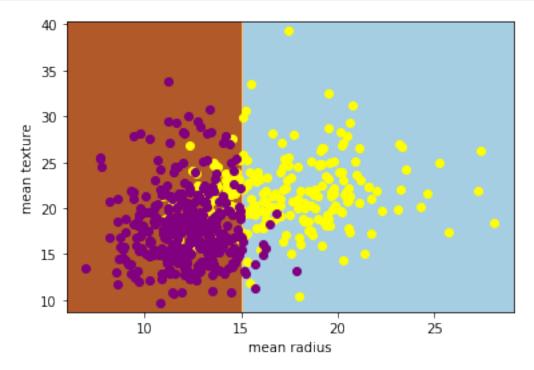
[14]: plot_regions (clf_a)



```
[15]: min_alpha = alpha_list[np.argmax(acc_test)]

clf_min_b = tree.DecisionTreeClassifier(ccp_alpha=min_alpha)
```

```
clf_min_b = clf_min_b.fit(X_train, y_train)
plot_regions(clf_min_b)
```

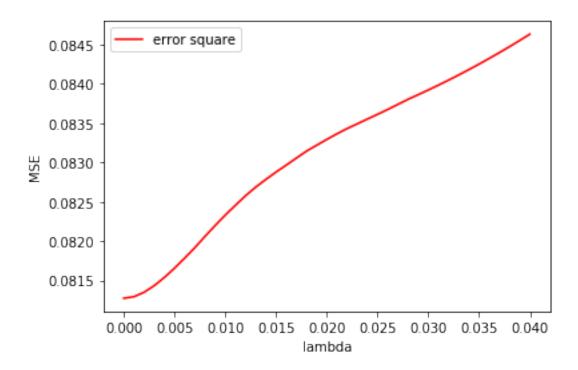


1.3 Question 3

1.3.1 Part A

```
[51]: import numpy as np
      np . random . seed (0)
      N_fold = 10
      N_{test} = 500
      N_{train} = 1000
      N = N_{test} + N_{train}
      \# Specify feature dimensions of X and Y
      X_dim = 20
      Y_dim = 10
      X = np.random.randn (N , X_dim )
      \# Only have 10 non - zero entries in beta ,
      nnz = 10
      beta = np . zeros (( X_dim * Y_dim ))
      nnz_idx = np.random.choice( X_dim * Y_dim , nnz , replace = False )
      beta [ nnz_idx ] = np.random.randn ( nnz ) * 2
      beta = beta . reshape ( X_dim , Y_dim )
```

```
xb = X @ beta
      Y = xb + np \cdot random \cdot rand (N , Y_dim)
      # Split training and testing set
      X_test = X [: N_test ]
      Y_test = Y [: N_test ]
      X_train = X [ N_test :]
      Y_train = Y [ N_test :]
[52]: def lasso_prediction_insample(X_train, X_test, Y_train, Y_test, alpha_c):
          clf = sklearn.linear_model.Lasso(alpha=alpha_c,max_iter=20000)
          clf = clf.fit(X_train,Y_train)
          y_hat = clf.predict(X_train)
          from sklearn.metrics import mean_squared_error
          ee = mean_squared_error(Y_train, y_hat)
          return ee
[53]: rv = []
      lambda_list = []
      for alpha in range(0,41):
          ee_i = lasso_prediction_insample(X_train, X_test, Y_train, Y_test, alpha/1000)
          rv.append(ee_i)
          lambda_list.append(alpha/1000)
      plt.plot(lambda_list,rv,color='r', label='error square')
      plt.legend()
      plt.xlabel('lambda')
      plt.ylabel('MSE')
```



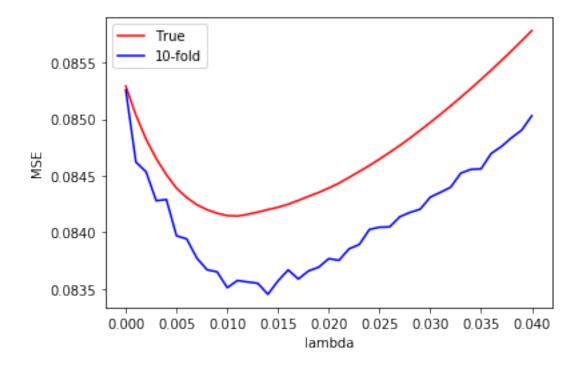
1.3.2 Part B

```
[54]: import random
      import numpy as np
      \#data=np.stack(Y\_test, X\_test,axis=0)
      # Split a dataset into k folds
      def cross_validation_split(dataset_X, dataset_Y, folds=3):
          dataset_split_X = list()
          dataset_split_Y = list()
          dataset_copy_X = list(dataset_X)
          dataset_copy_Y = list(dataset_Y)
          fold_size = int(len(dataset_X) / folds)
          for i in range(folds):
              fold_X = list()
              fold_Y = list()
              while len(fold_X) < fold_size:</pre>
                  index = random.randrange(len(dataset_copy_X))
                  fold_X.append(dataset_copy_X.pop(index))
                  fold_Y.append(dataset_copy_Y.pop(index))
              dataset_split_X.append(fold_X)
              dataset_split_Y.append(fold_Y)
          return dataset_split_X, dataset_split_Y
```

```
[55]: X_cv,Y_cv = cross_validation_split(X_train,Y_train,10)
[56]: | (np.concatenate(X_cv, axis=0)).shape
[56]: (1000, 20)
[57]: def lasso_prediction_outsample(X_train, X_test, Y_train, Y_test, alpha_c):
          clf = sklearn.linear_model.Lasso(alpha=alpha_c,max_iter=20000)
          clf = clf.fit(X_train,Y_train)
          y_hat = clf.predict(X_test)
          from sklearn.metrics import mean squared error
          ee = mean_squared_error(Y_test, y_hat)
          return ee
      def k_fold(fold, X, Y, alpha_c):
          X_cv,Y_cv = cross_validation_split(X,Y,fold)
          mean_ee = []
          for i in range(10):
              Y_test_iter = Y_cv[i]
              X_test_iter = X_cv[i]
              Y_train_iter = np.concatenate(Y_cv[:i]+Y_cv[i+1:], axis = 0)
              X_train_iter = np.concatenate(X_cv[:i]+X_cv[i+1:], axis= 0)
              ee =
       →lasso_prediction_outsample(X_train_iter, X_test_iter, Y_train_iter, Y_test_iter, alpha_c)
              mean ee.append(ee)
          return np.array(mean_ee).mean()
      #k_fold(10, X_train, Y_train, .2)
      rv_fold = []
      lambda_list_fold = []
      for alpha in range(0,41):
          ee_i = k_fold(10, X_train, Y_train,alpha/1000)
          rv_fold.append(ee_i)
          lambda_list_fold.append(alpha/1000)
[58]: rv_cv = []
      lambda_cv_list = []
      for alpha in range(0,41):
          ee_i = lasso_prediction_outsample(X_train, X_test, Y_train, Y_test, alpha/1000)
          rv_cv.append(ee_i)
          lambda_cv_list.append(alpha/1000)
```

```
plt.plot(lambda_cv_list,rv_cv,color='r', label='True')
plt.plot(lambda_cv_list,rv_fold,color='b', label='10-fold')
plt.legend()
plt.xlabel('lambda')
plt.ylabel('MSE')
```

[58]: Text(0, 0.5, 'MSE')



[]: