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(a)

These are the datasets for the first question.

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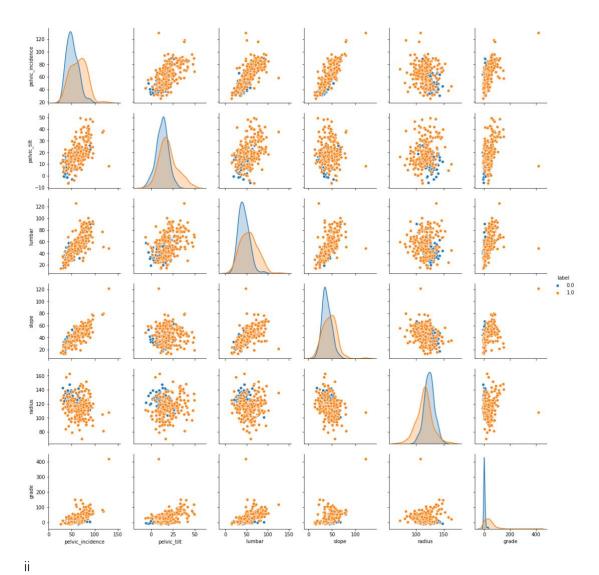
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(b)
Here are the codes and graphs of (i), (ii) and (iii).
trykNN.py
##
import pandas as pd
import numpy as np
import math
import seaborn as sns
from sklearn.model_selection import train_test_split
                        pd.read_table('vertebral_column_data/column_2C.dat',sep
users
'::',header=None,engine='python')
users=users[0]
#loading the data and get into the list
datalist=np.zeros((len(users),7))
for idx in range (len(users)):
    Str=users[idx]
    Str=Str.split()
    for i in range(len(Str)):
         element=Str[i]
         if i < (len(Str)-1):
              datalist[idx][i]=float(element)
         else:
              if element=='AB':
                  datalist[idx][i]=1
              else:
                  datalist[idx][i]=0
newdf=pd.DataFrame({"label":datalist[:,6],"pelvic_incidence":datalist[:,0],"pelvic_tilt":datalist[:,
1],"lumbar":datalist[:,2],"slope":datalist[:,3],"radius":datalist[:,4],"grade":datalist[:,5]})
##corresponding to b
sns.pairplot(data=newdf,
                           hue='label',
                                          vars=['pelvic_incidence',
                                                                     'pelvic_tilt',
                                                                                    'lumbar',
'slope', 'radius', 'grade'])
###
Here are the graphs for the questions:
```

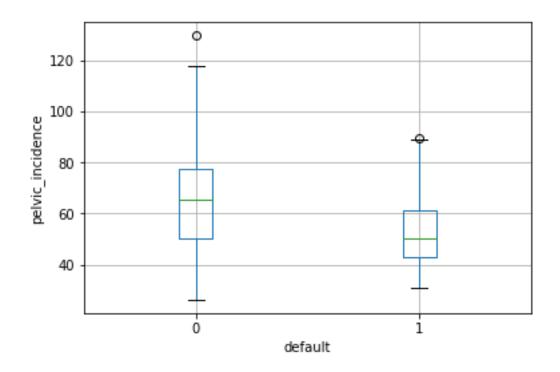
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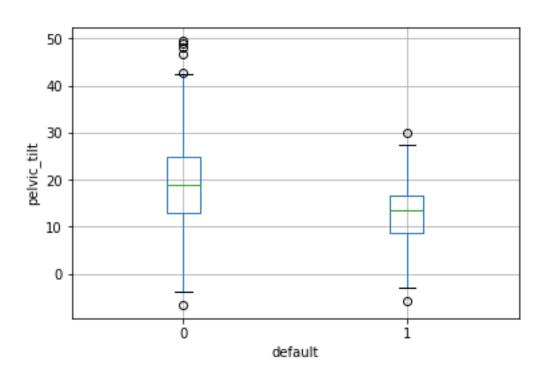


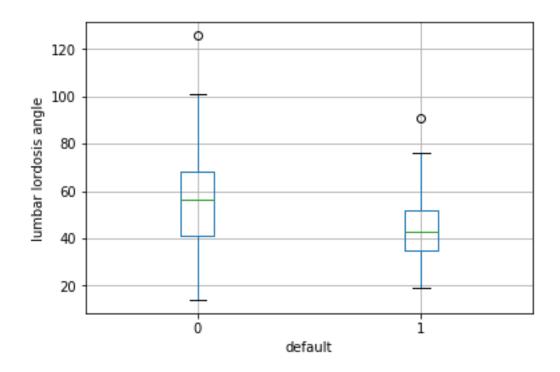
Make boxplots: the following code is about pelvic incidence variables, the others boxplots are made in a similar way.

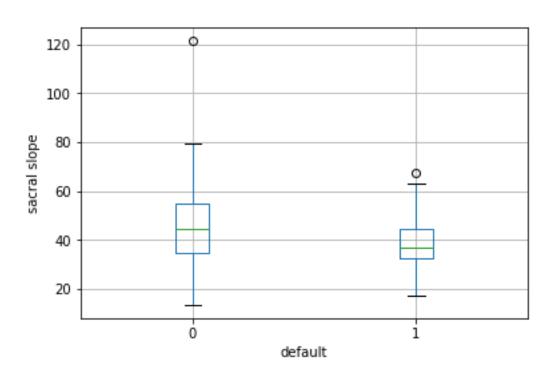
```
##
pel_1=newdf_1['pelvic_incidence']
pel_1=pel_1.rename(columns={'pelvic_incidence':'1'},inplace=True)
pel_0=newdf_0['pelvic_incidence']
pel_0=pel_0.rename(columns={'pelvic_incidence':'0'},inplace=True)

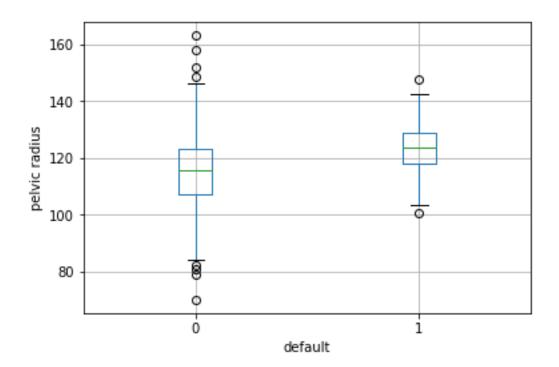
a=pd.concat([pel_1,pel_0],axis=1)
#plt.boxplot([pel_1,pel_0],patch_artist=True, boxprops={'color'})
a.boxplot()
plt.ylabel('pelvic_incidence')
plt.xlabel('default')
plt.show()
##
```

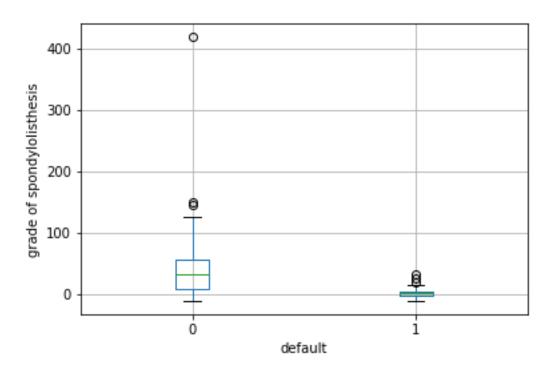












iii

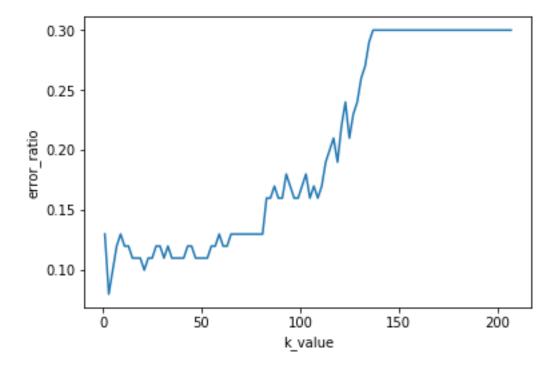
##

X=newdf[['pelvic_incidence','pelvic_tilt','lumbar','slope','radius','grade']]

y=newdf['label']

$$x0=X[y==0]$$

```
x1=X[y==1]
x0_{train}=x0.head(70)
y0_{train}=y[y==0].head(70)
x0_{\text{test}} = x0[70:\text{len}(x0)]
y0_{test}=y[y==0].head(x0_{test.shape}[0])
x1_train=x1.head(140)
y1_{train}=y[y==1].head(140)
x1_{\text{test}} = x1[140:\text{len}(x1)]
y1_test=y[y==1].head(x1_test.shape[0])
x_train=x0_train.append(x1_train)
y_train=y0_train.append(y1_train)
x_test=x0_test.append(x1_test)
y_test=y0_test.append(y1_test)
##
(c)
Here are the codes for the k-nearest neighbors with the Euclidean metric.
def Euclidean(single_data,norm_x_train):
     diffmat=np.tile(single_data,(norm_x_train.shape[0],1))-norm_x_train
     sq_diff=diffmat**2
     distance=(sq_diff.sum(axis=1))**0.5
     return distance #Euclidean metric
##
(ii)
In this question...
trykNN3.py
k values: {1, 3, 5, .....209}
```



When K=3, the minimum error rate is 0.08.

```
Calculating the confusion matrix
##
tp=0
fn=0
fp=0
tn=0
for row in range (test_num):
    compare=best_compare[row,:]
    y_real=compare[0]
    y_pre=compare[1]
    if y_real==1 and y_pre==1:
        tp+=1
    else:
        if y_real==1 and y_pre==0:
             fn+=1
        else:
             if y_real==0 and y_pre==1:
                 fp+=1
             else:
                 tn+=1
```

```
confusion_matrix=[[tp,fp,tp+fp],[fn,tn,fn+tn],[tp+fn,fp+tn,tp+fn+fp+tn]]
##
```

Now, we will get the results like this...

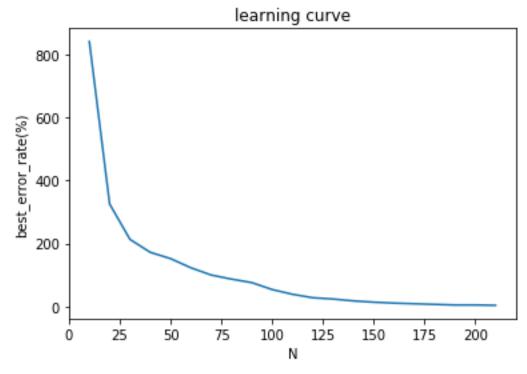
Confusion matrix

		True status		
		1	0	total
Predicted status	1	69	7	76
	0	1	23	24
	total	70	30	100

Precision=0.908 Fscore=0.945

(iii)

learning_curve.py



The codes are here:

##

 $N_{range} = np.arange(10,220,10)$

N_best_error_ratio=np.zeros_like(N_range)

for N in N_range:

```
x_train,y_train,x_test,y_test=prepare_data(x0,x1,y,N)
norm_x_train=norm2(x_train)
norm_x_test=norm2(x_test)
```

```
test_num=norm_x_test.shape[0]
y_train_dist=np.zeros((norm_x_train.shape[0],2))
```

compare=np.zeros((test_num,2))

```
k_{value} = np.arange(1, min(N, 196 + 5), 5)
    error_ratio=∏
    for k in k_value:
         error_sum=0
compare_result,error_sum_result=kNN_compare(k,norm_x_test,norm_x_train,y_test,test_num)
         error=error_sum_result/test_num
         error_ratio.append(error)
    min_index=np.array(error_ratio).argsort()
    k_best=k_value[min_index[0]]
best_compare,best_error=kNN_compare(k_best,norm_x_test,norm_x_train,y_test,test_num)
    idx_e=int(N/10-1)
    N_best_error_ratio[idx_e]=best_error/N*100
plt.figure(1)
plt.xlabel('N')
plt.ylabel('best_error_rate(%)')
plt.title('learning curve')
plt.plot(N_range,N_best_error_ratio)
##
(d)
    (I)
    The definition of Minkovski distance:
    def Minkow(single_data,norm_x_train,pvalue):
         distance=np.zeros((norm_x_train.shape[0]))
         x1=single_data
         for row in range(norm_x_train.shape[0]):
              x2=norm_x_train[row,:]
              dist=DistanceMetric.get_metric('minkowski',p=pvalue)
              xmat=[x1,x2]
              distance[row]=dist.pairwise(xmat)[0][1]
         return distance
    #
       A. When p=1, it will be the Manhattan Distance.
       The file is knn_d.py
       When using Minkow function, make the p-value to 1.
```

```
In the kNN_compare function, the minimum value of the error rate in
      distance=Minkow(single_data,norm_x_train,1)
      Is 0.11, the k is 6
      B. For the (list) q_value, with log10(p)
      The corresponding file is knn_d2.py
      # q_value=np.arange(0.1,1.1,0.1)
      ##
      for q in q_value:
           pvalue=10**q
           error_sum=0
      compare_result,error_sum_result=kNN_compare_p(k,norm_x_test,norm_x_train,y_test,
      test_num,pvalue)
           error=error_sum_result/test_num
           error_ratio.append(error)
     ##
    distance=Minkow(single_data,norm_x_train,pvalue)
After the calculation,
The lowest error rate is 0.1, and for the corresponding log10(p) is 0.6.
С.
# distance=chebdist(single_data,norm_x_train)
def chebdist(single_data,norm_x_train):
    distance=np.zeros((norm_x_train.shape[0]))
    x1=single_data
    for row in range(norm_x_train.shape[0]):
         x2=norm_x_train[row,:]
         dist=DistanceMetric.get_metric('chebyshev')
         xmat=[x1,x2]
         distance[row]=dist.pairwise(xmat)[0][1]
    return distance
##
As the result, the smallest error rate is 0.1 and the k will be 16.
    (ii)
    Mahalanobis distance
    #distance=Minkow(single_data,norm_x_train,pvalue)
    def mahadist(single_data,norm_x_train):
         distance=np.zeros((norm_x_train.shape[0]))
         x1=single_data
         for row in range(norm_x_train.shape[0]):
             x2=norm_x_train[row,:]
```

```
X=np.vstack([x1,x2])
XT=X.T
S=np.cov(X)
SI=np.linalg.inv(S)
n=XT.shape[0]

for i in range(0,n):
    for j in range(i+1,n):
        delta=XT[i]-XT[j]
        d=np.sqrt(np.dot(np.dot(delta,SI),delta.T))
        distance[row]=d
```

For the return distance, the smallest error rate is 0.1 and the k will be 21.

(e) ##

count_class={}
for ik in range (k):
 vote_y=y_train.iloc[sort_dist[ik]]

count_class[vote_y]=count_class.get(vote_y,0)+float(1/distance[sort_dist[ik]]) #

##

	Euclidean	Manhattan	Chebyshev
Majority vote	K=16	K=6	K=16
	Error rate=0.1	Error rate=0.11	Error rate=0.07
Weight	K=6	K=26	K=16
	Error rate=0.1	Error rate=0.1	Error rate=0.1