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In [20]:

```
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
from pandas import read csv
#path = r"C:\pima-indians-diabetes.csv"
path=r"C:\myprogram\python\machine_learning\data\pima-indians-diabetes.csv"
headernames = ['preg', 'plas', 'pres', 'skin', 'test', 'mass', 'pedi', 'age', 'class']
data = read csv(path, names = headernames)
array = data.values
X = array[:,0:8]
Y = array[:,8]
print(data.shape)
\#(768, 9)
print(data.head())
patient data = data.iloc[:, 3:5].values
import scipy.cluster.hierarchy as shc
plt.figure(figsize = (10, 7))
plt.title("Patient Dendograms")
dend = shc.dendrogram(shc.linkage(data, method = 'ward'))
from sklearn.cluster import AgglomerativeClustering
cluster = AgglomerativeClustering(n_clusters = 4, affinity = 'euclidean', linkage = 'ward'
)
cluster.fit predict(patient data)
plt.figure(figsize = (10, 7))
plt.scatter(patient_data[:,0], patient_data[:,1], c = cluster.labels_, cmap = 'rainbow')
```

localhost:8888/lab

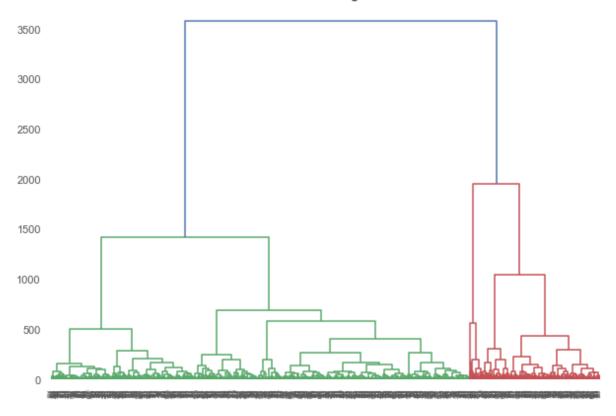
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| (768, 9) | | | | | | | | | |
|----------|------|------|------|------|------|------|-------|-----|-------|
| | preg | plas | pres | skin | test | mass | pedi | age | class |
| 0 | 6 | 148 | 72 | 35 | 0 | 33.6 | 0.627 | 50 | 1 |
| 1 | 1 | 85 | 66 | 29 | 0 | 26.6 | 0.351 | 31 | 0 |
| 2 | 8 | 183 | 64 | 0 | 0 | 23.3 | 0.672 | 32 | 1 |
| 3 | 1 | 89 | 66 | 23 | 94 | 28.1 | 0.167 | 21 | 0 |
| 4 | 0 | 137 | 40 | 35 | 168 | 43.1 | 2.288 | 33 | 1 |

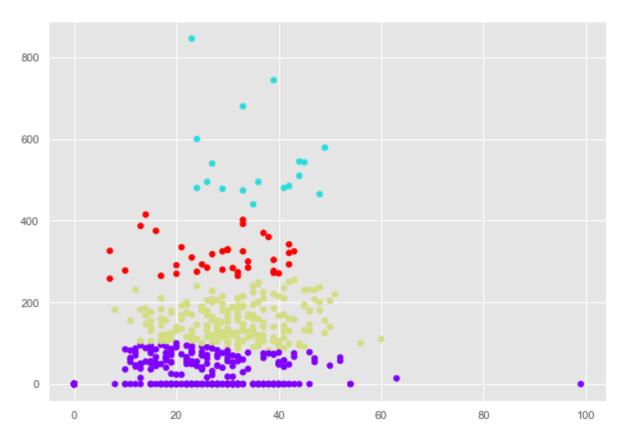
Out[20]:

<matplotlib.collections.PathCollection at 0x18474c992c8>

Patient Dendograms



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In []:

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