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## Assignment 1 - Introduction to Machine Learning

For this assignment, you will be using the Breast Cancer Wisconsin (Diagnostic) Database to create a classifier that can help diagnose patients. First, read through the description of the dataset (below).

```
In [2]: import numpy as np
import pandas as pd
from sklearn.datasets import load_breast_cancer

cancer = load_breast_cancer()

#print(cancer.DESCR) # Print the data set description
```

The object returned by `load_breast_cancer()` is a scikit-learn Bunch object, which is similar to a dictionary.

```
In [2]: cancer.keys()
```

```
Out[2]: dict_keys(['feature_names', 'DESCR', 'target', 'target_names', 'data'])
```

In [3]: `print(cancer.DESCR)`

## Breast Cancer Wisconsin (Diagnostic) Database

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## Notes

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## Data Set Characteristics:

:Number of Instances: 569

:Number of Attributes: 30 numeric, predictive attributes and the class

## :Attribute Information:

- radius (mean of distances from center to points on the perimeter)
- texture (standard deviation of gray-scale values)
- perimeter
- area
- smoothness (local variation in radius lengths)
- compactness ( $\text{perimeter}^2 / \text{area} - 1.0$ )
- concavity (severity of concave portions of the contour)
- concave points (number of concave portions of the contour)
- symmetry
- fractal dimension ("coastline approximation" - 1)

The mean, standard error, and "worst" or largest (mean of the three largest values) of these features were computed for each image, resulting in 30 features. For instance, field 3 is Mean Radius, field

d

13 is Radius SE, field 23 is Worst Radius.

## - class:

- WDBC-Malignant
- WDBC-Benign

## :Summary Statistics:

=====	=====	=====
	Min	Max
=====	=====	=====
radius (mean):	6.981	28.11
texture (mean):	9.71	39.28
perimeter (mean):	43.79	188.5
area (mean):	143.5	2501.0
smoothness (mean):	0.053	0.163
compactness (mean):	0.019	0.345
concavity (mean):	0.0	0.427
concave points (mean):	0.0	0.201
symmetry (mean):	0.106	0.304
fractal dimension (mean):	0.05	0.097
radius (standard error):	0.112	2.873
texture (standard error):	0.36	4.885
perimeter (standard error):	0.757	21.98
area (standard error):	6.802	542.2
smoothness (standard error):	0.002	0.031
compactness (standard error):	0.002	0.135
concavity (standard error):	0.0	0.396
concave points (standard error):	0.0	0.053
symmetry (standard error):	0.008	0.079
fractal dimension (standard error):	0.001	0.03

radius (worst):	7.93	36.04
texture (worst):	12.02	49.54
perimeter (worst):	50.41	251.2
area (worst):	185.2	4254.0
smoothness (worst):	0.071	0.223
compactness (worst):	0.027	1.058
concavity (worst):	0.0	1.252
concave points (worst):	0.0	0.291
symmetry (worst):	0.156	0.664
fractal dimension (worst):	0.055	0.208
=====	=====	=====

:Missing Attribute Values: None

:Class Distribution: 212 - Malignant, 357 - Benign

:Creator: Dr. William H. Wolberg, W. Nick Street, Olvi L. Mangasarian

:Donor: Nick Street

:Date: November, 1995

This is a copy of UCI ML Breast Cancer Wisconsin (Diagnostic) datasets.  
<https://goo.gl/U2Uwz2>

Features are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image.

Separating plane described above was obtained using Multisurface Method-Tree (MSM-T) [K. P. Bennett, "Decision Tree Construction Via Linear Programming." Proceedings of the 4th Midwest Artificial Intelligence and Cognitive Science Society, pp. 97-101, 1992], a classification method which uses linear programming to construct a decision tree. Relevant features were selected using an exhaustive search in the space of 1-4 features and 1-3 separating planes.

The actual linear program used to obtain the separating plane in the 3-dimensional space is that described in:  
 [K. P. Bennett and O. L. Mangasarian: "Robust Linear Programming Discrimination of Two Linearly Inseparable Sets", Optimization Methods and Software 1, 1992, 23-34].

This database is also available through the UW CS ftp server:

ftp ftp.cs.wisc.edu  
 cd math-prog/cpo-dataset/machine-learn/WDBC/

#### References

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- W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction for breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on Electronic Imaging: Science and Technology, volume 1905, pages 861-870, San Jose, CA, 1993.
- O.L. Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis

and  
prognosis via linear programming. Operations Research, 43(4), pages 570-577, July-August 1995.  
- W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine learning techniques to diagnose breast cancer from fine-needle aspirates. Cancer Letters 77 (1994) 163-171.

## Question 0 (Example)

How many features does the breast cancer dataset have?

*This function should return an integer.*

```
In [3]: # You should write your whole answer within the function provided. The autograder will call
# this function and compare the return value against the correct solution value
def answer_zero():
    # This function returns the number of features of the breast cancer dataset, which is an integer.
    # The assignment question description will tell you the general format the autograder is expecting
    return len(cancer['feature_names'])

# You can examine what your function returns by calling it in the cell. If you have questions
# about the assignment formats, check out the discussion forums for any FAQs
answer_zero()
```

Out[3]: 30

## Question 1

Scikit-learn works with lists, numpy arrays, scipy-sparse matrices, and pandas DataFrames, so converting the dataset to a DataFrame is not necessary for training this model. Using a DataFrame does however help make many things easier such as munging data, so let's practice creating a classifier with a pandas DataFrame.

Convert the sklearn.dataset cancer to a DataFrame.

*This function should return a (569, 31) DataFrame with*

*columns =*

```
['mean radius', 'mean texture', 'mean perimeter', 'mean area',  
'mean smoothness', 'mean compactness', 'mean concavity',  
'mean concave points', 'mean symmetry', 'mean fractal dimension',  
'radius error', 'texture error', 'perimeter error', 'area error',  
'smoothness error', 'compactness error', 'concavity error',  
'concave points error', 'symmetry error', 'fractal dimension error',  
'worst radius', 'worst texture', 'worst perimeter', 'worst area',  
'worst smoothness', 'worst compactness', 'worst concavity',  
'worst concave points', 'worst symmetry', 'worst fractal dimension',  
'target']
```

*and index =*

```
RangeIndex(start=0, stop=569, step=1)
```

```
In [3]: def answer_one():  
  
        df = pd.DataFrame(data=cancer.data, columns=cancer.feature_names)  
        df['target'] = cancer.target  
  
        return df  
  
#answer_one()
```

Out[3]:

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points
0	17.990	10.38	122.80	1001.0	0.11840	0.27760	0.300100	0.147100
1	20.570	17.77	132.90	1326.0	0.08474	0.07864	0.086900	0.070170
2	19.690	21.25	130.00	1203.0	0.10960	0.15990	0.197400	0.127900
3	11.420	20.38	77.58	386.1	0.14250	0.28390	0.241400	0.105200
4	20.290	14.34	135.10	1297.0	0.10030	0.13280	0.198000	0.104300
5	12.450	15.70	82.57	477.1	0.12780	0.17000	0.157800	0.080890
6	18.250	19.98	119.60	1040.0	0.09463	0.10900	0.112700	0.074000
7	13.710	20.83	90.20	577.9	0.11890	0.16450	0.093660	0.059850
8	13.000	21.82	87.50	519.8	0.12730	0.19320	0.185900	0.093530
9	12.460	24.04	83.97	475.9	0.11860	0.23960	0.227300	0.085430
10	16.020	23.24	102.70	797.8	0.08206	0.06669	0.032990	0.033230
11	15.780	17.89	103.60	781.0	0.09710	0.12920	0.099540	0.066060
12	19.170	24.80	132.40	1123.0	0.09740	0.24580	0.206500	0.111800
13	15.850	23.95	103.70	782.7	0.08401	0.10020	0.099380	0.053640
14	13.730	22.61	93.60	578.3	0.11310	0.22930	0.212800	0.080250
15	14.540	27.54	96.73	658.8	0.11390	0.15950	0.163900	0.073640
16	14.680	20.13	94.74	684.5	0.09867	0.07200	0.073950	0.052590
17	16.130	20.68	108.10	798.8	0.11700	0.20220	0.172200	0.102800
18	19.810	22.15	130.00	1260.0	0.09831	0.10270	0.147900	0.094980
19	13.540	14.36	87.46	566.3	0.09779	0.08129	0.066640	0.047810
20	13.080	15.71	85.63	520.0	0.10750	0.12700	0.045680	0.031100
21	9.504	12.44	60.34	273.9	0.10240	0.06492	0.029560	0.020760
22	15.340	14.26	102.50	704.4	0.10730	0.21350	0.207700	0.097560
23	21.160	23.04	137.20	1404.0	0.09428	0.10220	0.109700	0.086320
24	16.650	21.38	110.00	904.6	0.11210	0.14570	0.152500	0.091700
25	17.140	16.40	116.00	912.7	0.11860	0.22760	0.222900	0.140100
26	14.580	21.53	97.41	644.8	0.10540	0.18680	0.142500	0.087830
27	18.610	20.25	122.10	1094.0	0.09440	0.10660	0.149000	0.077310
28	15.300	25.27	102.40	732.4	0.10820	0.16970	0.168300	0.087510
29	17.570	15.05	115.00	955.1	0.09847	0.11570	0.098750	0.079530
...	...	...	...	...	...	...	...	...



	mean radius	mean texture	mean perimeter	mean area	mean smoothness	mean compactness	mean concavity	mean concave points
539	7.691	25.44	48.34	170.4	0.08668	0.11990	0.092520	0.013640
540	11.540	14.44	74.65	402.9	0.09984	0.11200	0.067370	0.025940
541	14.470	24.99	95.81	656.4	0.08837	0.12300	0.100900	0.038900
542	14.740	25.42	94.70	668.6	0.08275	0.07214	0.041050	0.030270
543	13.210	28.06	84.88	538.4	0.08671	0.06877	0.029870	0.032750
544	13.870	20.70	89.77	584.8	0.09578	0.10180	0.036880	0.023690
545	13.620	23.23	87.19	573.2	0.09246	0.06747	0.029740	0.024430
546	10.320	16.35	65.31	324.9	0.09434	0.04994	0.010120	0.005495
547	10.260	16.58	65.85	320.8	0.08877	0.08066	0.043580	0.024380
548	9.683	19.34	61.05	285.7	0.08491	0.05030	0.023370	0.009615
549	10.820	24.21	68.89	361.6	0.08192	0.06602	0.015480	0.008160
550	10.860	21.48	68.51	360.5	0.07431	0.04227	0.000000	0.000000
551	11.130	22.44	71.49	378.4	0.09566	0.08194	0.048240	0.022570
552	12.770	29.43	81.35	507.9	0.08276	0.04234	0.019970	0.014990
553	9.333	21.94	59.01	264.0	0.09240	0.05605	0.039960	0.012820
554	12.880	28.92	82.50	514.3	0.08123	0.05824	0.061950	0.023430
555	10.290	27.61	65.67	321.4	0.09030	0.07658	0.059990	0.027380
556	10.160	19.59	64.73	311.7	0.10030	0.07504	0.005025	0.011160
557	9.423	27.88	59.26	271.3	0.08123	0.04971	0.000000	0.000000
558	14.590	22.68	96.39	657.1	0.08473	0.13300	0.102900	0.037360
559	11.510	23.93	74.52	403.5	0.09261	0.10210	0.111200	0.041050
560	14.050	27.15	91.38	600.4	0.09929	0.11260	0.044620	0.043040
561	11.200	29.37	70.67	386.0	0.07449	0.03558	0.000000	0.000000
562	15.220	30.62	103.40	716.9	0.10480	0.20870	0.255000	0.094290
563	20.920	25.09	143.00	1347.0	0.10990	0.22360	0.317400	0.147400
564	21.560	22.39	142.00	1479.0	0.11100	0.11590	0.243900	0.138900
565	20.130	28.25	131.20	1261.0	0.09780	0.10340	0.144000	0.097910
566	16.600	28.08	108.30	858.1	0.08455	0.10230	0.092510	0.053020
567	20.600	29.33	140.10	1265.0	0.11780	0.27700	0.351400	0.152000
568	7.760	24.54	47.92	181.0	0.05263	0.04362	0.000000	0.000000

569 rows × 31 columns

## Question 2

What is the class distribution? (i.e. how many instances of malignant (encoded 0) and how many benign (encoded 1)?)

*This function should return a Series named target of length 2 with integer values and index = ['malignant', 'benign']*

```
In [4]: def answer_two():
        cancerdf = answer_one()

        target = cancerdf.target.value_counts()
        target = pd.Series([target[1], target[0]], index=['benign', 'malignant'])

        return target

#answer_two()
```

```
Out[4]: benign      357
        malignant   212
        dtype: int64
```

## Question 3

Split the DataFrame into X (the data) and y (the labels).

*This function should return a tuple of length 2: (X, y), where*

- X, a pandas DataFrame, has shape (569, 30)
- y, a pandas Series, has shape (569,).

```
In [6]: def answer_three():  
        cancerdf = answer_one()  
        X = cancerdf.loc[:, cancerdf.columns != 'target']  
        y = cancerdf['target']  
        #df_tuple = (X,y)  
  
        return X, y  
#answer_three()
```

```

Out[6]: (
\
0      17.990      10.38      122.80      1001.0      0.11840
1      20.570      17.77      132.90      1326.0      0.08474
2      19.690      21.25      130.00      1203.0      0.10960
3      11.420      20.38       77.58       386.1      0.14250
4      20.290      14.34      135.10      1297.0      0.10030
5      12.450      15.70       82.57       477.1      0.12780
6      18.250      19.98      119.60      1040.0      0.09463
7      13.710      20.83       90.20       577.9      0.11890
8      13.000      21.82       87.50       519.8      0.12730
9      12.460      24.04       83.97       475.9      0.11860
10     16.020      23.24      102.70       797.8      0.08206
11     15.780      17.89      103.60       781.0      0.09710
12     19.170      24.80      132.40      1123.0      0.09740
13     15.850      23.95      103.70       782.7      0.08401
14     13.730      22.61       93.60       578.3      0.11310
15     14.540      27.54       96.73       658.8      0.11390
16     14.680      20.13       94.74       684.5      0.09867
17     16.130      20.68      108.10       798.8      0.11700
18     19.810      22.15      130.00      1260.0      0.09831
19     13.540      14.36       87.46       566.3      0.09779
20     13.080      15.71       85.63       520.0      0.10750
21       9.504      12.44       60.34       273.9      0.10240
22     15.340      14.26      102.50       704.4      0.10730
23     21.160      23.04      137.20      1404.0      0.09428
24     16.650      21.38      110.00       904.6      0.11210
25     17.140      16.40      116.00       912.7      0.11860
26     14.580      21.53       97.41       644.8      0.10540
27     18.610      20.25      122.10      1094.0      0.09440

```

28	15.300	25.27	102.40	732.4	0.10820
29	17.570	15.05	115.00	955.1	0.09847
..	...	...	...	...	...
539	7.691	25.44	48.34	170.4	0.08668
540	11.540	14.44	74.65	402.9	0.09984
541	14.470	24.99	95.81	656.4	0.08837
542	14.740	25.42	94.70	668.6	0.08275
543	13.210	28.06	84.88	538.4	0.08671
544	13.870	20.70	89.77	584.8	0.09578
545	13.620	23.23	87.19	573.2	0.09246
546	10.320	16.35	65.31	324.9	0.09434
547	10.260	16.58	65.85	320.8	0.08877
548	9.683	19.34	61.05	285.7	0.08491
549	10.820	24.21	68.89	361.6	0.08192
550	10.860	21.48	68.51	360.5	0.07431
551	11.130	22.44	71.49	378.4	0.09566
552	12.770	29.43	81.35	507.9	0.08276
553	9.333	21.94	59.01	264.0	0.09240
554	12.880	28.92	82.50	514.3	0.08123
555	10.290	27.61	65.67	321.4	0.09030
556	10.160	19.59	64.73	311.7	0.10030
557	9.423	27.88	59.26	271.3	0.08123
558	14.590	22.68	96.39	657.1	0.08473
559	11.510	23.93	74.52	403.5	0.09261
560	14.050	27.15	91.38	600.4	0.09929
561	11.200	29.37	70.67	386.0	0.07449
562	15.220	30.62	103.40	716.9	0.10480
563	20.920	25.09	143.00	1347.0	0.10990

564	21.560	22.39	142.00	1479.0	0.11100
565	20.130	28.25	131.20	1261.0	0.09780
566	16.600	28.08	108.30	858.1	0.08455
567	20.600	29.33	140.10	1265.0	0.11780
568	7.760	24.54	47.92	181.0	0.05263

	mean compactness	mean concavity	mean concave points	mean symmetry \
0	0.27760	0.300100	0.147100	0.2419
1	0.07864	0.086900	0.070170	0.1812
2	0.15990	0.197400	0.127900	0.2069
3	0.28390	0.241400	0.105200	0.2597
4	0.13280	0.198000	0.104300	0.1809
5	0.17000	0.157800	0.080890	0.2087
6	0.10900	0.112700	0.074000	0.1794
7	0.16450	0.093660	0.059850	0.2196
8	0.19320	0.185900	0.093530	0.2350
9	0.23960	0.227300	0.085430	0.2030
10	0.06669	0.032990	0.033230	0.1528
11	0.12920	0.099540	0.066060	0.1842
12	0.24580	0.206500	0.111800	0.2397
13	0.10020	0.099380	0.053640	0.1847
14	0.22930	0.212800	0.080250	0.2069
15	0.15950	0.163900	0.073640	0.2303
16	0.07200	0.073950	0.052590	0.1586
17	0.20220	0.172200	0.102800	0.2164
18	0.10270	0.147900	0.094980	0.1582
19	0.08129	0.066640	0.047810	0.1885
20	0.12700	0.045680	0.031100	0.1967
21	0.06492	0.029560	0.020760	0.1815
22	0.21350	0.207700	0.097560	0.2521
23	0.10220	0.109700	0.086320	0.1769
24	0.14570	0.152500	0.091700	0.1995
25	0.22760	0.222900	0.140100	0.3040
26	0.18680	0.142500	0.087830	0.2252
27	0.10660	0.149000	0.077310	0.1697
28	0.16970	0.168300	0.087510	0.1926
29	0.11570	0.098750	0.079530	0.1739
..	...	...	...	...
539	0.11990	0.092520	0.013640	0.2037
540	0.11200	0.067370	0.025940	0.1818
541	0.12300	0.100900	0.038900	0.1872
542	0.07214	0.041050	0.030270	0.1840
543	0.06877	0.029870	0.032750	0.1628
544	0.10180	0.036880	0.023690	0.1620
545	0.06747	0.029740	0.024430	0.1664
546	0.04994	0.010120	0.005495	0.1885
547	0.08066	0.043580	0.024380	0.1669
548	0.05030	0.023370	0.009615	0.1580
549	0.06602	0.015480	0.008160	0.1976
550	0.04227	0.000000	0.000000	0.1661
551	0.08194	0.048240	0.022570	0.2030
552	0.04234	0.019970	0.014990	0.1539

553	0.05605	0.039960	0.012820	0.1692
554	0.05824	0.061950	0.023430	0.1566
555	0.07658	0.059990	0.027380	0.1593
556	0.07504	0.005025	0.011160	0.1791
557	0.04971	0.000000	0.000000	0.1742
558	0.13300	0.102900	0.037360	0.1454
559	0.10210	0.111200	0.041050	0.1388
560	0.11260	0.044620	0.043040	0.1537
561	0.03558	0.000000	0.000000	0.1060
562	0.20870	0.255000	0.094290	0.2128
563	0.22360	0.317400	0.147400	0.2149
564	0.11590	0.243900	0.138900	0.1726
565	0.10340	0.144000	0.097910	0.1752
566	0.10230	0.092510	0.053020	0.1590
567	0.27700	0.351400	0.152000	0.2397
568	0.04362	0.000000	0.000000	0.1587

	mean fractal dimension	...	worst radius \
0	0.07871	...	25.380
1	0.05667	...	24.990
2	0.05999	...	23.570
3	0.09744	...	14.910
4	0.05883	...	22.540
5	0.07613	...	15.470
6	0.05742	...	22.880
7	0.07451	...	17.060
8	0.07389	...	15.490
9	0.08243	...	15.090
10	0.05697	...	19.190
11	0.06082	...	20.420
12	0.07800	...	20.960
13	0.05338	...	16.840
14	0.07682	...	15.030
15	0.07077	...	17.460
16	0.05922	...	19.070
17	0.07356	...	20.960
18	0.05395	...	27.320
19	0.05766	...	15.110
20	0.06811	...	14.500
21	0.06905	...	10.230
22	0.07032	...	18.070
23	0.05278	...	29.170
24	0.06330	...	26.460
25	0.07413	...	22.250
26	0.06924	...	17.620
27	0.05699	...	21.310
28	0.06540	...	20.270
29	0.06149	...	20.010
..	...	...	...
539	0.07751	...	8.678
540	0.06782	...	12.260
541	0.06341	...	16.220
542	0.05680	...	16.510
543	0.05781	...	14.370
544	0.06688	...	15.050
545	0.05801	...	15.350
546	0.06201	...	11.250

547	0.06714	...	10.830
548	0.06235	...	10.930
549	0.06328	...	13.030
550	0.05948	...	11.660
551	0.06552	...	12.020
552	0.05637	...	13.870
553	0.06576	...	9.845
554	0.05708	...	13.890
555	0.06127	...	10.840
556	0.06331	...	10.650
557	0.06059	...	10.490
558	0.06147	...	15.480
559	0.06570	...	12.480
560	0.06171	...	15.300
561	0.05502	...	11.920
562	0.07152	...	17.520
563	0.06879	...	24.290
564	0.05623	...	25.450
565	0.05533	...	23.690
566	0.05648	...	18.980
567	0.07016	...	25.740
568	0.05884	...	9.456

	worst texture	worst perimeter	worst area	worst smoothness \
0	17.33	184.60	2019.0	0.16220
1	23.41	158.80	1956.0	0.12380
2	25.53	152.50	1709.0	0.14440
3	26.50	98.87	567.7	0.20980
4	16.67	152.20	1575.0	0.13740
5	23.75	103.40	741.6	0.17910
6	27.66	153.20	1606.0	0.14420
7	28.14	110.60	897.0	0.16540
8	30.73	106.20	739.3	0.17030
9	40.68	97.65	711.4	0.18530
10	33.88	123.80	1150.0	0.11810
11	27.28	136.50	1299.0	0.13960
12	29.94	151.70	1332.0	0.10370
13	27.66	112.00	876.5	0.11310
14	32.01	108.80	697.7	0.16510
15	37.13	124.10	943.2	0.16780
16	30.88	123.40	1138.0	0.14640
17	31.48	136.80	1315.0	0.17890
18	30.88	186.80	2398.0	0.15120
19	19.26	99.70	711.2	0.14400
20	20.49	96.09	630.5	0.13120
21	15.66	65.13	314.9	0.13240
22	19.08	125.10	980.9	0.13900
23	35.59	188.00	2615.0	0.14010
24	31.56	177.00	2215.0	0.18050
25	21.40	152.40	1461.0	0.15450
26	33.21	122.40	896.9	0.15250
27	27.26	139.90	1403.0	0.13380
28	36.71	149.30	1269.0	0.16410
29	19.52	134.90	1227.0	0.12550
..	...	...	...	...
539	31.89	54.49	223.6	0.15960
540	19.68	78.78	457.8	0.13450



541	31.73	113.50	808.9	0.13400
542	32.29	107.40	826.4	0.10600
543	37.17	92.48	629.6	0.10720
544	24.75	99.17	688.6	0.12640
545	29.09	97.58	729.8	0.12160
546	21.77	71.12	384.9	0.12850
547	22.04	71.08	357.4	0.14610
548	25.59	69.10	364.2	0.11990
549	31.45	83.90	505.6	0.12040
550	24.77	74.08	412.3	0.10010
551	28.26	77.80	436.6	0.10870
552	36.00	88.10	594.7	0.12340
553	25.05	62.86	295.8	0.11030
554	35.74	88.84	595.7	0.12270
555	34.91	69.57	357.6	0.13840
556	22.88	67.88	347.3	0.12650
557	34.24	66.50	330.6	0.10730
558	27.27	105.90	733.5	0.10260
559	37.16	82.28	474.2	0.12980
560	33.17	100.20	706.7	0.12410
561	38.30	75.19	439.6	0.09267
562	42.79	128.70	915.0	0.14170
563	29.41	179.10	1819.0	0.14070
564	26.40	166.10	2027.0	0.14100
565	38.25	155.00	1731.0	0.11660
566	34.12	126.70	1124.0	0.11390
567	39.42	184.60	1821.0	0.16500
568	30.37	59.16	268.6	0.08996

	worst compactness	worst concavity	worst concave points	worst symmetr
y \				
0	0.66560	0.71190	0.26540	0.460
1				
1	0.18660	0.24160	0.18600	0.275
0				
2	0.42450	0.45040	0.24300	0.361
3				
3	0.86630	0.68690	0.25750	0.663
8				
4	0.20500	0.40000	0.16250	0.236
4				
5	0.52490	0.53550	0.17410	0.398
5				
6	0.25760	0.37840	0.19320	0.306
3				
7	0.36820	0.26780	0.15560	0.319
6				
8	0.54010	0.53900	0.20600	0.437
8				
9	1.05800	1.10500	0.22100	0.436
6				
10	0.15510	0.14590	0.09975	0.294
8				
11	0.56090	0.39650	0.18100	0.379
2				
12	0.39030	0.36390	0.17670	0.317
6				

13	0.19240	0.23220	0.11190	0.280
9				
14	0.77250	0.69430	0.22080	0.359
6				
15	0.65770	0.70260	0.17120	0.421
8				
16	0.18710	0.29140	0.16090	0.302
9				
17	0.42330	0.47840	0.20730	0.370
6				
18	0.31500	0.53720	0.23880	0.276
8				
19	0.17730	0.23900	0.12880	0.297
7				
20	0.27760	0.18900	0.07283	0.318
4				
21	0.11480	0.08867	0.06227	0.245
0				
22	0.59540	0.63050	0.23930	0.466
7				
23	0.26000	0.31550	0.20090	0.282
2				
24	0.35780	0.46950	0.20950	0.361
3				
25	0.39490	0.38530	0.25500	0.406
6				
26	0.66430	0.55390	0.27010	0.426
4				
27	0.21170	0.34460	0.14900	0.234
1				
28	0.61100	0.63350	0.20240	0.402
7				
29	0.28120	0.24890	0.14560	0.275
6				
..	...	...	...	
...				
539	0.30640	0.33930	0.05000	0.279
0				
540	0.21180	0.17970	0.06918	0.232
9				
541	0.42020	0.40400	0.12050	0.318
7				
542	0.13760	0.16110	0.10950	0.272
2				
543	0.13810	0.10620	0.07958	0.247
3				
544	0.20370	0.13770	0.06845	0.224
9				
545	0.15170	0.10490	0.07174	0.264
2				
546	0.08842	0.04384	0.02381	0.268
1				
547	0.22460	0.17830	0.08333	0.269
1				
548	0.09546	0.09350	0.03846	0.255
2				
549	0.16330	0.06194	0.03264	0.305

9				
550	0.07348	0.00000	0.00000	0.245
8				
551	0.17820	0.15640	0.06413	0.316
9				
552	0.10640	0.08653	0.06498	0.240
7				
553	0.08298	0.07993	0.02564	0.243
5				
554	0.16200	0.24390	0.06493	0.237
2				
555	0.17100	0.20000	0.09127	0.222
6				
556	0.12000	0.01005	0.02232	0.226
2				
557	0.07158	0.00000	0.00000	0.247
5				
558	0.31710	0.36620	0.11050	0.225
8				
559	0.25170	0.36300	0.09653	0.211
2				
560	0.22640	0.13260	0.10480	0.225
0				
561	0.05494	0.00000	0.00000	0.156
6				
562	0.79170	1.17000	0.23560	0.408
9				
563	0.41860	0.65990	0.25420	0.292
9				
564	0.21130	0.41070	0.22160	0.206
0				
565	0.19220	0.32150	0.16280	0.257
2				
566	0.30940	0.34030	0.14180	0.221
8				
567	0.86810	0.93870	0.26500	0.408
7				
568	0.06444	0.00000	0.00000	0.287
1				

worst fractal dimension

0	0.11890
1	0.08902
2	0.08758
3	0.17300
4	0.07678
5	0.12440
6	0.08368
7	0.11510
8	0.10720
9	0.20750
10	0.08452
11	0.10480
12	0.10230
13	0.06287
14	0.14310
15	0.13410

16	0.08216
17	0.11420
18	0.07615
19	0.07259
20	0.08183
21	0.07773
22	0.09946
23	0.07526
24	0.09564
25	0.10590
26	0.12750
27	0.07421
28	0.09876
29	0.07919
..	...
539	0.10660
540	0.08134
541	0.10230
542	0.06956
543	0.06443
544	0.08492
545	0.06953
546	0.07399
547	0.09479
548	0.07920
549	0.07626
550	0.06592
551	0.08032
552	0.06484
553	0.07393
554	0.07242
555	0.08283
556	0.06742
557	0.06969
558	0.08004
559	0.08732
560	0.08321
561	0.05905
562	0.14090
563	0.09873
564	0.07115
565	0.06637
566	0.07820
567	0.12400
568	0.07039

[569 rows x 30 columns], 0      0

1	0
2	0
3	0
4	0
5	0
6	0
7	0
8	0
9	0
10	0

11	0
12	0
13	0
14	0
15	0
16	0
17	0
18	0
19	1
20	1
21	1
22	0
23	0
24	0
25	0
26	0
27	0
28	0
29	0

..

539	1
540	1
541	1
542	1
543	1
544	1
545	1
546	1
547	1
548	1
549	1
550	1
551	1
552	1
553	1
554	1
555	1
556	1
557	1
558	1
559	1
560	1
561	1
562	0
563	0
564	0
565	0
566	0
567	0
568	1

Name: target, dtype: int64)

## Question 4

Using `train_test_split`, split `X` and `y` into training and test sets (`X_train`, `X_test`, `y_train`, and `y_test`).

**Set the random number generator state to 0 using `random_state=0` to make sure your results match the autograder!**

*This function should return a tuple of length 4: (`X_train`, `X_test`, `y_train`, `y_test`), where*

- `X_train` has shape (426, 30)
- `X_test` has shape (143, 30)
- `y_train` has shape (426,)
- `y_test` has shape (143,)

```
In [16]: from sklearn.model_selection import train_test_split

def answer_four():
    X, y = answer_three()

    X_train, X_test, y_train, y_test = train_test_split(X, y, random_state=0)
    y_train = y_train.astype(np.float)
    y_test = y_test.astype(np.float)

    return X_train, X_test, y_train, y_test
```

## Question 5

Using `KNeighborsClassifier`, fit a k-nearest neighbors (knn) classifier with `X_train`, `y_train` and using one nearest neighbor (`n_neighbors = 1`).

*This function should return a `sklearn.neighbors.classification.KNeighborsClassifier`.*

```
In [19]: from sklearn.neighbors import KNeighborsClassifier

def answer_five():
    X_train, X_test, y_train, y_test = answer_four()
    knn = KNeighborsClassifier(n_neighbors = 1)
    knn.fit(X_train, y_train)

    return knn
```

## Question 6

Using your knn classifier, predict the class label using the mean value for each feature.

Hint: You can use `cancerdf.mean()[:-1].values.reshape(1, -1)` which gets the mean value for each feature, ignores the target column, and reshapes the data from 1 dimension to 2 (necessary for the `predict` method of `KNeighborsClassifier`).

*This function should return a numpy array either `array([ 0.])` or `array([ 1.])`*

```
In [20]: def answer_six():
          cancerdf = answer_one()
          knn = answer_five()
          means = cancerdf.mean()[:-1].values.reshape(1, -1)
          knn.predict(means)

          return knn.predict(means)

          #answer_six()
```

Out[20]: `array([ 1.])`

## Question 7

Using your knn classifier, predict the class labels for the test set `X_test`.

*This function should return a numpy array with shape `(143,)` and values either `0.0` or `1.0`.*

```
In [21]: def answer_seven():
          X_train, X_test, y_train, y_test = answer_four()
          knn = answer_five()
          p = knn.predict(X_test)

          return p
          answer_seven()
```

Out[21]: `array([ 1., 1., 1., 0., 1., 1., 1., 1., 1., 1., 0., 1., 1.,`  
`1., 0., 0., 1., 0., 0., 0., 0., 1., 1., 1., 0., 1.,`  
`1., 1., 1., 0., 1., 0., 1., 0., 1., 0., 1., 0., 1.,`  
`0., 0., 1., 0., 1., 0., 0., 1., 1., 1., 0., 0., 1.,`  
`0., 1., 1., 1., 1., 1., 1., 0., 0., 0., 1., 1., 0.,`  
`1., 0., 0., 0., 1., 1., 0., 1., 1., 0., 1., 1., 1.,`  
`1., 1., 0., 0., 0., 1., 0., 1., 1., 1., 0., 0., 1.,`  
`0., 1., 0., 1., 1., 0., 1., 1., 1., 1., 1., 1., 1.,`  
`0., 1., 0., 1., 0., 1., 1., 0., 0., 1., 1., 1., 0.,`  
`1., 1., 1., 1., 1., 1., 1., 0., 1., 1., 1., 1., 1.,`  
`0., 1., 1., 1., 1., 1., 1., 0., 0., 1., 1., 1., 0.]`

## Question 8

Find the score (mean accuracy) of your knn classifier using `X_test` and `y_test`.

*This function should return a float between 0 and 1*

```
In [22]: def answer_eight():
          X_train, X_test, y_train, y_test = answer_four()
          knn = answer_five()

          return knn.score(X_test, y_test)

          answer_eight()
```

```
Out[22]: 0.91608391608391604
```

## Optional plot

Try using the plotting function below to visualize the different prediction scores between training and test sets, as well as malignant and benign cells.



```

In [ ]: def accuracy_plot():
    import matplotlib.pyplot as plt

    %matplotlib notebook

    X_train, X_test, y_train, y_test = answer_four()

    # Find the training and testing accuracies by target value (i.e. malignant, benign)
    mal_train_X = X_train[y_train==0]
    mal_train_y = y_train[y_train==0]
    ben_train_X = X_train[y_train==1]
    ben_train_y = y_train[y_train==1]

    mal_test_X = X_test[y_test==0]
    mal_test_y = y_test[y_test==0]
    ben_test_X = X_test[y_test==1]
    ben_test_y = y_test[y_test==1]

    knn = answer_five()

    scores = [knn.score(mal_train_X, mal_train_y), knn.score(ben_train_X, ben_train_y),
               knn.score(mal_test_X, mal_test_y), knn.score(ben_test_X, ben_test_y)]

    plt.figure()

    # Plot the scores as a bar chart
    bars = plt.bar(np.arange(4), scores, color=['#4c72b0', '#4c72b0', '#55a868', '#55a868'])

    # directly label the score onto the bars
    for bar in bars:
        height = bar.get_height()
        plt.gca().text(bar.get_x() + bar.get_width()/2, height*.90, '{0:.1f}'.format(height, 2),
                        ha='center', color='w', fontsize=11)

    # remove all the ticks (both axes), and tick labels on the Y axis
    plt.tick_params(top='off', bottom='off', left='off', right='off', labelleft='off', labelbottom='on')

    # remove the frame of the chart
    for spine in plt.gca().spines.values():
        spine.set_visible(False)

    plt.xticks([0,1,2,3], ['Malignant\nTraining', 'Benign\nTraining', 'Malignant\nTest', 'Benign\nTest'], alpha=0.8);
    plt.title('Training and Test Accuracies for Malignant and Benign Cells', alpha=0.8)

```

Uncomment the plotting function to see the visualization.

**Comment out** the plotting function when submitting your notebook for grading.

```
In [ ]: #accuracy_plot()
```

```
In [ ]:
```