

Breast Cancer Wisconsin (Diagnostic)

January 12, 2022

```
[1]: import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
import seaborn as sns # data visualization library
import matplotlib.pyplot as plt
import time
```

```
[2]: data = pd.read_csv('data.csv')
```

```
[3]: data.head()
```

```
[3]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	\
0	842302	M	17.99	10.38	122.80	1001.0	
1	842517	M	20.57	17.77	132.90	1326.0	
2	84300903	M	19.69	21.25	130.00	1203.0	
3	84348301	M	11.42	20.38	77.58	386.1	
4	84358402	M	20.29	14.34	135.10	1297.0	

	smoothness_mean	compactness_mean	concavity_mean	concave	points_mean	\
0	0.11840	0.27760	0.3001		0.14710	
1	0.08474	0.07864	0.0869		0.07017	
2	0.10960	0.15990	0.1974		0.12790	
3	0.14250	0.28390	0.2414		0.10520	
4	0.10030	0.13280	0.1980		0.10430	

	...	texture_worst	perimeter_worst	area_worst	smoothness_worst	\
0	...	17.33	184.60	2019.0	0.1622	
1	...	23.41	158.80	1956.0	0.1238	
2	...	25.53	152.50	1709.0	0.1444	
3	...	26.50	98.87	567.7	0.2098	
4	...	16.67	152.20	1575.0	0.1374	

	compactness_worst	concavity_worst	concave	points_worst	symmetry_worst	\
0	0.6656	0.7119		0.2654	0.4601	
1	0.1866	0.2416		0.1860	0.2750	
2	0.4245	0.4504		0.2430	0.3613	
3	0.8663	0.6869		0.2575	0.6638	
4	0.2050	0.4000		0.1625	0.2364	

```

    fractal_dimension_worst  Unnamed: 32
0          0.11890          NaN
1          0.08902          NaN
2          0.08758          NaN
3          0.17300          NaN
4          0.07678          NaN

```

[5 rows x 33 columns]

```

[4]: col = data.columns
     print(col)

```

```

Index(['id', 'diagnosis', 'radius_mean', 'texture_mean', 'perimeter_mean',
      'area_mean', 'smoothness_mean', 'compactness_mean', 'concavity_mean',
      'concave points_mean', 'symmetry_mean', 'fractal_dimension_mean',
      'radius_se', 'texture_se', 'perimeter_se', 'area_se', 'smoothness_se',
      'compactness_se', 'concavity_se', 'concave points_se', 'symmetry_se',
      'fractal_dimension_se', 'radius_worst', 'texture_worst',
      'perimeter_worst', 'area_worst', 'smoothness_worst',
      'compactness_worst', 'concavity_worst', 'concave points_worst',
      'symmetry_worst', 'fractal_dimension_worst', 'Unnamed: 32'],
      dtype='object')

```

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[5]: y = data.diagnosis
     drop_cols = ['Unnamed: 32', 'id', 'diagnosis']
     x = data.drop(drop_cols, axis=1)
     x.head()

```

```

[5]:   radius_mean  texture_mean  perimeter_mean  area_mean  smoothness_mean  \
0         17.99         10.38         122.80        1001.0         0.11840
1         20.57         17.77         132.90        1326.0         0.08474
2         19.69         21.25         130.00        1203.0         0.10960
3         11.42         20.38          77.58         386.1         0.14250
4         20.29         14.34         135.10        1297.0         0.10030

      compactness_mean  concavity_mean  concave points_mean  symmetry_mean  \
0          0.27760         0.3001         0.14710         0.2419
1          0.07864         0.0869         0.07017         0.1812
2          0.15990         0.1974         0.12790         0.2069
3          0.28390         0.2414         0.10520         0.2597
4          0.13280         0.1980         0.10430         0.1809

      fractal_dimension_mean  ...  radius_worst  texture_worst  perimeter_worst  \
0          0.07871  ...         25.38         17.33         184.60
1          0.05667  ...         24.99         23.41         158.80
2          0.05999  ...         23.57         25.53         152.50

```

3	0.09744	...	14.91	26.50	98.87
4	0.05883	...	22.54	16.67	152.20

	area_worst	smoothness_worst	compactness_worst	concavity_worst	\
0	2019.0	0.1622	0.6656	0.7119	
1	1956.0	0.1238	0.1866	0.2416	
2	1709.0	0.1444	0.4245	0.4504	
3	567.7	0.2098	0.8663	0.6869	
4	1575.0	0.1374	0.2050	0.4000	

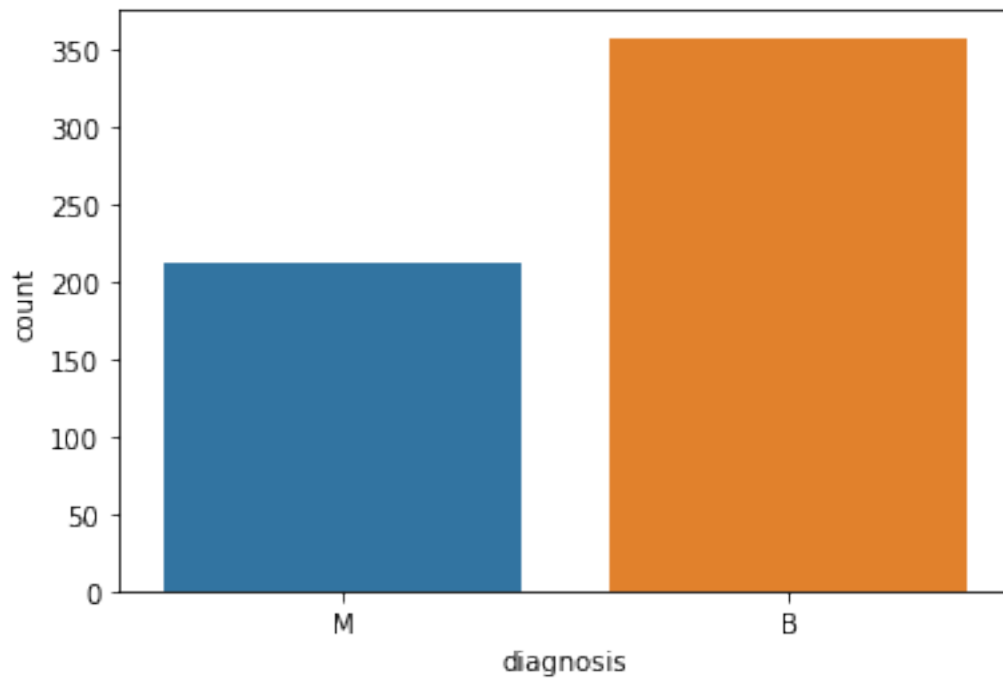
	concave points_worst	symmetry_worst	fractal_dimension_worst
0	0.2654	0.4601	0.11890
1	0.1860	0.2750	0.08902
2	0.2430	0.3613	0.08758
3	0.2575	0.6638	0.17300
4	0.1625	0.2364	0.07678

[5 rows x 30 columns]

```
[6]: ax = sns.countplot(y, label="Count")
      B, M = y.value_counts()
      print('Number of Benign Tumors', B)
      print('Number of Malignant Tumors', M)
```

Number of Benign Tumors 357

Number of Malignant Tumors 212



```
[7]: x.describe()
```

```
[7]:
```

	radius_mean	texture_mean	perimeter_mean	area_mean	\
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	

	smoothness_mean	compactness_mean	concavity_mean	concave points_mean	\
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	

	symmetry_mean	fractal_dimension_mean	...	radius_worst	\
count	569.000000	569.000000	...	569.000000	
mean	0.181162	0.062798	...	16.269190	
std	0.027414	0.007060	...	4.833242	
min	0.106000	0.049960	...	7.930000	
25%	0.161900	0.057700	...	13.010000	
50%	0.179200	0.061540	...	14.970000	
75%	0.195700	0.066120	...	18.790000	
max	0.304000	0.097440	...	36.040000	

	texture_worst	perimeter_worst	area_worst	smoothness_worst	\
count	569.000000	569.000000	569.000000	569.000000	
mean	25.677223	107.261213	880.583128	0.132369	
std	6.146258	33.602542	569.356993	0.022832	
min	12.020000	50.410000	185.200000	0.071170	
25%	21.080000	84.110000	515.300000	0.116600	
50%	25.410000	97.660000	686.500000	0.131300	
75%	29.720000	125.400000	1084.000000	0.146000	
max	49.540000	251.200000	4254.000000	0.222600	

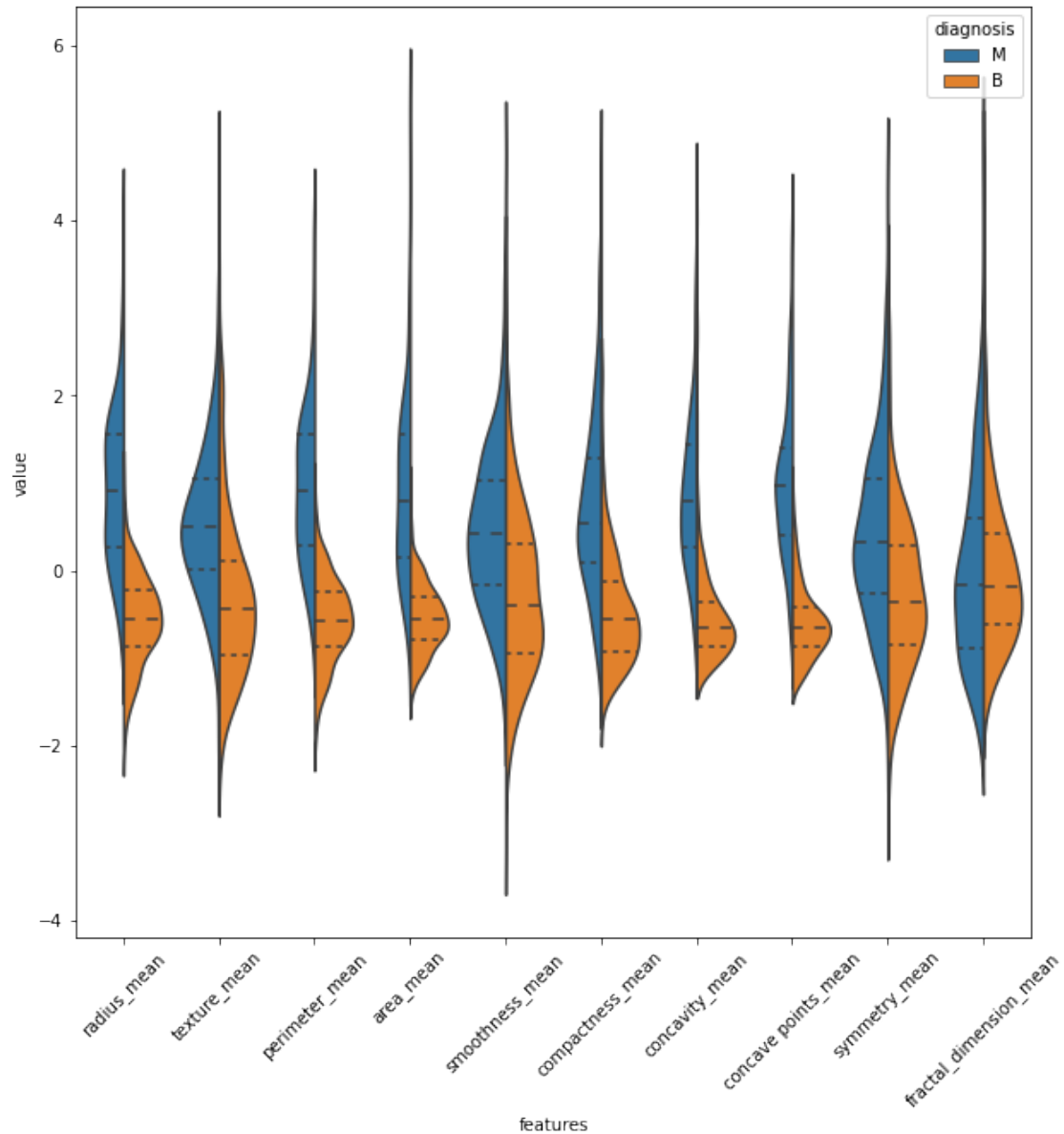
	compactness_worst	concavity_worst	concave points_worst	\
count	569.000000	569.000000	569.000000	
mean	0.254265	0.272188	0.114606	
std	0.157336	0.208624	0.065732	

min	0.027290	0.000000	0.000000
25%	0.147200	0.114500	0.064930
50%	0.211900	0.226700	0.099930
75%	0.339100	0.382900	0.161400
max	1.058000	1.252000	0.291000

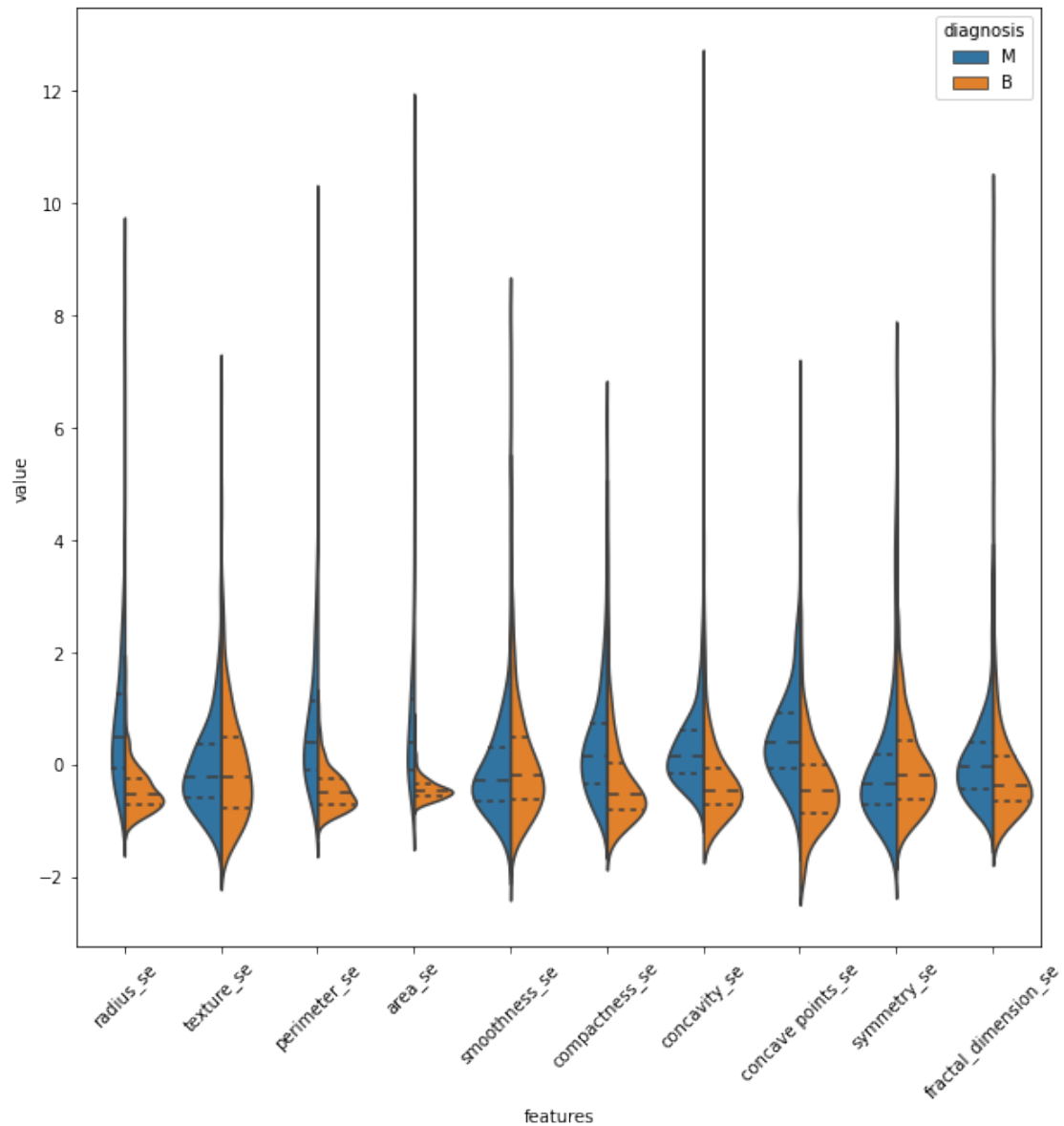
	symmetry_worst	fractal_dimension_worst
count	569.000000	569.000000
mean	0.290076	0.083946
std	0.061867	0.018061
min	0.156500	0.055040
25%	0.250400	0.071460
50%	0.282200	0.080040
75%	0.317900	0.092080
max	0.663800	0.207500

[8 rows x 30 columns]

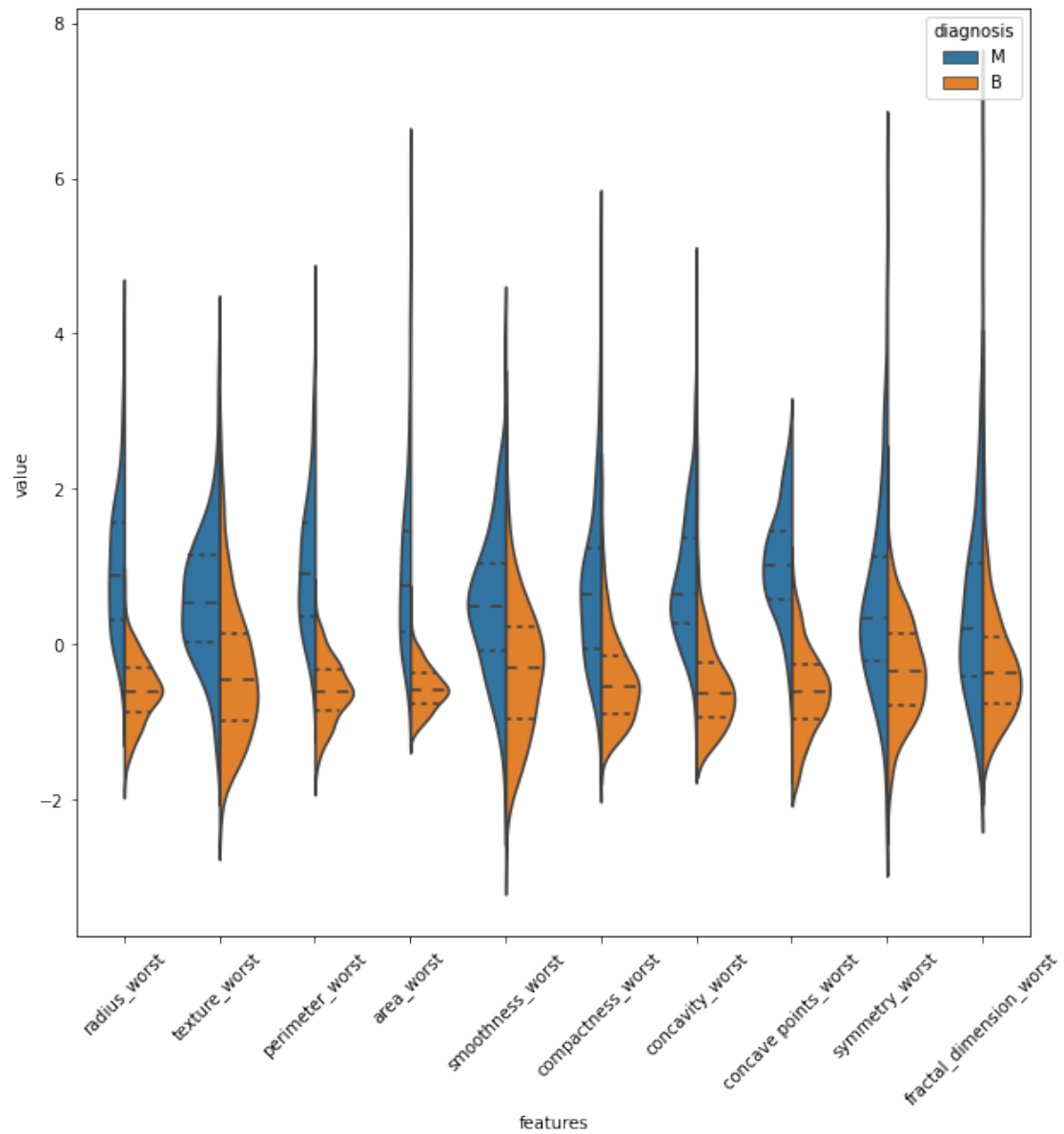
```
[8]: data = x
data_std = (data - data.mean()) / data.std()
data = pd.concat([y, data_std.iloc[:,0:10]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.violinplot(x='features', y='value', hue='diagnosis', data=data, split=True,
               inner='quart')
plt.xticks(rotation=45);
```



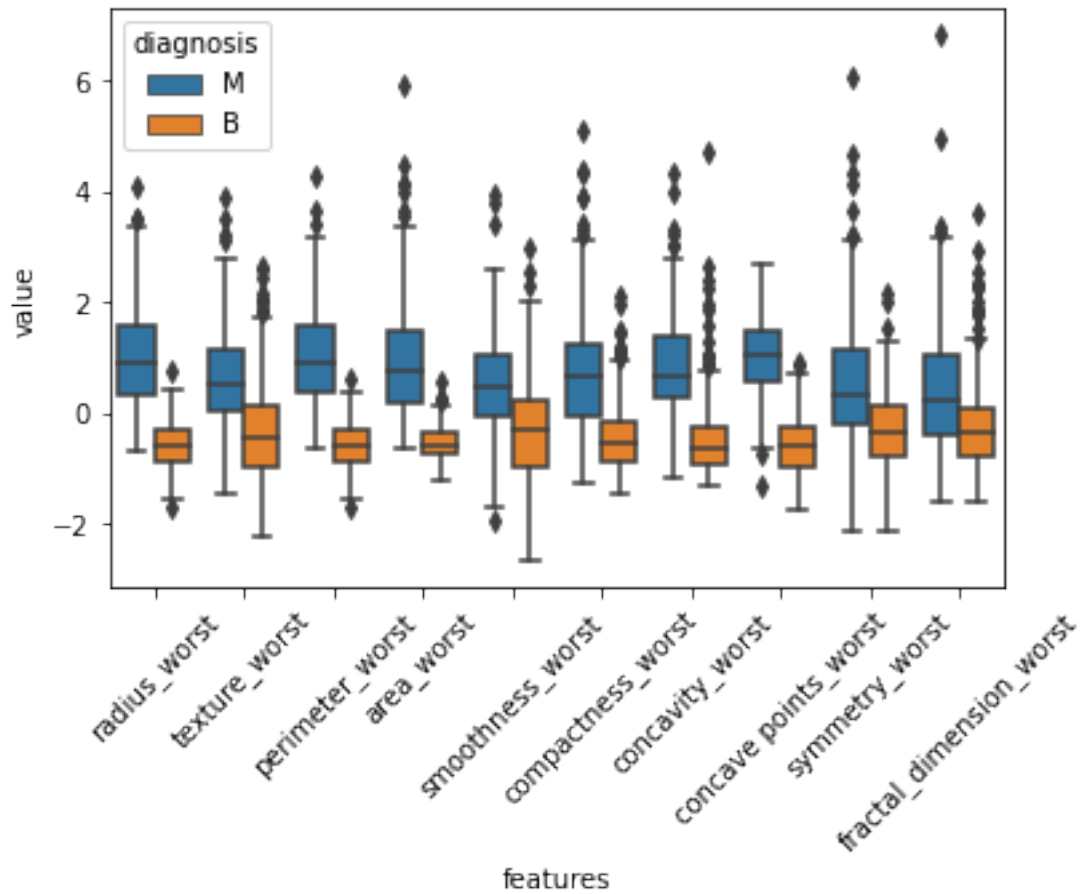
```
[9]: data = pd.concat([y, data_std.iloc[:,10:20]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.violinplot(x='features', y='value', hue='diagnosis', data=data, split=True,
               inner='quart')
plt.xticks(rotation=45);
```



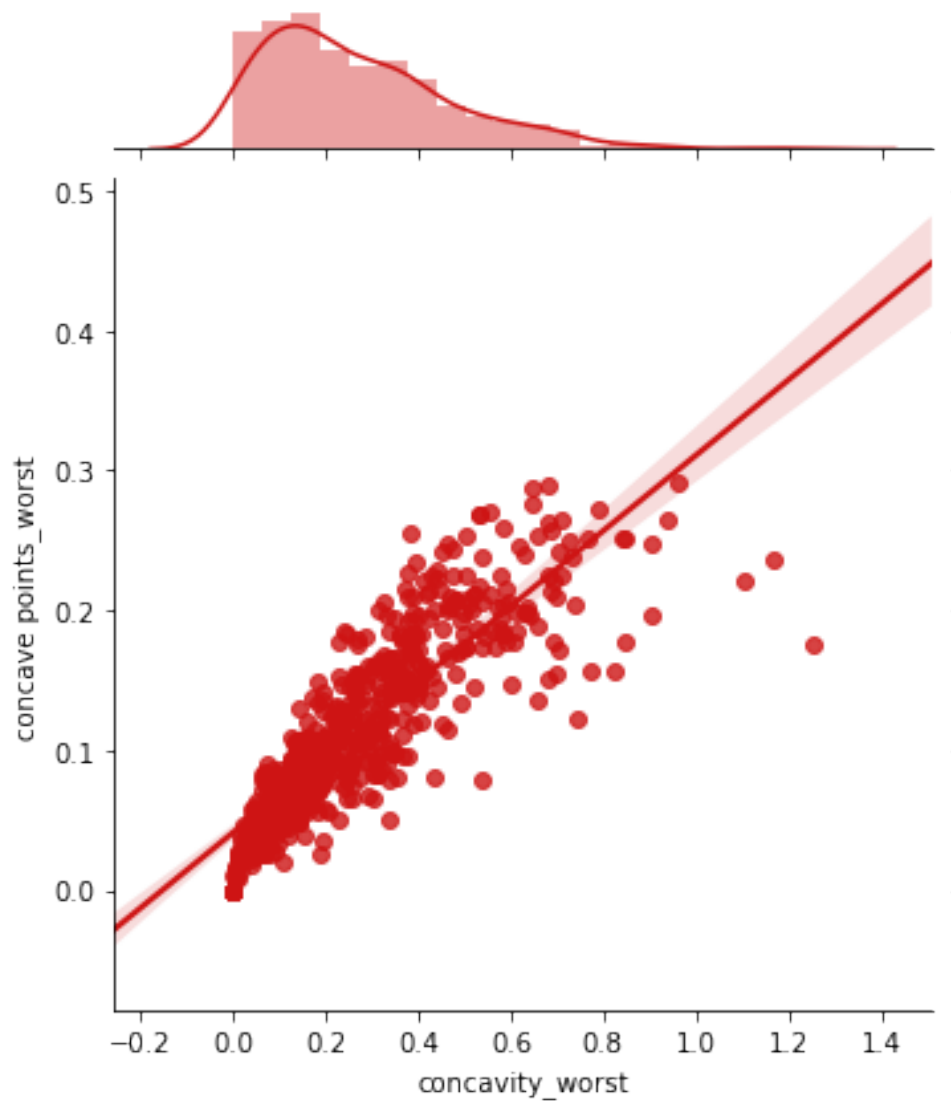
```
[10]: data = pd.concat([y, data_std.iloc[:,20:30]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.violinplot(x='features', y='value', hue='diagnosis', data=data, split=True,
               inner='quart')
plt.xticks(rotation=45);
```



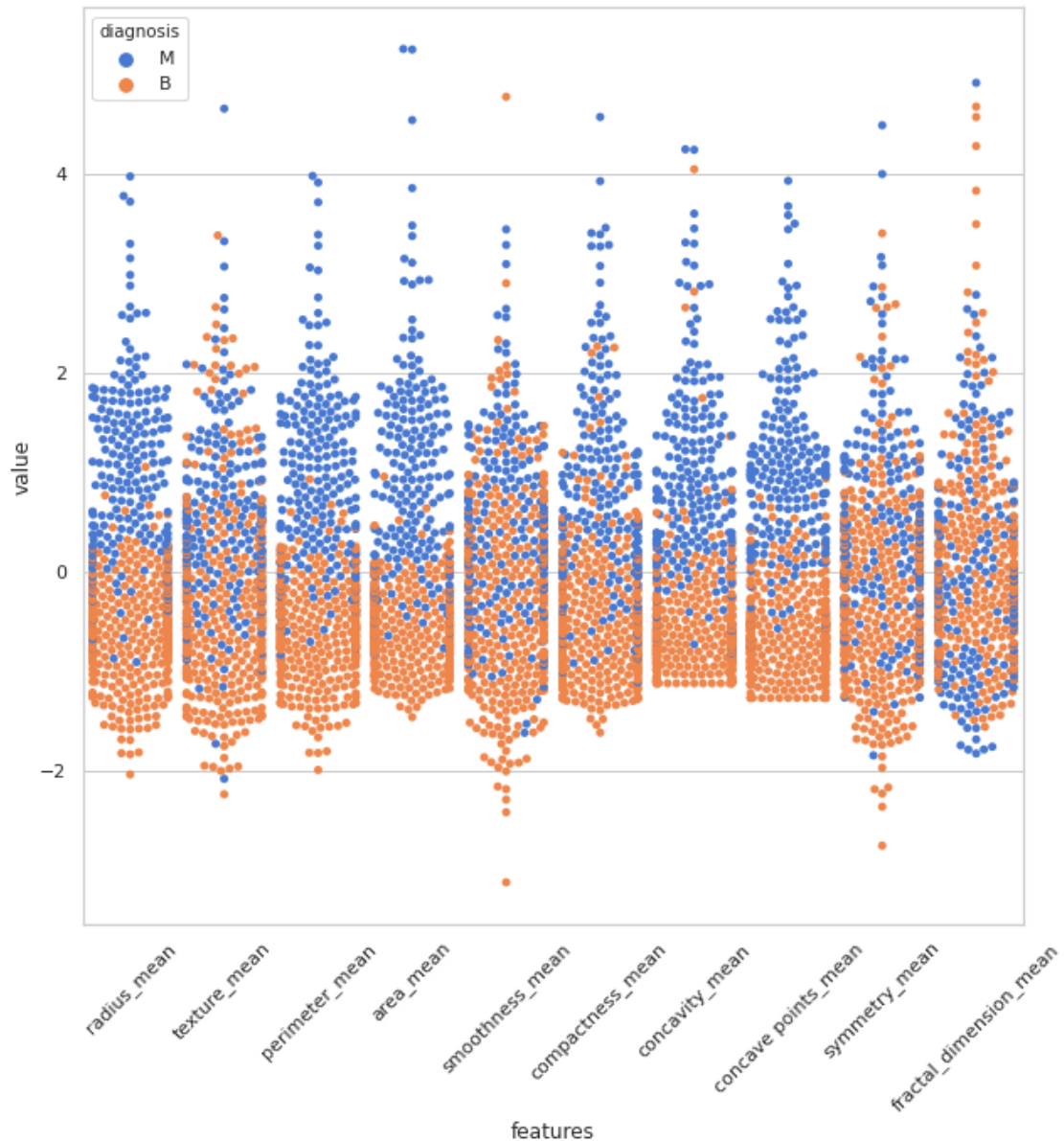
```
[11]: sns.boxplot(x='features', y='value', hue='diagnosis', data=data)
plt.xticks(rotation=45);
```

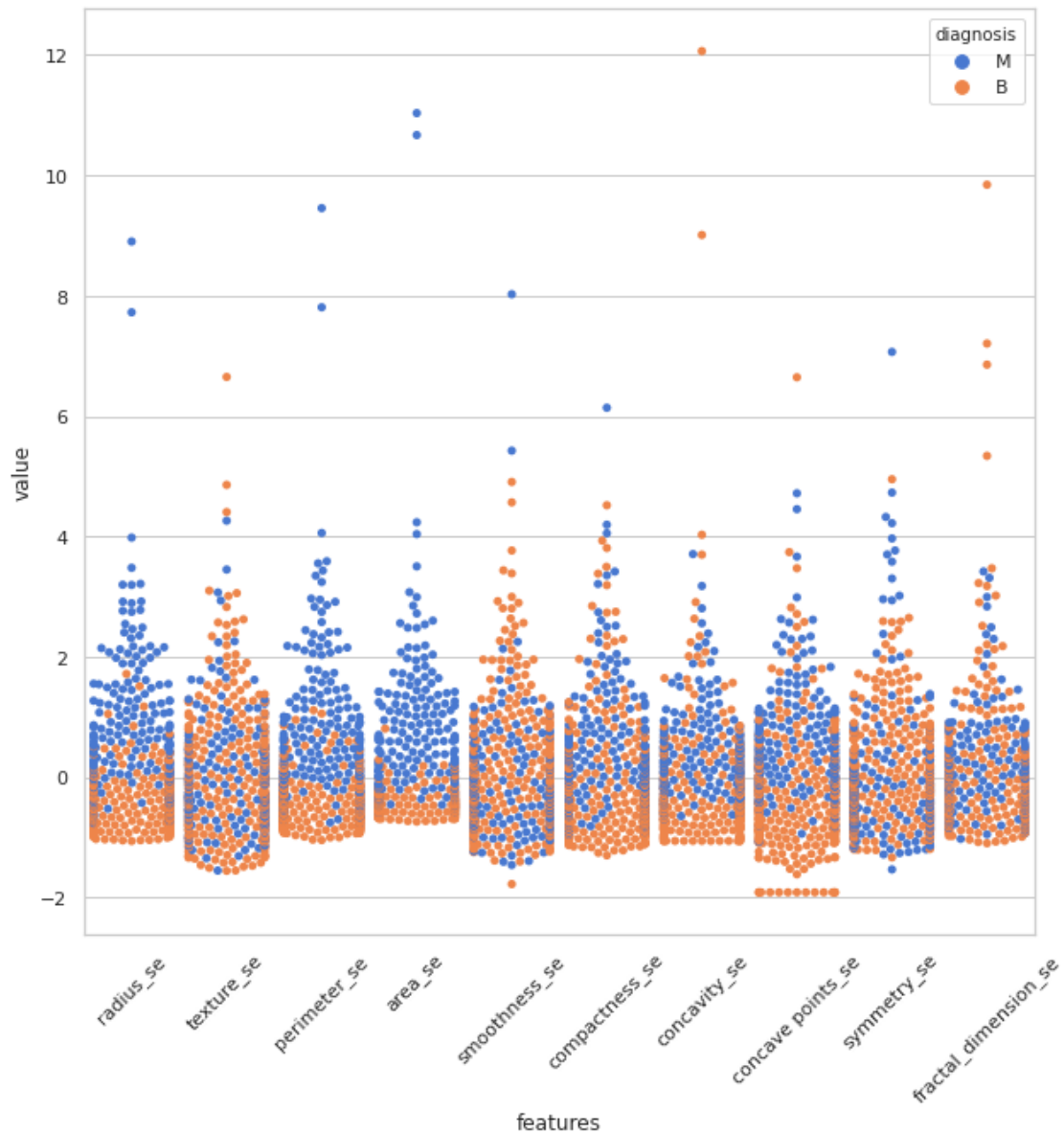
```
[12]: sns.jointplot(x.loc[:, 'concavity_worst'],
                  x.loc[:, 'concave points_worst'],
                  kind='regg',
                  color='#ce1414');
```



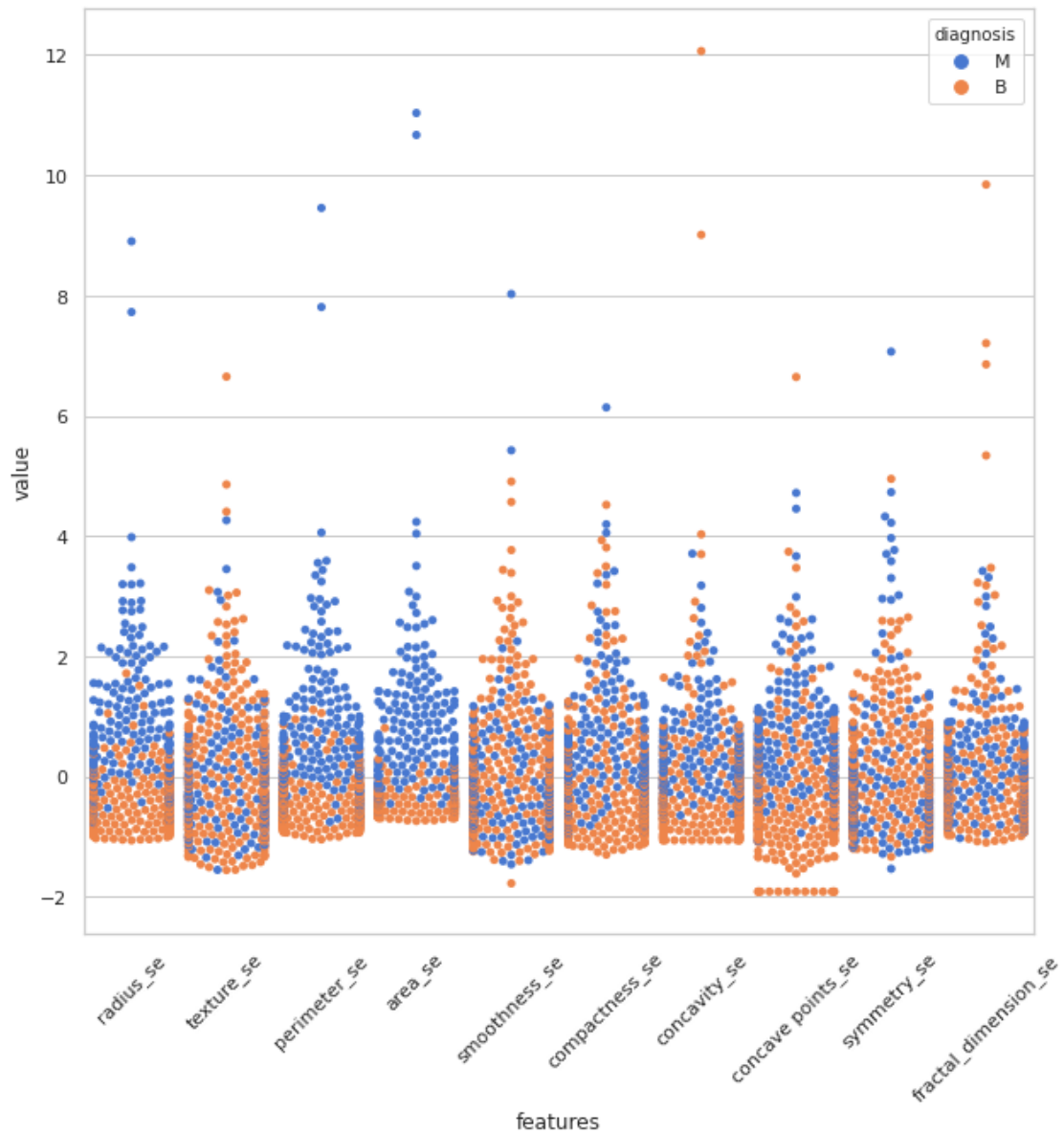
```
[13]: sns.set(style='whitegrid', palette='muted')
data = x
data_std = (data - data.mean()) / data.std()
data = pd.concat([y, data_std.iloc[:,0:10]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.swarmplot(x='features', y='value', hue='diagnosis', data=data)
plt.xticks(rotation=45);
```



```
[14]: sns.set(style='whitegrid', palette='muted')
data = x
data_std = (data - data.mean()) / data.std()
data = pd.concat([y, data_std.iloc[:,10:20]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.swarmplot(x='features', y='value', hue='diagnosis', data=data)
plt.xticks(rotation=45);
```

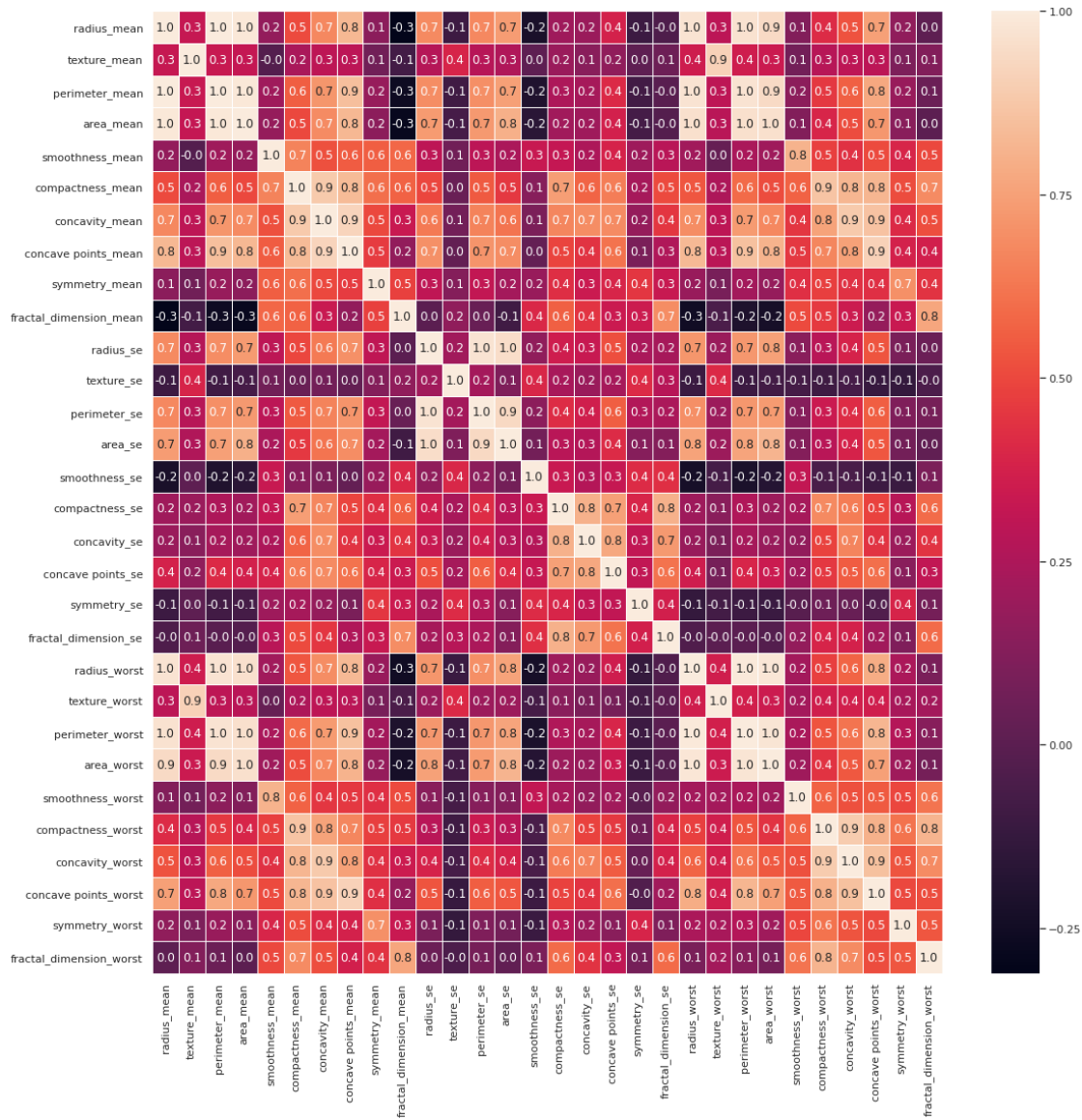


```
[15]: sns.set(style='whitegrid', palette='muted')
data = x
data_std = (data - data.mean()) / data.std()
data = pd.concat([y, data_std.iloc[:,10:20]], axis=1)
data = pd.melt(data, id_vars='diagnosis',
               var_name='features',
               value_name='value')
plt.figure(figsize=(10,10))
sns.swarmplot(x='features', y='value', hue='diagnosis', data=data)
plt.xticks(rotation=45);
```



```
[16]: f, ax = plt.subplots(figsize=(18, 18))
      sns.heatmap(x.corr(), annot=True, linewidth=.5, fmt='.1f', ax=ax)
```

```
[16]: <AxesSubplot:>
```



[]: