

# Breast Cancer Prediction with SVM

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## Breast Cancer Prediction using Support Vector Machines

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Importing Libraries

```
[4]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
```

Getting the data

```
[5]: from sklearn.datasets import load_breast_cancer
```

```
[6]: cancer = load_breast_cancer()
```

```
[8]: cancer.keys()
```

```
[8]: dict_keys(['data', 'target', 'frame', 'target_names', 'DESCR', 'feature_names',
'filename'])
```

```
[10]: df = pd.DataFrame(cancer['data'], columns=cancer['feature_names'])
```

```
[11]: df.head()
```

```
[11]:
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	\
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	

  

	mean compactness	mean concavity	mean concave points	mean symmetry	\
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

  

	mean fractal dimension	...	worst radius	worst texture	worst perimeter \
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

  

	worst area	worst smoothness	worst compactness	worst concavity \
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

  

	worst concave points	worst symmetry	worst fractal dimension
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]

```
[12]: cancer['target_names']
```

```
[12]: array(['malignant', 'benign'], dtype='<U9')
```

## Exploratory Data Analysis

Due to the lack of domain knowledge, I am skipping this step as I am incapable of interpreting the visual comparison between features,

## Train Test Split

```
[13]: from sklearn.model_selection import train_test_split
```

```
[14]: X = df
      y = cancer['target']

      X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.30,
      ↪random_state=101)
```

## Training Support Vector Classifier Model

```
[15]: from sklearn.svm import SVC
```

<frozen importlib.\_bootstrap>:219: RuntimeWarning: numpy.ufunc size changed, may

indicate binary incompatibility. Expected 192 from C header, got 216 from PyObject

```
[16]: model = SVC()
```

```
[17]: model.fit(X_train, y_train)
```

```
[17]: SVC()
```

```
[18]: predict = model.predict(X_test)
```

```
[19]: from sklearn.metrics import classification_report, confusion_matrix
```

```
[20]: confusion_matrix(y_test, predict)
```

```
[20]: array([[ 56,  10],
        [   3, 102]], dtype=int64)
```

```
[21]: print(classification_report(y_test, predict))
```

	precision	recall	f1-score	support
0	0.95	0.85	0.90	66
1	0.91	0.97	0.94	105
accuracy			0.92	171
macro avg	0.93	0.91	0.92	171
weighted avg	0.93	0.92	0.92	171

Gridsearch

```
[23]: from sklearn.model_selection import GridSearchCV
```

```
[25]: pgrid = {'C':[0.1,1,10,100,1000], 'gamma':[1,0.1,0.01,0.001,0.0001]}
```

```
[26]: grid = GridSearchCV(SVC(), pgrid, verbose=5)
```

```
[27]: grid.fit(X_train, y_train)
```

Fitting 5 folds for each of 25 candidates, totalling 125 fits

```
[CV] C=0.1, gamma=1 ...
[CV] ... C=0.1, gamma=1, score=0.637, total= 0.0s
[CV] C=0.1, gamma=1 ...
[CV] ... C=0.1, gamma=1, score=0.637, total= 0.0s
[CV] C=0.1, gamma=1 ...
[CV] ... C=0.1, gamma=1, score=0.625, total= 0.0s
[CV] C=0.1, gamma=1 ...
[CV] ... C=0.1, gamma=1, score=0.633, total= 0.0s
[CV] C=0.1, gamma=1 ...
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```

[CV] ... C=0.1, gamma=1, score=0.633, total= 0.0s
[CV] C=0.1, gamma=0.1 ...
[CV] ... C=0.1, gamma=0.1, score=0.637, total= 0.0s
[CV] C=0.1, gamma=0.1 ...

[Parallel(n_jobs=1)]: Using backend SequentialBackend with 1 concurrent workers.
[Parallel(n_jobs=1)]: Done 1 out of 1 | elapsed: 0.0s remaining: 0.0s
[Parallel(n_jobs=1)]: Done 2 out of 2 | elapsed: 0.0s remaining: 0.0s
[Parallel(n_jobs=1)]: Done 3 out of 3 | elapsed: 0.0s remaining: 0.0s
[Parallel(n_jobs=1)]: Done 4 out of 4 | elapsed: 0.0s remaining: 0.0s

[CV] ... C=0.1, gamma=0.1, score=0.637, total= 0.0s
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[CV] C=1, gamma=1 ...

```

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 [CV] C=1000, gamma=0.0001 ...  
 [CV] ... C=1000, gamma=0.0001, score=0.938, total= 0.0s  
 [CV] C=1000, gamma=0.0001 ...



```

[CV] ... C=1000, gamma=0.0001, score=0.912, total= 0.0s
[CV] C=1000, gamma=0.0001 ...
[CV] ... C=1000, gamma=0.0001, score=0.963, total= 0.0s
[CV] C=1000, gamma=0.0001 ...
[CV] ... C=1000, gamma=0.0001, score=0.924, total= 0.0s
[CV] C=1000, gamma=0.0001 ...
[CV] ... C=1000, gamma=0.0001, score=0.962, total= 0.0s

[Parallel(n_jobs=1)]: Done 125 out of 125 | elapsed: 3.1s finished

```

```

[27]: GridSearchCV(estimator=SVC(),
                  param_grid={'C': [0.1, 1, 10, 100, 1000],
                              'gamma': [1, 0.1, 0.01, 0.001, 0.0001]},
                  verbose=5)

```

```

[28]: #best combination
      grid.best_params_

```

```

[28]: {'C': 1, 'gamma': 0.0001}

```

```

[29]: grid.best_estimator_

```

```

[29]: SVC(C=1, gamma=0.0001)

```

```

[30]: gpred = grid.predict(X_test)

```

```

[31]: confusion_matrix(y_test, gpred)

```

```

[31]: array([[ 59,   7],
           [  4, 101]], dtype=int64)

```

```

[32]: print(classification_report(y_test, gpred))

```

	precision	recall	f1-score	support
0	0.94	0.89	0.91	66
1	0.94	0.96	0.95	105
accuracy			0.94	171
macro avg	0.94	0.93	0.93	171
weighted avg	0.94	0.94	0.94	171

```

[ ]:

```