

```
In [ ]: from sklearn.model_selection import train_test_split
        from sklearn.svm import SVC
        from sklearn import datasets
        from sklearn.preprocessing import StandardScaler
        #import gridsearchcv
        from sklearn.model_selection import GridSearchCV
```

```
In [ ]: bc = datasets.load_breast_cancer()
        X = bc.data
        y = bc.target

        # Create training and test split
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=42)
        sc = StandardScaler()
        sc.fit(X_train)
        X_train_std = sc.transform(X_train)
        X_test_std = sc.transform(X_test)

        params = {'C': [0.01, 0.1, 1, 10, 100], 'kernel': ['linear', 'rbf', 'poly', 'sigmoid']}

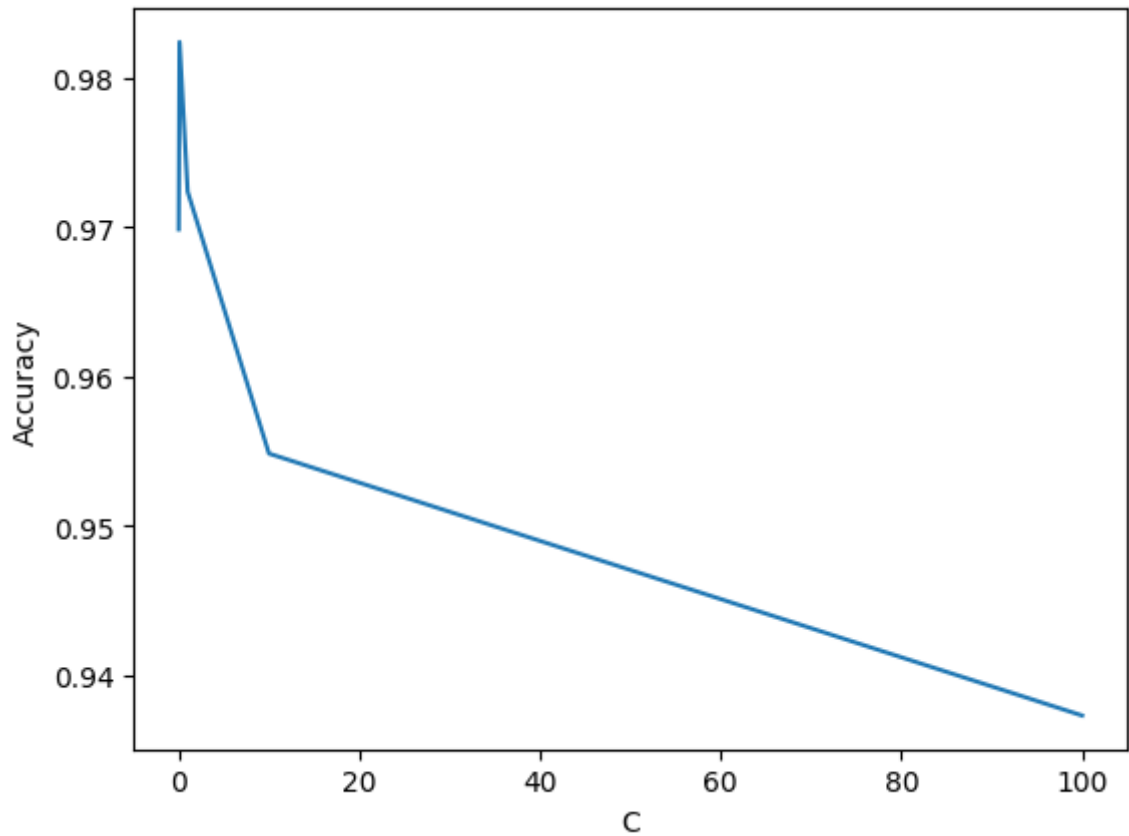
        # Instantiate the Support Vector Classifier (SVC)
        svc = SVC( random_state=1,)
        cv = GridSearchCV(svc, params, cv=5)
        cv.fit(X_train_std, y_train)

        print('Best parameter: ', cv.best_params_)
        print('Best score: ', cv.best_score_)
        print('Test score: ', cv.score(X_test_std, y_test))

        Best parameter: {'C': 0.1, 'kernel': 'linear'}
        Best score: 0.9824367088607595
        Test score: 0.9707602339181286
```

```
In [ ]: # plot the validation curve
        import matplotlib.pyplot as plt
        %matplotlib inline

        plt.plot(params['C'], cv.cv_results_['mean_test_score'])
        plt.xlabel('C')
        plt.ylabel('Accuracy')
        plt.show()
```



```
In [ ]: # kernel vs accuracy
import seaborn as sns
sns.set_style('whitegrid')
sns.barplot(x='param_kernel', y='mean_test_score', data=cv.cv_results_)
plt.xlabel('kernel')
plt.ylabel('Accuracy')
plt.show()
```

