0.5

0.0

1.0

1.5

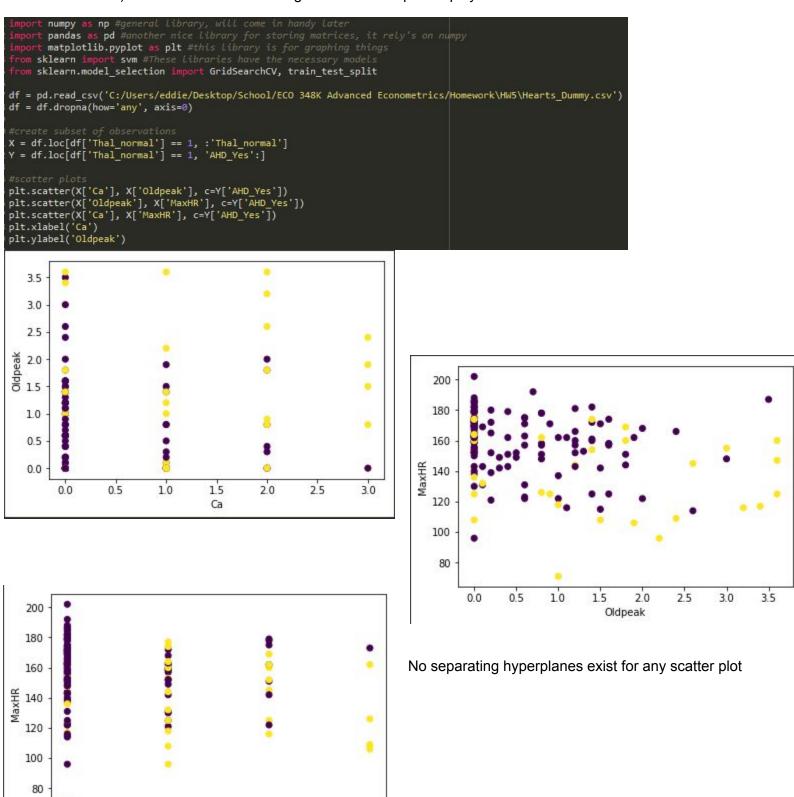
Ca

2.0

2.5

3.0

1) The axis labels will change based on the plot displayed so don't mind the last 2 lines



2)

```
% = discass of observations | X = df.loc(df('Thal_normal') = 1, :'Thal_normal') | Y = df.loc(df('Thal_normal') == 1, 'AHD_Yes':] |
% create training set and test set from subset |
Xtrain, Xtest, Ytrain, Ytest = train_test_split(X, Y, test_size=0.5, random_state=0) |
% create test observation from assignment |
iobs = [[0, 1, 2.5, 150, 1]] |
% begin model creation, fitting, and cross-validation |
% model = swm.SVC() |
% model.fit(Xtrain, Ytrain.values.ravel()) |
% print(') |
% print('insting data score w/ default parameters:', model.fit(Xtrain, Ytrain.values.ravel()).score(Xtrain, Ytrain.values.ravel())) |
% print('insting data score w/ default parameters:', model.fit(Xtrain, Ytrain.values.ravel()).score(Xtest, Ytest.values.ravel())) |
% parameters = [
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']]] |
% (C': [0.01, .1, 1, 10, 100], 'gamma': [0.01, 0.1, 1, 10, 100], 'kernel': ['rbf']] |
% (C': [0.0
```

```
Default parameters are C=1, kernel=RBF, and gamma=auto
Training data score w/ default parameters: 1.0
Test data score w/ default parameters: 0.7560975609756098

CV parameters are: {'C': 1, 'kernel': 'linear'}
Training data w/ CV parameters: 0.8780487804878049
Test data score w/ CV parameters: 0.7926829268292683

Test observation is class: [1]
```

C is the tolerance for which we allow observations to cross the margin or be on the wrong side. Lower values of C allow less error on the soft margin meaning the model will try to overfit the data resulting in low bias. As C increases, the bias will increase but there will be less variance.

In a sense, gamma determines how far an observation has influence. Lower values of gamma will result in data that has higher peaks to adjust for observations further away. As gamma increases, the bias will increase but there will be less variance.