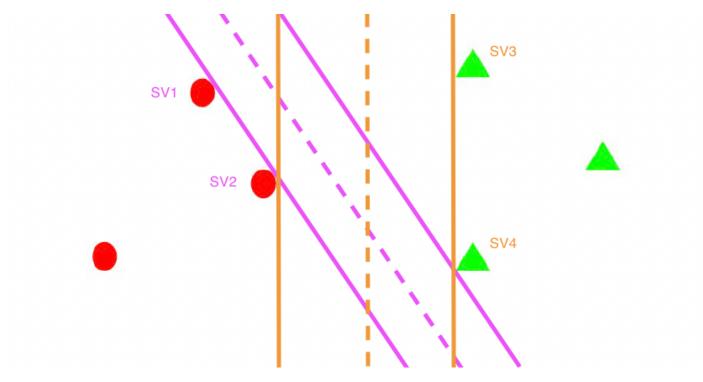
3.a

a. Draw the possible decision boundary learned by svm model, and point out all the support vectors.



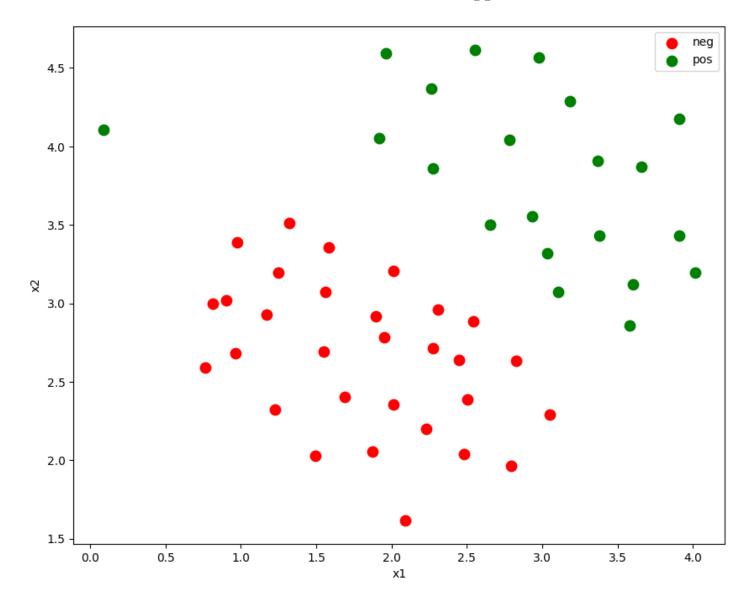
b. Does changing the support vectors change the decision boundary? Does changing the non support vectors change the decision boundary?

Yes, We can see here that changing the suppport vectors to SV1, SV2 and SV4 would result in a smaller decision boundry with less margin.

```
import random
import numpy as np
import matplotlib.pyplot as plt

%matplotlib inline
plt.rcParams['figure.figsize'] = (10.0, 8.0) # set default size of plots
plt.rcParams['image.interpolation'] = 'nearest'
plt.rcParams['image.cmap'] = 'gray'

%load_ext autoreload
%autoreload 2
```

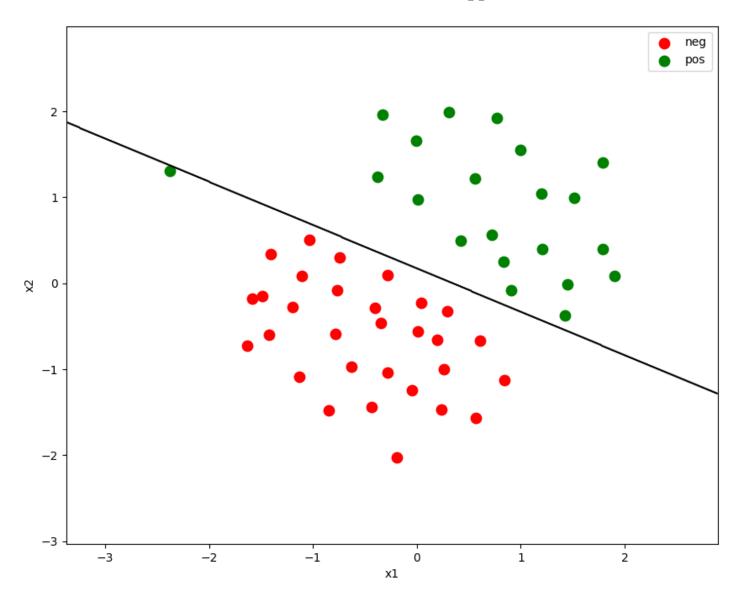


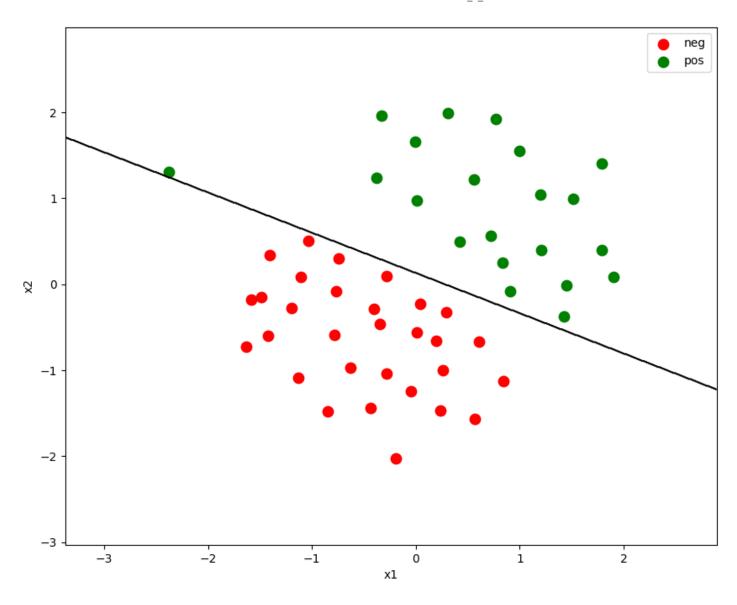
Question 1 Implement SVM with sklearn

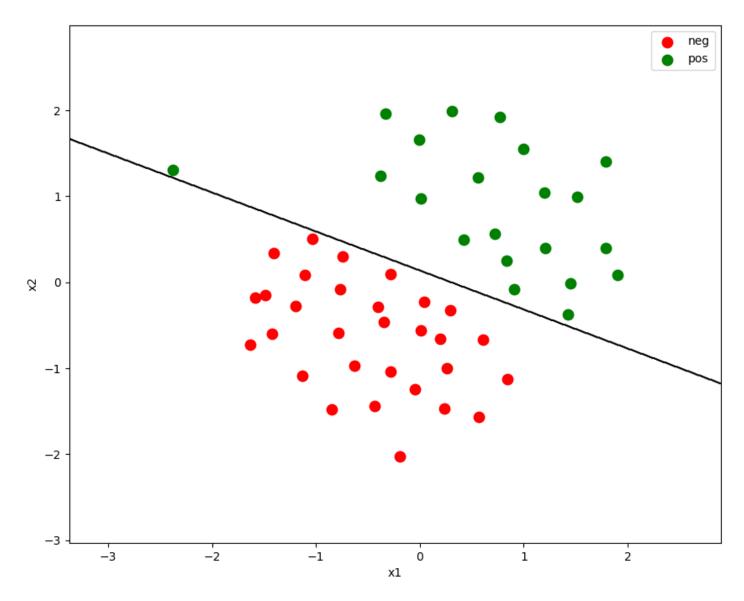
Implement SVM classifier to classify the dataset, and vary the value of C. What do you observe?

```
# scale the data
scaler = preprocessing.StandardScaler().fit(X)
scaleX = scaler.transform(X)
# add an intercept term and convert y values from [0,1] to [-1,1]
XX = np.array([(1,x1,x2) for (x1,x2) in scaleX])
yy = np.ones(y.shape)
vv[v == 0] = -1
yy[y == 0] = -1
# Training linear SVM
# Train a linear SVM on the data set and the plot the learned
# decision boundary
# TODO :: You will change this line below to vary C.
C \text{ list} = [1, 5, 10]
for C in C list:
   # TODO :: define your svm classifier by using sklearn LinearSVC; expect 1 line of code
   svm = LinearSVC(C=C, dual=False, random state=42)
   # TODO :: fit on your training data; expect 1 line of code
   svm.fit(XX, yy)
   # TODO :: classify the training data; expect 1 line of code
   y pred = svm.predict(XX)
   print(f"Accuracy on training data, C={C} = %.3f" %metrics.accuracy_score(yy,y_pred))
   # visualize the decision boundarFy
   utils.plot_decision_boundary(scaleX,y,svm,'x1','x2',['neg','pos'])
Accuracy on training data, C=1 = 0.980
```

Accuracy on training data, C=1 = 0.980Accuracy on training data, C=5 = 1.000Accuracy on training data, C=10 = 1.000







AS we increase the value of C it seems that the model is attempting to distinctly seperate the data. This can cause overfitting.

Question 2 Add PolynomialFeatures

Add polynomial features to the data and fit the LinearSVC with the new dataset. Tune the degree of the feature interaction to make the model correctly classify all the data in training set.

```
In []: # TODO :: expect 13 - 15 lines of code
        from sklearn.pipeline import Pipeline
        from sklearn.preprocessing import PolynomialFeatures
        from sklearn.svm import SVC
        polynomial svm clf = Pipeline([
                ("poly features", PolynomialFeatures(degree=3)),
                ("scaler", preprocessing.StandardScaler()),
                ("svm clf", LinearSVC(C=.25, loss="hinge", random_state=42))
            ])
        polynomial svm clf.fit(XX, yy)
        y pred poly = polynomial svm clf.predict(XX)
        poly_kernel_svm_clf = Pipeline([
                ("scaler", preprocessing.StandardScaler()),
                ("svm clf", SVC(kernel="poly", degree=5, coef0=1, C=1, random state=42))
            1)
        poly kernel svm clf.fit(XX, yy)
        y_pred_kernel = poly_kernel_svm_clf.predict(XX)
        print("Accuracy on training Polynomial SVM data = %.3f" %metrics.accuracy score(yy,y pred poly))
        print("Accuracy on training Kernel Poly SVM data = %.3f" %metrics.accuracy score(yy,y pred kernel))
        print('plot 1 = polynomial svm')
        utils.plot decision boundary(scaleX,y,polynomial svm clf,'x1','x2',['neq','pos'])
        print('plot 2 = polynomial kernel svm')
        utils.plot decision boundary(scaleX,y,poly kernel svm clf,'x1','x2',['neq','pos'])
       /Library/Frameworks/Python.framework/Versions/3.10/lib/python3.10/site-packages/sklearn/svm/ classes.py:32: FutureWarning: The def
       ault value of `dual` will change from `True` to `'auto'` in 1.5. Set the value of `dual` explicitly to suppress the warning.
         warnings.warn(
       Accuracy on training Polynomial SVM data = 1.000
       Accuracy on training Kernel Poly SVM data = 1.000
       plot 1 = polvnomial svm
       plot 2 = polynomial kernel svm
```

