

```
In [1]: import numpy as np
import pandas as pd
from sklearn.datasets import load_wine
from sklearn.model_selection import train_test_split

import matplotlib.pyplot as plt
from sklearn.svm import SVC
import seaborn as sns
```

Question 1.1

```
In [2]: # Read the wine dataset
dataset = load_wine()
df = pd.DataFrame(data=dataset['data'], columns=dataset['feature_names'])
df = df.assign(target=pd.Series(dataset['target']).values)

# Filter the irrelevant columns
df = df[['alcohol', 'magnesium', 'target']]
# Filter the irrelevant label
df = df[df.target != 0]

train_df, val_df = train_test_split(df, test_size=30, random_state=3)

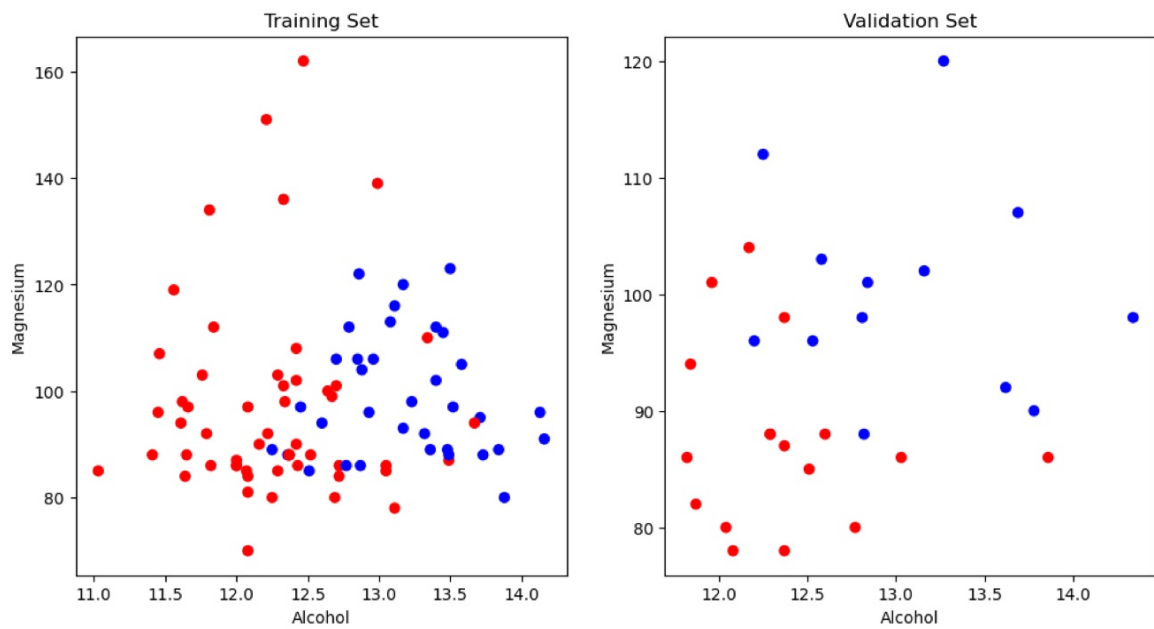
# Define colors for wineries
colors = {1: 'red', 2: 'blue'}

# Create scatter plots
fig, axes = plt.subplots(1, 2, figsize=(12, 6))

# Training set scatter plot
axes[0].scatter(train_df['alcohol'], train_df['magnesium'],
                c=train_df['target'].map(colors))
axes[0].set_title('Training Set')
axes[0].set_xlabel('Alcohol')
axes[0].set_ylabel('Magnesium')

# Validation set scatter plot
axes[1].scatter(val_df['alcohol'], val_df['magnesium'],
                c=val_df['target'].map(colors))
axes[1].set_title('Validation Set')
axes[1].set_xlabel('Alcohol')
axes[1].set_ylabel('Magnesium')

plt.show()
```



As we can see in the above plot, the data cannot be linearly separated in perfect manner. If the data is not linearly separable, running Hard-SVM will result in the model failing to find a suitable separating hyperplane. This is because Hard-SVM requires the data to be perfectly

Question 1.2

```
In [3]: from sklearn.svm import SVC
```

```

In [4]: # Function to plot the decision boundary
def plot_svc_decision_function(model, X, y, ax=None, plot_support=True):
    """Plot the decision function for a 2D SVC"""
    if ax is None:
        ax = plt.gca()
    xlim = ax.get_xlim()
    ylim = ax.get_ylim()

    # Create grid to evaluate model
    x = np.linspace(xlim[0], xlim[1], 30)
    y = np.linspace(ylim[0], ylim[1], 30)
    Y, X_grid = np.meshgrid(y, x)
    xy = np.vstack([X_grid.ravel(), Y.ravel()]).T
    xy_df = pd.DataFrame(xy, columns=['alcohol', 'magnesium'])
    P = model.decision_function(xy_df).reshape(X_grid.shape)

    # Plot decision boundary and margins
    ax.contour(X_grid, Y, P, colors='k',
               levels=[-1, 0, 1], alpha=0.5,
               linestyles=['--', '-', '--'])

    # Plot support vectors
    if plot_support:
        ax.scatter(model.support_vectors_[0],
                   model.support_vectors_[1],
                   s=100, linewidth=1, facecolors='none', edgecolors='k')
    ax.set_xlim(xlim)
    ax.set_ylim(ylim)

    # Train SVM models with different C values
    C_values = [0.1, 0.05, 0.01]
    models = [SVC(kernel='linear', C=C) for C in C_values]

    # Fit the models
    for model in models:
        model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

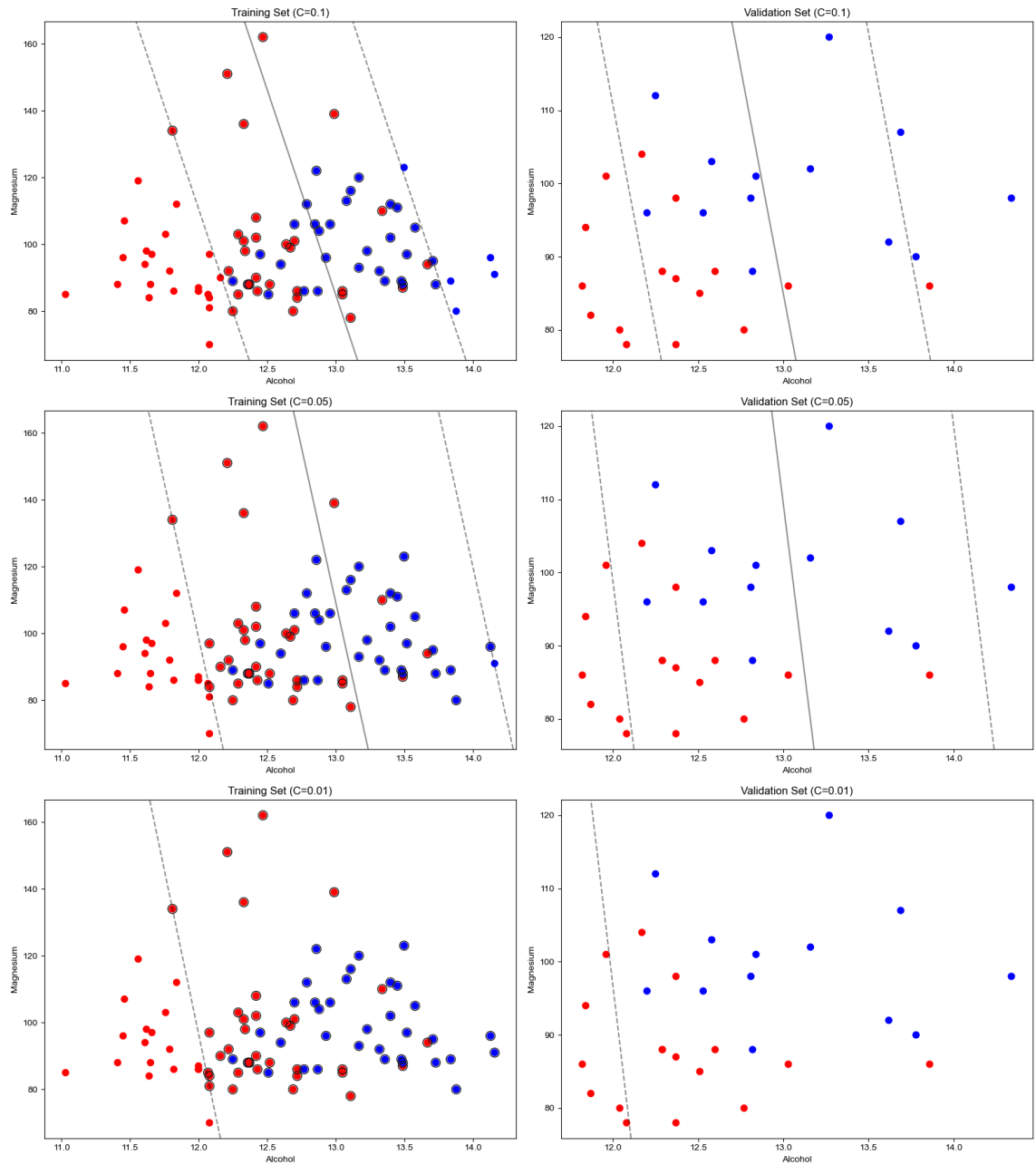
    # Create plots
    fig, axes = plt.subplots(3, 2, figsize=(16, 18))
    sns.set()

    for i, (model, C) in enumerate(zip(models, C_values)):
        # Training set scatter plot
        ax = axes[i, 0]
        ax.scatter(train_df['alcohol'], train_df['magnesium'],
                   c=train_df['target'].map(colors), s=50)
        plot_svc_decision_function(model,
                                   train_df[['alcohol', 'magnesium']], train_df
                                   ax=ax, plot_support=True)
        ax.set_title(f'Training Set (C={C})')
        ax.set_xlabel('Alcohol')
        ax.set_ylabel('Magnesium')

        # Validation set scatter plot
        ax = axes[i, 1]
        ax.scatter(val_df['alcohol'], val_df['magnesium'],
                   c=val_df['target'].map(colors), s=50)
        plot_svc_decision_function(model, val_df[['alcohol', 'magnesium']],
                                   val_df['target'], ax=ax, plot_support=False)
        ax.set_title(f'Validation Set (C={C})')
        ax.set_xlabel('Alcohol')
        ax.set_ylabel('Magnesium')

```

```
plt.tight_layout()  
plt.show()
```



Question 1.3

כדי שראינו בהרצאה, בעזרת האופטימיזציה הינה:

$$\min_w \|w\|^2 \quad \text{s.t.} \quad \forall i, \gamma_i \langle w, x_i \rangle \geq 1$$

כאשר ה-margin מוגדר γ_i :

$$\Rightarrow \min_{i \in [m]} |\langle \hat{w}, x_i \rangle| = \min_{i \in [m]} \left| \langle \frac{w_0}{\|w_0\|}, x_i \rangle \right| = \frac{1}{\|w_0\|} \min_{i \in [m]} |\langle w_0, x_i \rangle|$$

$\hat{w} = \frac{w_0}{\|w_0\|}$ מנורמל ↑ ↑
סלר הריבוע

וקטור $S = (x_i, \gamma_i)$ הנקרא ה support vector לפי ההצדקה.

ראינו בהרצאה כי צמד support vectors מקיימים $|\langle w_0, x_i \rangle| \geq 1$.
 שכן הם נמצאים במרחק המינימלי מ-0 אשר הגדול 1.
 מכאן שמתקיים:

$$\min_{i \in [m]} |\langle \hat{w}, x_i \rangle| = \frac{1}{\|w_0\|} \min_{i \in [m]} |\langle w_0, x_i \rangle| = \frac{1}{\|w_0\|} |\langle w_0, x_i \rangle| = \frac{1}{\|w_0\|}$$

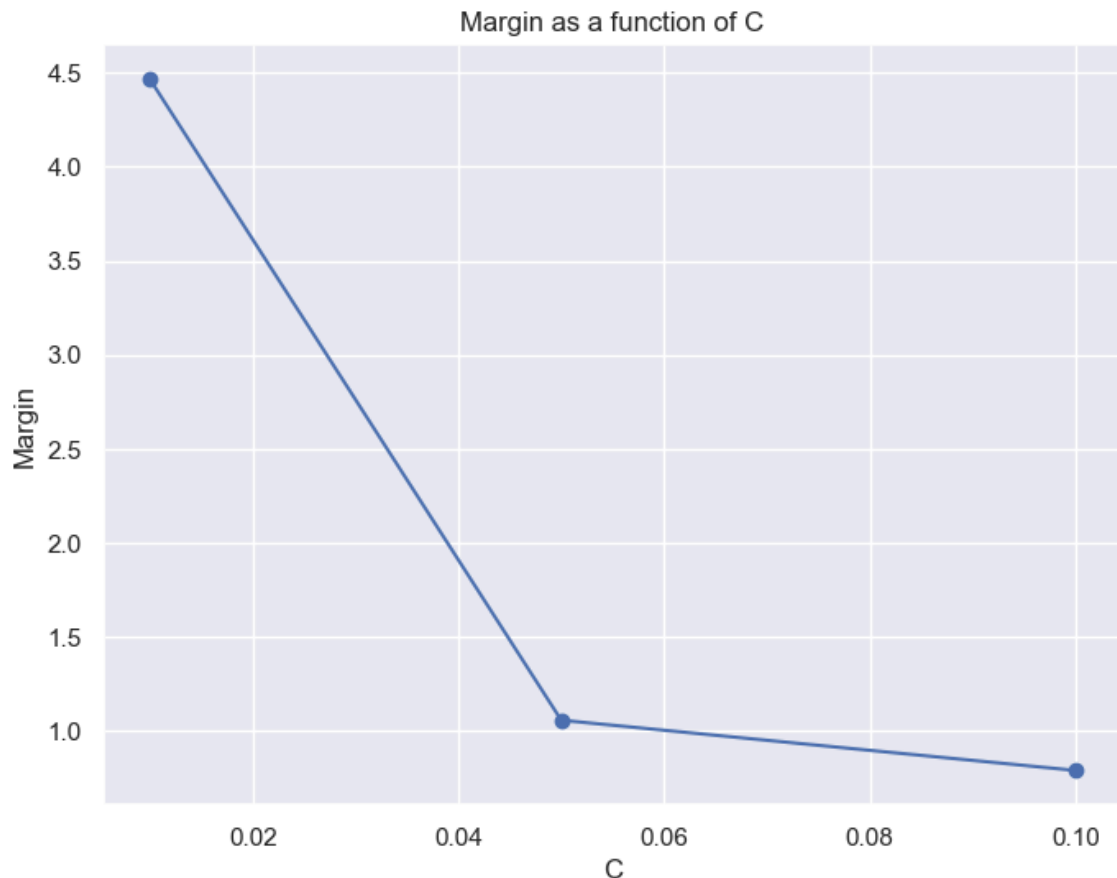
Question 1.4

```
In [5]: # Fit the models
for model in models:
    model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

# Function to calculate margin from SVC model
def calculate_margin(model):
    w_norm = np.linalg.norm(model.coef_)
    return 1 / w_norm

# Calculate margins for each model
margins = [calculate_margin(model) for model in models]

# Plot margin as a function of C
plt.figure(figsize=(8, 6))
plt.plot(C_values, margins, marker='o', linestyle='--', color='b')
plt.xlabel('C')
plt.ylabel('Margin')
plt.title('Margin as a function of C')
plt.grid(True)
plt.show()
```



We can see from the plot above that while C grows the margin narrows. In the optimization problem we have two parameters:

1. $\|W\|$ which affects the width of the margins.
2. The hinge loss * C. when C grows we would want to have a less training errors, since we give more weight to each mistake. Therefore the margin narrows and allow for less mistakes to be made. On the other hand, as C gets smaller - the margin widens and the model generalized better.

Question 1.5:

```
In [6]: from sklearn.metrics import accuracy_score

# Train SVM models with different C values
C_values = [0.1, 0.05, 0.01]
models = [SVC(kernel='linear', C=C) for C in C_values]

# Initialize lists to store errors
train_errors = []
val_errors = []

# Fit the models and compute errors
for model in models:
    model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

    # Predictions on training set
    train_pred = model.predict(train_df[['alcohol', 'magnesium']])
    train_accuracy = accuracy_score(train_df['target'], train_pred)
    train_error = 1.0 - train_accuracy
    train_errors.append(train_error)

    # Predictions on validation set
    val_pred = model.predict(val_df[['alcohol', 'magnesium']])
    val_accuracy = accuracy_score(val_df['target'], val_pred)
    val_error = 1.0 - val_accuracy
    val_errors.append(val_error)

# Plot errors as a function of C
plt.figure(figsize=(8, 6))
plt.plot(C_values, train_errors, marker='o', linestyle='--', color='b',
         label='Training Error')
plt.plot(C_values, val_errors, marker='o', linestyle='--', color='r',
         label='Validation Error')
plt.xlabel('C')
plt.ylabel('Error')
plt.title('Training and Validation Error as a function of C')
plt.grid(True)
plt.legend()
plt.show()
```



As we can observe in the plot above, both training and validation error are decreasing until $C=0.05$, moreover - the error value of the validation set is higher than the training set. for C larger than 0.05 we can see that as we expected for the last question (1.4) the margin narrows which may result in overfitting of the training set which in turn causes a higher error values in the validation set.

Question 1.6

```
In [7]: import numpy as np
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score

# Define the degree range and model parameter C
degrees = range(2, 9)
C = 1

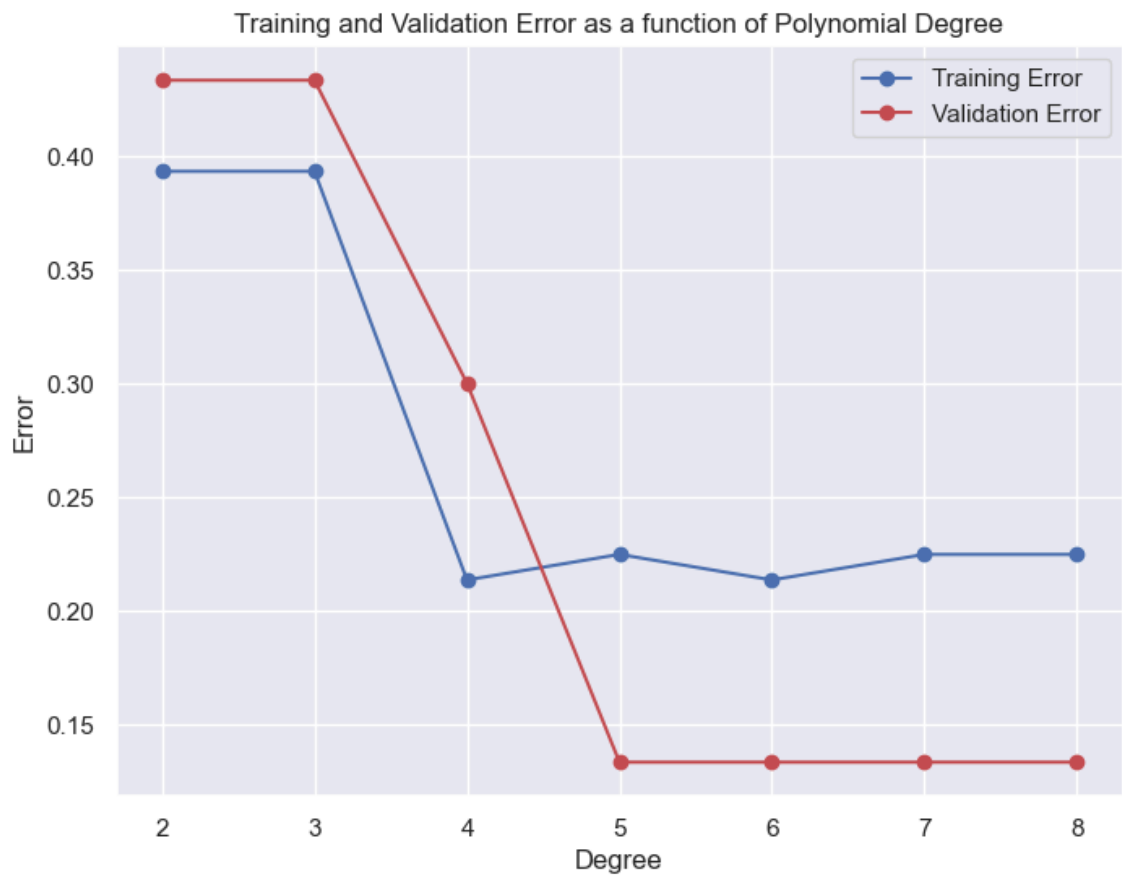
# Initialize lists to store errors
train_errors = []
val_errors = []

# Fit the models and compute errors
for degree in degrees:
    model = SVC(kernel='poly', degree=degree, C=C)
    model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

    # Predictions on training set
    train_pred = model.predict(train_df[['alcohol', 'magnesium']])
    train_accuracy = accuracy_score(train_df['target'], train_pred)
    train_error = 1.0 - train_accuracy
    train_errors.append(train_error)

    # Predictions on validation set
    val_pred = model.predict(val_df[['alcohol', 'magnesium']])
    val_accuracy = accuracy_score(val_df['target'], val_pred)
    val_error = 1.0 - val_accuracy
    val_errors.append(val_error)

# Plot errors as a function of polynomial degree
plt.figure(figsize=(8, 6))
plt.plot(degrees, train_errors, marker='o', linestyle='--', color='b',
         label='Training Error')
plt.plot(degrees, val_errors, marker='o', linestyle='--', color='r',
         label='Validation Error')
plt.xlabel('Degree')
plt.ylabel('Error')
plt.title('Training and Validation Error as a function of Polynomial Degree')
plt.legend()
plt.show()
```



As observed in the plot above, when moving to a higher dimension the error value decreases for both training and validation sets, and specifically for the validation set - starting from 5D we can assume that the data is almost linearly separable as the error value is the lowest.

Question 1.7

```

In [12]: # Function to plot the decision boundary
def plot_svc_decision_function(model, X, y, ax=None, plot_support=True):
    if ax is None:
        ax = plt.gca()
    xlim = ax.get_xlim()
    ylim = ax.get_ylim()

    # Create grid to evaluate model
    x = np.linspace(xlim[0], xlim[1], 30)
    y = np.linspace(ylim[0], ylim[1], 30)
    Y, X_grid = np.meshgrid(y, x)
    xy = np.vstack([X_grid.ravel(), Y.ravel()]).T
    xy_df = pd.DataFrame(xy, columns=['alcohol', 'magnesium'])
    P = model.decision_function(xy_df).reshape(X_grid.shape)

    # Plot decision boundary and margins
    ax.contour(X_grid, Y, P, colors='k', levels=[-1, 0, 1], alpha=0.5, line

    # Plot support vectors
    if plot_support:
        ax.scatter(model.support_vectors_[0],
                   model.support_vectors_[1],
                   s=100, linewidth=1, facecolors='none', edgecolors='k')
    ax.set_xlim(xlim)
    ax.set_ylim(ylim)

# Train SVM model with polynomial kernel of degree 3
model = SVC(kernel='poly', degree=3, C=1)
model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

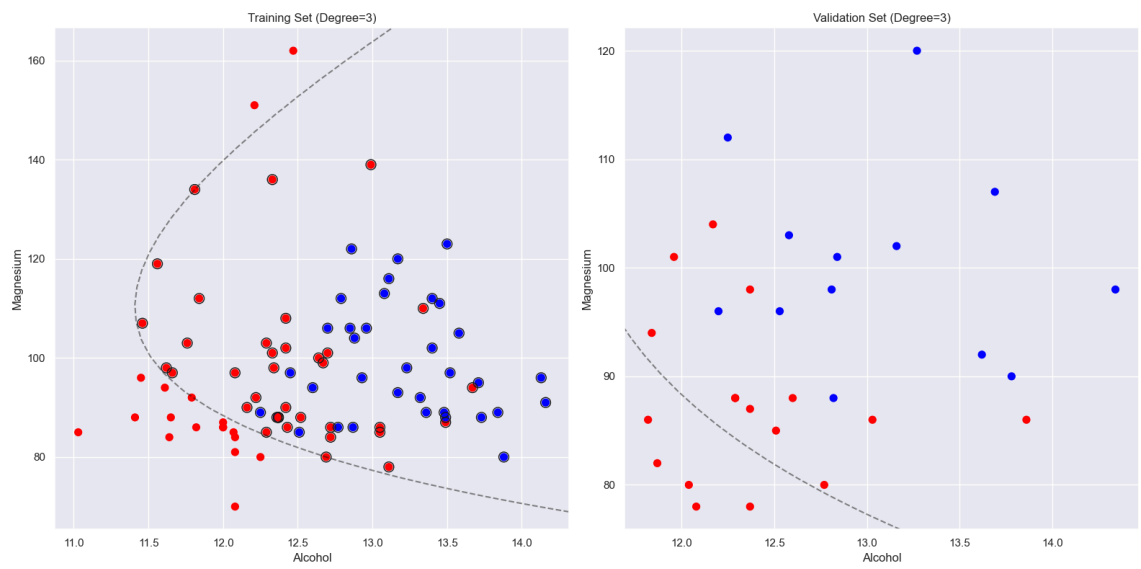
# Create plots
fig, axes = plt.subplots(1, 2, figsize=(16, 8))
sns.set()

# Training set scatter plot
ax = axes[0]
ax.scatter(train_df['alcohol'], train_df['magnesium'],
           c=train_df['target'].map(colors), s=50)
plot_svc_decision_function(model, train_df[['alcohol', 'magnesium']],
                           train_df['target'], ax=ax, plot_support=True)
ax.set_title('Training Set (Degree=3)')
ax.set_xlabel('Alcohol')
ax.set_ylabel('Magnesium')

# Validation set scatter plot
ax = axes[1]
ax.scatter(val_df['alcohol'], val_df['magnesium'],
           c=val_df['target'].map(colors), s=50)
plot_svc_decision_function(model, val_df[['alcohol', 'magnesium']],
                           val_df['target'], ax=ax, plot_support=False)
ax.set_title('Validation Set (Degree=3)')
ax.set_xlabel('Alcohol')
ax.set_ylabel('Magnesium')

plt.tight_layout()
plt.show()

```



```

In [13]: # Function to plot the decision boundary
def plot_svc_decision_function(model, X, y, ax=None, plot_support=True):
    if ax is None:
        ax = plt.gca()
    xlim = ax.get_xlim()
    ylim = ax.get_ylim()

    # Create grid to evaluate model
    x = np.linspace(xlim[0], xlim[1], 30)
    y = np.linspace(ylim[0], ylim[1], 30)
    Y, X_grid = np.meshgrid(y, x)
    xy = np.vstack([X_grid.ravel(), Y.ravel()]).T
    xy_df = pd.DataFrame(xy, columns=['alcohol', 'magnesium'])
    P = model.decision_function(xy_df).reshape(X_grid.shape)

    # Plot decision boundary and margins
    ax.contour(X_grid, Y, P, colors='k', levels=[-1, 0, 1],
               alpha=0.5, linestyles=['--', '-', '--'])

    # Plot support vectors
    if plot_support:
        ax.scatter(model.support_vectors_[0],
                   model.support_vectors_[1],
                   s=100, linewidth=1, facecolors='none', edgecolors='k')
    ax.set_xlim(xlim)
    ax.set_ylim(ylim)

    # Train SVM model with polynomial kernel of degree 6
    model = SVC(kernel='poly', degree=6, C=1)
    model.fit(train_df[['alcohol', 'magnesium']], train_df['target'])

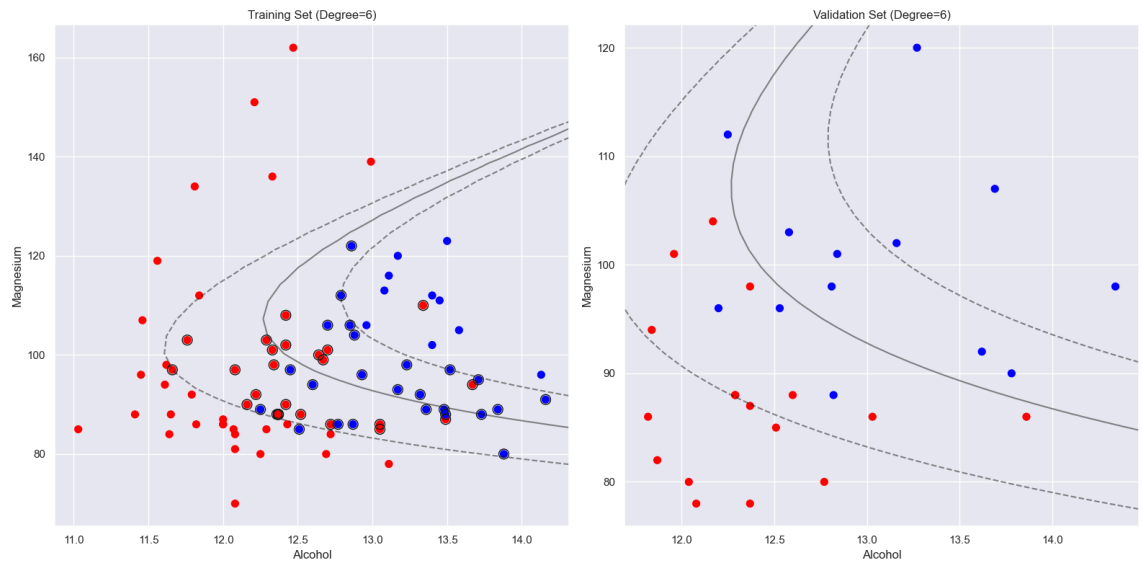
    # Create plots
    fig, axes = plt.subplots(1, 2, figsize=(16, 8))
    sns.set()

    # Training set scatter plot
    ax = axes[0]
    ax.scatter(train_df['alcohol'], train_df['magnesium'],
               c=train_df['target'].map(colors), s=50)
    plot_svc_decision_function(model, train_df[['alcohol', 'magnesium']],
                              train_df['target'], ax=ax, plot_support=True)
    ax.set_title('Training Set (Degree=6)')
    ax.set_xlabel('Alcohol')
    ax.set_ylabel('Magnesium')

    # Validation set scatter plot
    ax = axes[1]
    ax.scatter(val_df['alcohol'], val_df['magnesium'],
               c=val_df['target'].map(colors), s=50)
    plot_svc_decision_function(model, val_df[['alcohol', 'magnesium']],
                              val_df['target'], ax=ax, plot_support=False)
    ax.set_title('Validation Set (Degree=6)')
    ax.set_xlabel('Alcohol')
    ax.set_ylabel('Magnesium')

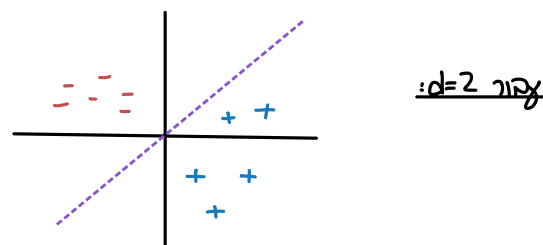
    plt.tight_layout()
    plt.show()

```



שאלה 3:

1. נתון דיגמה באשר $bias=0$. במקרה זה, המפריד הליניארי יצבור בהאשית הצירים, כלומר הפרדה ליניארית של הדאטה צריכה להתקיים בהיחס להאשית הצירים.



2. נמאן: $x_1 = (p, 0), y_1 = -1$ $x_2 = (0, q), y_2 = 1$

$$y_1 < w \cdot (p, 0) = -w_1 p \geq 1 \Rightarrow w_1 p \leq -1$$

מהאילווצים נקבל:

$$y_2 < w \cdot (0, q) = w_2 \cdot q \geq 1 \Rightarrow w_2 q \geq 1$$

נשים לב ש על מנת שהדאטה יהיה פריד ליניארית בהיחס להאשית הצירים, נרצה שאל אחת מהנקודות לא תמצא ב- (סיס), ואכן לכל מקרה בו $p \cdot q = 0$, הדאטה לא יהיה פריד ליניארית, ואז לא יהיה פתרון ל-hard SVM.

נחלק למקרים:

$$|w_2| \geq \left| \frac{1}{q} \right| \Leftrightarrow \begin{cases} w_2 \geq \frac{1}{q} & : q > 0 \\ w_2 \leq -\frac{1}{q} & : q < 0 \end{cases} \quad |w_1| \geq \left| \frac{-1}{p} \right| \Leftrightarrow \begin{cases} w_1 \leq -\frac{1}{p} & : p > 0 \\ w_1 \geq -\frac{1}{p} & : p < 0 \end{cases}$$

$$w_1 \text{ ו } w_2$$

ראינו בהרצאה כי:

$$\|w\|^2 = w_1^2 + w_2^2 = |w_1|^2 + |w_2|^2 \geq \left| \frac{-1}{p} \right|^2 + \left| \frac{1}{q} \right|^2 = \frac{1}{p^2} + \frac{1}{q^2}$$

$$y_1 \langle \tilde{w}, (p, 0) \rangle = -(-\frac{1}{p}) \cdot p = 1 \geq 1 \quad \checkmark$$

נבחר $\tilde{w} = (\frac{1}{p}, -\frac{1}{q})$. אזווי הנציה מתקיימת, שכן:

$$y_2 \langle \tilde{w}, (0, q) \rangle = \frac{1}{q} \cdot q = 1 \geq 1 \quad \checkmark$$

בנוסף, $\|\tilde{w}\|^2 = \tilde{w}_1^2 + \tilde{w}_2^2 = \frac{1}{p^2} + \frac{1}{q^2}$, ולכן \tilde{w} מקיים את המרחק התחתון ומקיים $\tilde{w} = \argmin_w \|w\|^2$

אם \tilde{w} הוא פתרון של hard SVM.

3. המרחב אולי הינו מקביל לא היה משתנה, היות שה-margin נקבע לפי ה-Support vectors, וכל הנקודות של y=1 אלוה נמצאות מעבר ל-margin.

4. נשים לב שבכל שנגדל את λ , ניתן חשיבות גבוהה יותר ל-margin ונעשה פחות בטעויות הסיווג. אכן, ככל ש- λ גדלה יותר, נשאף ל-margin ולהפכה טובה יותר של λ .

$\lambda = 200$

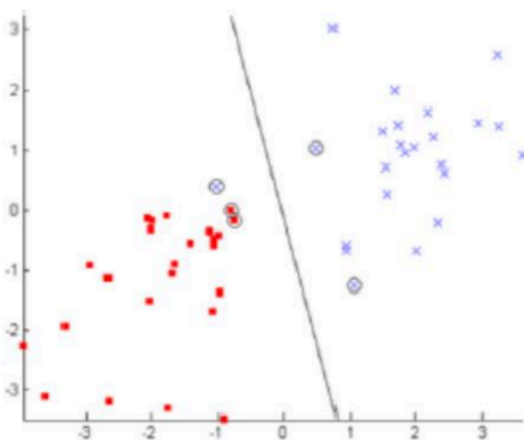


Figure 1

$\lambda = 2$

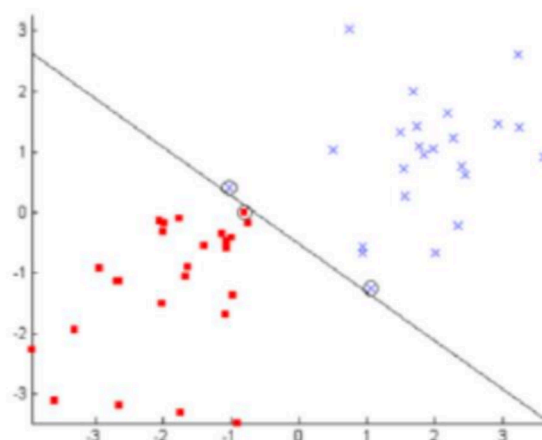


Figure 2

שאלה 4:

1. a) הפלט של האלגוריתם יהיה סקלרים $1 \leq i \leq m$, המייצגים את סמות הקיבועים שעשיר על הדוגמה i .

b) נגזר את פלט המודל, $\hat{y}(x) = \text{Sign} \left(\sum_{i=1}^m \alpha_i y_i K(x, x_i) \right)$

הפסאודו קוד:

$$\forall i \quad \alpha_i = 0$$

for $t = 1, 2, \dots$

if $(\exists i \text{ s.t. } y_i \neq \hat{y}(x_i))$

$$\alpha_i = \alpha_i + 1$$

else

return $[\alpha_i]_{i=1}^m$

2. נסמן: $w = \sum_{i=1}^m \alpha_i \Psi(x_i)$, וכאשר נשתמש ב- w באלגוריתם הפספסטרון, נקבל את האלגוריתם החדש שבענין.

Ψ לא ידועה, ולכן נאלץ להשתמש ב- w כשהוא נמצא בתוך מכלול פנימיות עם דוגמה אחת ו- K נתון.

לדונך הוכחה כי השתמשנו ב- w במכלול פנימיות של w עם וקטורים אחרים, ולכן בהינתן K , נוכל לחשב גם

את המכלול הולו. ניתן אף לשלוח את המכלול הולו Ψ וקטור החדש w של w . לפיכך, נמנהג אלגוריתם

הפספסטרון מתקיים כשהאלגוריתם המוצג מתקנס. $\Leftrightarrow S^\Psi$ פהיז אינארת ב- F .