

# svm\_homework

April 11, 2025

## 1 Worked with:

### 1.1 Trevor Mathisen

### 1.2 Viet Nguyen

```
[1]: from numpy import set_printoptions, logspace, mean, std
import matplotlib.pyplot as plt
import pandas as pd
from pandas import set_option
from pandas import read_csv
from pandas.plotting import scatter_matrix

from sklearn.preprocessing import StandardScaler, Normalizer, LabelEncoder
from sklearn.model_selection import KFold, GridSearchCV, cross_val_score

import seaborn as sns
```

### 1.3 3. Use the Wine dataset (check under ML dataset Module on Canvas)

```
[2]: filename = 'wine.csv'
# Column names added to csv including 'Class' as the first column/output_
↳variable
data = read_csv(filename)
```

### 1.4 Exploratory Data Analysis

```
[3]: set_printoptions(precision=3)
print(data.head(5))
print(data.isnull().sum())
print(f'{data.shape[0]} rows and {data.shape[1]} columns')
for column in data.columns:
    print(column)
    print(data[column].unique())
```

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash	Magnesium \
0	1	14.23	1.71	2.43	15.6	127
1	1	13.20	1.78	2.14	11.2	100

2	1	13.16	2.36	2.67	18.6	101
3	1	14.37	1.95	2.50	16.8	113
4	1	13.24	2.59	2.87	21.0	118

	Total phenols	Flavanoids	Nonflavanoid phenols	Proanthocyanins \
0	2.80	3.06	0.28	2.29
1	2.65	2.76	0.26	1.28
2	2.80	3.24	0.30	2.81
3	3.85	3.49	0.24	2.18
4	2.80	2.69	0.39	1.82

	Color intensity	Hue	OD280/OD315 of diluted wines	Proline
0	5.64	1.04	3.92	1065
1	4.38	1.05	3.40	1050
2	5.68	1.03	3.17	1185
3	7.80	0.86	3.45	1480
4	4.32	1.04	2.93	735

Class	0
Alcohol	0
Malic acid	0
Ash	0
Alcalinity of ash	0
Magnesium	0
Total phenols	0
Flavanoids	0
Nonflavanoid phenols	0
Proanthocyanins	0
Color intensity	0
Hue	0
OD280/OD315 of diluted wines	0
Proline	0

dtype: int64

178 rows and 14 columns

Class

[1 2 3]

Alcohol

[14.23 13.2 13.16 14.37 13.24 14.2 14.39 14.06 14.83 13.86 14.1 14.12  
13.75 14.75 14.38 13.63 14.3 13.83 14.19 13.64 12.93 13.71 12.85 13.5  
13.05 13.39 13.3 13.87 14.02 13.73 13.58 13.68 13.76 13.51 13.48 13.28  
13.07 14.22 13.56 13.41 13.88 14.21 13.9 13.94 13.82 13.77 13.74 13.29  
13.72 12.37 12.33 12.64 13.67 12.17 13.11 13.34 12.21 12.29 13.49 12.99  
11.96 11.66 13.03 11.84 12.7 12. 12.72 12.08 12.67 12.16 11.65 11.64  
12.69 11.62 12.47 11.81 12.6 12.34 11.82 12.51 12.42 12.25 12.22 11.61  
11.46 12.52 11.76 11.41 11.03 12.77 11.45 11.56 11.87 12.07 12.43 11.79  
12.04 12.86 12.88 12.81 12.53 12.84 13.36 13.52 13.62 12.87 13.32 13.08  
12.79 13.23 12.58 13.17 13.84 12.45 14.34 12.36 13.69 12.96 13.78 13.45  
12.82 13.4 12.2 14.16 13.27 14.13]

Malic acid

[1.71 1.78 2.36 1.95 2.59 1.76 1.87 2.15 1.64 1.35 2.16 1.48 1.73 1.81  
 1.92 1.57 1.59 3.1 1.63 3.8 1.86 1.6 2.05 1.77 1.72 1.9 1.68 1.5  
 1.66 1.83 1.53 1.8 1.65 3.99 3.84 1.89 3.98 4.04 3.59 2.02 1.75 1.67  
 1.7 1.97 1.43 0.94 1.1 1.36 1.25 1.13 1.45 1.21 1.01 1.17 1.19 1.61  
 1.51 1.09 1.88 0.9 2.89 0.99 3.87 0.92 3.86 0.89 0.98 2.06 1.33 2.83  
 1.99 1.52 2.12 1.41 1.07 3.17 2.08 1.34 2.45 2.55 1.29 3.74 2.43 2.68  
 0.74 1.39 1.47 3.43 2.4 4.43 5.8 4.31 2.13 4.3 2.99 2.31 3.55 1.24  
 2.46 4.72 5.51 2.96 2.81 2.56 4.95 3.88 3.57 5.04 4.61 3.24 3.9 3.12  
 2.67 3.3 5.19 4.12 3.03 3.83 3.26 3.27 3.45 2.76 4.36 3.7 3.37 2.58  
 4.6 2.39 2.51 5.65 3.91 4.28 4.1 ]

#### Ash

[2.43 2.14 2.67 2.5 2.87 2.45 2.61 2.17 2.27 2.3 2.32 2.41 2.39 2.38  
 2.7 2.72 2.62 2.48 2.56 2.28 2.65 2.36 2.52 3.22 2.8 2.21 2.84 2.55  
 2.1 2.51 2.31 2.12 2.59 2.29 2.44 2.4 2.04 2.6 2.42 2.68 2.25 2.46  
 1.36 2.02 1.92 2.16 2.53 1.7 1.75 2.24 1.71 2.23 1.95 2. 2.2 2.58  
 2.26 2.22 2.74 1.98 1.9 1.88 1.94 1.82 2.92 1.99 2.19 3.23 2.73 2.13  
 2.78 2.54 2.64 2.35 2.15 2.75 2.69 2.86 2.37]

#### Alcalinity of ash

[15.6 11.2 18.6 16.8 21. 15.2 14.6 17.6 14. 16. 18. 11.4 12. 17.2  
 20. 16.5 16.6 17.8 25. 16.1 17. 19.4 22.5 19.1 19.5 19. 20.5 15.5  
 13.2 16.2 18.8 15. 17.5 18.9 17.4 12.4 17.1 16.4 16.3 16.7 10.6 18.1  
 19.6 20.4 24. 30. 14.8 23. 22.8 26. 21.6 23.6 18.5 22. 20.7 21.5  
 20.8 28.5 26.5 24.5 23.5 25.5 27. ]

#### Magnesium

[127 100 101 113 118 112 96 121 97 98 105 95 89 91 102 120 115 108  
 116 126 124 93 94 107 106 104 132 110 128 117 90 103 111 92 88 87  
 78 151 86 139 136 85 99 84 70 81 80 162 134 119 82 122 123]

#### Total phenols

[2.8 2.65 3.85 3.27 2.5 2.6 2.98 2.95 2.2 3.1 3.3 2.85 2.7 3.  
 2.41 2.61 2.48 2.53 2.63 2.4 2.86 2.42 2.35 2.45 3.15 3.25 2.64 2.75  
 2.88 2.72 3.88 2.96 3.2 3.4 1.98 2.05 2.02 2.1 3.5 1.89 2.11 1.85  
 1.1 1.88 3.38 1.61 1.95 1.72 1.9 2.83 2. 1.65 1.78 1.92 1.6 1.45  
 1.38 3.02 2.55 3.52 2.23 2.56 1.68 2.36 2.74 3.18 1.75 2.46 1.63 2.9  
 2.62 2.13 2.22 1.51 1.3 1.15 1.7 1.62 1.79 2.32 1.54 1.4 1.55 1.5  
 0.98 1.93 1.41 1.48 1.8 1.74 2.3 1.83 1.39 1.35 1.28 1.25 1.59]

#### Flavanoids

[3.06 2.76 3.24 3.49 2.69 3.39 2.52 2.51 2.98 3.15 3.32 2.43 3.69 3.64  
 2.91 3.14 3.4 3.93 3.03 3.17 2.41 2.88 2.37 2.61 2.68 2.94 2.19 2.97  
 2.33 3.25 3.19 2.74 2.53 2.64 3.04 3.29 3.56 2.63 3. 2.65 2.92 3.54  
 3.27 2.99 3.74 2.79 2.9 2.78 3.23 3.67 0.57 1.09 1.41 1.79 3.1 1.75  
 3.18 2. 1.3 1.28 1.02 2.86 1.84 2.89 2.14 1.57 2.03 1.32 1.85 2.55  
 2.26 1.58 1.59 2.21 1.94 1.69 1.61 1.5 1.25 1.46 2.25 2.27 0.99 2.5  
 3.75 2.17 1.36 2.11 1.64 1.92 1.76 2.04 2.58 2.01 2.29 1.6 2.09 5.08  
 2.13 2.24 2.45 1.22 1.2 0.58 0.66 0.47 0.6 0.48 0.5 0.52 0.8 0.78  
 0.55 0.34 0.65 0.76 1.39 0.83 0.63 1.31 1.1 0.92 0.56 0.7 0.68 0.84  
 0.96 0.49 0.51 0.61 0.75 0.69]

#### Nonflavanoid phenols

[0.28 0.26 0.3 0.24 0.39 0.34 0.31 0.29 0.22 0.43 0.33 0.4 0.32 0.17

0.25 0.27 0.47 0.37 0.42 0.5 0.2 0.21 0.19 0.63 0.53 0.45 0.55 0.14  
0.13 0.35 0.61 0.48 0.52 0.58 0.66 0.6 0.41 0.44 0.56]

#### Proanthocyanins

[2.29 1.28 2.81 2.18 1.82 1.97 1.98 1.25 1.85 2.38 1.57 1.81 2.96 1.46  
1.72 1.86 1.66 2.1 1.69 1.92 1.45 1.35 1.76 1.95 1.54 1.36 1.44 1.37  
2.08 2.34 1.48 1.7 2.03 2.19 2.14 2.91 1.87 1.68 1.62 2.45 2.04 0.42  
0.41 0.62 0.73 1.03 2.28 1.04 2.5 1.96 1.65 1.15 0.95 2.76 1.43 1.77  
1.4 2.35 1.56 1.34 1.38 1.64 1.63 1.99 3.28 1.31 1.42 2.49 3.58 1.22  
1.05 2.01 1.53 1.61 0.83 1.83 1.71 1.9 0.94 0.84 0.8 1.1 0.88 0.81  
0.75 0.64 0.55 1.02 1.14 1.3 0.68 0.86 1.26 1.55 2.7 0.96 0.97 1.11  
1.24 1.06 1.41]

#### Color intensity

[ 5.64 4.38 5.68 7.8 4.32 6.75 5.25 5.05 5.2 7.22 5.75 5.  
5.6 5.4 7.5 7.3 6.2 6.6 8.7 5.1 5.65 4.5 3.8 3.93  
3.52 3.58 4.8 3.95 4.7 5.7 6.9 3.84 4.2 4.6 4.25 3.7  
6.13 4.28 5.43 4.36 5.04 5.24 4.9 6.1 8.9 7.2 7.05 6.3  
5.85 6.25 6.38 6. 6.8 1.95 3.27 4.45 2.95 5.3 4.68 3.17  
2.85 3.05 3.38 3.74 3.35 3.21 2.65 3.4 2.57 2.5 3.9 2.2  
2.62 2.45 2.6 2.8 1.74 2.4 3.6 2.15 3.25 2.9 2.3 3.3  
2.06 2.94 2.7 2. 3.08 1.9 1.28 2.08 2.76 3.94 3. 2.12  
4.1 5.45 7.1 3.85 4.92 4.35 4.4 8.21 4. 7.65 8.42 9.4  
8.6 10.8 10.52 7.6 7.9 9.01 13. 11.75 5.88 5.58 5.28 9.58  
6.62 10.68 10.26 8.66 8.5 5.5 9.9 9.7 7.7 10.2 9.3 9.2 ]

#### Hue

[1.04 1.05 1.03 0.86 1.02 1.06 1.08 1.01 1.25 1.17 1.15 1.2  
1.28 1.07 1.13 1.23 0.96 1.09 1.11 1.12 0.92 1.19 1.1 1.18  
0.89 0.95 0.91 0.88 0.82 0.87 1.24 0.98 0.94 1.22 1.45 0.906  
1.36 1.31 0.99 1.38 1.16 0.84 0.79 1.33 1. 1.42 1.27 0.8  
0.75 0.9 0.93 1.71 0.7 0.73 0.69 0.97 0.76 0.74 0.66 0.78  
0.81 0.77 0.65 0.6 0.58 0.54 0.55 0.57 0.59 0.48 0.61 0.56  
0.67 0.68 0.85 0.72 0.62 0.64 ]

#### OD280/OD315 of diluted wines

[3.92 3.4 3.17 3.45 2.93 2.85 3.58 3.55 2.82 2.9 2.73 3. 2.88 2.65  
2.57 3.36 3.71 3.52 4. 3.63 3.82 3.2 3.22 2.77 3.59 2.71 2.87 3.47  
2.78 2.51 2.69 3.53 3.38 3.56 3.35 3.33 3.44 2.75 3.1 2.91 3.37 3.26  
3.03 3.31 2.84 1.82 1.67 1.59 2.46 2.23 2.3 3.18 3.48 1.93 3.07 3.16  
3.5 3.13 2.14 2.48 2.52 2.31 3.12 3.14 2.72 2.01 3.08 2.26 3.21 2.27  
2.06 3.3 2.96 2.63 2.74 2.83 2.44 3.57 2.42 3.02 2.81 2.5 3.19 2.12  
3.05 3.39 3.69 3.64 3.28 1.29 1.42 1.36 1.51 1.58 1.27 1.69 2.15 2.47  
2.05 2. 1.68 1.33 1.86 1.62 1.3 1.47 1.55 1.48 1.64 1.73 1.96 1.78  
2.11 1.75 1.56 1.8 1.92 1.83 1.63 1.71 1.74 1.6 ]

#### Proline

[1065 1050 1185 1480 735 1450 1290 1295 1045 1510 1280 1320 1150 1547  
1310 1130 1680 845 780 770 1035 1015 830 1195 1285 915 1515 990  
1235 1095 920 880 1105 1020 760 795 680 885 1080 985 1060 1260  
1265 1190 1375 1120 970 1270 520 450 630 420 355 678 502 510  
750 718 870 410 472 886 428 392 500 463 278 714 515 495  
562 625 480 290 345 937 660 406 710 438 415 672 315 488

```

312 325 607 434 385 407 372 564 465 365 380 378 352 466
342 580 530 560 600 650 695 720 590 550 855 425 675 640
725 620 570 615 685 470 740 835 840]

```

```

[4]: def data_info(_data):
    print(_data.describe())
    _data.hist()
    plt.tight_layout()
    plt.show()
    plt.figure() # new plot
    plt.tight_layout()
    corMat = _data.corr(method='pearson')
    print(corMat)
    ## plot correlation matrix as a heat map
    sns.heatmap(corMat, square=True)
    plt.yticks(rotation=0)
    plt.xticks(rotation=90)
    plt.title(f"CORRELATION MATRIX USING HEAT MAP")
    plt.show()

    ## scatter plot of all _data
    plt.figure()
    # # The output overlaps itself, resize it to display better (w padding)
    scatter_matrix(_data)
    plt.tight_layout(pad=0.1)
    plt.show()

data_info(data)

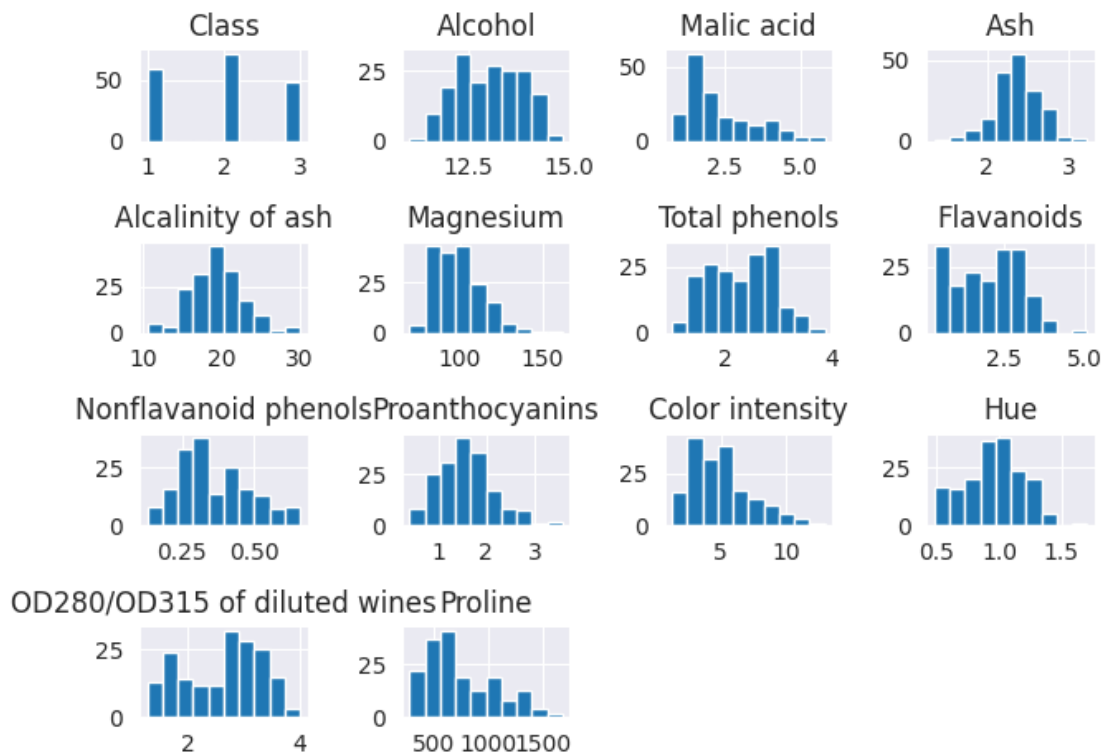
```

	Class	Alcohol	Malic acid	Ash	Alcalinity of ash \
count	178.000000	178.000000	178.000000	178.000000	178.000000
mean	1.938202	13.000618	2.336348	2.366517	19.494944
std	0.775035	0.811827	1.117146	0.274344	3.339564
min	1.000000	11.030000	0.740000	1.360000	10.600000
25%	1.000000	12.362500	1.602500	2.210000	17.200000
50%	2.000000	13.050000	1.865000	2.360000	19.500000
75%	3.000000	13.677500	3.082500	2.557500	21.500000
max	3.000000	14.830000	5.800000	3.230000	30.000000

	Magnesium	Total phenols	Flavanoids	Nonflavanoid phenols \
count	178.000000	178.000000	178.000000	178.000000
mean	99.741573	2.295112	2.029270	0.361854
std	14.282484	0.625851	0.998859	0.124453
min	70.000000	0.980000	0.340000	0.130000
25%	88.000000	1.742500	1.205000	0.270000
50%	98.000000	2.355000	2.135000	0.340000
75%	107.000000	2.800000	2.875000	0.437500
max	162.000000	3.880000	5.080000	0.660000

	Proanthocyanins	Color intensity	Hue \
count	178.000000	178.000000	178.000000
mean	1.590899	5.058090	0.957449
std	0.572359	2.318286	0.228572
min	0.410000	1.280000	0.480000
25%	1.250000	3.220000	0.782500
50%	1.555000	4.690000	0.965000
75%	1.950000	6.200000	1.120000
max	3.580000	13.000000	1.710000

	OD280/OD315 of diluted wines	Proline
count	178.000000	178.000000
mean	2.611685	746.893258
std	0.709990	314.907474
min	1.270000	278.000000
25%	1.937500	500.500000
50%	2.780000	673.500000
75%	3.170000	985.000000
max	4.000000	1680.000000



	Class	Alcohol	Malic acid	Ash \
Class	1.000000	-0.328222	0.437776	-0.049643

Alcohol	-0.328222	1.000000	0.094397	0.211545
Malic acid	0.437776	0.094397	1.000000	0.164045
Ash	-0.049643	0.211545	0.164045	1.000000
Alcalinity of ash	0.517859	-0.310235	0.288500	0.443367
Magnesium	-0.209179	0.270798	-0.054575	0.286587
Total phenols	-0.719163	0.289101	-0.335167	0.128980
Flavanoids	-0.847498	0.236815	-0.411007	0.115077
Nonflavanoid phenols	0.489109	-0.155929	0.292977	0.186230
Proanthocyanins	-0.499130	0.136698	-0.220746	0.009652
Color intensity	0.265668	0.546364	0.248985	0.258887
Hue	-0.617369	-0.071747	-0.561296	-0.074667
OD280/OD315 of diluted wines	-0.788230	0.072343	-0.368710	0.003911
Proline	-0.633717	0.643720	-0.192011	0.223626

	Alcalinity of ash	Magnesium	Total phenols	\
Class	0.517859	-0.209179	-0.719163	
Alcohol	-0.310235	0.270798	0.289101	
Malic acid	0.288500	-0.054575	-0.335167	
Ash	0.443367	0.286587	0.128980	
Alcalinity of ash	1.000000	-0.083333	-0.321113	
Magnesium	-0.083333	1.000000	0.214401	
Total phenols	-0.321113	0.214401	1.000000	
Flavanoids	-0.351370	0.195784	0.864564	
Nonflavanoid phenols	0.361922	-0.256294	-0.449935	
Proanthocyanins	-0.197327	0.236441	0.612413	
Color intensity	0.018732	0.199950	-0.055136	
Hue	-0.273955	0.055398	0.433681	
OD280/OD315 of diluted wines	-0.276769	0.066004	0.699949	
Proline	-0.440597	0.393351	0.498115	

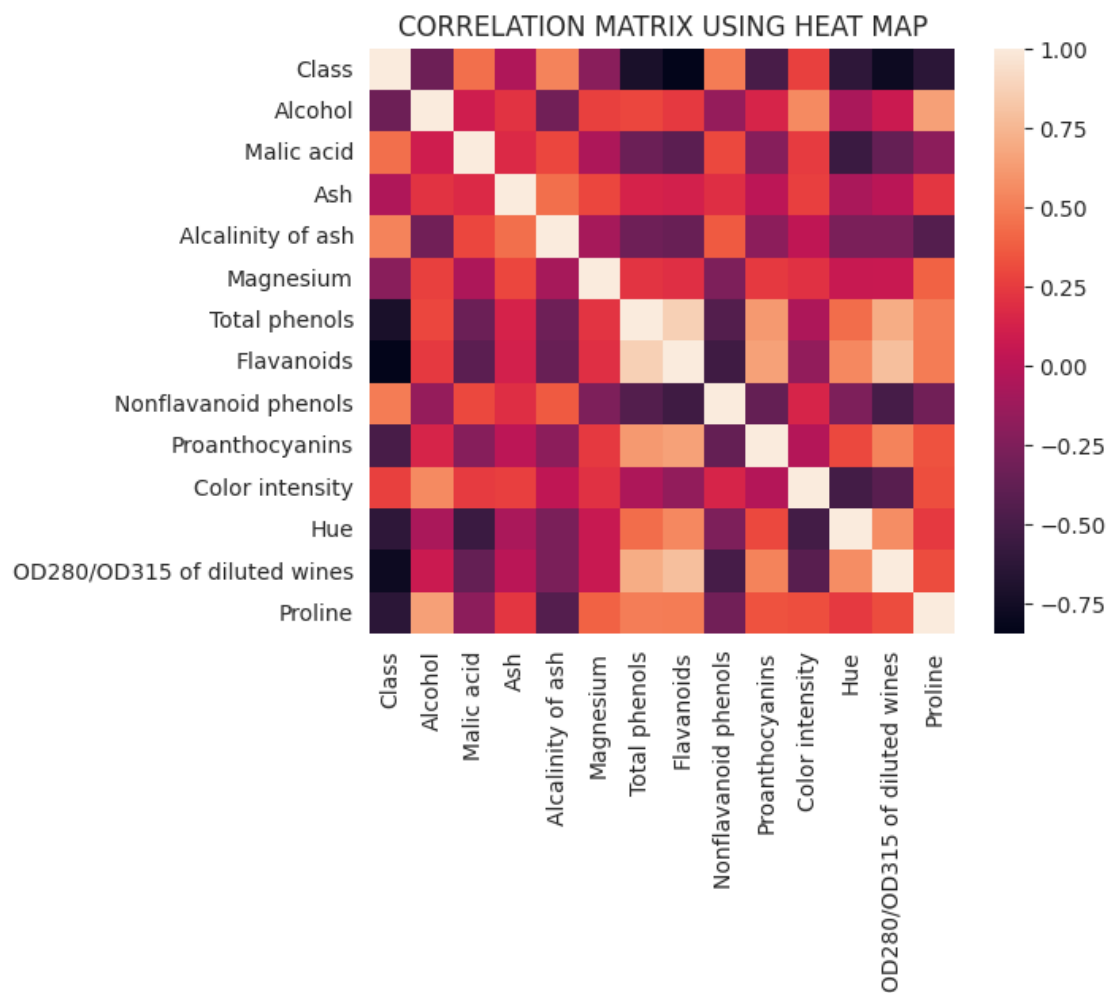
	Flavanoids	Nonflavanoid phenols	\
Class	-0.847498	0.489109	
Alcohol	0.236815	-0.155929	
Malic acid	-0.411007	0.292977	
Ash	0.115077	0.186230	
Alcalinity of ash	-0.351370	0.361922	
Magnesium	0.195784	-0.256294	
Total phenols	0.864564	-0.449935	
Flavanoids	1.000000	-0.537900	
Nonflavanoid phenols	-0.537900	1.000000	
Proanthocyanins	0.652692	-0.365845	
Color intensity	-0.172379	0.139057	
Hue	0.543479	-0.262640	
OD280/OD315 of diluted wines	0.787194	-0.503270	
Proline	0.494193	-0.311385	

	Proanthocyanins	Color intensity	Hue	\
Class	-0.499130	0.265668	-0.617369	

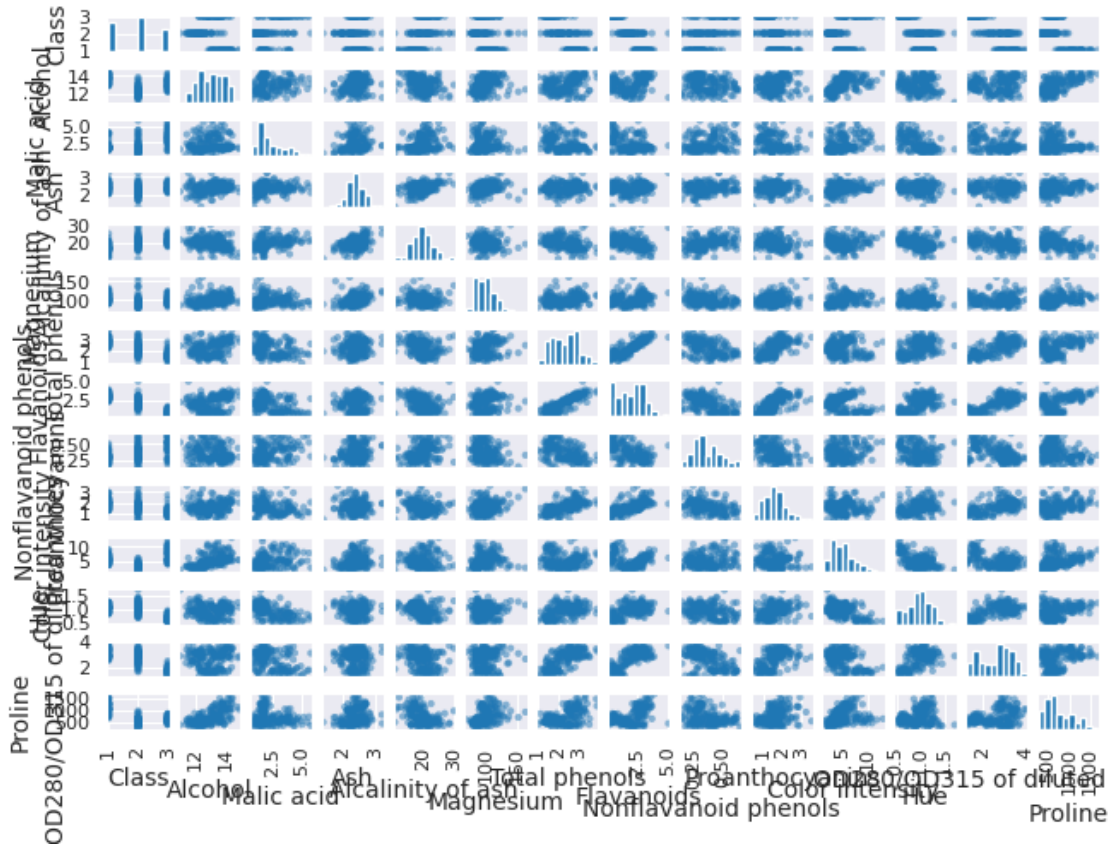
Alcohol	0.136698	0.546364	-0.071747
Malic acid	-0.220746	0.248985	-0.561296
Ash	0.009652	0.258887	-0.074667
Alcalinity of ash	-0.197327	0.018732	-0.273955
Magnesium	0.236441	0.199950	0.055398
Total phenols	0.612413	-0.055136	0.433681
Flavanoids	0.652692	-0.172379	0.543479
Nonflavanoid phenols	-0.365845	0.139057	-0.262640
Proanthocyanins	1.000000	-0.025250	0.295544
Color intensity	-0.025250	1.000000	-0.521813
Hue	0.295544	-0.521813	1.000000
OD280/OD315 of diluted wines	0.519067	-0.428815	0.565468
Proline	0.330417	0.316100	0.236183

	OD280/OD315 of diluted wines	Proline
Class	-0.788230	-0.633717
Alcohol	0.072343	0.643720
Malic acid	-0.368710	-0.192011
Ash	0.003911	0.223626
Alcalinity of ash	-0.276769	-0.440597
Magnesium	0.066004	0.393351
Total phenols	0.699949	0.498115
Flavanoids	0.787194	0.494193
Nonflavanoid phenols	-0.503270	-0.311385
Proanthocyanins	0.519067	0.330417
Color intensity	-0.428815	0.316100
Hue	0.565468	0.236183
OD280/OD315 of diluted wines	1.000000	0.312761
Proline	0.312761	1.000000





<Figure size 640x480 with 0 Axes>



### 1.5 Standardize the data as all are continuous variables and appear mostly gaussian

```
[5]: Y1 = data['Class']
data_stand = data.copy()
X1_stand = data.drop(columns=['Class'])
X1_stand = StandardScaler().fit(X1_stand).transform(X1_stand)
```

```
[6]: from matplotlib import pyplot

def evaluate_each_model_in_turn(models, X, Y):
    results = []
    names = []
    scoring = 'accuracy'
    for name, model in models:
        kfold = KFold(n_splits=10, random_state=7, shuffle=True)
        cv_results = cross_val_score(model, X, Y, cv=kfold, scoring=scoring)
        results.append(cv_results)
        names.append(name)
        msg = "%s: %f (%f)" % (name, cv_results.mean(), cv_results.std())
```

```

    print(msg)
    ## boxplot algorithm comparison
    fig = pyplot.figure()
    fig.suptitle('Algorithm Comparison')
    ax = fig.add_subplot(111)
    pyplot.boxplot(results)
    ax.set_xticklabels(names)
    pyplot.show()

```

1.6 3.

1.7 add an SV Classifier(SVC: use the default settings given in the sample code, use RFB kernel with  $C = 1.0$ )

1.8 a random forest classifier with a depth of 2

1.9 and an Adaboost classifier

1.10 and compare them using kfold cross validation with  $k=10$ .

```

[7]: # add an SV Classifier(SVC), a random forest classifier with a depth of 2 and
    ↪ an Adaboost
    # classifier and compare them using kfold cross validation with k=10. For the
    ↪ SVC, use
    # the default settings given in the sample code, use RFB kernel with C = 1.0
    models = []
    from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier
    from sklearn.svm import SVC
    svc_model = ('SVC', SVC(kernel='rbf', C=1.0, random_state=1, probability=True))
    models.append(svc_model)
    models.append(('Random Forest', RandomForestClassifier(max_depth=2,
    ↪ random_state=1)))
    models.append(('AdaBoost', AdaBoostClassifier(n_estimators=100,
    ↪ random_state=1)))

    from sklearn.model_selection import train_test_split
    X_train, X_test, Y_train, Y_test = train_test_split(X1_stand, Y1, test_size=0.2)

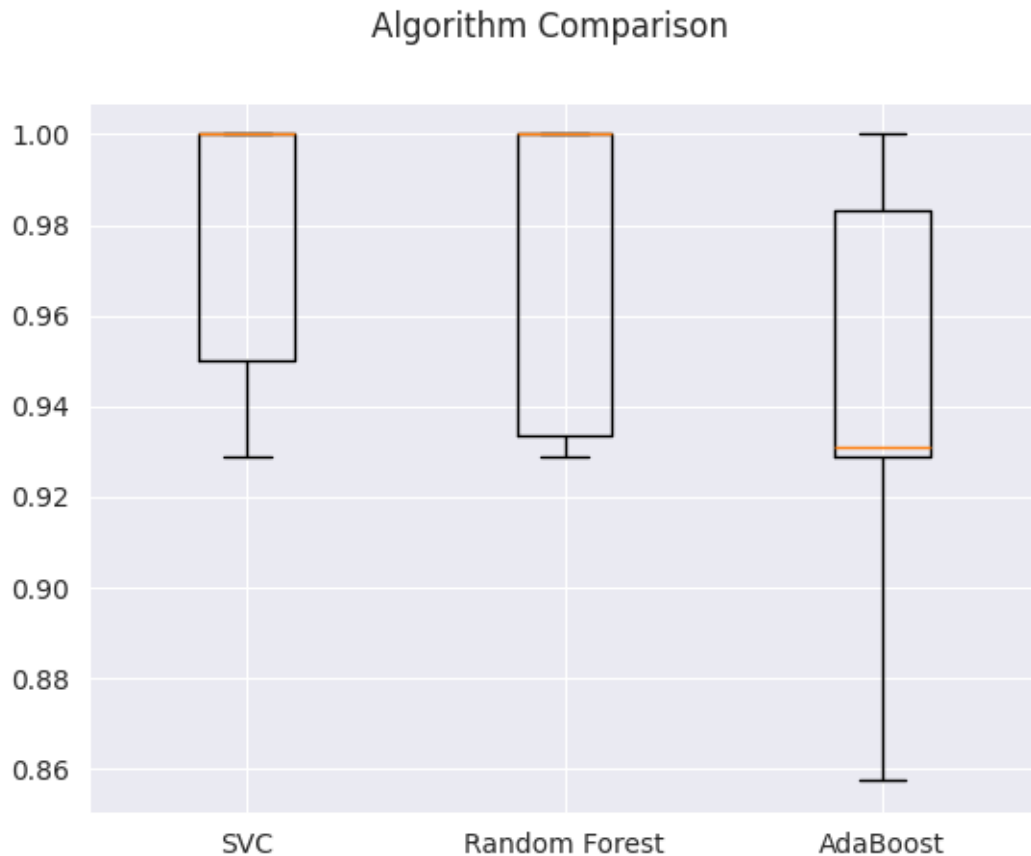
    evaluate_each_model_in_turn(models, X_train, Y_train)

```

SVC: 0.979048 (0.032029)

Random Forest: 0.972381 (0.033860)

AdaBoost: 0.936667 (0.049900)



1.11 4. Plot all the accuracy results vs. each model (model type on the x-axis and accuracy on the y-axis)

1.11.1 Results:

1.11.2 1. The SVC with RBF kernel has the highest accuracy.

1.11.3 2. The AdaBoost classifier has the lowest accuracy.

1.11.4 3. SVC also has the lowest standard deviation, indicating more stable performance.

1.11.5 4. In all runs, SVC continuously outperformed the other classifiers.

```
[8]: accuracy_dict = {}  
for name, model in models:  
    model.fit(X_train, Y_train)  
    y_pred = model.predict(X_test)  
    accuracy = model.score(X_test, Y_test)  
    accuracy_dict[name] = accuracy  
    print(f"{name} Accuracy: {accuracy:.4f}")
```

```

# Plotting the accuracies
import matplotlib.pyplot as plt
import numpy as np
import seaborn as sns

# Set the style
sns.set_theme(style="whitegrid")
methods = list(accuracy_dict.keys())
accuracies = list(accuracy_dict.values())

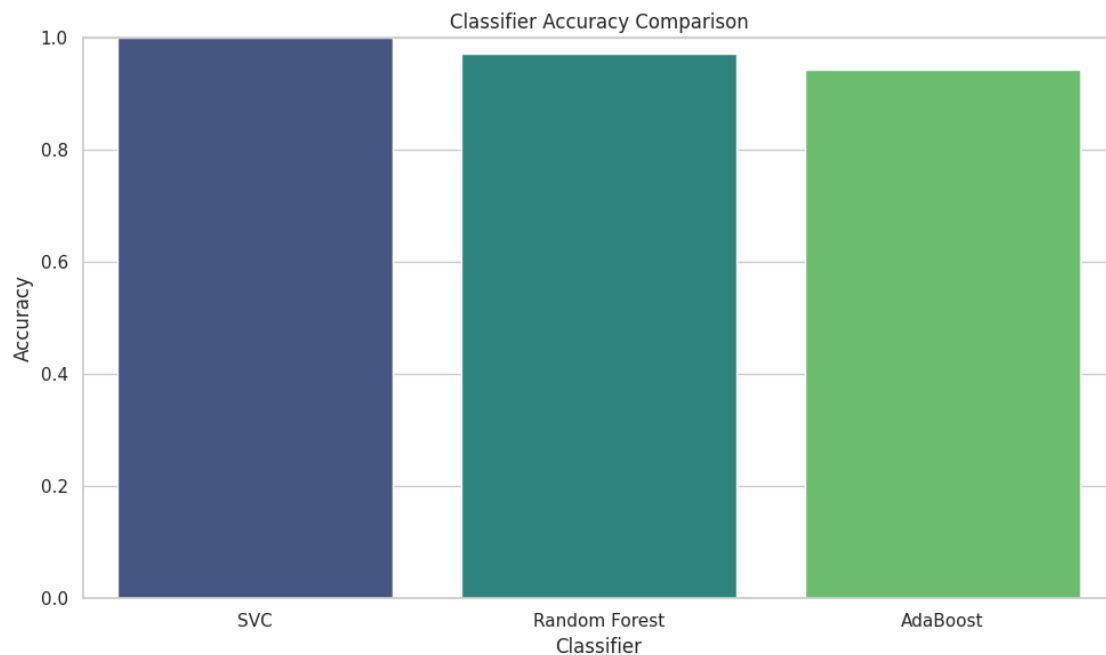
plt.figure(figsize=(10, 6))
sns.barplot(x=methods, y=accuracies, hue=methods, palette="viridis",
            legend=False)
plt.title('Classifier Accuracy Comparison')
plt.xlabel('Classifier')
plt.ylabel('Accuracy')
plt.ylim(0, 1) # Accuracy domain
plt.tight_layout()
plt.show()

```

SVC Accuracy: 1.0000

Random Forest Accuracy: 0.9722

AdaBoost Accuracy: 0.9444



1.12 5. Try a polynomial kernel by setting `kernel = 'poly'` and change the kernel degree from 2 – 5.

1.13 6. Compare the results with the RBF kernel and the same value of `C=1.0`

1.13.1 Results:

1.13.2 The RBF kernel significantly outperforms the polynomial kernel in accuracy. The polynomial kernel has both lower accuracy and higher variance in cross-validation scores, indicating less robust performance across different data subsets.

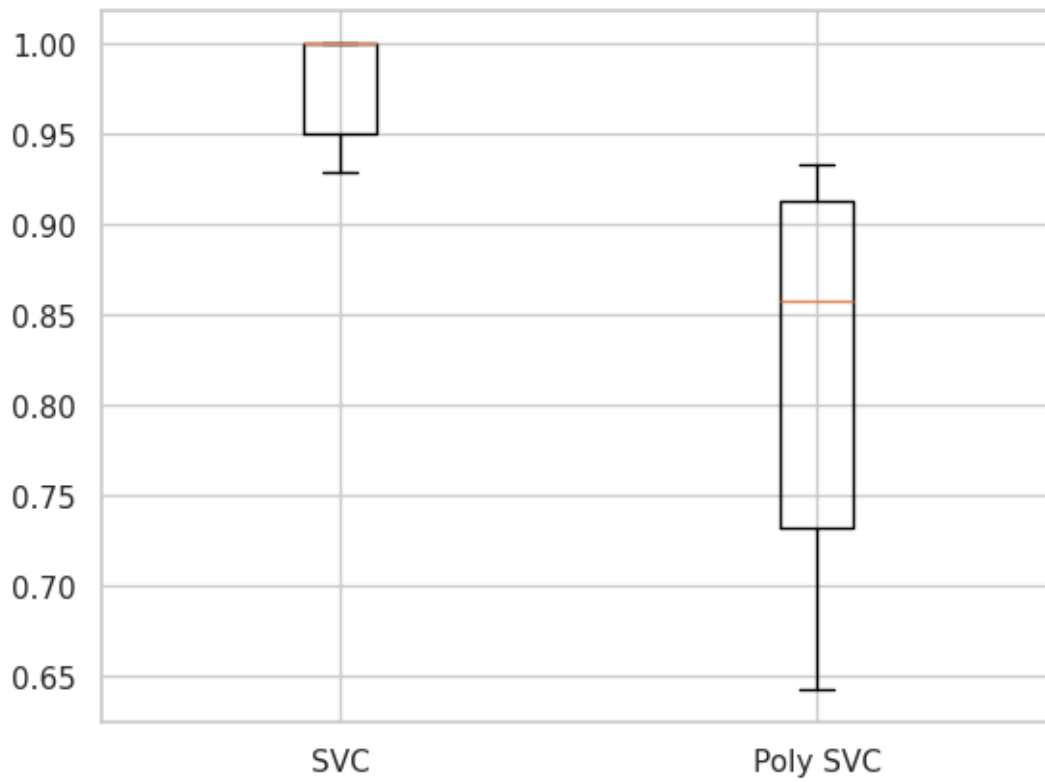
```
[9]: poly_svc_model = SVC(kernel='poly', degree=5, C=1.0, random_state=1,
    ↪probability=True)
    # fit and append
    poly_svc_model.fit(X_train, Y_train)
    y_pred_poly = poly_svc_model.predict(X_test)
    accuracy_poly = poly_svc_model.score(X_test, Y_test)
    poly_svc = ('Poly SVC', poly_svc_model)
    accuracy_dict[poly_svc[0]] = accuracy_poly
    models.append(poly_svc)

    compare_svc_models = [svc_model, poly_svc]
    evaluate_each_model_in_turn(compare_svc_models, X_train, Y_train)
```

SVC: 0.979048 (0.032029)

Poly SVC: 0.815714 (0.107835)

## Algorithm Comparison



1.14 7. Write down your observation on the comparison results.

1.14.1 Results:

1.14.2 1. The RBF kernel significantly outperforms the polynomial kernel in accuracy.

1.14.3 2. The polynomial kernel has both lower accuracy and higher variance in cross-validation scores, indicating less robust performance across different data subsets.

1.14.4 3. Despite the accuracy differences, the polynomial kernel still achieves excellent ROC AUC, suggesting good ranking capabilities even when making some classification errors.

1.15 8. Plot the multi-class ROC curve and use the `roc_auc_score` function to calculate ROC score.

1.15.1 Results:

1.15.2 1. Both SVC and Random Forest achieve perfect ROC AUC scores despite Random Forest having lower accuracy. This suggests Random Forest still ranks predictions correctly even when it makes some misclassifications.

1.15.3 2. AdaBoost performs slightly worse but still shows excellent performance

```
[12]: # Plot the multi-class ROC curve and use the roc_auc_score function to
      ↪ calculate ROC
      # AUC for each class.
      from sklearn.metrics import roc_curve, roc_auc_score

      # Now you can calculate and plot ROC curves
      from sklearn.metrics import roc_curve, auc
      from sklearn.preprocessing import label_binarize
      import numpy as np

      # For multi-class problems, you need to binarize the output
      y_test_bin = label_binarize(Y_test, classes=np.unique(Y_test))
      n_classes = y_test_bin.shape[1]

      # Plot ROC curves
      plt.figure(figsize=(10, 8))

      for name, model in models:
          # Getting probabilities
          y_score = model.predict_proba(X_test)

          # Compute ROC curve and ROC area for each class
          fpr = dict()
          tpr = dict()
          roc_auc = dict()
```



```

for i in range(n_classes):
    fpr[i], tpr[i], _ = roc_curve(y_test_bin[:, i], y_score[:, i])
    roc_auc[i] = auc(fpr[i], tpr[i])

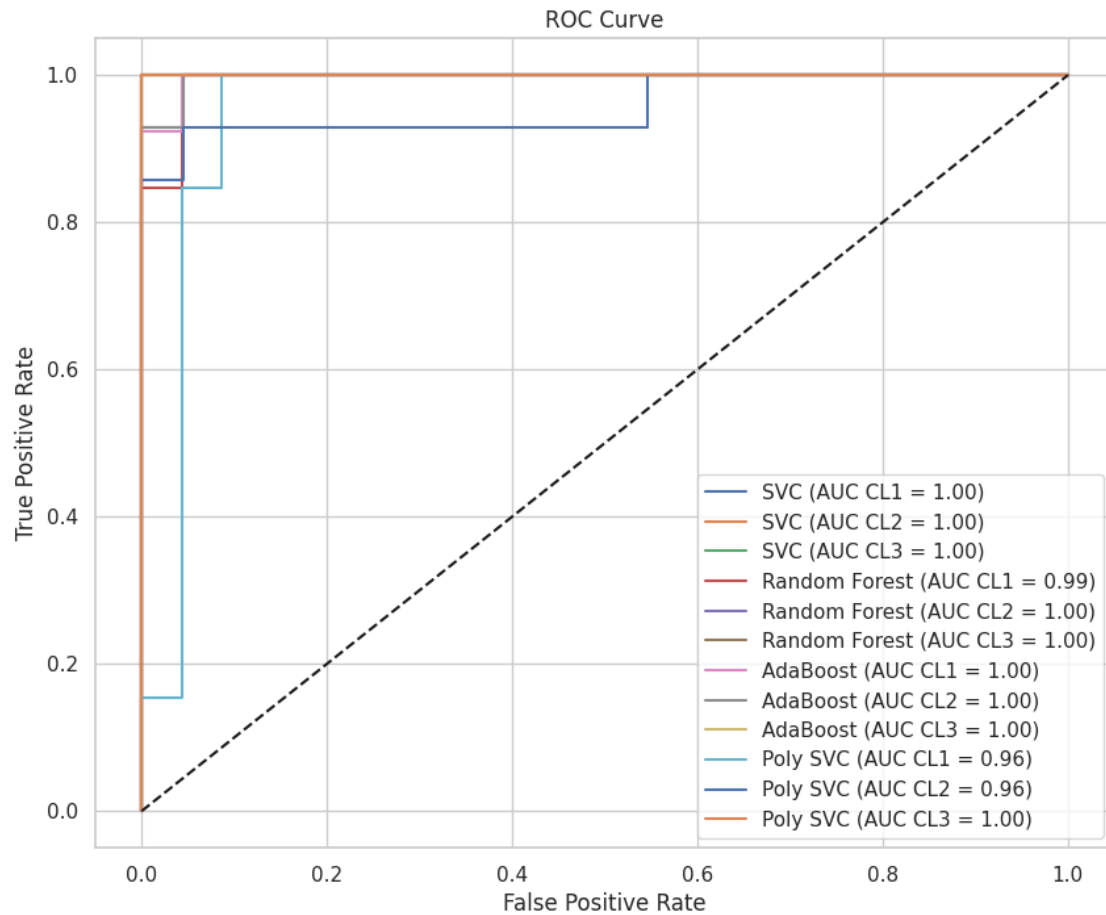
    # Plot ROC for the first class as an example
    # plt.plot(fpr[0], tpr[0], label=f'{name} (AUC = {roc_auc[0]:.2f})')
    # Plot ROC for all classes
    plt.plot(fpr[0], tpr[0], label=f'{name} (AUC CL1 = {roc_auc[0]:.2f})')
    plt.plot(fpr[1], tpr[1], label=f'{name} (AUC CL2 = {roc_auc[1]:.2f})')
    plt.plot(fpr[2], tpr[2], label=f'{name} (AUC CL3 = {roc_auc[2]:.2f})')

plt.plot([0, 1], [0, 1], 'k--')
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('ROC Curve')
plt.legend(loc='lower right')
plt.show()

# Calculate micro-average ROC curve and ROC area (will average performance
↳ across all classes)
for name, model in models:
    y_score = model.predict_proba(X_test)
    y_pred = model.predict(X_test)

    # Calculate and print micro-average ROC AUC score
    print(f"{name} - Micro-average ROC AUC: {roc_auc_score(Y_test, y_score,
↳ multi_class='ovo'):.4f}")

```



SVC - Micro-average ROC AUC: 1.0000  
 Random Forest - Micro-average ROC AUC: 0.9973  
 AdaBoost - Micro-average ROC AUC: 0.9982  
 Poly SVC - Micro-average ROC AUC: 0.9746