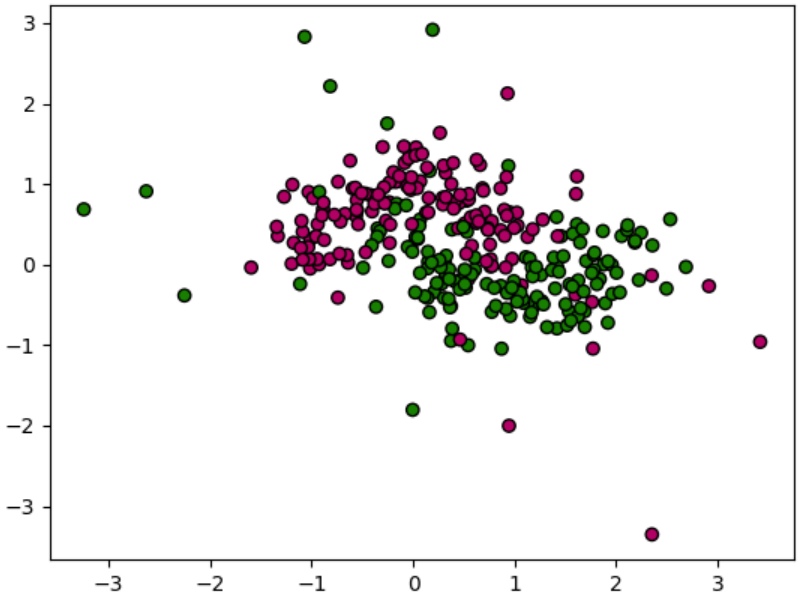
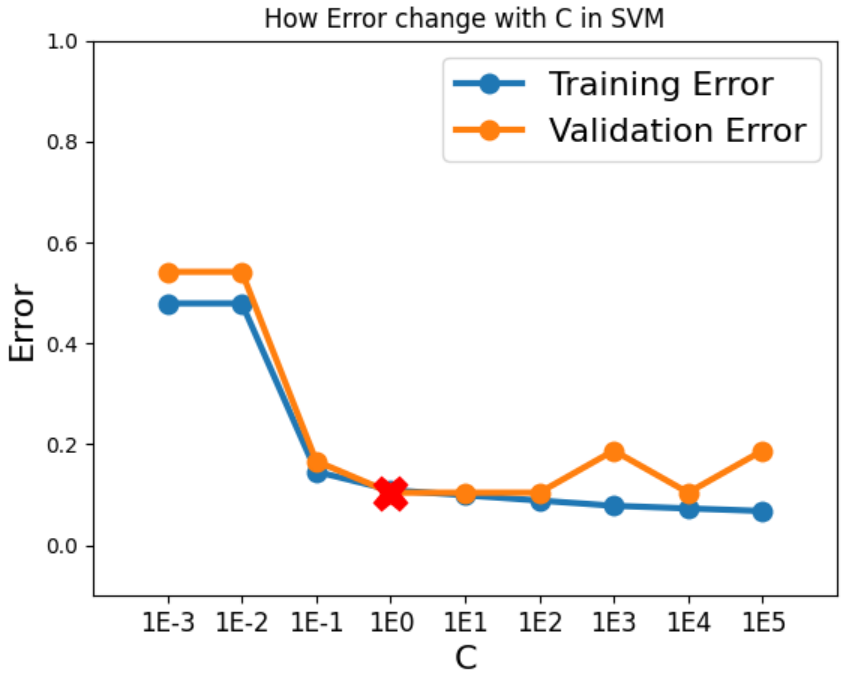
Assignment 4



1. Support Vector Machines with Synthetic Data

a. The effect of the regularization parameter C

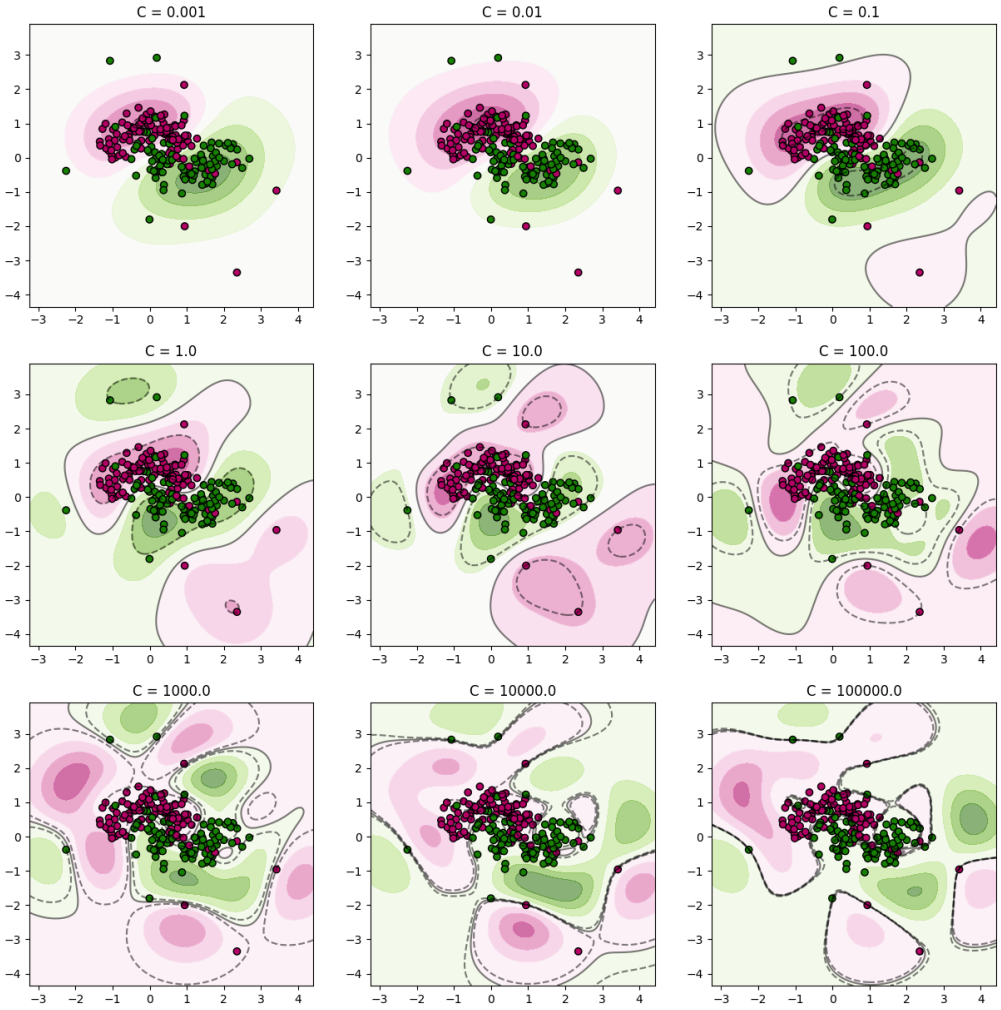
Plot:



Discussion:

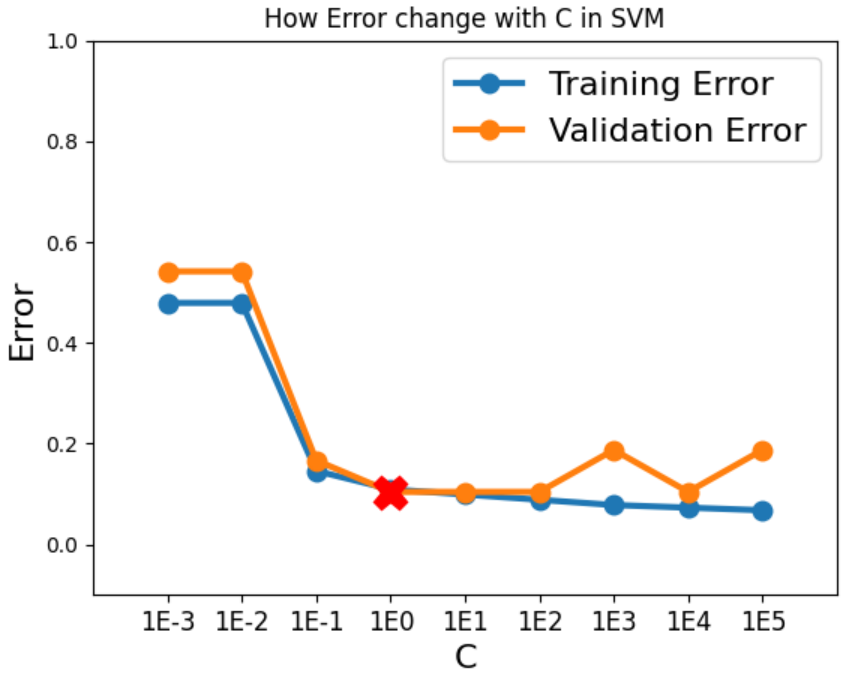
Training Error: Training Error monotonically decreases while C increases. It decreases quickly at first and slowly in the end.

Validation Error: Validation Error decreases when C <= 1 and generally increases afterwards (overfitting).



C is a trade-off between training error and flatness. Some people called C as Cross-validation parameter. While C increases, the model is softer, which means more slack are allowed and whole model is more precise and less misclassifying. Vice versa.

Final Model Selection:



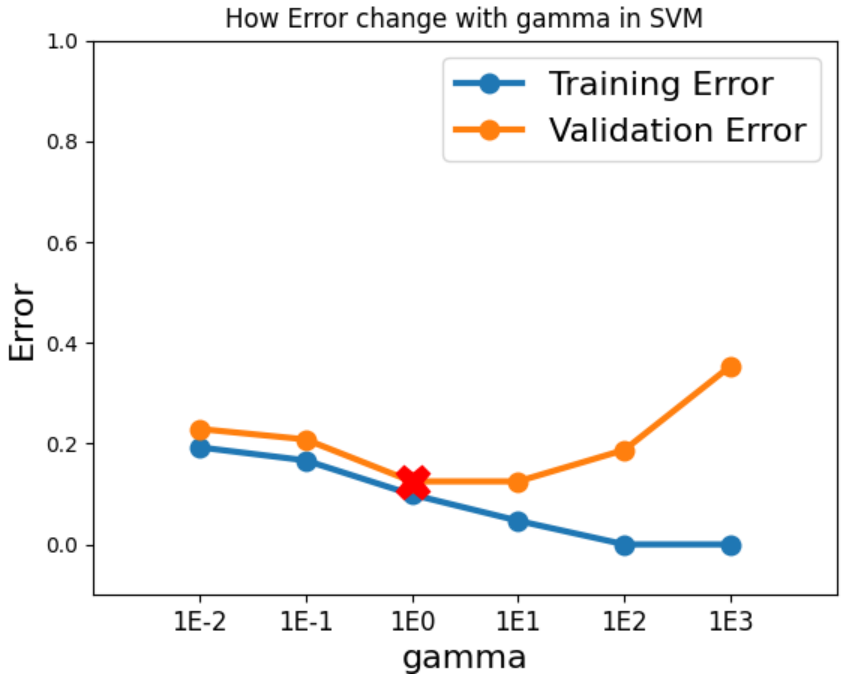
To have the least Validation Error, $C_{best} = 1$.

Console:

```
Part 1.a
Among all C values:
Best C: 1, Test Accuracy: 0.8333.
```

b. The effect of RBF kernel parameter γ

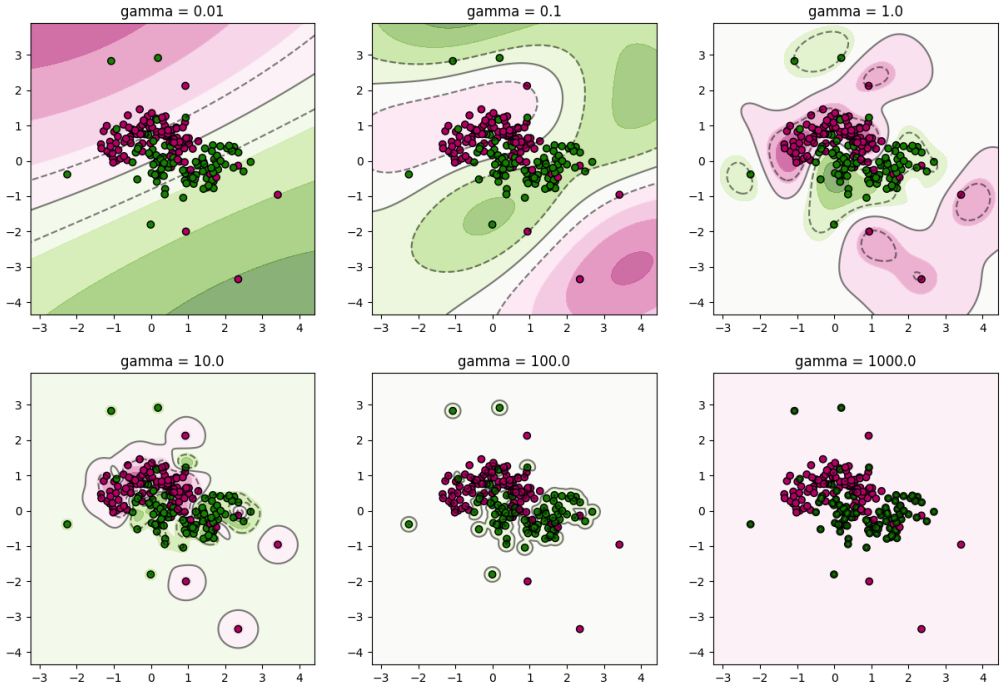
Plot:



Discussion:

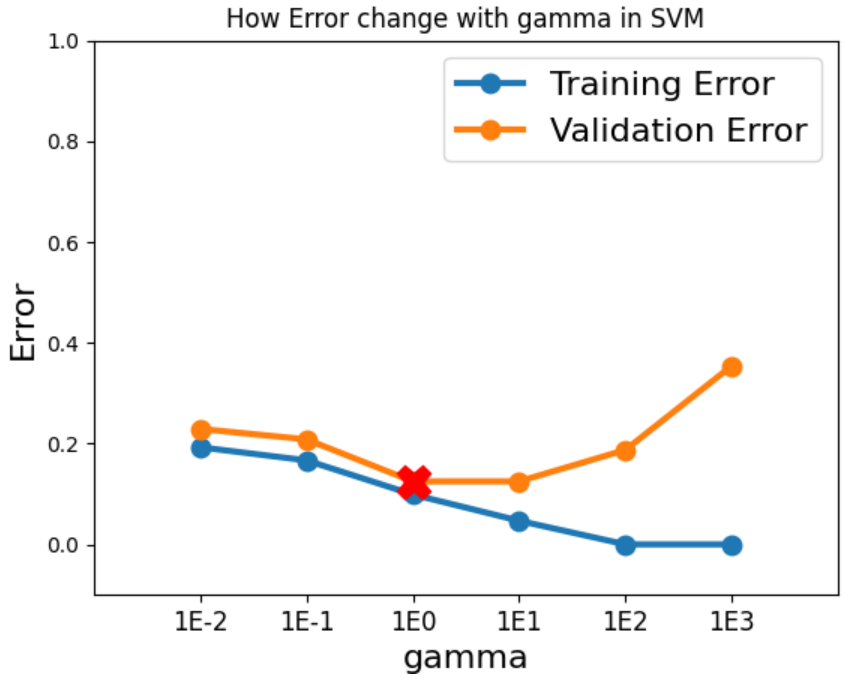
Training Error: Training Error monotonically decreases while γ increases. It decreases quickly at first and slowly in the end.

Validation Error: Validation Error decreases when γ <= 1 and generally increases afterwards (overfitting).



γ defines how far the influence of a single training example reaches. For a big **γ**, it will generate a sharp heap which will locate most of its contribution near the center. Hence, less constrain will cause the model loss the sense of the overall shape of data. When **γ** is large enough, the model's accuracy is close to 1 but useless for classification.

Final Model Selection:



To have the least Validation Error, $\gamma_{best} = 1$.

```
Part 1.b
Among all gamma values:
Best gamma: 1, Test Accuracy: 0.8333.
```

2. Breast Cancer Diagnosis with Support Vector Machines

Print Errors:

```
Training Errors:
gamma =     1E-3        1E-2        1E-1        1E0         1E1         1E2
C = 1E-2    0.371681    0.371681    0.371681    0.371681    0.371681    0.371681
C = 1E-1    0.306785    0.050147    0.035398    0.371681    0.371681    0.371681
C = 1E0     0.047198    0.029499    0.011799    0.000000    0.000000    0.000000
C = 1E1     0.026549    0.011799    0.000000    0.000000    0.000000    0.000000
C = 1E2     0.014749    0.002950    0.000000    0.000000    0.000000    0.000000
C = 1E3     0.005900    0.000000    0.000000    0.000000    0.000000    0.000000
C = 1E4     0.000000    0.000000    0.000000    0.000000    0.000000    0.000000
Validation Errors:
gamma =     1E-3        1E-2        1E-1        1E0         1E1         1E2
C = 1E-2    0.373913    0.373913    0.373913    0.373913    0.373913    0.373913
C = 1E-1    0.304348    0.069565    0.078261    0.373913    0.373913    0.373913
C = 1E0     0.060870    0.060870    0.043478    0.373913    0.373913    0.373913
C = 1E1     0.034783    0.043478    0.034783    0.373913    0.373913    0.373913
C = 1E2     0.034783    0.026087    0.034783    0.373913    0.373913    0.373913
C = 1E3     0.034783    (0.026087)<-best    0.034783    0.373913    0.373913    0.373913
C = 1E4     0.026087    0.026087    0.034783    0.373913    0.373913    0.373913
```

I use median to blur the matrix and find the final “best” (C, γ) pair.

```
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 1, 0, 0, 0, 0]
[0, 1, 0, 0, 0, 0]
[1, 1, 0, 0, 0, 0]
Blur to:
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 1, 0, 0, 0, 0]
[0, 1, 0, 0, 0, 0]
Blur to:
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
[0, 0, 0, 0, 0, 0]
(5, 1)
```

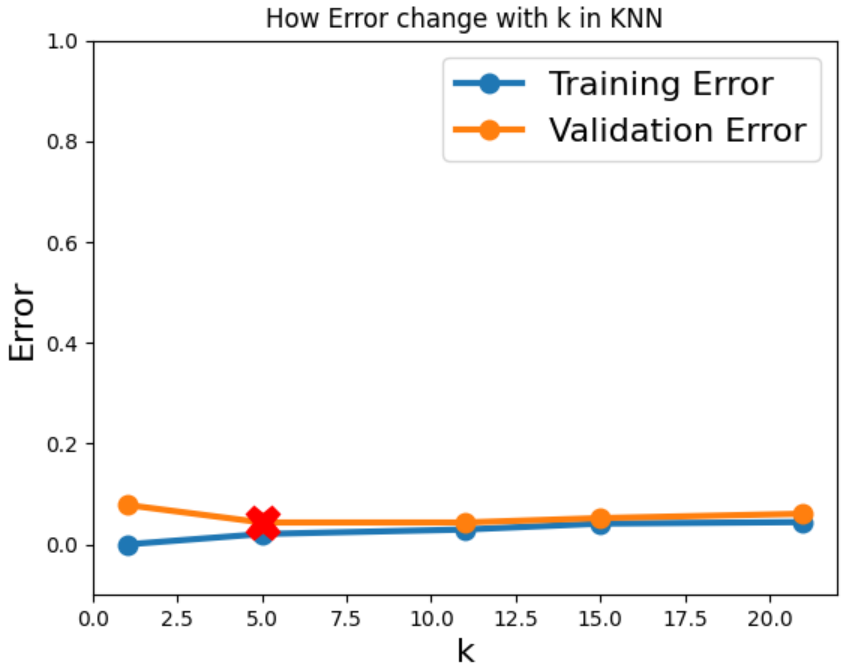
Final Model Selection:

To have the least Validation Error, C_{best} = 1000 and $\boldsymbol{\gamma}_{best}$ = 0.01.

```
Among all C and gamma value combinations:
Best C: 1000, Best gamma: 0.01, Test Accuracy: 0.9478.
```

3. Breast Cancer Diagnosis with k-Nearest Neighbors

Plot:



Final Model Selection:

To have the least Validation Error, k_{best} = 5.

```
Part 3
Among all k values:
Best k: 5, Test Accuracy: 0.9565.

Process finished with exit code 0
```

Discussion:

Depending on the result which I got, I will prefer to use kNN. For the test accuracy of kNN in k_{best} is larger than that of SVM in (C_{best}, $\boldsymbol{\gamma}_{best}$). But all these two is good for Breast Cancer Diagnosis:

Test Accuracy(SVMs(C_{best}=1000, $\boldsymbol{\gamma}_{best}$=0.01)) $= 0.9478 < 0.9565 =$ Test Accuracy(kNN(k_{best}=5))