Project 2 Report

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1 Member Roles

| | Train | MyClass | sifier | MyCrossValidate | TestMyClassifier | | | MyConfusionMatrix | |
|----------------|-------|---------|--------|-------------------|------------------|-----|-----|------------------------|--|
| | RVM | SVM | GPR | iviyCiOssvaiidate | RVM | SVM | GPR | iviy com asiom viacinx | |
| B. Poddar | | 1 | | | | 1 | | | |
| N. K. Sunki | | | 1 | | | | 1 | | |
| V. Chennapalli | 1 | | | | 1 | | | | |
| C. Luo | | | | 1 | | | | 1 | |

2 Observations

In this project, all the three algorithms were trained using all-pairs methods. Because of time limit, we performed 5-fold cross validation only on a random subset (5000 samples) of the dataset provided by Dr. Gader and compared the 3 algorithms based on this subset.

Note that the 'NaN' in confusion matrix represents there is no sample from the unseen class.

2.1 Support Vector Machines

| | C | C | C | C | C | Uns |
|-------|-------|-------|-------|-------|-------|-------|
| | C_1 | C_2 | C_3 | C_4 | C_5 | een |
| C_1 | 0.99 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| C_2 | 0.01 | 0.96 | 0.00 | 0.02 | 0.00 | 0.01 |
| C_3 | 0.00 | 0.00 | 0.98 | 0.00 | 0.00 | 0.01 |
| C_4 | 0.00 | 0.02 | 0.00 | 0.96 | 0.01 | 0.01 |
| C_5 | 0.00 | 0.00 | 0.00 | 0.01 | 0.98 | 0.00 |
| Uns | NaN | NaN | NaN | NaN | NaN | NaN |
| een | INCIN | INGIN | INGIN | INCIN | INGIN | INCIN |

| | # of support vectors |
|----------|----------------------|
| 1st Fold | 366 |
| 2nd Fold | 362 |
| 3rd Fold | 353 |
| 4th Fold | 355 |
| 5th Fold | 358 |

Table 1 Overall Confusion Matrix for SVM.

Overall Accuracy = 97.29%. (5,000 samples)

Table 2 Number of Support Vectors for Each Fold (5,000 samples)

2.2 Relevance Vector Machines

| | C_1 | \mathcal{C}_2 | \mathcal{C}_3 | C_4 | C_5 | Uns |
|-------|----------------|-----------------|-----------------|-------|----------------|-------|
| | ^L 1 | C ₂ | L3 | 64 | C ₅ | een |
| C_1 | 0.71 | 0.05 | 0.08 | 0.13 | 0.00 | 0.03 |
| C_2 | 0.11 | 0.68 | 0.03 | 0.13 | 0.01 | 0.04 |
| C_3 | 0.02 | 0.08 | 0.66 | 0.04 | 0.16 | 0.04 |
| C_4 | 0.10 | 0.14 | 0.07 | 0.61 | 0.03 | 0.05 |
| C_5 | 0.01 | 0.01 | 0.13 | 0.05 | 0.77 | 0.04 |
| Uns | NaN | NaN | NaN | NaN | NaN | NaN |
| een | INGIN | INGIN | INGIN | INGIN | INGIN | INGIN |

| | # of relevance |
|----------|----------------|
| | vectors |
| 1st Fold | |
| 2nd Fold | |
| 3rd Fold | |
| 4th Fold | |
| 5th Fold | |

Table 3 Overall Confusion Matrix for RVM.

Overall Accuracy = 68.41%. (5,000 samples)

Table 4 Number of Relevance Vectors for Each Fold (5,000 samples)

2.3 Gaussian Process Regression

| | _ | _ | _ | _ | _ | |
|-------|-------|-------|-------|-------|-------|------|
| | C_1 | C_2 | C_3 | C_4 | C_5 | Uns |
| | | | | | | een |
| C_1 | 1.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| C_2 | 0.00 | 0.96 | 0.01 | 0.03 | 0.00 | 0.00 |
| C_3 | 0.04 | 0.00 | 0.89 | 0.00 | 0.07 | 0.00 |
| C_4 | 0.00 | 0.01 | 0.00 | 0.95 | 0.03 | 0.00 |
| C_5 | 0.01 | 0.00 | 0.00 | 0.00 | 0.98 | 0.00 |
| Uns | NaN | NaN | NaN | NaN | NaN | NaN |
| een | | | | | | |

Table 5 Overall Confusion Matrix for GPR. Overall Accuracy = 95.63% (5,000 samples)

3 Conclusion

<u>Accuracy:</u> Based on 5000 random samples, the accuracy for SVM, RVM and GPR are respectively 97.29%, 68.41% and 95.63%. Thus, SVM is the best.

Speed: From both time complexity and our experiments, SVM is the fastest algorithm.

Hence, SVM is the best algorithm in terms of accuracy and speed. To end this report, we provided the results of SVM for the whole dataset (25,000 samples) below.

| | C_1 | \mathcal{C}_2 | \mathcal{C}_3 | C_4 | C_5 | Uns |
|-------|----------------|-----------------|-----------------|----------------|----------------|-------|
| | ι ₁ | C ₂ | υ ₃ | C ₄ | C ₅ | een |
| C_1 | 0.99 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| C_2 | 0.00 | 0.97 | 0.00 | 0.02 | 0.00 | 0.00 |
| C_3 | 0.01 | 0.00 | 0.98 | 0.00 | 0.01 | 0.00 |
| C_4 | 0.00 | 0.01 | 0.00 | 0.97 | 0.01 | 0.01 |
| C_5 | 0.00 | 0.00 | 0.00 | 0.00 | 0.99 | 0.00 |
| Uns | NaN | NaN | NaN | NaN | NaN | NaN |
| een | INGIN | INGIN | INGIN | INGIN | INGIN | INGIN |

Table 6 Overall Confusion Matrix for SVM.

Overall Accuracy = 98.18% (25,000 samples)

| | # of support vectors |
|----------------------|----------------------|
| 1 st Fold | 1143 |
| 2 nd Fold | 1201 |
| 3 rd Fold | 1182 |
| 4 th Fold | 1183 |
| 5 th Fold | 1182 |

Table 7 Number of Support Vectors for Each Fold. (25,000 samples)