Machine Learning, Machine Learning (extended)

5 – Supervised Learning: Evaluation Metrics for Classification Kashif Rajpoot

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Outline

- Performance evaluation
- 0/1 loss
- Classification accuracy
- Confusion matrix
 - True positive
 - True negative
 - False positive
 - False negative
- Sensitivity
- Specificity
- ROC analysis
 - Area under curve
- Hold-out validation
- Cross-validation
 - K-fold
 - Leave-one-out
- Repeated cross-validation

Performance evaluation

- How to assess a classification algorithm?
- How to choose?
 - Classification algorithm?
 - Algorithm parameters?

0/1 loss

- 0/1 loss: proportion of times classifier is wrong
- Consider a set of classifier label predictions $t_1, t_2, ..., t_N$ and ground truth target labels $t_1^*, t_2^*, ..., t_N^*$
- Mean 0/1 loss can be computed as:
 - $\frac{1}{N}\sum_{n=1}^{N}\delta(t_n\neq t_n^*)$
- For a particular test sample prediction t_n :
 - $\delta(t_n \neq t_n^*) = 1$
 - $\delta(t_n = t_n^*) = 0$
- The lower the 0/1 loss, the better
- Advantages
 - Simple
 - Suitable for binary or multi-class classification

0/1 loss

- Disadvantage
 - Suffers from class imbalance
- Imagine we're building a classifier to detect a rare disease
 - Consider only 1% of population is diseased
 - t = 1, for diseased
 - t = 0, for healthy
- What if algorithm always predicts t = 0?
- Accuracy will be 99%, but the classification algorithm is rubbish

Classification accuracy

- A slight variant of 0/1 loss, computed as percentage accuracy
- Consider a set of classifier label predictions $t_1, t_2, ..., t_N$ and ground truth target labels $t_1^*, t_2^*, ..., t_N^*$
- Mean classification accuracy can be computed as:

•
$$100 * \frac{1}{N} \sum_{n=1}^{N} \delta(t_n = t_n^*)$$

- For a particular test sample prediction t_n :
 - $\delta(t_n \neq t_n^*) = 0$
 - $\bullet \ \delta(t_n = t_n^*) = 1$
- The higher the classification accuracy, the better
- Disadvantage
 - Suffers from class imbalance, similar to 0/1 loss

Confusion matrix (CM)

- Shows the simple count of correctly (and wrongly) classified samples by the classifier
- Consider a binary classification problem with 80 diseased and 20 healthy test samples

| | | Actual cla | Total | | | |
|-----------------------|----------|------------|-----------------|----|--|--|
| | | Diseased | iseased Healthy | | | |
| Predicted class label | Diseased | 15 | 4 | 19 | | |
| | Healthy | 5 | 76 | 81 | | |
| | Total | 20 | 80 | | | |

- Variety of metrics can be estimated from CM
- Suitable for binary and multi-class classification
 - Particularly useful for multi-class problems
 - Specifically indicates problems with an individual class

Confusion matrix

- Example: classify ~7000 test documents in 20 classes (newsgroups data)
 - Too similar classes?
 - Need more data?

| 60000000000000000000000000000000000000 | Theed more data? | | | | | | | | | | | | | | | | | | | | |
|--|------------------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|
| | True class | | | | | | | | | | | | | | | | | | | | |
| | | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 | 15 | 16 | 18 | 18 | 19 | 20 |
| | 1 | 242 | 3 | 3 | 0 | 1 | 0 | 0 | 1 | 0 | 4 | 2 | 0 | 2 | 10 | 4 | 7 | 1 | 12 | 7 | 47 |
| | 2 | 0 | 296 | 33 | 8 | 8 | 42 | 9 | 1 | 1 | 0 | 0 | 4 | 18 | 7 | 8 | 2 | 0 | 1 | 1 | 3 |
| | 3 | 0 | 6 | 209 | 15 | 9 | 8 | 4 | 0 | 0 | 0 | 0 | 1 | 0 | 1 | 0 | 1 | 0 | 0 | 0 | 0 |
| | 4 | 0 | 12 | 60 | 303 | 36 | 12 | 46 | 2 | 0 | 1 | 0 | 1 | 28 | 3 | 0 | 0 | 0 | 0 | 0 | 0 |
| | 5 | 0 | 8 | 10 | 22 | 277 | 2 | 21 | 0 | 0 | 1 | 0 | 2 | 7 | 0 | 0 | 1 | 1 | 0 | 0 | 0 |
| | 6 | 1 | 21 | 30 | 2 | 2 | 304 | 0 | 1 | 0 | 3 | 0 | 1 | 3 | 0 | 1 | 2 | 0 | 0 | 1 | 0 |
| | 7 | 0 | 1 | 0 | 5 | 5 | 1 | 235 | 5 | 1 | 2 | 0 | 1 | 1 | 0 | 0 | 0 | 1 | 0 | 0 | 0 |
| class | 8 | 0 | 3 | 1 | 6 | 4 | 0 | 31 | 356 | 25 | 3 | 1 | 0 | 9 | 4 | 0 | 0 | 2 | 2 | 1 | 0 |
| cla | 9 | 0 | 2 | 2 | 0 | 1 | 2 | 5 | 4 | 353 | 1 | 0 | 0 | 2 | 0 | 1 | 0 | 1 | 1 | 0 | 1 |
| Predicted | 10 | 0 | 0 | 2 | 0 | 1 | 1 | 0 | 2 | 2 | 348 | 4 | 0 | 0 | 1 | 0 | 0 | 1 | 1 | 0 | 0 |
| ict | 11 | 1 | 0 | 1 | 1 | 0 | 0 | 1 | 0 | 0 | 16 | 382 | 0 | 1 | 0 | 1 | 0 | 1 | 1 | 0 | 0 |
| . eq | 12 | 1 | 16 | 16 | 5 | 4 | 10 | 3 | 1 | 1 | 2 | 0 | 360 | 45 | 0 | 4 | 1 | 3 | 4 | 3 | 1 |
| $P_{\mathbf{r}}$ | 13 | 1 | 4 | 1 | 24 | 16 | 0 | 9 | 5 | 1 | 2 | 0 | 3 | 260 | 3 | 4 | 0 | 0 | 0 | 0 | 0 |
| | 14 | 2 | 3 | 4 | 0 | 8 | 0 | 2 | 0 | 1 | 0 | 2 | 2 | 6 | 324 | 4 | 1 | 1 | 0 | 3 | 3 |
| | 15 | 3 | 7 | 4 | 1 | 2 | 3 | 3 | 2 | 0 | 0 | 1 | 0 | 4 | 3 | 336 | 0 | 2 | 0 | 7 | 5 |
| | 16 | 39 | 4 | 5 | 0 | 0 | 1 | 3 | 1 | 1 | 3 | 2 | 2 | 5 | 17 | 4 | 376 | 3 | 7 | 2 | 68 |
| | 17 | 4 | 0 | 0 | 0 | 3 | 1 | 1 | 5 | 4 | 1 | 0 | 9 | 0 | 3 | 1 | 3 | 325 | 3 | 95 | 19 |
| | 18 | 7 | 1 | 0 | 0 | 0 | 1 | 3 | 1 | 2 | 2 | 1 | 0 | 2 | 6 | 2 | 1 | 2 | 325 | 4 | 5 |
| | 19 | 7 | 2 | 9 | 0 | 6 | 2 | 5 | 8 | 5 | 8 | 4 | 8 | 0 | 10 | 21 | 1 | 16 | 19 | 185 | 7 |
| | 20 | 10 | 0 | 1 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | 1 | 0 | 1 | 1 | 2 | 4 | 0 | 1 | 92 |

Confusion matrix (CM)

| | | Actual cla | Total | | |
|-----------------------|----------|------------|---------|-------|--|
| | | Diseased | Healthy | TOTAL | |
| Predicted class label | Diseased | TP | FP | TP+FP | |
| | Healthy | FN | TN | FN+TN | |
| | Total | TP+FN | FP+TN | | |

- True positive (TP): diseased samples are classified as diseased
 - Number of test samples with $t_n^*=1$ that are classified as $t_n=1$
- True negative (TN): healthy samples are classified as healthy
 - Number of test samples with $t_n^*=0$ that are classified as $t_n=0$
- False positive (FP): healthy samples are classified as diseased
 - Number of test samples with $t_n^st=0$ that are classified as $t_n=1$
- False negative (FN): diseased samples are classified as healthy
 - Number of test samples with $t_n^*=1$ that are classified as $t_n=0$

Sensitivity and Specificity

- Sensitivity: proportion of diseased samples that are classified as diseased
 - The higher, the better

$$Sen = \frac{TP}{TP + FN} = \frac{TP}{All\ Positive}$$

- Specificity: the proportion of healthy samples that are classified as healthy
 - The higher, the better

$$Spec = \frac{TN}{TN + FP} = \frac{TN}{All\ Negative}$$

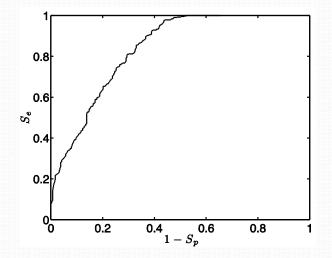
Sensitivity and Specificity

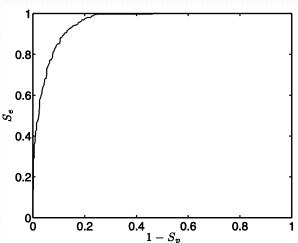
- Consider only 1% of population is diseased
- Let's say:
 - t = 1, for diseased
 - t = 0, for healthy
- What if algorithm always predicts t = 0?
 - Sen = ?
 - Sen = 0
 - *Spec* =?
 - Spec = 1
- We would like both Sen and Spec to be as high as possible
 - Often, increasing one will decrease the other
- In a disease diagnosis system:
 - We can probably tolerate a decrease in specificity (healthy samples classified as diseased), if it provides an increase in sensitivity (diseased samples classified as healthy)

Performance evaluation

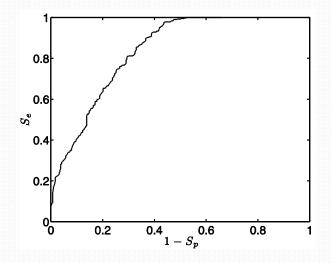
- Let's recall that often classification algorithms provide real-valued output which is then thresholded to assign a classification label
- Bayesian classifier
 - $P(t_{new} = 1 | \mathbf{x}_{new}, \mathbf{X}, t)$
- · SVM
 - $t_{new} = sign(\sum_{n=1}^{N} t_n \alpha_n k(\mathbf{x}_n, \mathbf{x}_{new}) + b)$
- How about perturbing this threshold value?

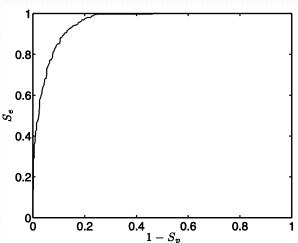
- Threshold is changed to evaluate classifier's performance
- Receiver operating characteristic (ROC) curve
 - Sensitivity (Sen) is plotted against the complementary specificity (1 – Spec)
 - Every point on the curve reflects classifier's performance at a particular threshold value



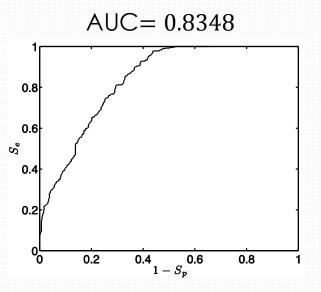


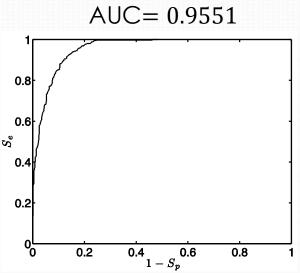
- We would like both Sen and Spec to be as high as possible
 - i.e. Sen = 1, Spec = 1, 1 Spec = 0
- Bottom left: every sample is classified as healthy (0)
- Top right: every sample is classified as diseased (1)
- Ideal: get the curve to the top left corner
 - Perfect classification: Sen = 1, Spec = 1

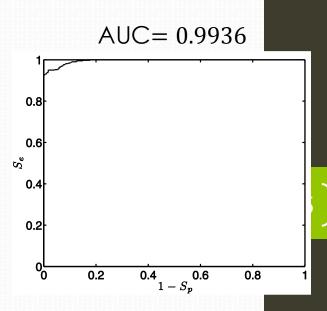




- Area under curve (AUC)
 - Quantify performance by estimating AUC
 - The higher, the better
- AUC is a better measure than 0/1 loss or classification accuracy
 - Considers class imbalance in data







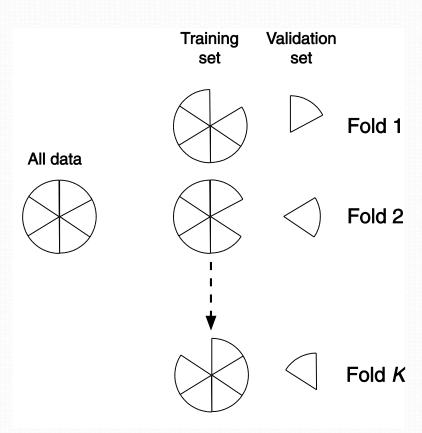
- Multi-class classification
 - Not naturally suitable
- One-against-all classification
- For example: for a 3 class problem, we will generate 3 ROC analyses
 - Each one looks at the binary classification results for class c against the rest

Hold-out validation

- We have earlier discussed the issues of overfitting and generalization
 - Ideally, we want a classifier that can generalize well i.e. on unseen data
 - Where to get unseen data?
- Hold-out validation
 - Partition observed data in to training and testing portions
 - For example: 80-20, or 50-50 split
- Disadvantages
 - Reduce training data
 - Representativeness i.e. validation is biased towards choice of data in validation set, particularly if data is small

Cross-validation

- K-fold cross-validation
 - Hold out Kth portion/fold for testing
 - Repeat the classifier training and testing for each fold
- Compute average error from all folds
- Leave-one-out cross-validation (LOOCV)
 - Extreme case of K-fold cross-validation
 - Computationally exhausting



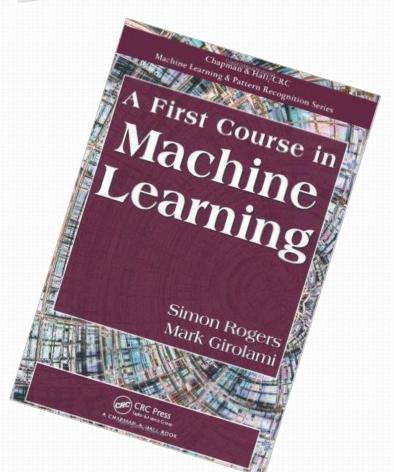
Repeated cross-validation

- The hold-out validation or k-fold cross-validation can be made more reliable by repeating it several times
 - With random selection of training and testing datasets
- Accuracies obtained from various repeats are averaged to indicate overall performance
- Computationally exhaustive

Summary

- 0/1 loss
- Classification accuracy
- Confusion matrix
- Sensitivity
- Specificity
- ROC analysis
- Assessing binary-class classification performance
- Assessing multi-class classification performance
- Cross-validation







Author's material (Simon Rogers)



Thankyou