

# Homework 6 for Pattern Recognition

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## Question 1.1

Note that

$$\begin{aligned}\varepsilon_B(x) &= h_B(x) - y(x) = \left[ \frac{1}{m} \sum_{i=1}^m h_i(x) \right] - y(x) \\ &= \frac{1}{m} \sum_{i=1}^m [h_i(x) - y(x)] = \frac{1}{m} \sum_{i=1}^m \varepsilon_i(x).\end{aligned}$$

It follows that

$$\begin{aligned}E_{h_B} &= \frac{1}{n} \sum_{j=1}^n [\varepsilon_B(x_j)]^2 = \frac{1}{m^2 n} \sum_{j=1}^n \left[ \sum_{i=1}^m \varepsilon_i(x_j) \right]^2 \\ &= \frac{1}{m^2 n} \sum_{j=1}^n \left[ \sum_{i=1}^m (\varepsilon_i(x_j))^2 + 2 \sum_{i=1}^m \sum_{k=1, k \neq i}^m \varepsilon_i(x_j) \varepsilon_k(x_j) \right] \\ &= \frac{1}{m^2 n} \left[ \sum_{i=1}^m \sum_{j=1}^n (\varepsilon_i(x_j))^2 + 2 \sum_{i=1}^m \sum_{k=1, k \neq i}^m \sum_{j=1}^n \varepsilon_i(x_j) \varepsilon_k(x_j) \right] \\ &= \frac{1}{m^2 n} \left[ \sum_{i=1}^m \sum_{j=1}^n (\varepsilon_i(x_j))^2 + 2 \sum_{i=1}^m \sum_{k=1, k \neq i}^m 0 \right] \\ &= \frac{1}{m^2 n} \left[ \sum_{i=1}^m \sum_{j=1}^n (\varepsilon_i(x_j))^2 \right] \\ &= \frac{1}{m} \frac{1}{m} \sum_{i=1}^m \left[ \frac{1}{n} \sum_{j=1}^n (\varepsilon_i(x_j))^2 \right] = \frac{1}{m} E_h.\end{aligned}$$

## Question 1.2

By Cauchy Inequality, we have

$$\left[ \sum_{i=1}^m (\varepsilon_i(x_j) \cdot 1) \right]^2 \leq \left[ \sum_{i=1}^m (\varepsilon_i(x_j))^2 \right] \cdot \left[ \sum_{i=1}^m 1^2 \right] = m \sum_{i=1}^m (\varepsilon_i(x_j))^2.$$

It follows that

$$\begin{aligned}
E_{h_B} &= \frac{1}{n} \sum_{j=1}^n [\varepsilon_B(x_j)]^2 = \frac{1}{m^2 n} \sum_{j=1}^n \left[ \sum_{i=1}^m \varepsilon_i(x_j) \right]^2 \\
&\leq \frac{1}{m^2 n} \sum_{j=1}^n \left[ m \sum_{i=1}^m (\varepsilon_i(x_j))^2 \right] \\
&= \frac{1}{mn} \sum_{j=1}^n \left[ \sum_{i=1}^m (\varepsilon_i(x_j))^2 \right] \\
&= \frac{1}{m} \sum_{i=1}^m \left[ \frac{1}{n} \sum_{j=1}^n (\varepsilon_i(x_j))^2 \right] = E_h.
\end{aligned}$$

## Question 2.1

**Lemma:** With all weights  $w_i$ ,  $d$  and  $j$  fixed, there exists  $k$  such that the following quantity

$$\sum_{i=1}^n w_i 1\{h_{a,d,j}(x_i) \neq y_i\},$$

as a function of  $a$ , is minimized at  $a = x_{k,j}$ , where  $x_{k,j}$  is the  $j$ -th feature of the  $k$ -th sample.

**Proof:** This quantity, with all weights  $w_i$ ,  $d$  and  $j$  fixed, is a linear combination of many sign functions

$$1\{h_{a,d,j}(x_i) \neq y_i\}$$

with respect to  $a$ . Therefore, it is a staircase function, which can be minimized at one of those steps. By definition, the threshold for  $1\{h_{a,d,j}(x_i) \neq y_i\}$  is  $a = x_{k,j}$ , and the lemma is thus proved.

With this lemma, we only need to find the optimal  $a$ ,  $d$ ,  $j$  from a finite set, i.e., to find  $k$ ,  $d$  and  $j$  such that

$$\begin{aligned}
a &= x_{k,j}, \\
d &= 1 \quad \text{or} \quad d = -1, \\
1 &\leq k \leq n, \\
1 &\leq j \leq p.
\end{aligned}$$

The size of this finite set is merely  $2np$ .

**Note:** For this dataset, the images are nearly binary: the gray-scale values are very close to either 0 or 1. In such cases, we can speed up calculations by neglecting the repeating values in  $X$ , i.e., doing a “unique” operation.

Iterations	Training error	Testing error
30.0000	0.0180	0.0753
100.0000	0	0.0653
200.0000	0	0.0598

Table 1: Error rates through training

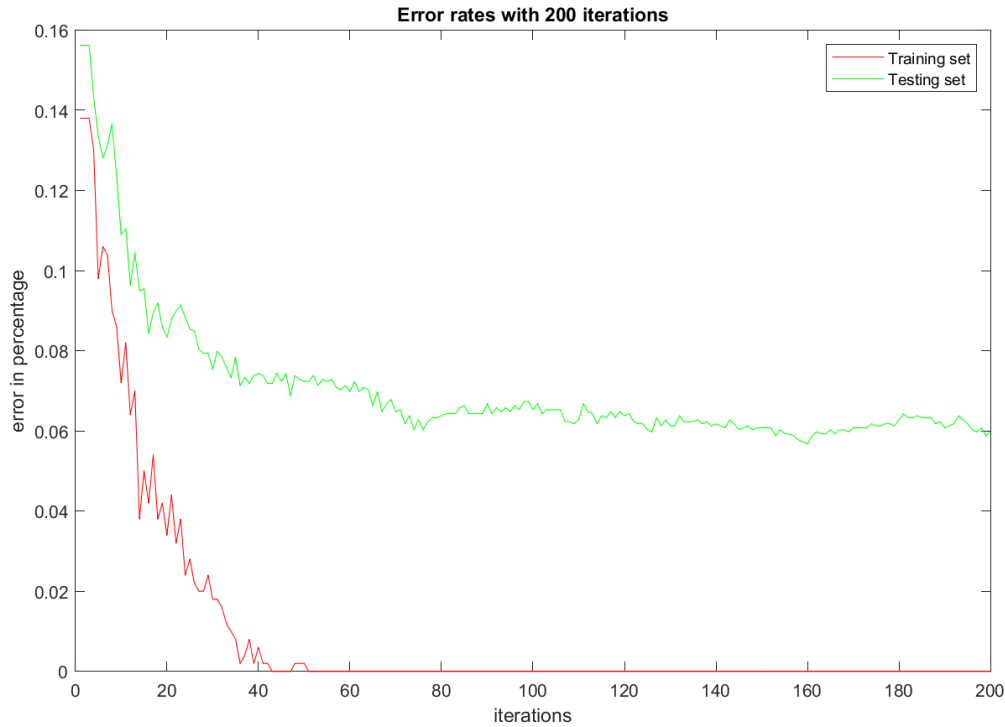


Figure 1: Error rates through training

## Questions 2.2 & 2.3

Keep using 5-fold cross validation through training.

Model	Validation error	Testing error
Fine Tree	0.122000	0.084380
Medium Tree	0.126000	0.084380
Coarse Tree	0.124000	0.098443
Bagged Trees 30	0.044000	0.047715
Bagged Trees 100	0.038000	0.037670
Bagged Trees 300	0.042000	0.035660

Table 2: Error rates of the 6 tree models

All 6 models have been saved to “models.mat” automatically.

For the 3 tree models, the FineTree (max division number = 4) and MediumTree (max division number = 20) give the best performance on both validation set and testing set, while the CoarseTree (max division number = 100) has much higher error rate, especially on the testing set.

For the 3 bagged trees, it attains the lowest error rate when the number of classifiers, the hyper-parameter, is tuned at 30.

## Question 2.4

**Briefly speaking:** “Boosting” and “random forest” are both useful workarounds to the problem that single decision trees are easily overfitting.

**Detailed comparison:**

- Judging from the output, ensembled models like bagged trees models attain the lowest error rate (around 4%), the AdaBoost the medium (around 6%), and single tree models the highest (around 12%). This means it is possible to improve the performance by ensembling multiple models. Ensembling contributes to better performance, makes the model robust against noise, and fits parallel computation as well.
- The idea of boosting is useful in that it assigns weights to different samples, and allows the classifier to concentrate on the samples in which it classifies wrong. It makes the training faster, and a bit more robust. The weights and the parameters are updated iteratively. Sometimes it also employs multiple trees and makes decision by taking a vote, as shown in AdaBoost.
- Hyperparameter tuning is critical to our model. This not only applies in a single decision tree (where we can adjust the max division number), but it is also seen in bagged trees (where we can adjust the number of classifiers). More divisions and more sub-classifiers do not always bring better performance, as shown in Table 2.
- It is necessary to identify overfitting. From Figure 1, we see it starts to overfit after 60 iterations. It is better to stop at 60 iterations than to continue training.

## Source Code

Please download the source code from [http://39.106.23.58/files/PR6\\_2015011506.7z](http://39.106.23.58/files/PR6_2015011506.7z)

For Question 2, please run “main.m”. It may take *less than a minute* to train the network, but the result is reproducible because of the random seed.

For each model, I clicked “Generate code” button to transcript my operations into MATLAB codes, and stored each of them in the corresponding “.m” file. These files include:

- trainClassifierFineTree.m
- trainClassifierMediumTree.m
- trainClassifierCoarseTree.m
- trainClassifierBaggedTrees\_30.m
- trainClassifierBaggedTrees\_100.m
- trainClassifierBaggedTrees\_300.m

Thus, the steps above can be easily reproduced without using the GUI of the toolbox.