## 1 Perceptron

Implement functions for simulating perceptron:

- p = perc\_create(n) creates and initializes randomly a perceptron with n inputs. The perceptron will be represented by an extended weight vector
   — a row vector of length n+1, where the last component is the bias of the perceptron with the opposite sign.
- c = perc\_recall(p,x) returns a row vector with the outputs of the perceptron p for a set of input (column) vectors stored in a matrix x.
- pn=perc\_update(p,x,c,lam) performs a single step of the learning algorithm for the perceptron p on a set of input vectors, where
  - $\mathbf{x}$  is a matrix with some k input vectors as columns,
  - ${\tt c}$  is a row vector of length k containing the desired outputs for the corresponding input vectors,

lam is a learning rate.

The function  $perc\_update$  performs the perceptron learning algorithm for each column x(i) with the desired output c(i), for i = 1, ..., k. Instead of adding or subtracting the vector x(i), the function adds or subtracts lam\*x(i).

- c = perc\_err(p,x,c) returns the error rate (between 0 and 1) of the perceptron p on the test set x with the correct answers c. The matrix x contains test set as column vectors, c is a row vector with the correct answers (0's and 1's).
- pn=perc\_learn(p,x,c,lam,maxit) makes at most maxit epochs of learning the perceptron p, where
  - x is a matrix of input vectors columns of the matrix,
  - c is a row vector of desired outputs of the perceptron,

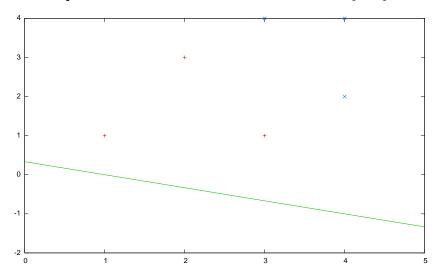
lam is a learning parameter.

The learning stops when the perceptron has a zero error on x or the number of epochs reaches maxit. One epoch consists of one step of perceptron learning performed on each input vector from x.

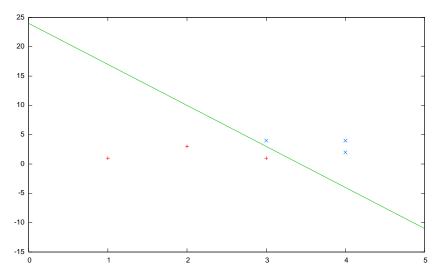
Test your implementation on the following example:

```
p=[1 \ 3 \ -1];
                              %manually initialized perceptron
xpos=[1 2 3; 1 3 1];
                              %training positive inputs, desired output 1
xneg=[3 4 4; 4 2 4];
                              %training negative inputs, desired output 0
x=[xpos xneg];
perm=[1 4 6 2 3 5];
x=x(:,perm)
                               %permute the training smaples
c=[ones(1,3) zeros(1,3)];
                               %permute also the desired outputs
c=c(perm)
xr=0:5:
                               %plot the samples and the separating
                               %hyperplane of the perceptron
```

The first plot-command illustrates the initial state of the perceptron.



The second plot draws the state of the learned perceptron.



What is the minimal number of epochs necessary to learn the perceptron to classify all training samples correctly?

# 2 Testing Machine Learning Algorithms

**Task:** Let X be a set, e.g., the set of all d-dimensional vectors with entries from the interval (0,1). Let  $f: X \to \{0,1\}$  be a target function. The goal of

a learning algorithm is to identify a function  $h: X \to \{0,1\}$ , called hypothesis, from some class of functions  $\mathcal{H}$  such that the function h is a good approximation of the target function f. The only information the algorithm can use is a sample  $S \subset X$  called training set together with the correct value f(x) for each  $x \in S$ . The sample S is a set of S elements from S randomly selected according to some probabilistic distribution S.

When the learning algorithm selects some function h, we can easily measure how well h approximates f on the training set by counting the number of elements from S for which both functions f and h have the same value. Then the error of the hypothesis h on the training set is

$$Error_S(h) = \frac{1}{n} \sum_{x \in S} \delta(f(x), h(x)) ,$$

where n is he size of S and  $\delta$  is the classification error function

$$\delta(\alpha, \beta) = \begin{cases} 1 & \text{for } \alpha \neq \beta \\ 0 & \text{for } \alpha = \beta \end{cases}$$

Hence,

$$\operatorname{Error}_S(h) = \frac{r}{n}$$
,

where r is the number of elements from S, which were incorrectly classified by h.

#### **Problems:**

- 1. How to estimate accuracy of h on any sample taken from X according to the same probability distribution D?
- 2. How good is this estimator what is the probable error of this estimator?

We want to know the error rate of the hypothesis h on the whole set X, i.e. the probability that an element  $x \in X$  selected according to the probability distribution D is classified incorrectly:

$$\mathsf{Error}_D = Pr_{x \in D}[f(x) \neq h(x)]$$
 .

Such error is called  $true\ error$ . It has a binomial distribution. An estimator of the error can be computed as the number of incorrectly classified samples divided by the number of samples. However, this estimator must be computed on a sample which is independent from the training set S! Hence:

an estimator of the error of a learning algorithm must be computed on a test set T ( $\subset X$ ) independent (disjunctive) from S.

The maximum likelihood estimate of the true error rate of the learning algorithm can be computed from the number of errors  $r_T$  on a test set T of size  $n_T = |T|$  as:

$$\operatorname{Error}_T(h) = \frac{r_T}{n_T}$$
.

Then the expected number of errors on  $n_T$  samples is  $r_T$ . Variance of the number of errors is according to the binomial distribution:

$$\mathsf{Var'}_T = n_T \cdot \frac{r_T}{n_T} \cdot \left(1 - \frac{r_T}{n_T}\right) = n_t \cdot \mathsf{Error}_T(h) \cdot \left(1 - \mathsf{Error}_T(h)\right) \ .$$

Also the standard deviation of the expected number of errors is:

$$\sigma_T' = \sqrt{n_t \cdot \mathsf{Error}_T(h) \cdot (1 - \mathsf{Error}_T(h))}$$
.

Hence he standard deviation of the estimator of error rate of hypothesis h on the test set T is

$$\sigma_T = rac{\sigma_T'}{n_T} = \sqrt{rac{\mathsf{Error}_T(h) \cdot (1 - \mathsf{Error}_T(h))}{n_T}}$$
 .

Using Matlab, it is easy to compute a confidence interval – the interval to which belongs the actual error rate of the hypothesis h on the whole set X with a prescribed probability. With the confidence  $\alpha$  the hypothesis h makes at most  $k=binoinv(\alpha, n_T, r_T/n_T)$  errors on a test set of size  $n_T$ . Therefore, the error of the hypothesis h on the whole X belongs to the interval  $<0, k/n_T>$  with the confidence  $\alpha$ .

For a sufficiently large  $n_T$  (e.g.,  $n_T \geq 30$ , or  $n_T \cdot \mathsf{Error}_T(h) \cdot (1 - \mathsf{Error}_T(h)) \geq 5$ , respectively) the binomial distribution can be approximated by the normal distribution with mean  $\mathsf{Error}_T(h)$  and standard deviation  $\sigma_T(h)$ . Consequently, the two-sided P%-confidence interval is given by the following formula

$$\mathsf{Error}_T(h) \pm z_P \cdot \sigma_T$$
,

where the values of  $z_P$  are called quantiles and can be found in statistical tables. In Matlab it is possible to compute  $z_P$  directly:

```
>> a=[80,90,95,98,99]
a =
    80    90    95    98    99
>> b=(100+a)/2/100
b =
    0.9000    0.9500    0.9750    0.9900    0.9950
>> norminv(b,0,1)
ans =
    1.2816    1.6449    1.9600    2.3263    2.5758
```

Hence, e.g.,  $z_{95} = 1.96$  and  $z_{99} = 2.5758$ .

In a single graph, plot the binomial probability distribution for n=40 samples with the probability of success p=0.1 (as this is a discrete probability distribution, your task is to plot the distribution function for arguments  $1, \ldots, n$ ) and the normal probability distribution with the same mean and standard deviation (in this case, it is a continuous probability distribution, hence you should plot its continuous density function).

### A general guidance for deriving a confidence interval

A N% confidence interval for a parameter p is an interval that is expected with probability N% to contain p.

Method:

- 1. Identify the parameter p, which should be estimated e.g., the error of a hypothesis  $\mathsf{Error}_D(h)$ .
- 2. Derive the estimate Y in our case,  $\mathsf{Error}_T(h)$ . If possible, use an unbiased estimate with minimal variance.
- 3. Determine the probability distribution  $D_Y$  for Y, including its mean and standard deviation.
- 4. Establish N% confidence interval i.e. find the limits L and U such that N% of samples selected according to the probability distribution  $D_Y$  are between L and U.

## 3 Comparing Learning Algorithms

Learning algorithms can be compared using methods for comparing statistical hypothesis, but better results can be obtained from pair tests or k-fold cross validation.

### 3.1 Difference in Error of Two Hypotheses

A hypothesis  $h_1$  is tested on a set  $T_1$  od size  $n_1$ , a hypothesis  $h_2$  is tested on a set  $T_2$  of size  $n_2$ . We want to estimate the difference  $d = \text{Error}_D(h_1) - \text{Error}_D(h_2)$  between their errors when the samples were drawn from the same underlying probability distribution D. Similarly to Section 2, we assume that the range of the target function is discrete and the error of the hypothesis can be computed as the fraction of incorrectly classified samples from the tested samples. An unbiased estimator for the difference of errors is

$$\widehat{d} = \mathsf{Error}_{T_1}(h_1) - \mathsf{Error}_{T_2}(h_2)$$
 .

This method has an disadvantage that the hypotheses have not only different errors but also variances. Hence the difference of the errors can be approximated by a normal distribution with the mean  $\hat{d}$  and variance

$$\sigma_{\widehat{d}}^2 = \frac{\mathsf{Error}_{T_1}(h_1) \cdot (1 - \mathsf{Error}_{T_1}(h_1))}{n_1} + \frac{\mathsf{Error}_{T_2}(h_2) \cdot (1 - \mathsf{Error}_{T_2}(h_2))}{n_2} \enspace .$$

Therefore N% confidence interval for the estimator  $\hat{d}$  is:

$$\hat{d} \pm z_N \sqrt{\sigma_{\hat{d}}^2}$$
.

#### 3.2 Paired Tests

We are interested in comparing not just two hypothesis, but two learning algorithms. We can decrease he variance of the difference in their errors by training both algorithms on the same training set a testing them on the same test set, The obtained confidence intervals are narrower, because the difference in performance of the learning algorithms are then caused by the algorithm themselves, not by different input data.

Let  $L_A$  and  $L_B$  be two learning algorithms learning the same target function f. For the comparison, we should compare  $L_A$  and  $L_B$  by learning on all subsets

of X of size n selected according to the same probabilistic distribution D. Then the expected value of the difference of error rates for the learning algorithms  $L_A$  and  $L_B$  is

$$E_{T \subset D}[\mathsf{Error}_D(L_A(T)) - \mathsf{Error}_D(L_B(T))]$$
,

where  $L_{\alpha}(T)$  denotes the hypothesis output by the learning algorithm  $L_{\alpha}$  after learning from the training set T ( $\alpha \in \{A, B\}$ ). The expected value is computed on subsets (of size n) of X selected according to the probability distribution D.

When comparing learning algorithms in practice, we have only a limited sample  $D_0$ . Therefore we split  $D_0$  randomly into a training set  $S_0$  and a test set  $T_0$ , which are disjoint. The training samples are used to train both  $L_A$  and  $L_B$ . The test data is used for comparing accuracy of the trained hypotheses – the obtained error rates are subtracted:

$$\mathsf{Error}_{T_0}(L_A(S_0)) - \mathsf{Error}_{T_0}(L_B(S_0)) \ . \tag{1}$$

Hence the difference in error rates for  $L_A$  and  $L_B$  is measured on a single training set  $S_0$  and not as the expected value of the difference on all samples S that might be drawn from the distribution D. If we have enough data, we can use them better.

### 3.3 k-Fold Cross-Validation

One way for improving the estimator given by Equation 1 is to partition the data  $D_0$  into disjoint training and test sets and to take the mean of the test errors. The procedure is as follows:

- 1. Partition the data  $D_0$  into k pairwise disjoint sets  $T_1, T_2, \ldots, T_k$  of equal size. The size of  $T_i$  should be at least 30!
- 2. For i = 1, ..., k do:

take  $T_i$  for the test set and the remaining data for the training set  $S_i$ 

$$\begin{split} S_i &:= D_0 \smallsetminus T_i \\ h_A &:= L_A(S_i) \\ h_B &:= L_B(S_i) \\ \delta_i &:= \mathsf{Error}_{T_i}(h_A) - \mathsf{Error}_{T_i}(h_B) \end{split}$$

3. The resulting value of the difference of errors of  $L_A$  a  $L_B$  is

$$\overline{\delta} = \frac{1}{k} \sum_{i=1}^{k} \delta_i .$$

The approximate N% confidence interval the real difference of errors is

$$\langle \overline{\delta} - t_{N,k-1} \cdot s_{\overline{\delta}}, \overline{\delta} + t_{N,k-1} \cdot s_{\overline{\delta}} \rangle$$

where  $s_{\overline{\delta}}$  is an estimate of the standard deviation

$$s_{\overline{\delta}} = \sqrt{\frac{1}{k(k-1)} \sum_{i=1}^{k} (\delta_i - \overline{\delta})^2}$$

and  $t_{N,k-1}$  is the  $\frac{N+100}{2}\%$  quantile of the Student t-distribution with k-1 degrees of freedom. In Matlab, we can compute  $t_{N,k-1}$  as

$$tinv((N+100)/200, k-1).$$