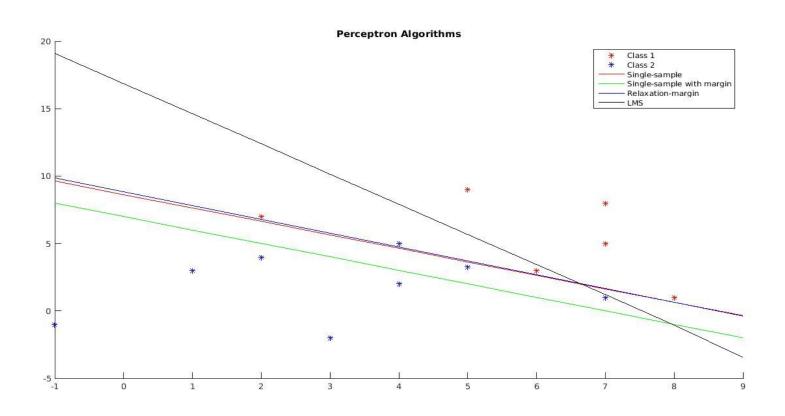
# Assignment #1: LDFs and Neural Networks

CSE471: SMAI

Implementation: Matlab

Problem 1:



Part ii.

Algorithm	Weight Vector	Convergence time(s)
Single-sample perceptron	[1,1,1]	0.0679
	[ 0.5 , 0.5 , 0.5 ]	0.0674
	[4,4,4]	0.0692
	[-1,-1,-1]	0.0691
	[1,2,3]	0.0680

Single-sample perceptron with margin	[1,1,1]	0.1323
withinargin	[ 0.5 , 0.5 , 0.5 ]	0.1341
	[4,4,4]	0.1381
	[-1,-1,-1]	0.1444
	[1,2,3]	0.1328
Relaxation algorithm with margin	[1,1,1]	1.4322
	[ 0.5 , 0.5 , 0.5 ]	1.3065
	[4,4,4]	1.4536
	[-1,-1,-1]	0.0256
	[1,2,3]	0.0262
Widrow-Hoff/LMS Rule	[1,1,1]	3.3214e-05
	[ 0.5 , 0.5 , 0.5 ]	2.2763e-05
	[4,4,4]	2.1751e-05
	[-1,-1,-1]	2.1852e-05
	[1,2,3]	2.2267e-05

There is no direct trend between higher or lower values of initial weights and convergence times, hence we can say starting with a random value of initial weights and then adjusting the initial parameters would be a good strategy.

## Part iii.

Single sample run with parameters :

init\_wts = [ones(1,dim)];

eta = 0.0005;

Relation : direct proportion

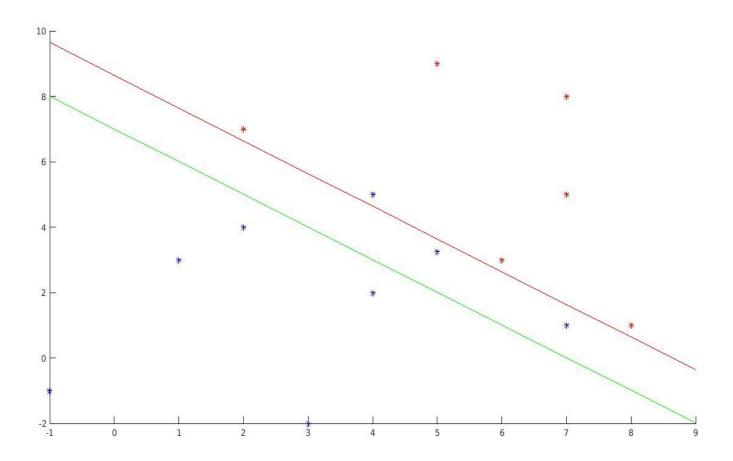
Perceptron Algorithm	Margin	Convergence Time (sec)
Single-sample perceptron with margin	0.001	2.3825e-04
	0.5	0.0038

	1	0.055
Relaxation algorithm with margin	0.1	0.340
	0.5	0.345.
	1	0.403

Hence we observe, If margin  $<< \eta(k)*norm(Yk)$ , the amount by which a is updated increases by a(k)\*y(k), which is a small amount. If it is  $>> \eta(k)*norm(Yk)$ , many corrections will be needed to satisfy the conditions  $a^T * y < b$ .

Part iii.

Varying parameters twice for non seperable point :



#### Part iv.

Initialization and augmentation of matrices:

Typical Perceptron Algorithm Run:

```
hold on
init_wts = [ones(1,dim)];
eta = 1;
res = single_sample(X,init_wts,eta);
[class1,class2] = classify(org_X,res,no_of_samples,dim);
%% plot solution a
Xpts = [-1:9];
Ypts =( res(1)* Xpts + res(3) )/ res(2) ;
plot(Xpts,-Ypts,'-r','MarkerSize',10); hold on;
```

The general approach for any perceptron algorithm that we apply is:

- 1. Preprocess the values by augmenting it and normalizing.
- 2. According to the appropriate criterion function that is minized if 'a' is solution vector, get value of 'a'
- 3. Classify using the new value of a.

#### A) Single-Sample and Single-Sample with margin :

```
function[res] = single sample(X,init wts,eta)
        [no of samples,dim] = size(X);
        a = init wts;
        prev a = zeros(1,dim);
        theta = 0.005;
        misclassified = 1;
        k = 1;
        counter = 0;
        while(1)
                misclassified = 0;
                while(k<=no of samples)
                        counter = counter + 1;
                        Y = X(k,:);
                        if a*Y' <0
                                prev a = a;
                                a = a + (eta.*Y);
                                misclassified = 1;
                                counter = 0;
                                break;
                        end
                        k = mod(k+1, no of samples);
                        if k == 0
                                k = 14:
                        end
                        if (counter==14\&misclassified==0)
                        end
                end
                if ((pdist([a;prev a])<theta)||(misclassified==0))</pre>
                        break;
                end
        end
        res = a;
end
```

The single sample algorithm, rather than testing a(k) on every sample, considers the samples in a sequence and modifies the weight vector when it misclassifies a single sample. The inner while loop runs a cycle for every updated a. The misclassified flag indicates if after the updation of a, a misclassified Yk still remains. The **single sample with margin** algorithm is exactly identical except for the condition: if  $a^TY < margin to decide whether a sample was misclassified or not.$ 

Result: In the graph obtained,

A = 0.9968 0.9984 0.9992

### C) Single Sample with Relaxation:

```
function[res] = relaxation margin(X,init wts,eta,margin,no of samples,dim)
        [no of samples,dim] = size(X);
        a = init_wts;
        prev a = zeros(1,dim);
        theta = 0.005;
        misclassified = 1;
        k = 1;
        counter = 0;
        while(1)
                misclassified = 0;
                while(k<=no of samples)
                        counter = counter + 1;
                        Y = X(k,:);
                        if a*Y' <0
                                prev a = a;
                                term = (margin - (a*Y')/norm(Y))*Y';
                                a = a + (eta.*term');
                                misclassified = 1;
                                counter = 0;
                                disp (Y)
                                break;
                        end
                        k = mod(k+1,no_of_samples);
                        if k == 0
                               k=14;
                        end
                        if (counter==14\&misclassified==0)
                                break;
                        end
                if ((pdist([a;prev a])<theta)||(misclassified==0))</pre>
                        break;
                end
        end
        res = a;
end
```

This code implements the single-sample relaxation rule. The value of 'term' in the code is the distance from a(k) to the hyperplane  $a^ty_k = b$ . The loop iterates over all samples once, as soon as it finds a misclassified sample, the value of a is updated using term. We start counting elements that are present in the cycle for the  $Y_k$ s that are present **AFTER** misclassified one and circle back till we find that all samples are correctly classified.

D) LMS:

```
function[a] = lms(X,a,b,theta,eta,no of samples,dim)
       k = 1;
       while(1)
               k = mod(k+1, no of samples);
               if k==0
                       k = 14;
               end
               Yk = X(k,:);
               Bk = b(k);
               term = (eta*(Bk - a*Yk')*Yk);
               a = a + term;
               if term'<theta
                       break
               end
       end
end
```

In this algorithm, we consider all sampled and not just the misclassified ones. Hence the code snippet doesn't have the a<sup>t</sup>y<margin as we observed before.

The error correction  $\eta$  (k)(b<sub>k</sub> - a<sup>t</sup>y<sub>k</sub>) \* y<sub>k</sub> <  $\theta$  is stored in the variable term. The loop is run until term < theta.

#### Problem 2:

