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Signal processing and identification in control of mechatronic devices

Topic: Classification and regression

## Introduction:

Classification is a predictive model that approximates a mapping function from input variables to identify discrete output variables, which can be labels or categories. The mapping function of classification algorithms is responsible for predicting the label or category of the given input variables. A classification algorithm can have both discrete and real-valued variables, but it requires that the examples be classified into one of two or more classes.

# Ex.1 Classification dataset generation

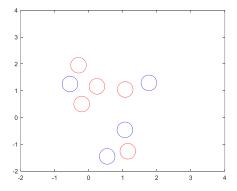
Classification of data is performed by the use of clastering. Process that requires assigning each sample with a cluster, using an unsupervised algorithm.

Code used for generating the clasters of data:

```
close all
rng('shuffle'); % To get different results each time
Clusters.ClustersA = 5; % How many clusters of data exist in class 1?
Clusters.ClustersB = 4; % How many clusters of class 2 exist?
% Definition of clusters centers
%Clusters.ACoordinates = randn(2,Clusters.ClustersA);
%Clusters.BCoordinates = randn(2,Clusters.ClustersB);
for k = 1:Clusters.ClustersA
plot(Clusters.ACoordinates(1,k),Clusters.ACoordinates(2,k),'or','MarkerSize',25); hold on
end
for k = 1:Clusters.ClustersB
plot(Clusters.BCoordinates(1,k),Clusters.BCoordinates(2,k),'ob','MarkerSize',25); hold on
end
xlim([-2 4]);
ylim([-2,4]);
```

Clusters that we will be using in the next exercises have to not overlap, should not be linearly separable.

## Example of valid result:



# Ex. 2 Generating and dividing dataset

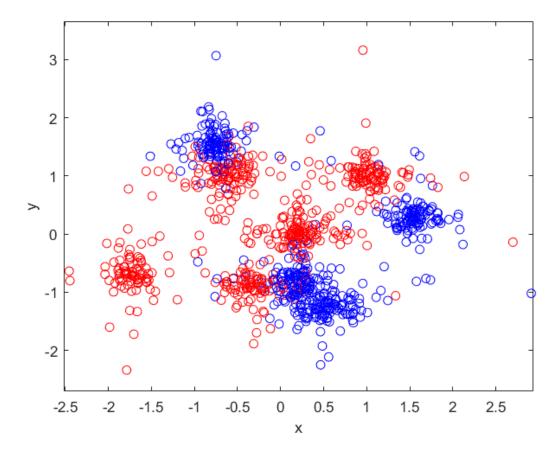
In machine learning to obtain proper results that secure us from overfitting (classifier or rergressor obtain high accuracy on training dataset, but much lower on a testing dataset), we devide set to subsets corresponding to testing, validation and training.

Train dataset: The actual dataset that we use to train the model.

Validation dataset: In this dataset we configure the classifier to obtain mataparameter values, which will improve our results.

Testing dataset: In this dataset we test the classifier estimating its future accuracy. It is only used once a model is completely trained(using the train and validation sets)

```
close all
Samples = 1000; % How many data samples there are?
DataDivision = 0.5; % How many data samples fall into which class?
v = 2; % v parameter of T Student's distribution
% Definition of data
for k = 1:Samples
if(rand()>DataDivision)
Ind = randi(Clusters.ClustersA);
DATA(2,k) = Clusters.ACoordinates(1,Ind)+random('T',v)*0.15;
 DATA(3,k) = Clusters.ACoordinates(2,Ind)+random('T',v)*0.15;
DATA(1,k) = 0;
 Ind = randi(Clusters.ClustersB);
DATA(2,k) = Clusters.BCoordinates(1,Ind)+random('T',v)*0.15;
 DATA(3,k) = Clusters.BCoordinates(2,Ind)+random('T',v)*0.15;
for k = 1:Samples
if(DATA(1,k) == 1)
 plot(DATA(2,k),DATA(3,k),'or'); hold on
plot(DATA(2,k),DATA(3,k),'ob'); hold on
end
xlabel('x');
ylabel('y');
vlim([-3 4])
```



The code above fills our clusters with data.

Worth mentioning is v value that determines how heavy are tails of the data distribution.

```
Indices = randperm(length(DATA1));

DATA_permutated = DATA1(:,Indices)

TR_number = ceil(length(DATA1)*0.5);

VA_number = ceil(length(DATA1)*0.25);

TE_number = ceil(length(DATA1)*0.25);

TE_number = ceil(length(DATA1)*0.25);

TR_DATA = DATA_permutated(:,1:TR_number);

VA_DATA = DATA_permutated(:,TR_number+VA_number);

TE_DATA = DATA_permutated(:,TR_number+VA_number);

save TR_DATA TR_DATA
save VA_DATA VA_DATA
save VA_DATA VA_DATA
```

The code above creates sets for training, validation and testing. In our case it would equal 500, 250 and 250 samples respectively.

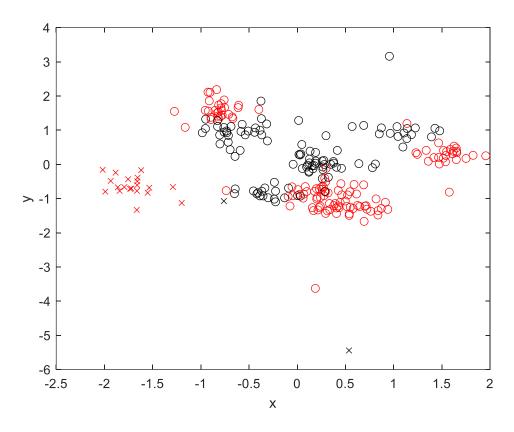
# Ex.3-6

A classifier in machine learning is an algorithm that automatically orders or categorizes data into one or more of a set of "classes." In this exercise we will desing simple linear classifier, even though our data is not linearly separable.

```
function [ClassLabel] = InitialClassifier(x,y,Parameters)
if(Parameters.W1*x + Parameters.W2*y + Parameters.B > 0)
ClassLabel = 1;
else
ClassLabel = 0;
end
end
```

#### Classification of our data:

```
load VA_DATA
Parameters.W1 = 1;
Parameters.W2 = 0.3;
Parameters.B = 1;
ErrorsA = 0;
ErrorsB = 0;
for k = 1:length(VA_DATA)
if(InitialClassifier(VA_DATA(2,k),VA_DATA(3,k),Parameters) == 1)
% Data point classified as A
if(VA_DATA(1,k) == 1)
% Data point classified correctly!
plot(VA_DATA(2,k),VA_DATA(3,k),'ok'); hold on
plot(VA_DATA(2,k),VA_DATA(3,k),'or'); hold on
ErrorsA = ErrorsA + 1;
end
% Data point classified as B
if(VA_DATA(1,k) == 0)
% Data point classified correctly!
plot(VA_DATA(2,k),VA_DATA(3,k),'xk'); hold on
plot(VA\_DATA(2,k),VA\_DATA(3,k),'xr'); hold on
ErrorsB = ErrorsB + 1;
 end
xlabel('x');
ylabel('y');
ErrorsA
ErrorsB
ErrorsA+ErrorsB
```



#### Results:

ErrorsA = 126

ErrorsB = 20

ans = 146

Obtained results are far form optimal. We have to optimize our algorithm, Using scripts developed in class 1 and 2 to optimize parameters of my linear classifier. I will use a grid search algorithm.

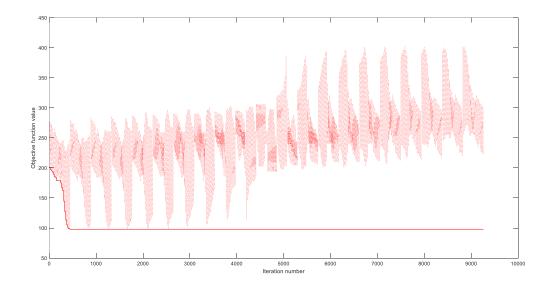
#### Grid code:

```
% A simple random optimization algorithm. It tries new locations until it
% runs out of time. Delay serves as a way of slowing FunctionPlot.
\% It requires a function for optimization (any function from folder
% "FunctionsForOptimization"
addpath FunctionsForOptimization
%% Optimization task:
FunctionForOptimization = str2func('zadanie3_TR');
%% Adjustable parameters:
MaxRangeX = [-10 \ 10]; % Range of parameters for optimization
MaxRangeY = [-10 10];
MaxRangeZ = [-10 10];
MaxSteps = 100;
                       % How many iterations do we perform?
FunctionPlot = 0;
                       % change to 0 If you want to get rid of the underlying function plot
                       \ensuremath{\mbox{\%}} Change to 0 if you want to get rid of the visualization
ConvergenceColor = 'r'; % Change color of the convergence curve here
%close all % Comment this out if you want to have many convergence curves plotted
```

```
ViewVect = [0,90];
                           % Initial viewpoint
Delay = 0.001;
                           % Inter-loop delay - to slow down the visualization
FunctionPlotQuality = 0.05; % Quality of function interpolation. Lower for a quicker run
%% Map initialization
InitialRangeX = MaxRangeX;
                            % This is the range from which we can draw points.
InitialRangeY = MaxRangeY;
InitialRangeZ = MaxRangeZ;
%% Map visualization (this code is not used for problem solving)
if(FunctionPlot == 1)
   figure(1);
   clf
        vectX = [MaxRangeX(1):FunctionPlotQuality:MaxRangeX(2)];
        vectY = [MaxRangeY(1):FunctionPlotQuality:MaxRangeY(2)];
       vectZ = [MaxRangeZ(1):FunctionPlotQuality:MaxRangeZ(2)];
        [X,Y,Z] = meshgrid(vectX,vectY,vectZ); indx = 1; indy = 1;
       for x = vectX
           indy = 1;
           for y = vectY
               Val(indx,indy) = FunctionForOptimization([x,y]);
               indy = indy + 1;
            end
            indx = indx + 1;
        mesh(X,Y,Val); surf(X,Y,Val,'LineStyle','none');
        view(ViewVect); colormap(bone); hold on
else end
%% Storing of a best solution
   CurrentMin = 50000;
   ResultX = 1;
    ResultY = 1;
    ResultZ = 1:
%% The main optimization loop
    EndingCondition = 0;
    iter = 0;
   tic;
    step x = 1;
    for NewX = MaxRangeX(1):step_x:MaxRangeX(2)
        for NewY = MaxRangeY(1):step_x:MaxRangeY(2)
           for NewZ = MaxRangeZ(1):step_x:MaxRangeZ(2)
\mbox{\ensuremath{\mbox{\$}}} If you'd like to provide function as a 2D image or use here any other objective function,
\% following line needs to be modified. The \theta passed to the function denotes the fact,
\ensuremath{\mathrm{\%}} that the function is constant in time.
        CurrentValue = FunctionForOptimization(NewX,NewY,NewZ);
        if(CurrentValue < CurrentMin)</pre>
           CurrentMin = CurrentValue; % Storing of a historically best result
            ResultX = NewX;
            ResultY = NewY;
            ResultZ = NewZ
           % FunctionPlot (green, if we have a new minimum):
           if(PointPlot == 1)
         figure(1); plot3(NewY, NewX, NewZ, CurrentValue, '.g'); hold on
```

```
end
   else
        \% FunctionPlot (red, if we don't have a new minimum):
       if(PointPlot == 1)
           figure(1); plot3(NewY, NewX, NewZ, CurrentValue, '.r'); hold on
   end
   \ensuremath{\mathrm{\%}} Command-window stuff for monitoring of algorithm's progress:
   SimTime = toc;
   clc
   fprintf('\nCurrent best: %f',CurrentMin);
   fprintf('\nCurrent:
                              %f',CurrentValue);
   fprintf('\n\n\n');
   fprintf('\nIteration: %d',iter);
   fprintf('\nTime:
                             %d',SimTime);
   BestHistory(iter) = CurrentMin;
                                          \% Here we store our historically best result
   CurrentHistory(iter) = CurrentValue;  % Here we store our currently investigated result
   \ensuremath{\mathrm{\%}} If we'd like to slow down the simulation - this line is where it
   % is done:
   pause(Delay);
end
    end
figure(2);
plot(BestHistory, 'Color', ConvergenceColor); hold on
\verb|plot(CurrentHistory, `Color', Convergence Color, `Line Style', `:'); | hold on |
xlabel('Iteration number');
ylabel('Objective function value');
```

# Grid search optimization:



### Code for genetic algorithm without stop condition:

```
\ensuremath{\mathrm{\%}} A simple random optimization algorithm. It tries new locations until it
% runs out of time. Delay serves as a way of slowing FunctionPlot.
% It requires a function for optimization (any function from folder
% "FunctionsForOptimization"
%% Optimization task:
FunctionForOptimization = str2func('zadanie6_TR');
%% Adjustable parameters:
InitialStep = 3; % Exploration/exploitation balance parameters:
P2 = 2:
lines=10;
P_size = 40; % Population size
n = 10: % Parameter n for n best succession
Step = 0.1; % Mutation range
MaxRangeX = [-10 \ 10]; \% Range of parameters for optimization
MaxRangeY = [-10 10];
MaxRangeZ = [-10 10];
MaxSteps = 20; % How many iterations do we perform?
FunctionPlot = 0; % change to 0 If you want to get rid of the underlying function plot
PointPlot = 0; % Change to 0 if you want to get rid of the visualization
ConvergenceColor = 'r'; % Change color of the convergence curve here
%close all % Comment this out if you want to have many convergence curves plotted
ViewVect = [0,90]; % Initial viewpoint
Delay =0.001; % Inter-loop delay - to slow down the visualization
\textbf{FunctionPlotQuality = 0.05; \% Quality of function interpolation. Lower for a quicker runchion of the property of the prope
%% Map initialization
InitialRangeX = MaxRangeX; % This is the range from which we can draw points.
InitialRangeY = MaxRangeY;
InitialRangeZ = MaxRangeZ;
\ensuremath{\text{\%\%}} Map visualization (this code is not used for problem solving)
TimePercent = 0:
if(FunctionPlot == 1)
  figure(1);
  vectX = [MaxRangeX(1):FunctionPlotQuality:MaxRangeX(2)];
  vectY = [MaxRangeY(1):FunctionPlotQuality:MaxRangeY(2)];
  [X,Y] = meshgrid(vectX,vectY); indx = 1; indy = 1;
  for x = vectX
  indy = 1;
  {\tt Val(indx,indy) = FunctionForOptimization([x,y]);}
  indy = indy + 1;
  indx = indx + 1;
  mesh(X,Y,Val); surf(X,Y,Val,'LineStyle','none');
  view(ViewVect); colormap(bone); hold on
%% Storing of a best solution
```

```
ResultX = 1:
 ResultY = 1;
%% The main optimization loop
 EndingCondition = 0;
 iter = 0;
  tic;
        for k = 1:P_size
                  Population(k).OF = Inf;
                  for line=1:lines
                  Population(k).Parameters.W1(line) = InitialRangeX(1) + ...
                  rand()*(InitialRangeX(2) - InitialRangeX(1));
                  Population(k).Parameters.W2(line) = InitialRangeY(1) + ...
                  rand()*(InitialRangeY(2) - InitialRangeY(1));
                  Population(k).Parameters.B(line) = InitialRangeY(1) + \dots
                  rand()*(InitialRangeY(2) - InitialRangeY(1));
        end
  while(EndingCondition == 0);
        iter = iter +1;
        Step(iter) = InitialStep * (1/(1+exp((iter-(MaxSteps/P1))/P2)));
        for k = 1:P_size
                        Population(k).OF = FunctionForOptimization(Population(k).Parameters.W1,Population(k).Parameters.W2,Population(k).Parameters.B); \\
        [~,Indices] = sortrows([Population(:).OF]');
        Population(Indices(1))
        if(FunctionPlot == 1)
               figure(1);
               clf
                surf(X,Y,Z,Val, 'LineStyle', 'none');
               view(ViewVect)
               colormap(bone)
                hold on
        else
        if(PointPlot == 1)
        for k = 1:1:P size
                \verb|plot3([Population(k).Parameters(2)], [Population(k).Parameters(1)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(2)]}, [Population(k).Parameters(1)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(2)]}, [Population(k).Parameters(1)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(2)]}, [Population(k).Parameters(3)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(3)]}, [Population(k).Parameters(3)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(3)]}, [Population(k).Parameters(3)], [Population(k).OF], '.r'); \\ \  \  \text{hold on the plot3} \\ \  \  \text{Population(k).Parameters(3)]}, [Population(k).Parameters(3)], [Population(k).Parameters(3)
        end
        BestHistory(iter) = Population(Indices(1)).OF;
        CurrentHistory(iter) = Population(Indices(floor(P_size/2))).0F;
        BestIndividualGenome(iter) = Population(Indices(1));
        NewPopulation(1) = Population(Indices(1));
        for k = 2:1:P_size
               ind1 = randi(n);
               ind2 = randi(n);
                NewPopulation(k) = Population(Indices(ind1));
              NewPopulation(k).Parameters.W1(2) = Population(Indices(ind2)).Parameters.W1(2);\\
                \label{eq:NewPopulation} NewPopulation(k). Parameters. W2(2) = Population(Indices(ind2)). Parameters. W2(2);
                  NewPopulation(k).Parameters.B(2) = Population(Indices(ind2)).Parameters.B(2);\\
                NewPopulation(k). Parameters. \\ \texttt{W1} = NewPopulation(k). Parameters. \\ \texttt{W1} + Step(iter)*randn(size(NewPopulation(k). Parameters. \\ \texttt{W1}));
                NewPopulation(k).Parameters.W2 = NewPopulation(k).Parameters.W2 + Step(iter)*randn(size(NewPopulation(k).Parameters.W2));\\
                NewPopulation(k).Parameters.B = NewPopulation(k).Parameters.B + Step(iter)*randn(size(NewPopulation(k).Parameters.B)); \\
                NewPopulation(k).OF = Inf;
```

```
Population = NewPopulation;
   clc
   fprintf('\nCurrent best: %f',BestHistory(end));
   fprintf('\nTime: %d',SimTime);
   if(iter > MaxSteps)
   EndingCondition = 1;
   else
   pause(Delay);
end
plot(BestHistory, 'Color', ConvergenceColor); hold on
plot(CurrentHistory, 'Color', ConvergenceColor, 'LineStyle', ':'); hold on
xlabel('Iteration number');
ylabel('Objective function value');
figure(4);
plot(Step)
xlabel('iteration');
ylabel('mutation step value');
```

If we want to obtain stop condition we have to include this fragment of code:

```
% with stop
testing(iter)=zadanie6_TE(Population(Indices(1)).Parameters.W1,Population(Indices(1)).Parameters.W2,Population(Indices(1)).Parameters.B);
[val,ind]=min(testing);
if(iter-ind>2)
    EndingCondition = 1;
    Population(Indices(1))=BestIndividualGenome(ind);
end
```

It works as following.

If the error value of the best individual is not improving in the testing set for longer than two iterations, we stop the training.

In the next exercises we repeated the procedure of training and validation and we obtained such results for corresponding algorithms.

	TR	VA
Grid	98	43
Genetic	97	44
3-line	81	41
Genetic		
5-line	83	41
Genetic		
7-line	42	19
Genetic		
10 line	5	11
genetic		
without		
stop		
condition		
10 line	8	6
genetic		
with stop		
condition		

As we can see if we increase the number of lines in genetic algorithm, we obtain better solution. Although when we decrease the size of dataset and increase the number of genetic lines without stoping condition we obtain worse solution than the one with stop condition. That is because, when we include stop condition we prevent overfiting, which is cricuial in obtaining satisfying solution.