## LABORATORY EXPERIENCE #1

# (Linear Regression)

#### Introduction

The first laboratory experience requests to perform **regression** algorithms in order to predict the grade of the **Parkinson disease** in a list of patients, starting from a dataset of measurements taken by the medical doctor.

The dataset that has been subject of study is public and can be downloaded at the following link https://archive.ics.uci.edu/ml/datasets/Parkinsons+Telemonitoring.

Each entry of the dataset is described by the following parameters, evaluated for 42 patients during a 6 months monitoring period:

- subject# Integer that uniquely identifies each subject
- age Subject age
- sex Subject gender '0' male, '1' female
- **test\_time** Time since recruitment into the trial. The integer part is the number of days since recruitment.
- motor\_UPDRS Clinician's motor UPDRS score, linearly interpolated
- total\_UPDRS Clinician's total UPDRS score, linearly interpolated
- Jitter (%) Several measures of variation in fundamental frequency
- Jitter (Abs)
- Jitter: RAPJitter: PP05
- Jittel. FFQ
- Jitter: DDP
- Shimmer Several measures of variation in amplitude
- Shimmer(dB)
- Shimmer: APQ3
- Shimmer: APQ5
- Shimmer: APO11
- Shimmer: DDA
- NHR
- HNR Two measures of ratio of noise to tonal components in the voice
- RPDE A nonlinear dynamical complexity measure
- DFA Signal fractal scaling exponent
- PPE A nonlinear measure of fundamental frequency variation

The parameter that indicates the grade of the Parkinson disease is the UPDRS, that stands for Unified Parkinson's Disease Rating Scale. In our case we will take into consideration the motor\_UPDRS parameter, that will be our regressand.

Regression will be performed through **Minimum Square Error (MSE) algorithm**, **gradient algorithm** and **steepest descent algorithm**.

### **Data preparation**

The first phase of the algorithm is dedicated to "clean" the input dataset, in order to ease the process of regression. In particular, for each patient, we aggregate the measurements taken

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# **Data preparation**

```
clear all
close all
load('updrs.mat')
A=parkinsonsupdrs;
num patients=max(A(:,1));
new_matrix=ones(1,22); % we define the matrix in which we insert
grouped values
for ind=1:num patients
    sub rows=find(A(:,1)==ind); % indexes of the matrix corresponding
to the 'ind' patient
   pat=A(sub_rows, :); % sub_matrix of 'ind' patient
    time=sort(abs(fix(pat(:,4)))); % we define an array of the days in
which measurements were taken for patient 'ind'
    % we had to take the integer part of that values and make the
 absolute
    % value because there were some negative time values. Then I
 ordered
    % them by increasing order
   time=unique(time); % we delete duplicates in the array of days
    for i=1:length(time) %we iterate within the days
        day=time(i);
        time rows=find(A(:,1)==ind & abs(fix(A(:,4)))==day); % we find
 the rows with all the measurements taken on patient 'ind' in the day
 'day'
        pat_time=A(time_rows, :); % we build a matrix of the
measurements of patient 'ind' on day 'day'
        pat_time(:,4)=abs(fix(pat_time(:,4))); % we adjust the time
 column in order to have integer day indexes
        mean row=mean(pat time,1);
        new matrix=[new matrix;mean row];
    end
end
pazienti=new matrix(2:end,:);
save('parkinson.mat','new_matrix');
```

```
%----- we build data train and data test matrices----
% data train=ones(1,22);
% data test=ones(1,22);
% for i=1:length(new matrix(:,1))
      if new matrix(i,1)<=36
용
          data train=[data train; new matrix(i,:)];
용
      else
          data test=[data test;new matrix(i,:)];
용
      end
% end
% data train=data train(2:end,:);
% data test=data test(2:end,:);
data train = pazienti(pazienti(:,1)<37,:);</pre>
data test = pazienti(pazienti(:,1)>36,:);
% ---- applying normalization to data train matrix
m data train=mean(data train,1);
v data train=var(data train,1);
s v data train=sqrt(v data train);
% data_train_norm=ones(length(data_train(:,1)),22);
% for i=1:length(data train(:,1))
      data train norm(i,1:4)=data train(i,1:4);
      for f=5:22
          data train norm(i,f)=(data train(i,f)-m data train(f))/
s v data train(f);
      end
% end
o = ones(size(data train,1),1);
data train norm = data train;
data_train_norm(:,5:end) = (data_train(:,5:end) -
 o*m data train(:,5:end)) ./ (o*s v data train(:,5:end));
mean train norm = mean(data train norm, 1); % verify that normalization
 is effective
var train norm = var(data train norm,1);
% ---- applying normalization to data test matrix
% data_test_norm=ones(length(data_test(:,1)),22);
% for i=1:length(data_test(:,1))
      data test norm(i,1:4)=data test(i,1:4);
      for f=5:22
          data test norm(i,f)=(data test(i,f)-m data train(f))/
s_v_data_train(f);
```

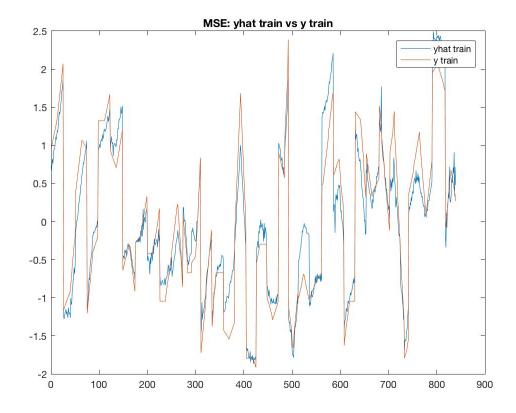
```
end
% end
o = ones(size(data test,1),1);
data_test_norm = data_test;
data_test_norm(:,5:end) = (data_test(:,5:end) -
 o*m data train(:,5:end)) ./ (o*s v data train(:,5:end));
mean test norm = mean(data train norm,1); % verify that normalization
 is effective
var_test_norm = var(data_train_norm,1);
% DEFINITION OF REGRESSOR AND REGRESSAND
F0=5; % we have to try with feature 7 and feature 5
y_train=data_train_norm(:,F0);
X train=data train norm;
X_train(:,F0)=[];
y_test=data_test_norm(:,F0);
X test=data test norm;
X_test(:,F0)=[];
```

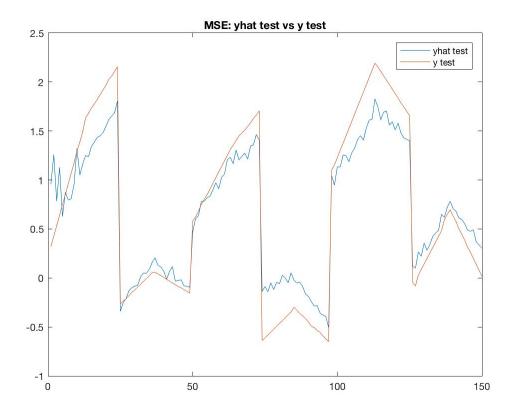
# Minimum Square Error (MSE)

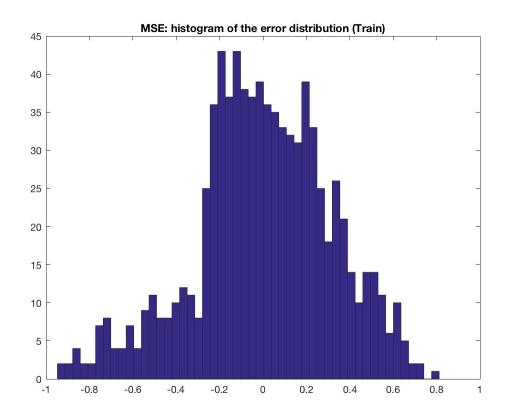
```
X_T=X_train(:,5:end).';
a hat=inv(X T*X train(:,5:end))*X T*y train;
yhat_train=X_train(:,5:end)*a_hat;
yhat_test=X_test(:,5:end)*a_hat;
error test=y test-yhat test;
error train=y train-yhat train;
figure
plot(yhat_train)
hold on
plot(y_train)
hold off
title('MSE: yhat train vs y train')
legend('yhat train','y train')
figure
plot(yhat test)
hold on
plot(y_test)
hold off
title('MSE: yhat test vs y test')
legend('yhat test','y test')
figure
```

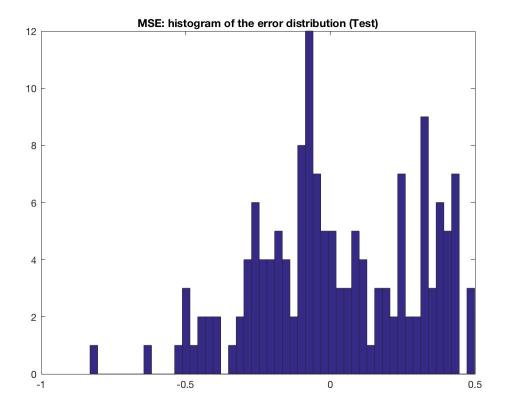
```
hist(error_train,50) % for the feature 5 the error is very variable,
  that means that feature 5 is less correlated to others
% for the feature 7 the error is distributed according something that
  is
% like a gaussian
title('MSE: histogram of the error distribution (Train)')

figure
hist(error_test,50) % for the feature 5 the error is very variable,
  that means that feature 5 is less correlated to others
% for the feature 7 the error is distributed according something that
  is
% like a gaussian
title('MSE: histogram of the error distribution (Test)')
```





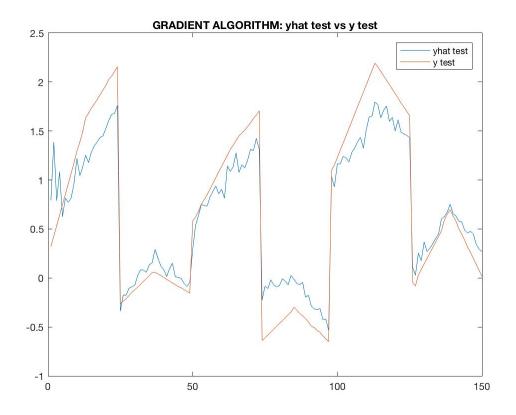


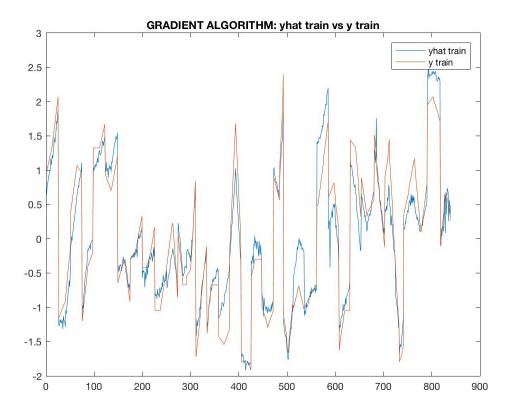


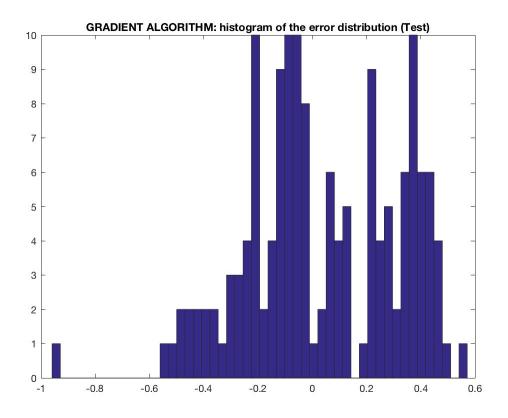
# **Gradient Algorithm**

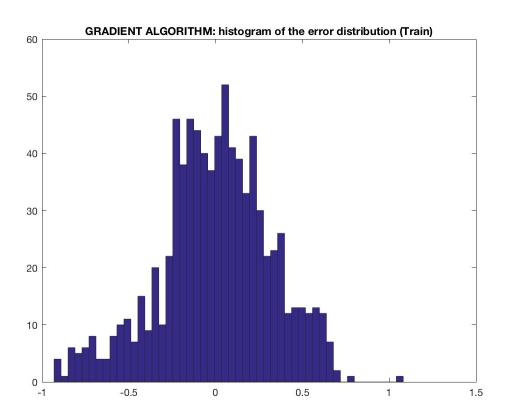
```
rng('default') % allows to have the same random vector each time we
 run the script
a i = rand(17,1); % generates a 21x1 matrix (a vector) of uniformly
 distributed random numbers
gamma=0.000001; % learning coefficient (usually 0.01/0.001/0.0001)
epsilon=0.00001; % stopping rule: a hat(i+1)-a hat(i)<epsilon
a old=zeros(17,1);
n=0;
a_story=a_i';
while (norm(a_i - a_old) > epsilon)
    grad= - 2 * X T * y train + 2 * X T * X train(:,5:end) * a i;
    a story=[a story;a i'];
    a old=a i;
    a i = a i - (gamma * grad);
    n=n+1;
end
yhat ga test=X test(:,5:end)*a i;
error_ga_test=y_test-yhat_ga_test;
yhat_ga_train=X_train(:,5:end)*a_i;
error ga train=y train-yhat ga train;
```

```
figure
plot(yhat_ga_test)
hold on
plot(y_test)
hold off
title('GRADIENT ALGORITHM: yhat test vs y test')
legend('yhat test','y test')
figure
plot(yhat_ga_train)
hold on
plot(y_train)
hold off
title('GRADIENT ALGORITHM: yhat train vs y train')
legend('yhat train','y train')
figure
hist(error_ga_test,50)
title('GRADIENT ALGORITHM: histogram of the error distribution
 (Test)')
figure
hist(error_ga_train,50)
title('GRADIENT ALGORITHM: histogram of the error distribution
 (Train)')
```





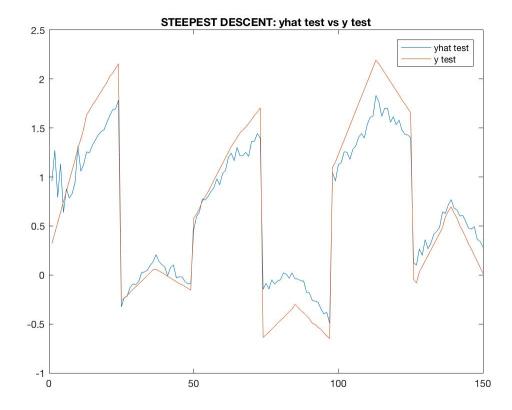


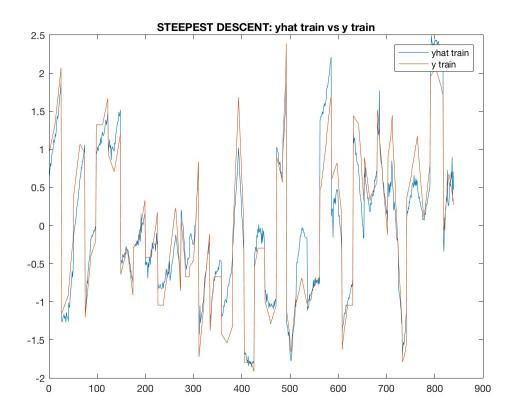


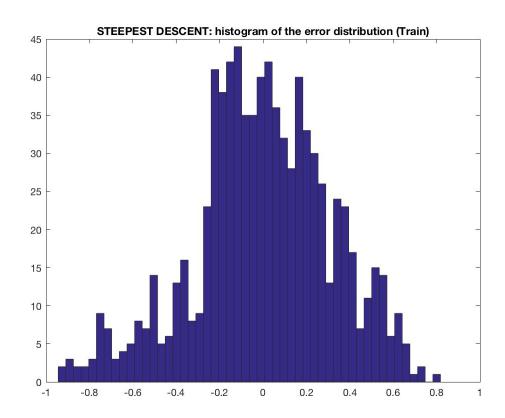
## **Steepest Descent Algorithm**

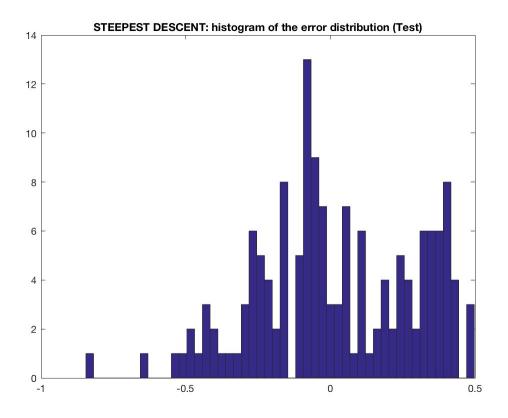
```
rng('default') % allows to have the same random vector each time we
 run the script
a i sd = rand(17,1); % generates a 21x1 matrix (a vector) of uniformly
 distributed random numbers
gamma=0.000001; % learning coefficient (usually 0.01/0.001/0.0001)
epsilon=0.00001; % stopping rule: a_hat(i+1)-a_hat(i)<epsilon
a old sd=zeros(17,1);
n=0;
a_story=a_i';
while (norm(a i sd - a old sd) > epsilon)
    grad sd= - 2 * X T * y train + 2 * X T * X train(:,5:end) *
 a i sd;
    hessian = 4 * X T * X train(:,5:end);
    a old sd = a i sd;
    a i sd = a i sd - ((norm(grad sd)^2 * grad sd)/(grad sd.' *
 hessian * grad sd));
end
yhat_sd_train=X_train(:,5:end)*a_i_sd;
error sd train=y train-yhat sd train;
yhat sd test=X test(:,5:end)*a i sd;
error_sd_test=y_test-yhat_sd test;
figure
plot(yhat sd test)
hold on
plot(y_test)
hold off
title('STEEPEST DESCENT: yhat test vs y test')
legend('yhat test','y test')
figure
plot(yhat_sd_train)
hold on
plot(y train)
hold off
title('STEEPEST DESCENT: yhat train vs y train')
legend('yhat train','y train')
figure
hist(error sd train,50)
title('STEEPEST DESCENT: histogram of the error distribution (Train)')
figure
hist(error sd test,50)
title('STEEPEST DESCENT: histogram of the error distribution (Test)')
```

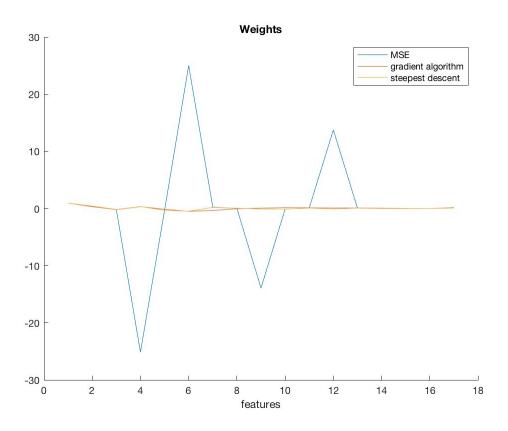
```
figure
hold on
plot(a_hat)
plot(a_i)
plot(a_i_sd)
hold off
title('Weights')
xlabel('features')
legend('MSE', 'gradient algorithm', 'steepest descent')
```













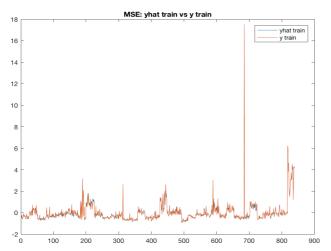
within each day evaluating the mean of the entries corresponding to that day. As some test\_time values were wrongly registered as negative numbers, the absolute value of that number was considered. Then, data is normalized and divide in two matrices: the first, containing the measurement of the first 36 patients, will be used as regressor in the training phase of the algorithms, while the second, corresponding to the patients from 37 to 42 will be used in the test phase.

## **Algorithms implementation and results**

The idea of linear regression starts from the hypothesis that we can obtain our regressand vector  $\mathbf{y}$  is a linear combination of our regressor matrix  $\mathbf{X}$ , properly weighted (coefficients vector  $\mathbf{a}$ ), plus a measurement error. The scope of the linear regression algorithm is to find the weights that better describe this relation between regressor and regressand.

**MSE algorithm** simply finds those weights as the values that minimizes the square error  $e(a) = \|y - Xa\|^2$  finding the values of a for which the gradient of e(a) is zero. This method allows to evaluate the unknown through a closed form expression, but requires the inversion of a matrix, that might be too complex in some cases. So, in those cases might be useful to perform iterative algorithms like the **gradient algorithm**, in which, starting from an initial guess of the weights, computes step by step the gradient of the error function and tries to "follow" the direction towards the minimum until a stopping function is reached. This algorithm requires the definition of a learning coefficient and of the stopping rule. **Steepest descent algorithm** allows to reach a faster convergence finding at each step the optimum value of the learning coefficient, instead of defining it a priori.

We are able to obtain in a similar way good performance for each of the three algorithms, especially if we try to predict feature 7 (Jitter (%)). Plotting the predicted values against the true ones we can see that all the algorithms are working quite well, both in the training phase and learning phase. Actually, plotting the histograms of the error distribution is it possible to observe a greater dispersion of the error in the test phase, as we could expect. The confrontation of the weights shows that the iterative algorithms find a different solution to the MSE algorithm, as well as the overall performance of the regression remains very good. Performing regression with feature 5 (motor\_UPDRS) we can see a slight decrease of the performances, expecially in the test phase. This could be due to the fact that Jitter is more related to the other features rather then the motor\_UPDRS, as the first one is, in line the other features, related to speech impairments, while the second one is dealing with motor impairments, and It is evaluated at discretion of the medical doctor.



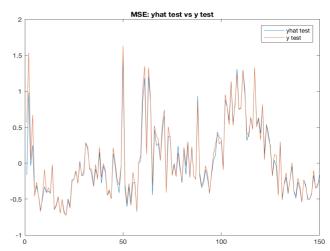


Figure 1: Regression results vs. real values in train and test phase - MSE feature 7

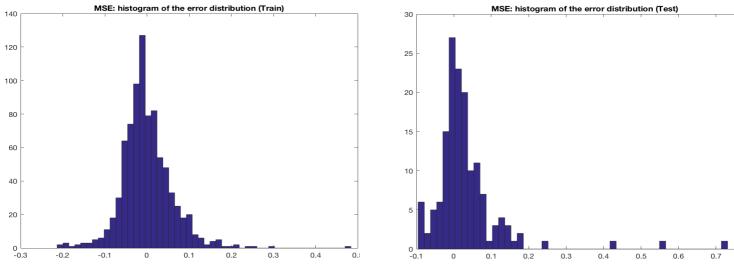


Figure 2: Comparison of error distribution in train and test phase - MSE feature 7

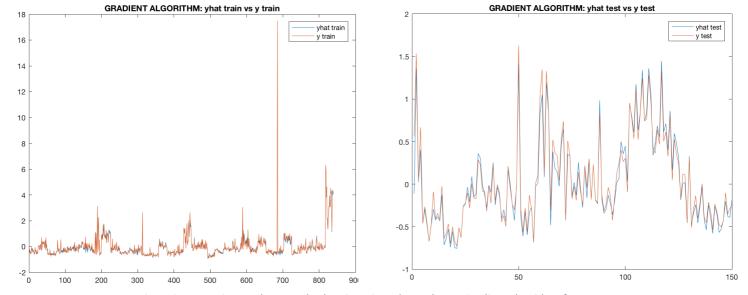


Figure 3: Regression results vs. real values in train and test phase - Gradient algorithm, feature 7

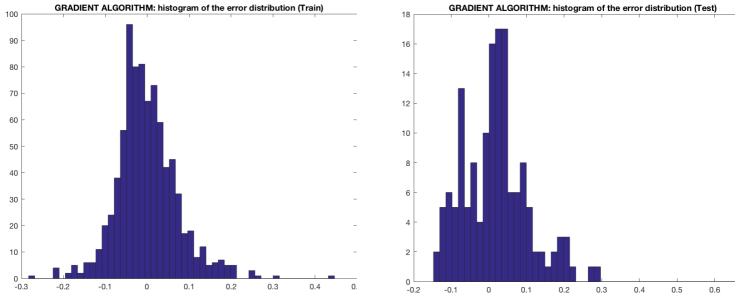
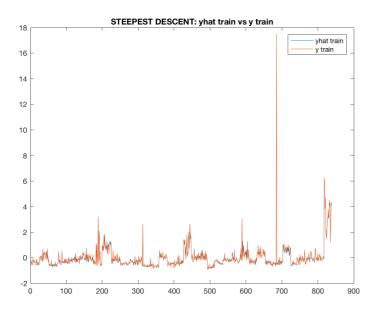


Figure 4: Comparison of error distribution in train and test phase - Gradient algorithm, feature 7



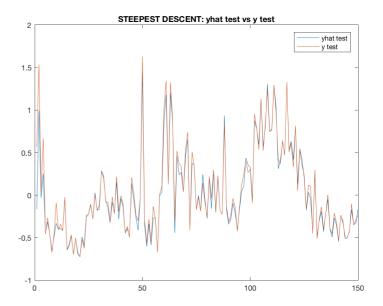
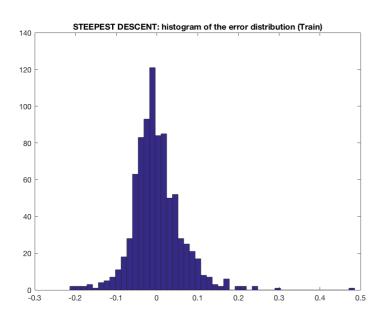


Figure 5: Regression results in train and test phase - Steepest descent, feature 7



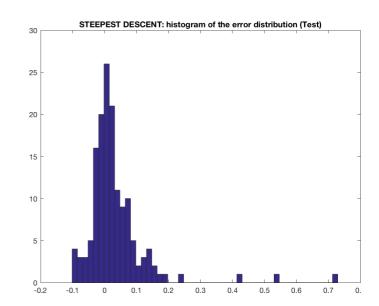
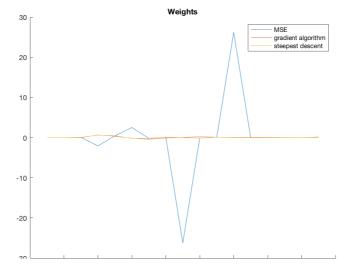


Figure 6: Comparison of error distribution in train and test phase - Steepest descent, feature 7



Pictures relative to regression with feature 5 are present in the publish of the Matlab scripts of Lab 01.

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# **Data preparation**

```
clear all
close all
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A=parkinsonsupdrs;
num patients=max(A(:,1));
new_matrix=ones(1,22); % we define the matrix in which we insert
grouped values
for ind=1:num patients
    sub rows=find(A(:,1)==ind); % indexes of the matrix corresponding
to the 'ind' patient
   pat=A(sub rows, :); % sub matrix of 'ind' patient
    time=sort(abs(fix(pat(:,4)))); % we define an array of the days in
which measurements were taken for patient 'ind'
    % we had to take the integer part of that values and make the
 absolute
    % value because there were some negative time values. Then I
 ordered
    % them by increasing order
   time=unique(time); % we delete duplicates in the array of days
    for i=1:length(time) %we iterate within the days
        day=time(i);
        time rows=find(A(:,1)==ind & abs(fix(A(:,4)))==day); % we find
 the rows with all the measurements taken on patient 'ind' in the day
 'day'
        pat_time=A(time_rows, :); % we build a matrix of the
measurements of patient 'ind' on day 'day'
        pat_time(:,4)=abs(fix(pat_time(:,4))); % we adjust the time
 column in order to have integer day indexes
        mean row=mean(pat time,1);
        new matrix=[new matrix;mean row];
    end
end
pazienti=new matrix(2:end,:);
save('parkinson.mat','new_matrix');
```

```
%----- we build data train and data test matrices----
% data train=ones(1,22);
% data test=ones(1,22);
% for i=1:length(new matrix(:,1))
      if new matrix(i,1)<=36
용
          data train=[data train; new matrix(i,:)];
용
      else
          data test=[data test;new matrix(i,:)];
용
      end
% end
% data train=data train(2:end,:);
% data test=data test(2:end,:);
data train = pazienti(pazienti(:,1)<37,:);</pre>
data test = pazienti(pazienti(:,1)>36,:);
% ---- applying normalization to data train matrix
m data train=mean(data train,1);
v data train=var(data train,1);
s v data train=sqrt(v data train);
% data_train_norm=ones(length(data_train(:,1)),22);
% for i=1:length(data train(:,1))
      data train norm(i,1:4)=data train(i,1:4);
      for f=5:22
          data train norm(i,f)=(data train(i,f)-m data train(f))/
s v data train(f);
      end
% end
o = ones(size(data train,1),1);
data train norm = data train;
data_train_norm(:,5:end) = (data_train(:,5:end) -
 o*m data train(:,5:end)) ./ (o*s v data train(:,5:end));
mean train norm = mean(data train norm, 1); % verify that normalization
 is effective
var train norm = var(data train norm,1);
% ---- applying normalization to data test matrix
% data_test_norm=ones(length(data_test(:,1)),22);
% for i=1:length(data_test(:,1))
      data test norm(i,1:4)=data test(i,1:4);
      for f=5:22
          data test norm(i,f)=(data test(i,f)-m data train(f))/
s_v_data_train(f);
```

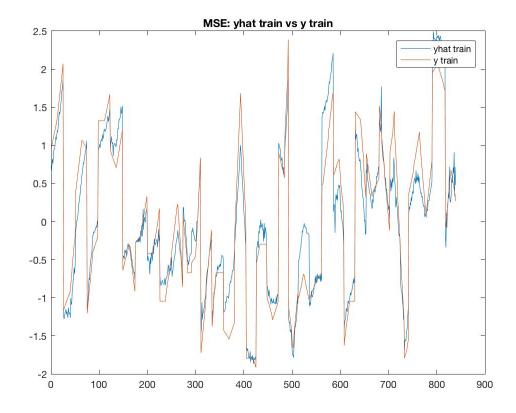
```
end
% end
o = ones(size(data test,1),1);
data_test_norm = data_test;
data_test_norm(:,5:end) = (data_test(:,5:end) -
 o*m data train(:,5:end)) ./ (o*s v data train(:,5:end));
mean test norm = mean(data train norm,1); % verify that normalization
 is effective
var_test_norm = var(data_train_norm,1);
% DEFINITION OF REGRESSOR AND REGRESSAND
F0=5; % we have to try with feature 7 and feature 5
y_train=data_train_norm(:,F0);
X train=data train norm;
X_train(:,F0)=[];
y_test=data_test_norm(:,F0);
X test=data test norm;
X_test(:,F0)=[];
```

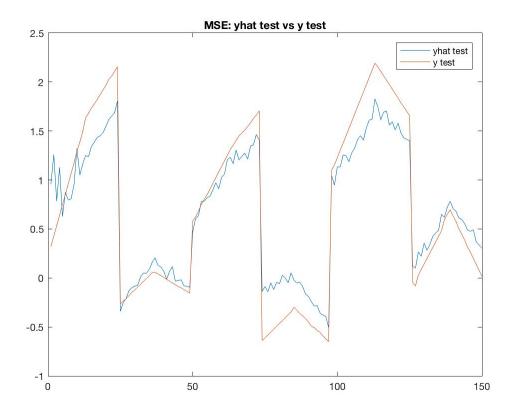
# Minimum Square Error (MSE)

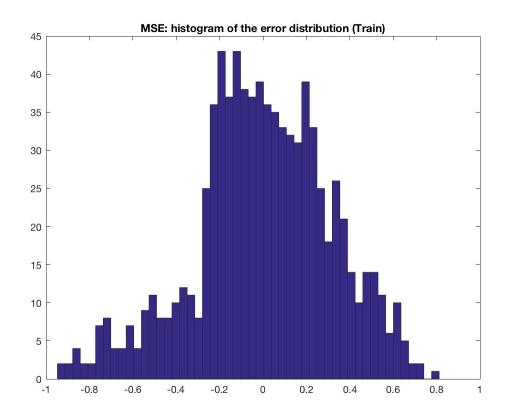
```
X_T=X_train(:,5:end).';
a hat=inv(X T*X train(:,5:end))*X T*y train;
yhat_train=X_train(:,5:end)*a_hat;
yhat_test=X_test(:,5:end)*a_hat;
error test=y test-yhat test;
error train=y train-yhat train;
figure
plot(yhat_train)
hold on
plot(y_train)
hold off
title('MSE: yhat train vs y train')
legend('yhat train','y train')
figure
plot(yhat test)
hold on
plot(y_test)
hold off
title('MSE: yhat test vs y test')
legend('yhat test','y test')
figure
```

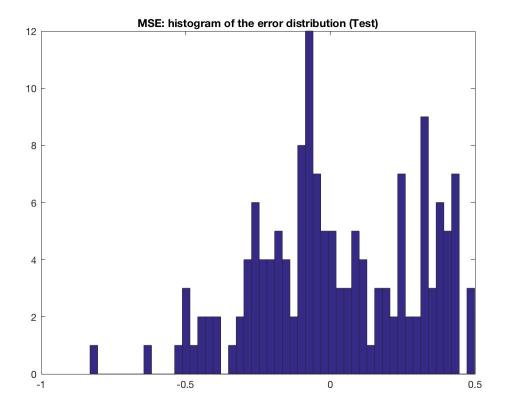
```
hist(error_train,50) % for the feature 5 the error is very variable,
  that means that feature 5 is less correlated to others
% for the feature 7 the error is distributed according something that
  is
% like a gaussian
title('MSE: histogram of the error distribution (Train)')

figure
hist(error_test,50) % for the feature 5 the error is very variable,
  that means that feature 5 is less correlated to others
% for the feature 7 the error is distributed according something that
  is
% like a gaussian
title('MSE: histogram of the error distribution (Test)')
```





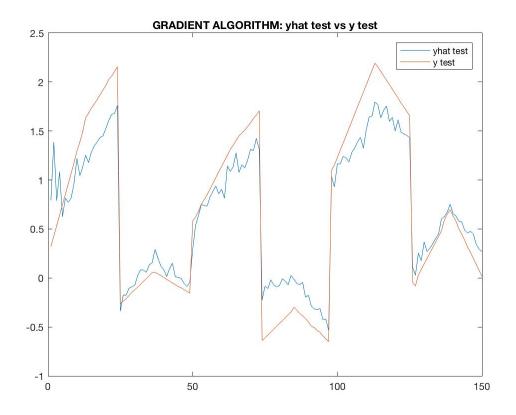


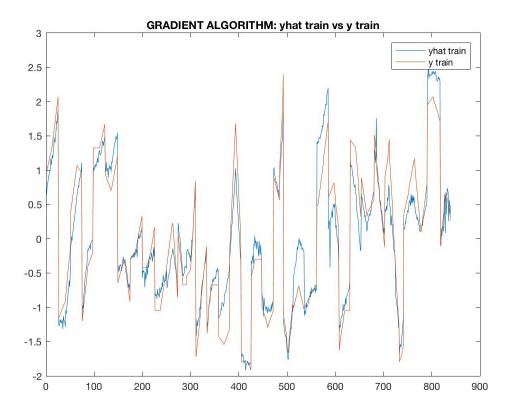


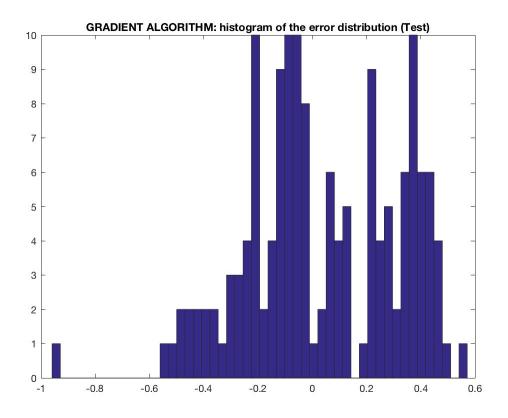
# **Gradient Algorithm**

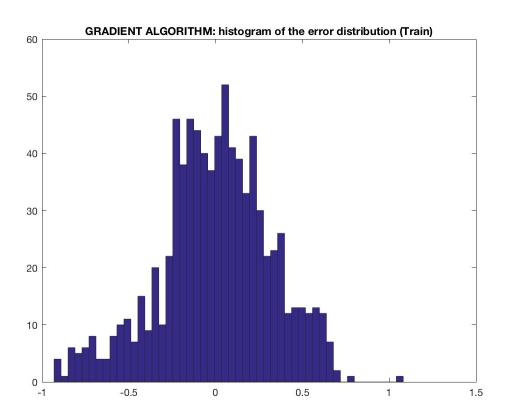
```
rng('default') % allows to have the same random vector each time we
 run the script
a i = rand(17,1); % generates a 21x1 matrix (a vector) of uniformly
 distributed random numbers
gamma=0.000001; % learning coefficient (usually 0.01/0.001/0.0001)
epsilon=0.00001; % stopping rule: a hat(i+1)-a hat(i)<epsilon
a old=zeros(17,1);
n=0;
a_story=a_i';
while (norm(a_i - a_old) > epsilon)
    grad= - 2 * X T * y train + 2 * X T * X train(:,5:end) * a i;
    a story=[a story;a i'];
    a old=a i;
    a i = a i - (gamma * grad);
    n=n+1;
end
yhat ga test=X test(:,5:end)*a i;
error_ga_test=y_test-yhat_ga_test;
yhat_ga_train=X_train(:,5:end)*a_i;
error ga train=y train-yhat ga train;
```

```
figure
plot(yhat_ga_test)
hold on
plot(y_test)
hold off
title('GRADIENT ALGORITHM: yhat test vs y test')
legend('yhat test','y test')
figure
plot(yhat_ga_train)
hold on
plot(y_train)
hold off
title('GRADIENT ALGORITHM: yhat train vs y train')
legend('yhat train','y train')
figure
hist(error_ga_test,50)
title('GRADIENT ALGORITHM: histogram of the error distribution
 (Test)')
figure
hist(error_ga_train,50)
title('GRADIENT ALGORITHM: histogram of the error distribution
 (Train)')
```









## **Steepest Descent Algorithm**

```
rng('default') % allows to have the same random vector each time we
 run the script
a i sd = rand(17,1); % generates a 21x1 matrix (a vector) of uniformly
 distributed random numbers
gamma=0.000001; % learning coefficient (usually 0.01/0.001/0.0001)
epsilon=0.00001; % stopping rule: a_hat(i+1)-a_hat(i)<epsilon
a old sd=zeros(17,1);
n=0;
a_story=a_i';
while (norm(a i sd - a old sd) > epsilon)
    grad sd= - 2 * X T * y train + 2 * X T * X train(:,5:end) *
 a i sd;
    hessian = 4 * X T * X train(:,5:end);
    a old sd = a i sd;
    a i sd = a i sd - ((norm(grad sd)^2 * grad sd)/(grad sd.' *
 hessian * grad sd));
end
yhat_sd_train=X_train(:,5:end)*a_i_sd;
error sd train=y train-yhat sd train;
yhat sd test=X test(:,5:end)*a i sd;
error_sd_test=y_test-yhat_sd test;
figure
plot(yhat sd test)
hold on
plot(y_test)
hold off
title('STEEPEST DESCENT: yhat test vs y test')
legend('yhat test','y test')
figure
plot(yhat_sd_train)
hold on
plot(y train)
hold off
title('STEEPEST DESCENT: yhat train vs y train')
legend('yhat train','y train')
figure
hist(error sd train,50)
title('STEEPEST DESCENT: histogram of the error distribution (Train)')
figure
hist(error sd test,50)
title('STEEPEST DESCENT: histogram of the error distribution (Test)')
```

```
figure
hold on
plot(a_hat)
plot(a_i)
plot(a_i_sd)
hold off
title('Weights')
xlabel('features')
legend('MSE', 'gradient algorithm', 'steepest descent')
```

