## **ABSTRACT**

Data Analysis for P300 data set including spatial averaging of parietal channels

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## First extract single trials after stimulus and Plot

In order to extract single trials after stimulus following steps were used:

- 1. Loading data set into workspace
- 2. Selecting channel 5 and Reducing Data dimension from 85x7794x64 in variable Signal to 26x1 using squeeze (Signal (1, 5, 39:end, 1)) %26 points from Subject A, channel 5, trial 3
- 3. Saving the reduced data in variable SingleStimulus
- 4. Plotting the signal
- 5. Plotting square Signal
- 6. Repeat steps 1-5 for each data set train/test
- 7. Repeat steps 1-5 for both subjects A and B

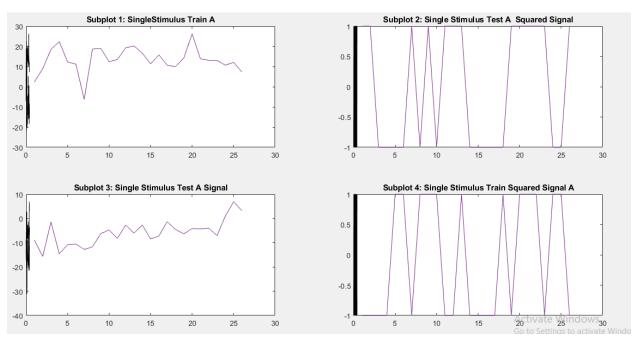
### Matlab Commands Subject A:

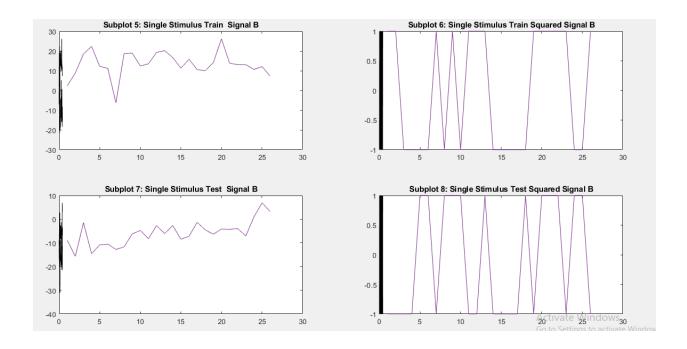
```
load('Subject A Train.mat')
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, channel 5, trial 3
singleStimulus Train A = squeeze(Signal(1,5,39:end,1))
N=length(singleStimulus Train A); % Get length of average per 5
t=(0:N-1)/frequency s; % Generating time vector
subplot(2,2,1)% Generating SingleStimulus Train A Plot
plot(t, singleStimulus Train A, 'k'); hold on;
title('Subplot 1: SingleStimulus Train A')
subplot(2,2,2)% Generating Single Stimulus Test A Squared Signal
Plot
plot(t, square(singleStimulus Train A), 'k'); hold on;
title ('Subplot 2: Single Stimulus Test A Squared Signal')
load('Subject A Test.mat')
singleStimulus Test A = squeeze(Signal(1,5,39:end,1))
N1=length(singleStimulus Test A); % Get length
t1=(0:N1-1)/frequency s; % Generating time vector
subplot(2,2,3)
plot(t1, singleStimulus Test A, 'k'); hold on;
title ('Subplot 3: Single Stimulus Test A Signal')
subplot(2,2,4)
plot(t1, square(singleStimulus Test A), 'k'); hold on;
title ('Subplot 4: Single Stimulus Train Squared Signal A')
```

## Matlab Commands Subject B:

```
load('Subject B Train.mat')
singleStimulus Train B = squeeze(Signal(1,5,39:end,1))
N2=length(singleStimulus Train B); % Get length
t2=(0:N2-1)/frequency s; % Generating time vector
subplot(2,2,1)
plot(t2, singleStimulus Train B, 'k'); hold on;
title('Subplot 5: Single Stimulus Train Signal B')
subplot(2,2,2)
plot(t2, square(singleStimulus Train B), 'k'); hold on;
title ('Subplot 6: Single Stimulus Train Squared Signal B')
load('Subject B Test.mat')
singleStimulus Test B = squeeze(Signal(1,5,39:end,1))
N3=length(singleStimulus Test B); % Get length
t3=(0:N3-1)/frequency s; % Generating time vector
subplot(2,2,3)
plot(t2, singleStimulus Test B, 'k'); hold on;
title('Subplot 7: Single Stimulus Test Signal B')
subplot(2,2,4)
plot(t2, square(singleStimulus Test B), 'k'); hold on;
title('Subplot 8: Single Stimulus Test Squared Signal B')
```

## Plots Subject A and Subject B





## 2. Take averaging of 5 trails and plot

In order to take average of 5 trials and plot following steps were used:

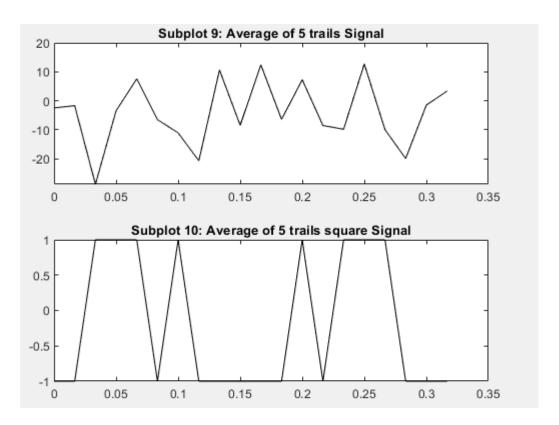
- 1. Loading data set into workspace
- 2. Selecting 5 trails by using values 1-5 and taking their mean, next using next 5 values and taking their mean and so on using reshape (column2(1:100), [], 5)
- 3. Saving the reduced data in variable out
- 4. Taking mean(out, 2)
- 5. Plotting the signal
- 6. Plotting square Signal
- 7. Repeat steps 1-4 for each data set train/test
- 8. Repeat steps 1-5 for both subjects A and B

## Matlab Commands:

```
column2 = Signal;
out = reshape(column2(1:100), [], 5)
N=length(out); % Get length of average per 5 trails
frequency_s=60; % Setting the frequency to 60 Hz

t=(0:N-1)/frequency_s; % Generating time vector
means5 = mean(out, 2)
subplot(2,1,1)
plot(t,means5,'k');hold on;
title('Subplot 9: Average of 5 trails Signal')
```

**Plots** 



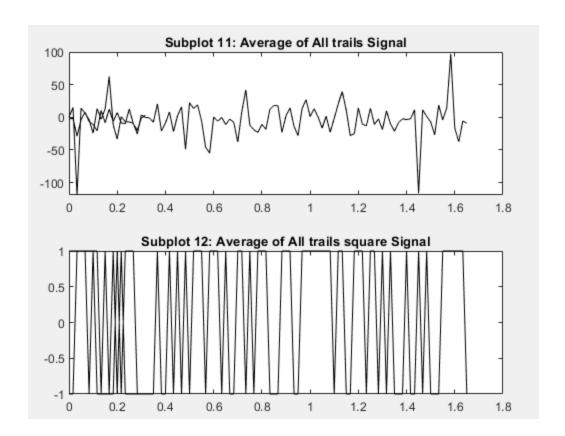
# 3. Take averaging to all trials from single channel and plot

### Matlab Commands:

```
column2 = Signal;
out = reshape(column2(1:100), [], 1)
N=length(out); % Get length of average per 5 trails
frequency_s=60; % Setting the frequency to 60 Hz

t=(0:N-1)/frequency_s; % Generating time vector
means5 = mean(out, 2)
subplot(2,1,1)
plot(t,means5,'k');hold on;
title('Subplot 11: Average of All trails Signal')
```

### **Plots**

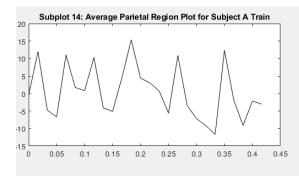


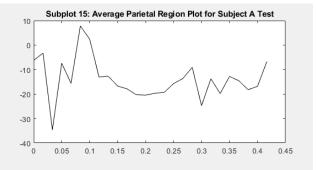
## Average the parietal region channels and find the spatial averaging of all trials.

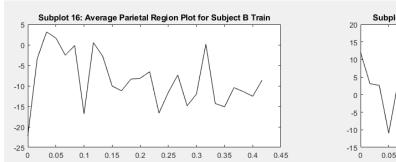
```
load('Subject A Train.mat')
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, channel 5, trial 3
48, 49, 53, 54
singleStimulus Train A P5 =
squeeze(Signal(1,48,39:end,1))
singleStimulus Train A P3 =
squeeze(Signal(1,49,39:end,1))
singleStimulus Train A P4 =
squeeze(Signal(1,53,39:end,1))
singleStimulus Train A P6 =
squeeze(Signal(1,54,39:end,1))
N=length(singleStimulus Train A P5); % Get length
t=(0:N-1)/frequency s; % Generating time vector
singleStimulus Train A parietal avg=(singleStimulus Tr
ain A P5+singleStimulus Train A P3+singleStimulus Trai
n_A_P4+singleStimulus_Train A_P6)/4
subplot(2,2,1)% Generating SingleStimulus Train A Plot
plot(t, singleStimulus Train A parietal avg, 'k'); hold
title ('Subplot 14: Average Parietal Region Plot for
Subject A Train')
load('Subject A Test.mat')
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, channel 5, trial 3
48, 49, 53, 54
singleStimulus Test A P5 =
squeeze(Signal(1,48,39:end,1))
singleStimulus Test A P3 =
squeeze (Signal (1, 49, 39:end, 1))
singleStimulus Test A P4 =
squeeze(Signal(1,53,39:end,1))
singleStimulus Test A P6 =
squeeze (Signal (1,54,39:end,1))
N=length(singleStimulus Test A P5); % Get length
t=(0:N-1)/frequency s; % Generating time vector
singleStimulus Test A parietal avg=(singleStimulus Tes
t_A_P5+singleStimulus_Test_A_P3+singleStimulus_Test_A_
P4+singleStimulus Test A P6)/4
subplot(2,2,2)% Generating SingleStimulus Train A Plot
plot(t, singleStimulus Test A parietal avg, 'k'); hold
title('Subplot 15: Average Parietal Region Plot for
Subject A Test')
```

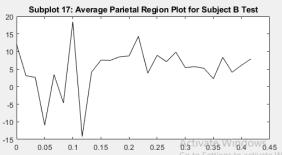
```
load('Subject B Train.mat')
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, channel 5, trial 3 48,49,53,54
singleStimulus Train B P5 = squeeze(Signal(1,48,39:end,1))
singleStimulus Train B P3 = squeeze(Signal(1,49,39:end,1))
singleStimulus Train B P4 = squeeze(Signal(1,53,39:end,1))
singleStimulus Train B P6 = squeeze(Signal(1,54,39:end,1))
N=length(singleStimulus Train B P5); % Get length
t=(0:N-1)/frequency s; % Generating time vector
singleStimulus Train B parietal avg=(singleStimulus Train B
P5+singleStimulus Train B P3+singleStimulus Train B P4+sin
gleStimulus Train B P6)/4
subplot(2,2,3)% Generating SingleStimulus Train A Plot
plot(t, singleStimulus Train B parietal avg, 'k'); hold on;
title ('Subplot 16: Average Parietal Region Plot for Subject
B Train')
load('Subject B Test.mat')
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, channel 5, trial 3 48,49,53,54
singleStimulus Test B P5 = squeeze(Signal(1,48,39:end,1))
singleStimulus Test B P3 = squeeze(Signal(1,49,39:end,1))
singleStimulus Test B P4 = squeeze(Signal(1,53,39:end,1))
singleStimulus Test B P6 = squeeze(Signal(1,54,39:end,1))
N=length(singleStimulus Test B P5); % Get length
t=(0:N-1)/frequency s; % Generating time vector
singleStimulus Test B parietal avg=(singleStimulus Test B P
5+singleStimulus Test B P3+singleStimulus Test B P4+singleS
timulus Test B P6)/4;
subplot(2,2,4)% Generating SingleStimulus Train A Plot
plot(t, singleStimulus_Test_B_parietal_avg, 'k'); hold on;
title('Subplot 17: Average Parietal Region Plot for Subject
B Test')
```

### Plots:





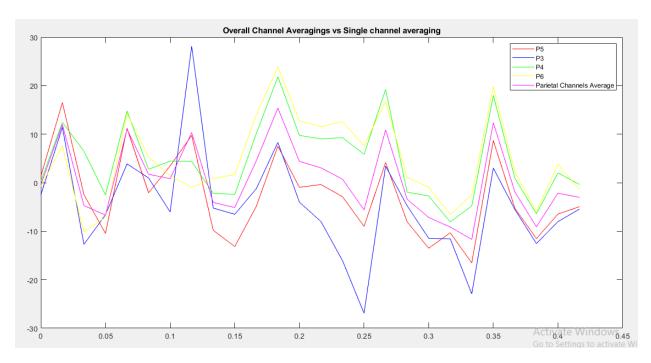




## Find if single channel averaging is better than averaging over channels.

#### Matlab Code

```
load('Subject A Train.mat')
x = linspace(0, 10, 50);
frequency s=60; % Setting the frequency to 60 Hz
% points from Subject A, Parietal channels 48,49,53,54
singleStimulus Train A P5 = squeeze(Signal(1,48,39:end,1))
singleStimulus Train A P3 = squeeze(Signal(1,49,39:end,1))
singleStimulus Train A P4 = squeeze(Signal(1,53,39:end,1))
singleStimulus Train A P6 = squeeze(Signal(1,54,39:end,1))
N=length(singleStimulus Train A P5); % Get length
t=(0:N-1)/frequency s; % Generating time vector
% Sp Averaging
singleStimulus Train A parietal avg=(singleStimulus Train A P
5+singleStimulus Train A P3+singleStimulus Train A P4+singleS
timulus Train A \overline{P6})/4
% Comparison Plot
plot(t, singleStimulus Train A P5, 'c'); hold on;
title('Overall Channel Averagings vs Single channel
averaging')
hold on
h1=plot(t,singleStimulus Train_A_P5,'r');hold on;
h2=plot(t, singleStimulus Train A P3, 'b'); hold on;
h3=plot(t,singleStimulus Train_A_P4,'g');hold on;
h4=plot(t, singleStimulus Train A P6, 'y'); hold on;
h5=plot(t, singleStimulus Train A parietal avg, 'm'); hold on;
legend([h1 h2 h3 h4 h5], 'P5', 'P3', 'P4', 'P6', 'Parietal
```



From the plot, the performance of the averaging over channels is significantly higher in comparison to individual channels. Standard deviation is lesser from the axis in cases of later

```
load('Subject_A_Train.mat')
% points from Subject A, channel 5, trial 3 48,49,53,54
singleStimulus_Train_A_P5 = squeeze(Signal(1,48,39:end,1))
singleStimulus_Train_A_P3 = squeeze(Signal(1,49,39:end,1))
singleStimulus_Train_A_P4 = squeeze(Signal(1,53,39:end,1))
singleStimulus_Train_A_P6 = squeeze(Signal(1,54,39:end,1))

% Standard deviation of individual vs average of parietal channels

Standard_Deviation_P5 = std(singleStimulus_Train_A_P5)
Standard_Deviation_P3 = std(singleStimulus_Train_A_P3)
Standard_Deviation_P4 = std(singleStimulus_Train_A_P4)
Standard_Deviation_P6 = std(singleStimulus_Train_A_P6)
Standard_Deviation_Average = std(singleStimulus_Train_A_Parietal_avg)
```

Standard Deviation P5 = 8.5473

Standard\_Deviation\_P3 =10.8791

Standard\_Deviation\_P4 = 8.0947

Standard\_Deviation\_P6 = 8.8262

Standard Deviation Average = 7.6199