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# 540.305 Problem Set 1: Vectors, Arrays, Data and Basic Plotting - Diego Alba

```
% Script will ask for some input and display all answers in the command window
clear
clc
addpath('functions','resources','figures')
```

## 1 Basic vector and array practice

```
disp('Basic vector and array practice')
% a
v = [54/(3+(4.2)^2),32,(6.3)^2-(7.2)^2,54,exp(3.7),
      sin(deg2rad(66))+cos(3*pi/8)];
disp(['1, (a) : ',num2str(v),']')

% b
x = 0.85;
y = 12.5;
w = [y,y^x,log(y/x),y+x,y+x]';
disp('1, (b) : ')
disp(num2str(w))
disp('                ]')

% c
a = repmat(7.5,1,9)-[repmat(7.5,1,8),0];
disp(['1, (c) : a = ',num2str(a),']')

% d
B = zeros(5,3);
B(:,1) = 1:5;
B(:,3) = repmat(3,1,5);
disp('1, (d) : B = ')
disp(num2str(B))
disp('                ]')

% e
```

---

```

N = [33 21 9 14 30; 30 18 6 18 34; 27 15 6 22 38; 24 12 10 26 42];
disp('1, (e) : A = [')
disp(num2str([N(1,1:4)', N(2,2:5)' ]))
disp('           ]')
disp(['1, (e) : B = [,num2str([N(:,3)' N(3,:)])],']'])
disp("1, (e) : C = Error: Expression or statement is incorrect--"
possibly unbalanced (, {, or [. ")
%-----
```

*Basic vector and array practice*

```

1, (a) : [2.61628          32          -12.15          54          40.4473
           1.29623]
1, (b) : [
           12.5
8.55802
2.68825
           13.35
           13.35
           ]
1, (c) : a = [0            0            0            0            0
           0            0            0            7.5]
1, (d) : B = [
1 0 3
2 0 3
3 0 3
4 0 3
5 0 3
           ]
1, (e) : A = [
33 18
21 6
9 18
14 34
           ]
1, (e) : B = [9 6 6 10 27 15 6 22 38]
1, (e) : C = Error: Expression or statement is incorrect--possibly
unbalanced (, {, or [. ].
```

## 2 Transforming vectors and arrays

```

disp('Transforming vectors and arrays')
% a
% function code in separate file
% b
% functions code in separate files
% c
seriesSum = sum(ones(1,10000)./(1:10000).*repmat([1,-1],1,10000/2));
disp(['2, (c): seriesSum = ',num2str(seriesSum)])
%-----
```

*Transforming vectors and arrays*

```

2, (c): seriesSum = 0.6931
```

---

## 3 Filtering, searching and summarizing arrays and vectors

```
disp('Filtering, searching and summarizing arrays and vectors')
% a
% function code in separate file
A=[1 5 3 4 2; 9 7 6 8 10; 11 14 13 12 15];
[v,w] = filterMat(A);
disp(['3, (a) : evens = [',num2str(v'),']'])
disp(['3, (a) : odds = [',num2str(w'),']'])
% b
% function code in separate file
evenA = odds2zero(A);
disp('3, (b) A without odds: ')
disp(num2str(evenA))
disp('          ')
%-----
```

*Filtering, searching and summarizing arrays and vectors*

```
3, (a) : evens = [14      6      4      8      12      2      10]
3, (a) : odds = [1      9      11      5      7      3      13      15]
3, (b) A without odds: [
  0      0      4      2
  0      6      8      10
  0     14      0     12      0
]
```

## 4 String handling

```
% a
hi = 'hello';
% b
strings = {'hello', 'goodbye', 'hello', 'hello', 'goodbye', 'goodbye'};
% c
n = sum(strcmp(hi, strings));
allstrings={};
for i = 1:length(strings)
    allstrings = [allstrings, strssplit(strings{i})];
end
m = sum(strcmp(hi,allstrings));

%
```

## 5 Processing EEG data

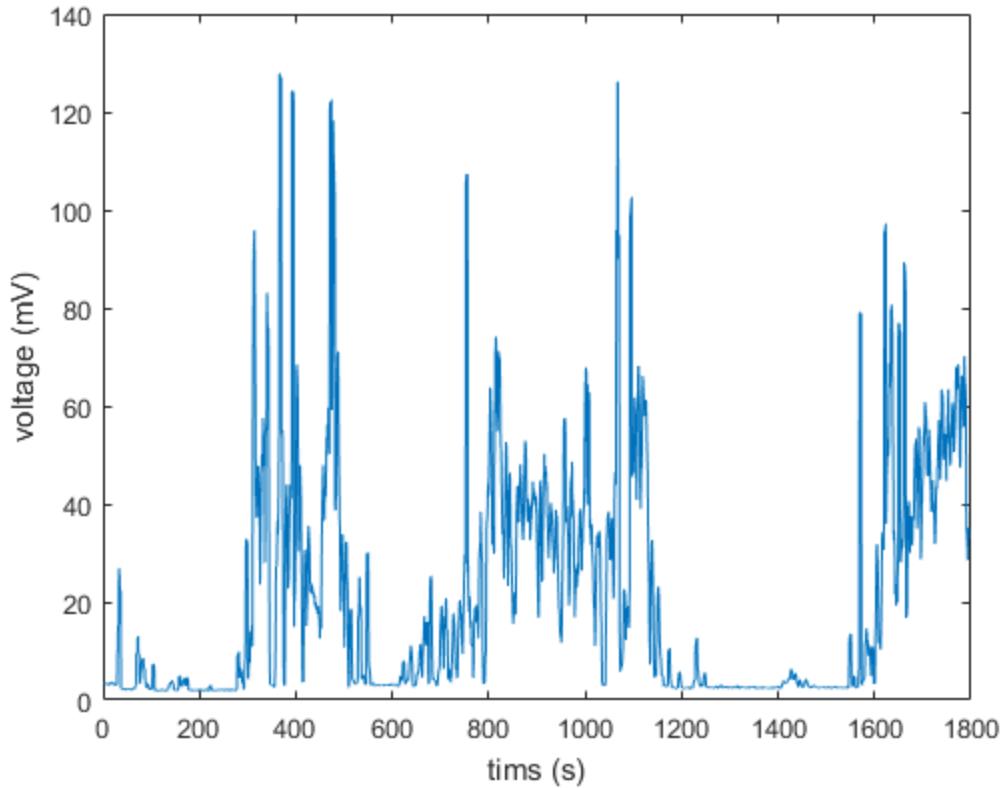
```
disp('Processing EEG data')
% a
load('acc')
load('time')
% b
```

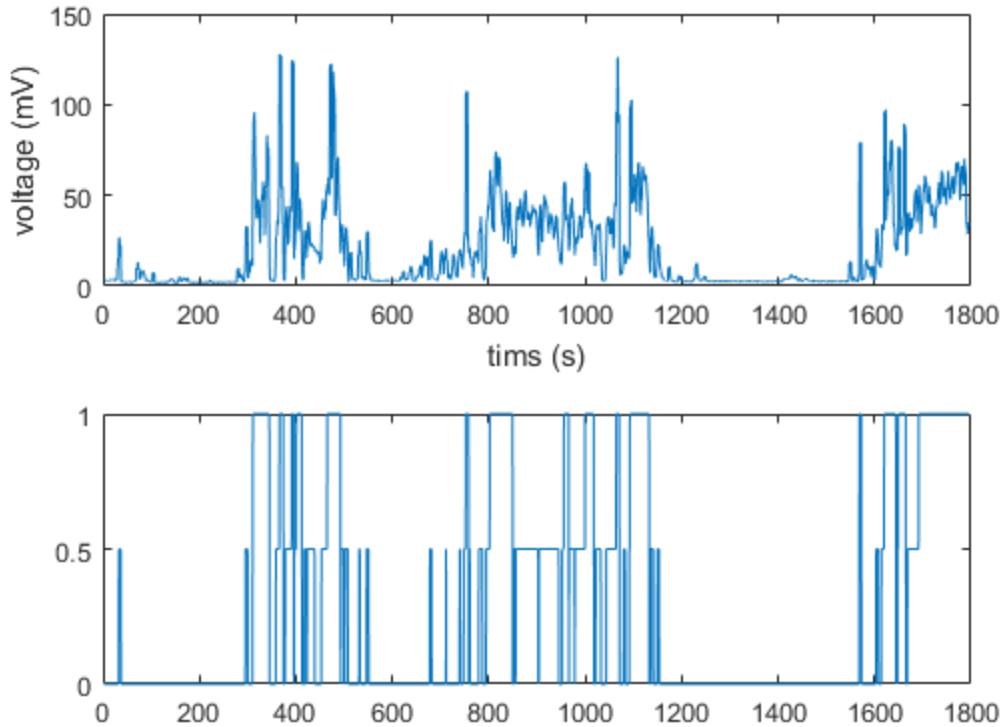
---

```
figure
plot(time,acc)
xlabel('tims (s)')
ylabel('voltage (mV)')
saveas(gcf,'figures/EEG.png')
%
detector = activity(acc, time);
%
figure
subplot(2,1,1)
plot(time,acc)
xlabel('tims (s)')
ylabel('voltage (mV)')
subplot(2,1,2)
plot(time,detector)
saveas(gcf,'figures/detector.png')

%
```

*Processing EEG data*





## 6 Processing PDB files

```

disp('Processing PDB files')
%
% a
% b
[~,~,coords] = xlsread('mystery_molecule.xls', 'G2:L660');
%
for i = 1:length(coords)
    coords{i,6} = double(coords{i,6});
end
coords = cell2mat(coords);
coords = sortrows(coords,6);
figure
plot3(coords(:,1),coords(:,2),coords(:,3))
saveas(gcf, 'figures/DNA.png')
%
C = sum(coords(:,6)==67); % black
N = sum(coords(:,6)==78)+C; % blue
O = sum(coords(:,6)==79)+N; % red
P = sum(coords(:,6)==80)+O; % green

figure
hold on
plot3(coords(1:C,1),coords(1:C,2),coords(1:C,3), 'black')
plot3(coords(C:N,1),coords(C:N,2),coords(C:N,3), 'blue')

```

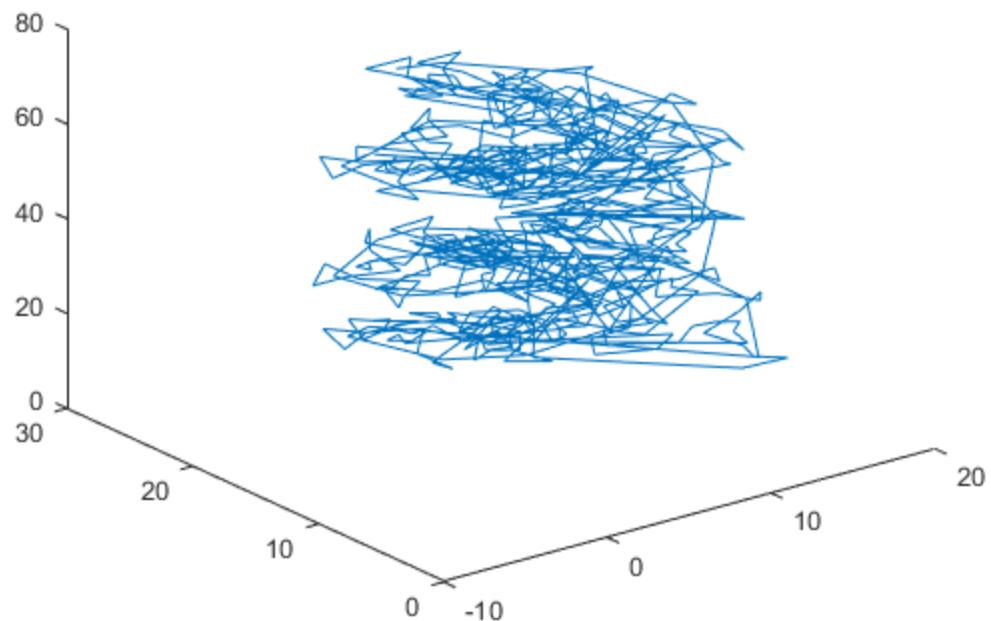
---

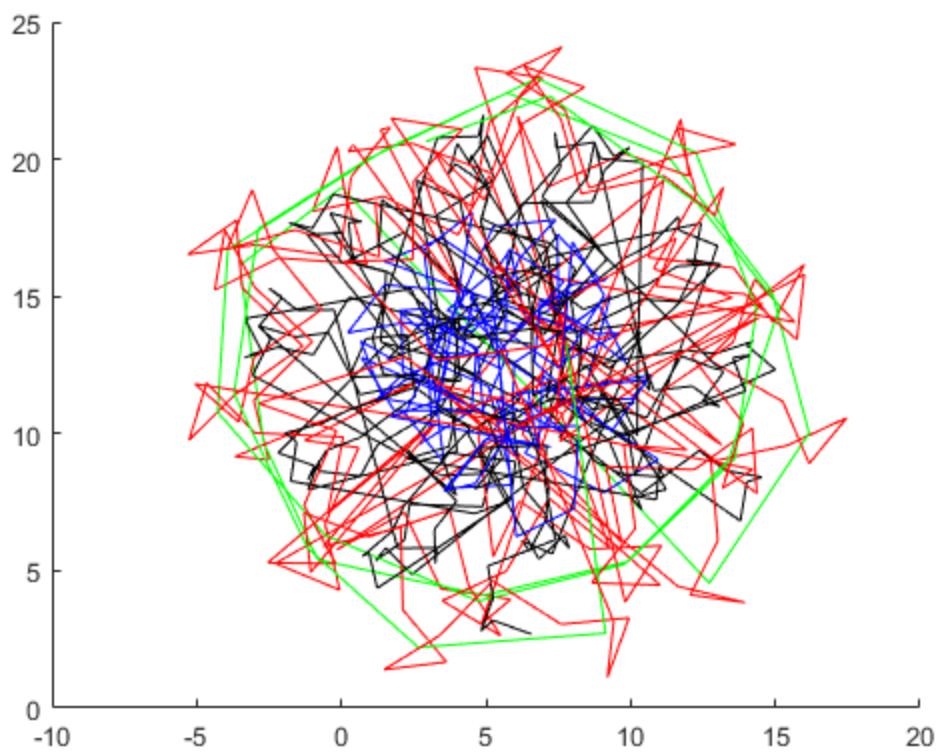
```
plot3(coords(N:O,1),coords(N:O,2),coords(N:O,3), 'red')
plot3(coords(O:P,1),coords(O:P,2),coords(O:P,3), 'green')
hold off
saveas(gcf, 'figures/colorDNA.png')

%-----
```

clear

*Processing PDB files*





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