

% Exercise 1.

```
N = 10;  
  
x = randn([1 N]);  
y = randn([1 N]);  
  
vector1 = x;  
vector2 = y;  
  
x_i = (vector1-mean(vector1))/std(vector1);  
y_i = (vector2-mean(vector2))/std(vector2);  
  
r = sum(x_i.*y_i)/N;  
  
disp(sprintf("Calculated correlation coefficient is %.4g",r));
```

Calculated correlation coefficient is 0.01154

```
disp(sprintf("Inbouded function correlation coefficient is %.4g",corr(x(:),y(:))));
```

Inbouded function correlation coefficient is 0.01282

% Exercise 2.

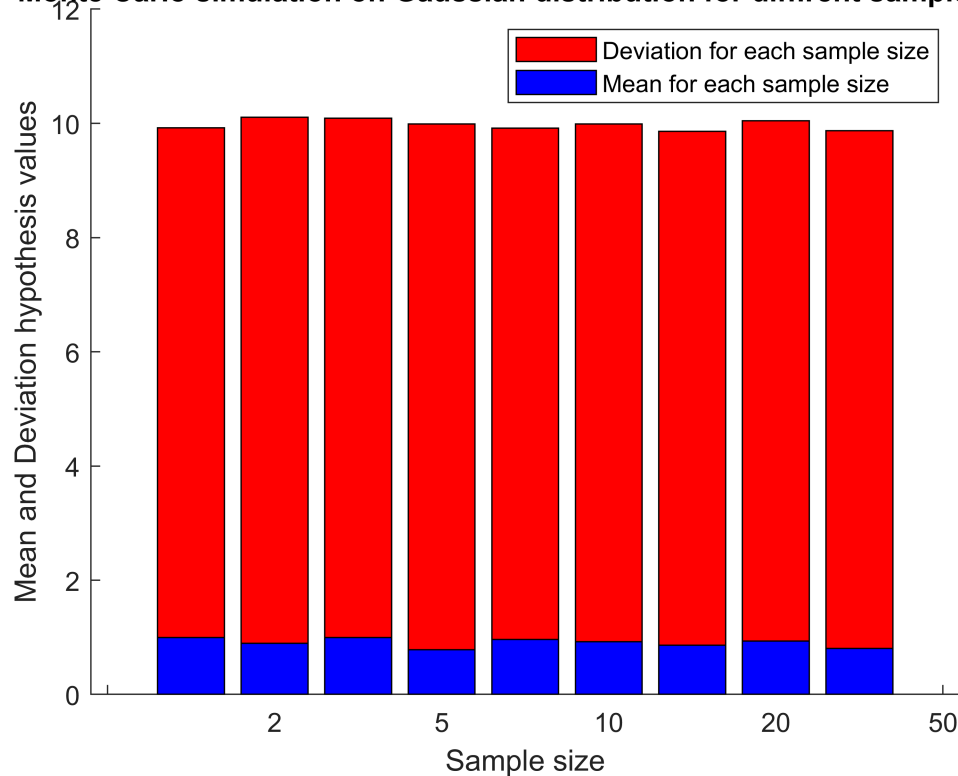
```
sample_sizes = [2 5 10 20 50 100 200 500 10000];  
iteration = 100;  
  
results = zeros(9,2);  
  
%results = zeros(size(sample_sizes(:)),iteration)  
index = 1;  
for r = sample_sizes  
  
    [mean_mean, std_std] = monte_carlo(5,iteration);  
  
    results(index,:) = [mean_mean std_std];  
    index = index +1;  
  
end  
  
% visualize the data  
  
names = {'','2', '5', '10', '20', '50', '100', '200', '500', '10000'};  
  
figure; hold on;  
h1 = bar(results(:,1), 'r');
```

```

h2 = bar(results(:,2), 'b');
legend([h1 h2], 'Deviation for each sample size', 'Mean for each sample size');
title("Monte Carlo simulation on Gaussian distribution for diffirent sample size")
set(gca, 'xticklabel', names);
xlabel('Sample size');
ylabel('Mean and Deviation hypothesis values');

```

Monte Carlo simulation on Gaussian distribution for diffirent sample size



% Exercise 3

```
x = rand(1,10);
```

```
bootstrapped = bootstrap(x);
```

```
% visualize the original sample and bootstraped
```

```
figure; hold on;
```

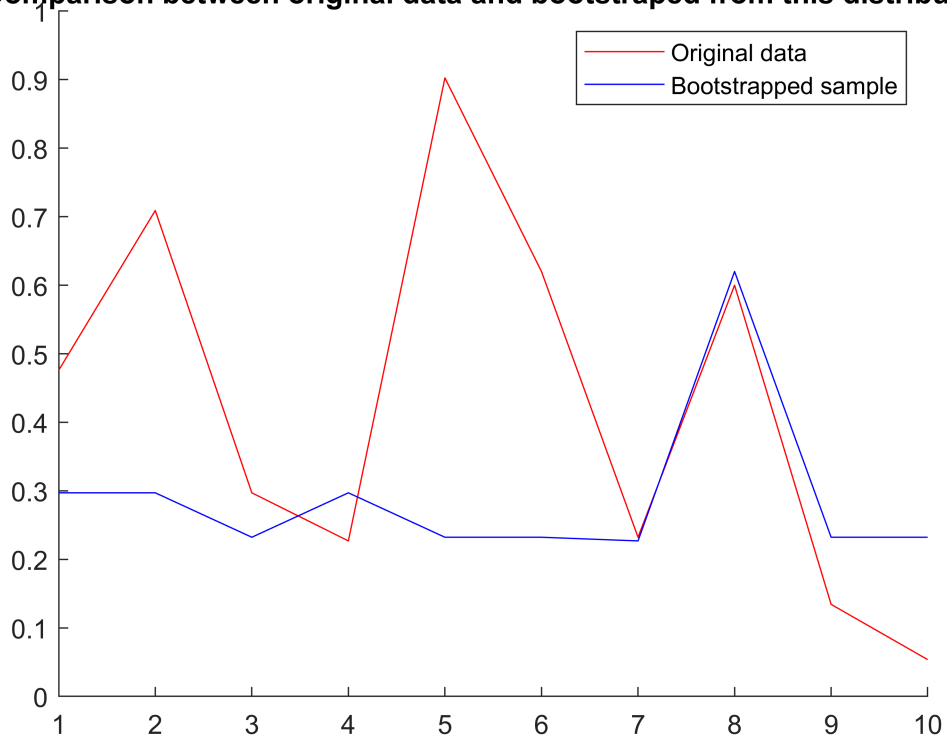
```
h1 = plot(x, 'r');
```

```
h2 = plot(bootstrapped, 'b');
```

```
title('Comparison between original data and bootstraped from this distribution')
```

```
legend([h1 h2], 'Original data', 'Bootstrapped sample');
```

Comparison between original data and bootstrapped from this distribution



% Exercise 4

```
bootstrapped_array = ex4_return_2d(1000,10);
```

```
% Visualize different samples
```

```
%
```

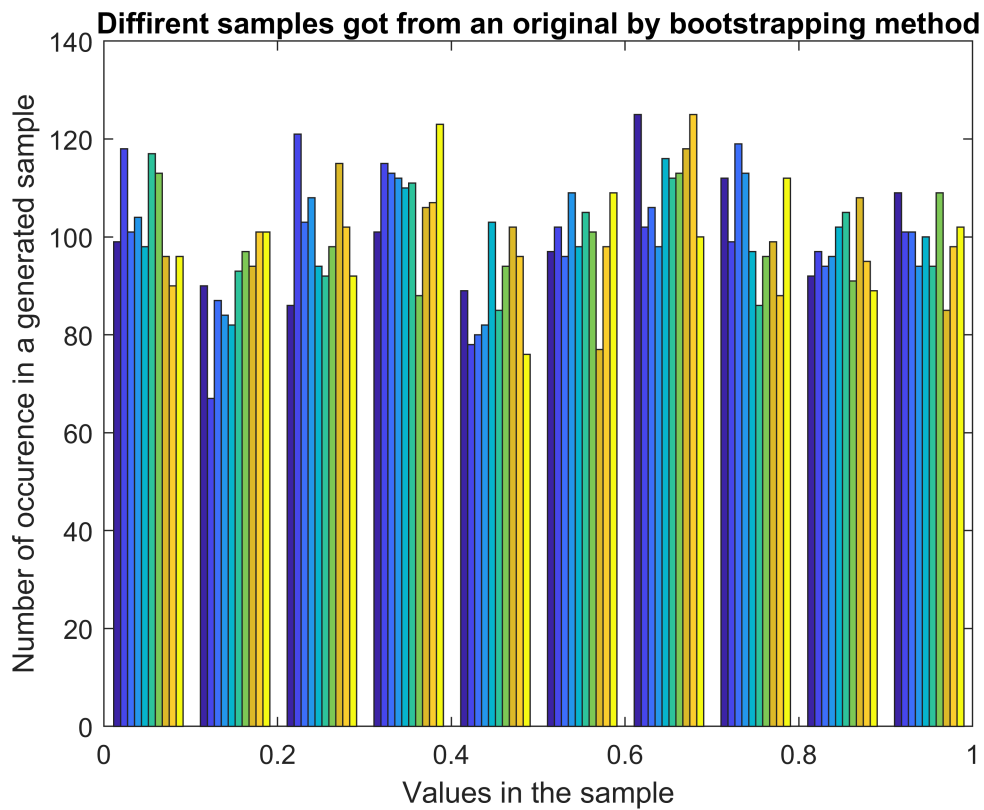
```
figure;
```

```
hist(bootstrapped_array');
```

```
title('Different samples got from an original by bootstrapping method');
```

```
xlabel('Values in the sample');
```

```
ylabel('Number of occurrence in a generated sample')
```



% Exercise 5

% X is for the speed of cars

```
x = [10 20 30 23 10 190 45 34 23 1 10 90 40 70 24 41 4 74 2 4]';
```

% Y is for the amount of accident for cars at specific speed

```
y = [ 2 5 4 2 5 19 8 3 2 0 0 14 7 10 4 7 1 9 2 1]';
```

%Visualize the dependency

```
figure;
```

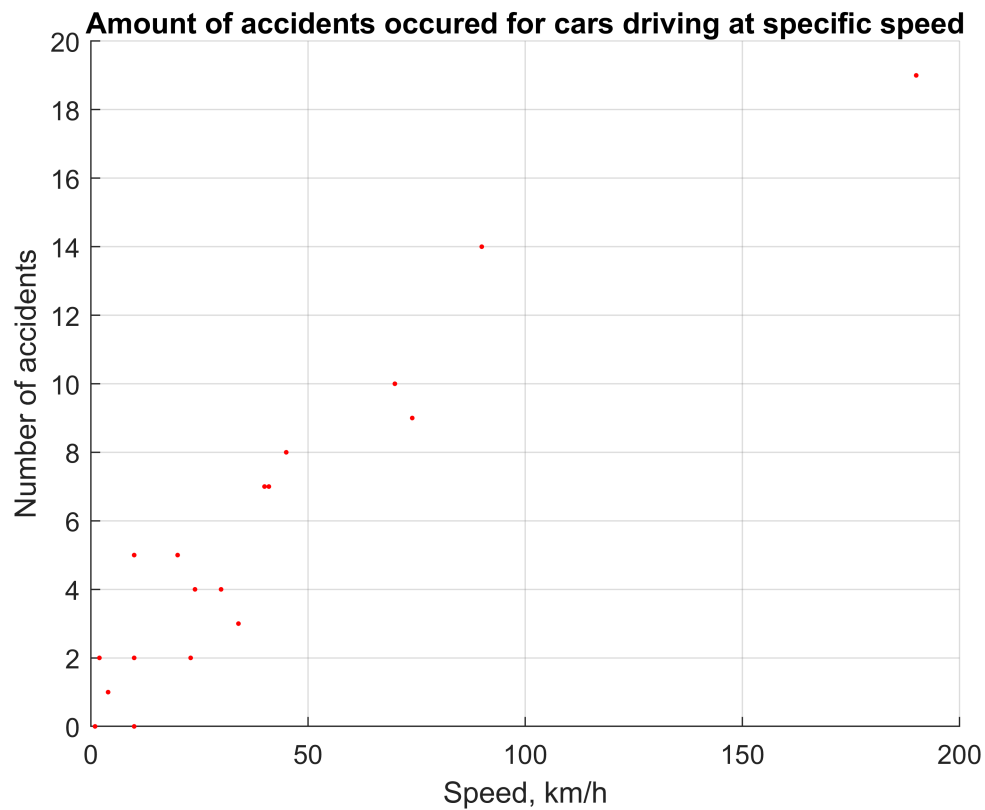
```
scatter(x,y,'r.');
```

```
title('Amount of accidents occured for cars driving at specific speed');
```

```
grid;
```

```
xlabel('Speed, km/h');
```

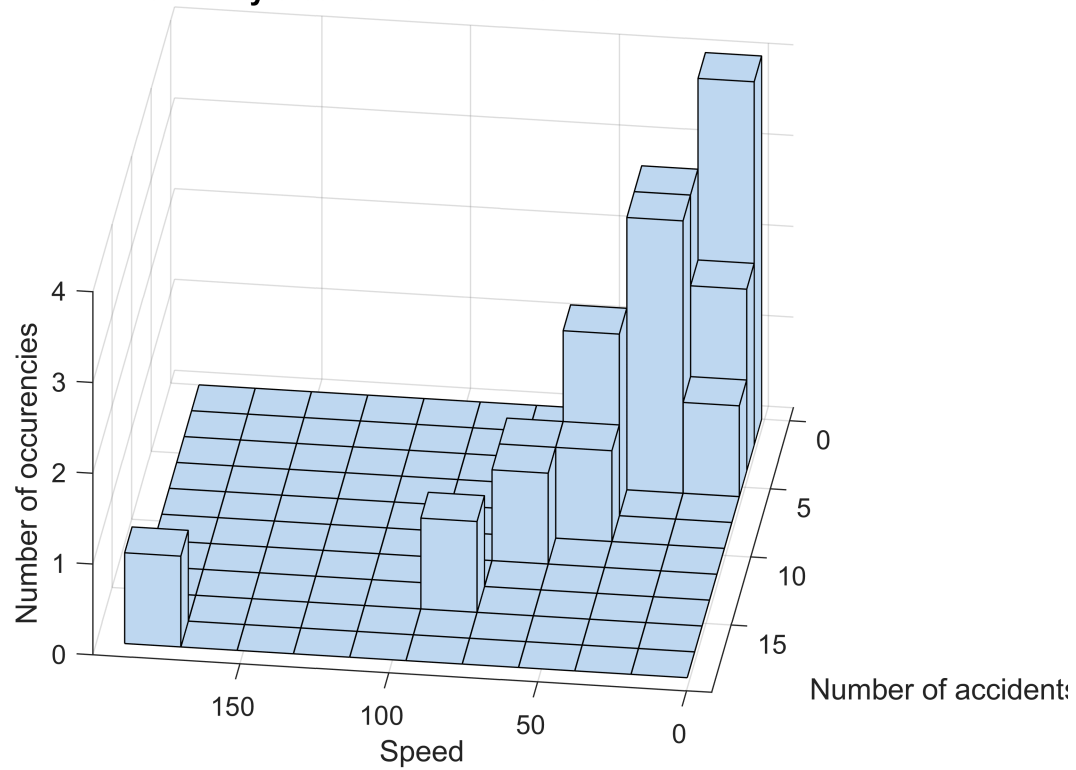
```
ylabel('Number of accidents');
```



```
% Plot 2D histogram. On x-axis - values from the first distribution. On  
% y-axis - values from the second distribution. On z-axis amount of values  
% took specific value.
```

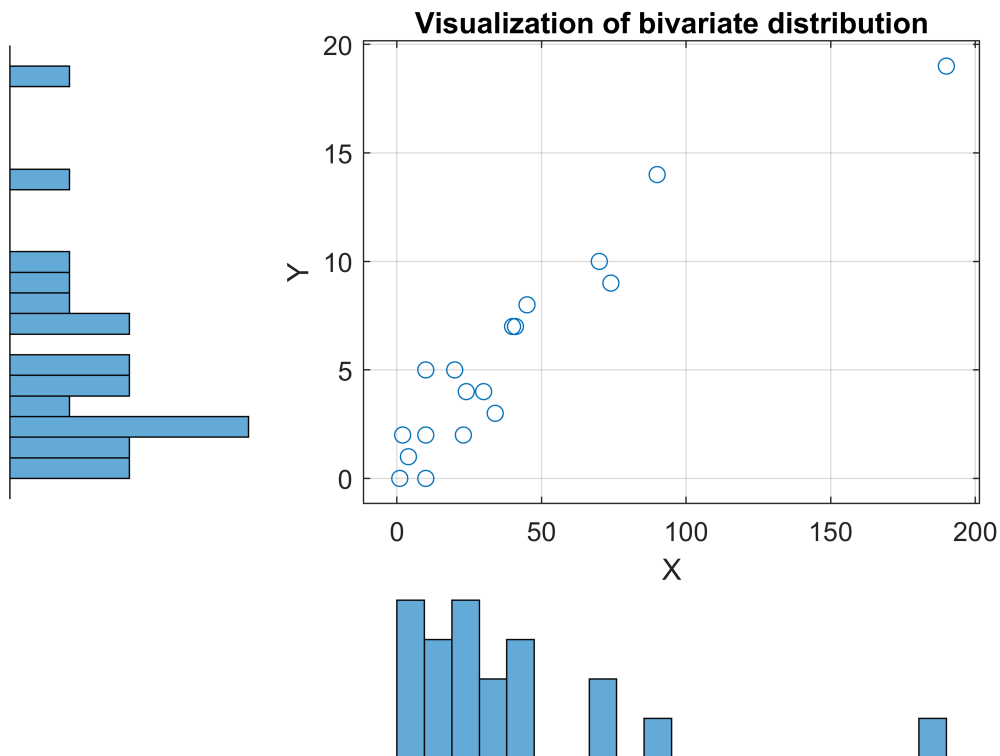
```
figure;  
hist3([x y],[10, 10]);  
title("Probability distribution for 2 Random Variables");  
xlabel('Speed');  
ylabel('Number of accidents');  
zlabel('Number of occurrences');
```

Probability distribution for 2 Random Variables



```
% Calculate marginal probability for X (stands for speed)
```

```
scatterhist(x,y,20);grid;  
title('Visualization of bivariate distribution');  
xlabel('X')  
ylabel('Y')
```



```
% use filter to sort out unnesessary dots for X variable
```

```
x_filter_lower = 100;  
x_filter_higher = 200;
```

```
% Apply a filter on the X
```

```
x_i = x( x_filter_lower<x .* (x<=x_filter_higher));
```

```
number_of_occurencies = length(x_i);
```

```
number_of_occurencies = 1
```

```
% What we got:
```

```
% N(0<x<=50) = 16
```

```
% N(50<x<100) = 3
```

```
% N(100<x<200) = 1
```

```
% use filter to sort out unnesessary dots for Y variable
```

```
y_filter_lower = 15;  
y_filter_higher = 20;
```

```
% Apply a filter on the Y
```

```
y_i = y( y_filter_lower<y .* (y<=y_filter_higher));
```

```
number_of_occurencies = length(y_i);
```

```
number_of_occurencies = 1
```

```
% What we got:
```

```
% N(0<y<=5) = 11
```

```
% N(5<y<10) = 5
```

```
% N(10<y<15) = 1
```

```
% N(15<y<20) = 1
```

```
% Finally, we got the matrix with occurancies for X and Y.
```

```
% Given that we can calculate the total lenght of samples, we are able to
```

```
% calculate the frequency of each random variables separately
```

```
N = length(x);
```

```
occurencies_x = [16 3 1] ./N
```

```
occurencies_x = 1×3
```

```
0.8000 0.1500 0.0500
```

```
occurencies_y = [11 5 1 1] ./N
```

```
occurencies_y = 1×4
```

```
0.5500 0.2500 0.0500 0.0500
```

```
% This is marginal distribution for P(x) and P(y) respectively
```

```
% Obviously, the data can be presented in the following table
```

% N	x	P(0<x<=50 y)	P(50<x<100 y)	P(100<x<200 y)	P(Y)
% P(0<y<=5 x)					55%
% P(5<y<10 x)					25%
% P(10<y<15 x)					5%
% P(15<y<20 x)					5%
% P(X)		80%	15%	5%	100%

```
% Since we expect the variables are independent, then the joint probability
```

```
% P(XY) = P(X)*P(Y), where P(X),P(Y) - marginal distribution for X,Y in
```

```
% probability space.
```

```
joint_probability_matrix = occurencies_x' * occurencies_y
```

```
joint_probability_matrix = 3×4
```

```
0.4400 0.2000 0.0400 0.0400
```

```
0.0825 0.0375 0.0075 0.0075
```


0.0275 0.0125 0.0025 0.0025

```
x_names = {'P(0<x<=50|y)'; 'P(50<x<100|y)'; 'P(100<x<200|y)'; ''};
y_names = {'P(0<y<=5)'; 'P(10<y<15|x)'; 'P(10<y<15|x)'; 'P(15<y<20|x)'};

s = mesh(joint_probability_matrix);
colormap(jet)
set(gca,'xticklabel',x_names);
set(gca,'yticklabel',y_names);
legend(s,'P(XY)');
title("Joint probability for independent random variables X and Y");
xlabel("P(X|Y)")
ylabel("P(Y|X)")
```

