Bangladesh University of Engineering and Technology



Numerical Technique Laboratory

EEE 212

Experiment No.: 04

Name of the Experiment: Curve Fitting

Department: EEE

Section: C1

Group: 01

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Curve Fitting

Polynomial Regression for nth Order

Code:

```
clearall ;
clc ,clf , close all ;
% Curve Fitting
x = [0 \ 1 \ 2 \ 3 \ 4 \ 5] ;
y = [2.1 7.7 13.6 27.2 40.9 61.1];
% plotting x and y as points
plot(x,y,'0','MarkerSize',6,'MarkerFaceColor','w','MarkerEdgeColor',...
'b','LineWidth',2) ;
holdon ;
% taking order of polynomial as input
input value= input('Enter the order of polynomial : ') ;
order = 1 ;
% plotting curve from order 1 to input value
while order <= input value + 1</pre>
power = 0;
for row = 1 : order
for col = 1 : order
A(row,col) = sum(x .^ power) ;
power = power + 1 ;
              % as every time power of x start from the previous
power = row ;
% row value
for row = 1 : order
B(row,1) = sum((x.^(row-1)) .* y);
end
coefficient = A \ ;
total = 0;
for i = 1 : length(coefficient) - 1
total = total + coefficient(i+1) * (x.^i) ;
new y = total + coefficient(1) ;
plot(x,new y) ;
% changing order
order = order + 1 ;
end
```



Output:

Command window output:

```
Command Window

Enter the order of polynomial : 6

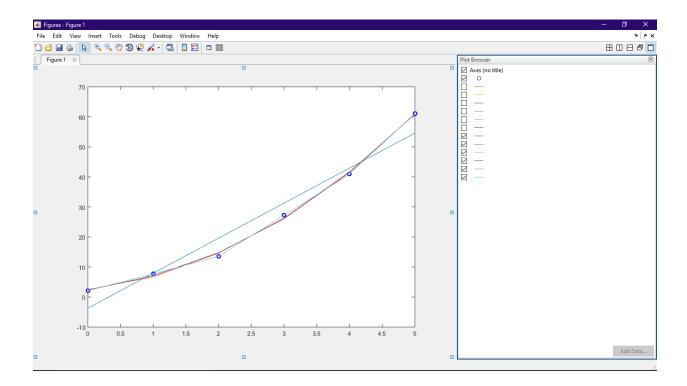
f_X >>

script

Ln 46 Col 1
```

Output:

Output of polynomial regression for order 1 to 6 in a single plot

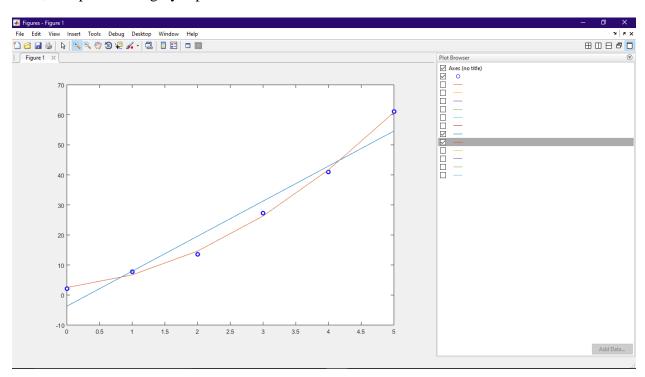




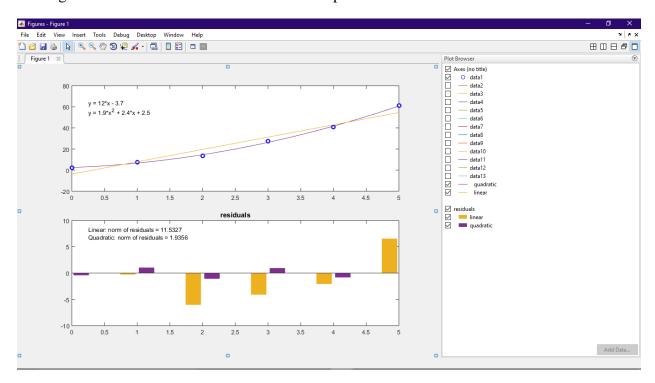
Comments:

Output plot for order 1 & 2

Here, two plots are highly separated.



Finding residuals and norm of the residuals from plot window:



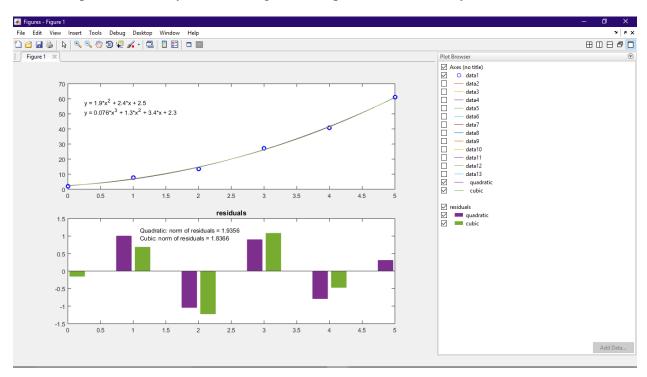


Residual is the difference between the sample and the estimated function value. If the points in a residual plot are randomly dispersed around the horizontal axis, a linear regression model is appropriate for the data otherwise a non-linear model is more appropriate. Here, residuals value is not randomly dispersed and non-random residuals indicate a non-linear better fitted curve.

So, a non-linear curve would be better fitted for the given points.

Output plot for order 2 &3:

Here, two plots are closely located. [Higher order plots are also closely located.]



Norm of the residuals is a measure of the deviation between the correlation and data. A lower norm of the residuals means a better fit. Here, norm of the residuals are:

Order 1: norm of residuals = 11.5327

Order 2: norm of residuals = 1.9356

Order 3: norm of residuals = 1.8366

The difference between the norm of residuals for order 1 & 2 is: 9.5971

The difference between the norm of residuals for order 2 & 3 is: .099

That is why plot for order 1 & 2 are highly separated and plot for order 2 & 3 are closely located.



Curve Fitting Of a Noisy Sin Wave

Code:

```
clearall ;
clc ,clf , close all ;
x = 0:360;
y = sind(x);
plot(x,y,'LineWidth',2) ;
axis([0 360 -1.5 1.5]);
title('Actual Sin wave', 'LineWidth',2) ;
xlabel('x','LineWidth',2);
ylabel('sin(x)','LineWidth',2) ;
figure ;
min = -.1;
max = .1;
n = length(x);
% Random number generator within limit
noise = min + rand(1,n) * (max-min);
y = y + noise ;
plot(x,y,'LineWidth',1.5) ;
axis([0 360 -1.5 1.5]);
title('Sin wave with noise', 'LineWidth', 2);
xlabel('x','LineWidth',2);
ylabel('sin(x) + Noise', 'LineWidth',2) ;
figure ;
new x = [] ;
new_y = [] ;
for i = 1: 15 : 361
new x = [new xx(i)];
new_y = [new_yy(i)];
end
plot(new x,new y,'0','Linewidth',1.5) ;
axis([0 \ \overline{3}60 \ -1.5 \ 1.5]);
holdon ;
% Curve Fitting
n = length(new x);
order = 8;
power = 0;
```

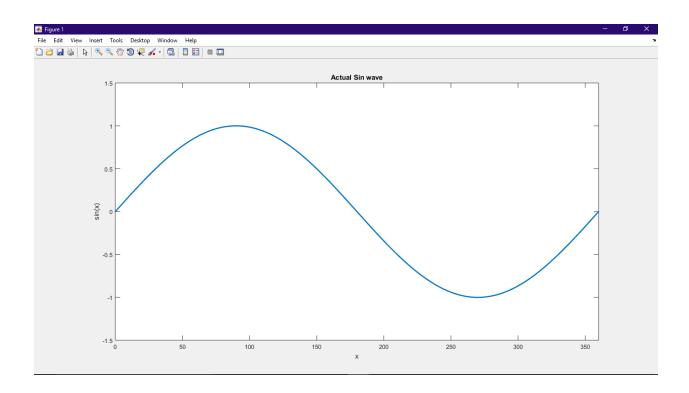


```
for row = 1 : order
for col = 1 : order
A(row,col) = sum(new x .^ power) ;
power = power + 1 ;
end
power = row ;
end
for row = 1 : order
B(row,1) = sum((new_x.^(row-1)) .* new_y);
end
coefficient = A \ ;
total = 0 ;
for i = 1 : length(coefficient) - 1
total = total + coefficient(i+1) * (new_x.^i) ;
end
estimated y = total + coefficient(1) ;
plot(new_x,estimated_y,'Linewidth',2) ;
title('Estimated Sin wave', 'LineWidth',2) ;
axis([0 360 -1.5 1.5]);
x = 0:360;
y = sind(x);
actual y = [] ;
for i = 1: 15 : 361
actual_y = [actual_yy(i)] ;
end
error = (actual_y - estimated_y) ./ actual_y ;
error = abs(error .* 100) ;
% Removing Inf and NaN from error matrix
error(~isfinite(error)) = 0;
figure ;
% plotting error
plot(error, 'LineWidth', 1.5) ;
title('Percentage of error between original sin wave and estimated sin
wave',...
'LineWidth',2);
ylabel('Error','LineWidth',2) ;
```

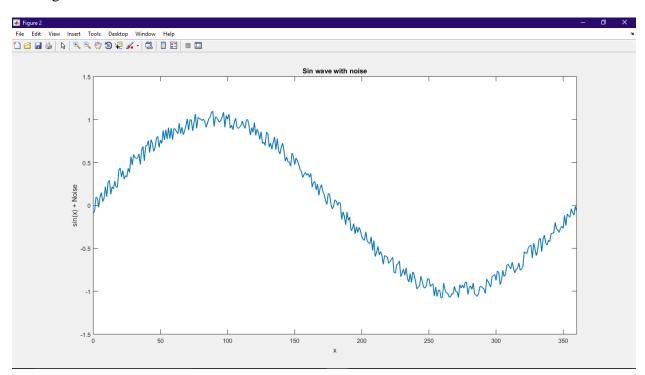


Output:

Actual Sin Wave:

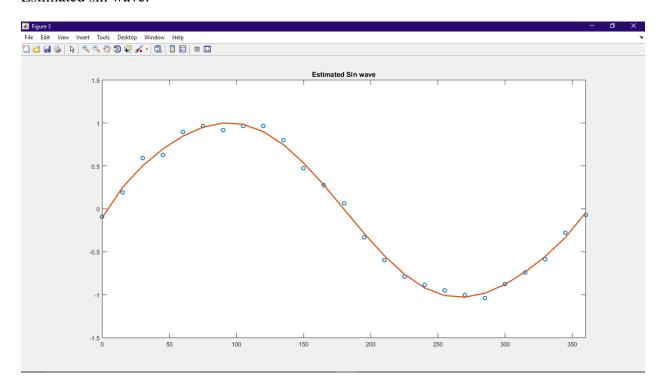


Adding noise to the sin wave:





Estimated sin wave:



Percentage of error:

Calculated as, percentage of error =
$$\frac{\text{Actual value - Estimated value}}{\text{Actual value}} \times 100\%$$

