
Interpolation i Matlab

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```
clear all, close all
```

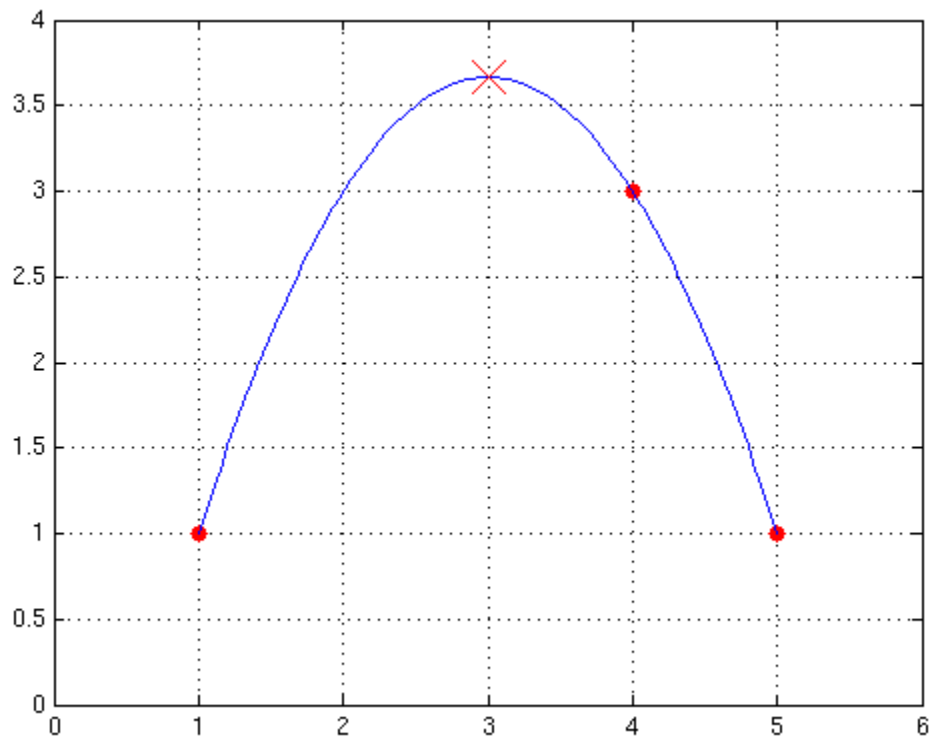
```
X = [1 4 5]';  
Y = [1 3 1]';
```

```
% ekvationssystemet  
c = [ones((size(X))) X X.^2]\Y  
f = @(x) c(1) + c(2)*x + c(3)*x.^2;
```

```
% derivera: df = c(2) + 2*c(3)*x = 0 =>  
xmax = -c(2)/(2*c(3))
```

```
x = linspace(1,5,100);  
plot(X,Y,'r.',x,f(x),'b',xmax,f(xmax),'rx','MarkerSize',20),  
axis([0 6 0 4]), grid on
```

```
c =  
-2.333333333333333  
 4.000000000000000  
-0.666666666666667  
xmax =  
 3
```



Med inbyggda funktioner

```
X = [1 4 5 7 8]';
Y = [1 3 1 4 5]';

% anpassa polynom
c = polyfit(X,Y,4)
x = linspace(1,8,100);
y = polyval(c,x); % evaluera polynomet i punkterna x

figure()
plot(X,Y,'r.',x,y,'MarkerSize',20), grid on

% styckvis kubisk
help pchip

figure()
c = pchip(X,Y)
y = ppval(c,x);
plot(X,Y,'r.',x,y,'MarkerSize',20), grid on

c =
Columns 1 through 3
-0.091269841269840    1.857142857142832   -12.757936507936355
Columns 4 through 5
33.214285714285381   -21.2222222222222076
PCHIP Piecewise Cubic Hermite Interpolating Polynomial.
PP = PCHIP(X,Y) provides the piecewise polynomial form of a certain
shape-preserving piecewise cubic Hermite interpolant, to the values
Y at the sites X, for later use with PPVAL and the spline utility UNMKPP.
X must be a vector.
If Y is a vector, then Y(j) is taken as the value to be matched at X(j),
hence Y must be of the same length as X.
If Y is a matrix or ND array, then Y(:,...,:),j) is taken as the value to
be matched at X(j), hence the last dimension of Y must equal length(X).

YY = PCHIP(X,Y,XX) is the same as YY = PPVAL(PCHIP(X,Y),XX), thus
providing, in YY, the values of the interpolant at XX.

The PCHIP interpolating function, p(x), satisfies:
On each subinterval,  $X(k) \leq x \leq X(k+1)$ , p(x) is the cubic Hermite
interpolant to the given values and certain slopes at the two endpoints.
Therefore, p(x) interpolates Y, i.e.,  $p(X(j)) = Y(:,j)$ , and
the first derivative,  $Dp(x)$ , is continuous, but
 $D^2p(x)$  is probably not continuous; there may be jumps at the  $X(j)$ .
The slopes at the  $X(j)$  are chosen in such a way that
p(x) is "shape preserving" and "respects monotonicity". This means that,
on intervals where the data is monotonic, so is p(x);
at points where the data have a local extremum, so does p(x).

Comparing PCHIP with SPLINE:
The function s(x) supplied by SPLINE is constructed in exactly the same way,
except that the slopes at the  $X(j)$  are chosen differently, namely to make
even  $D^2s(x)$  continuous. This has the following effects.
SPLINE is smoother, i.e.,  $D^2s(x)$  is continuous.
SPLINE is more accurate if the data are values of a smooth function.
PCHIP has no overshoots and less oscillation if the data are not smooth.
PCHIP is less expensive to set up.
The two are equally expensive to evaluate.

Example:

x = -3:3;
```

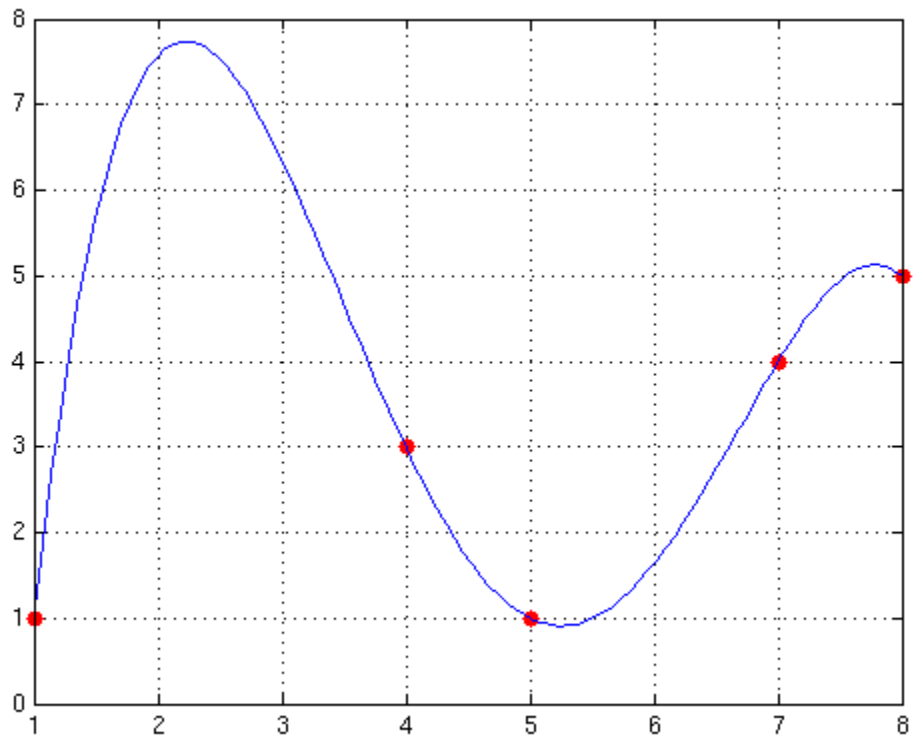
```
y = [-1 -1 -1 0 1 1 1];  
t = -3:.01:3;  
plot(x,y,'o',t,[pchip(x,y,t); spline(x,y,t)])  
legend('data','pchip','spline',4)
```

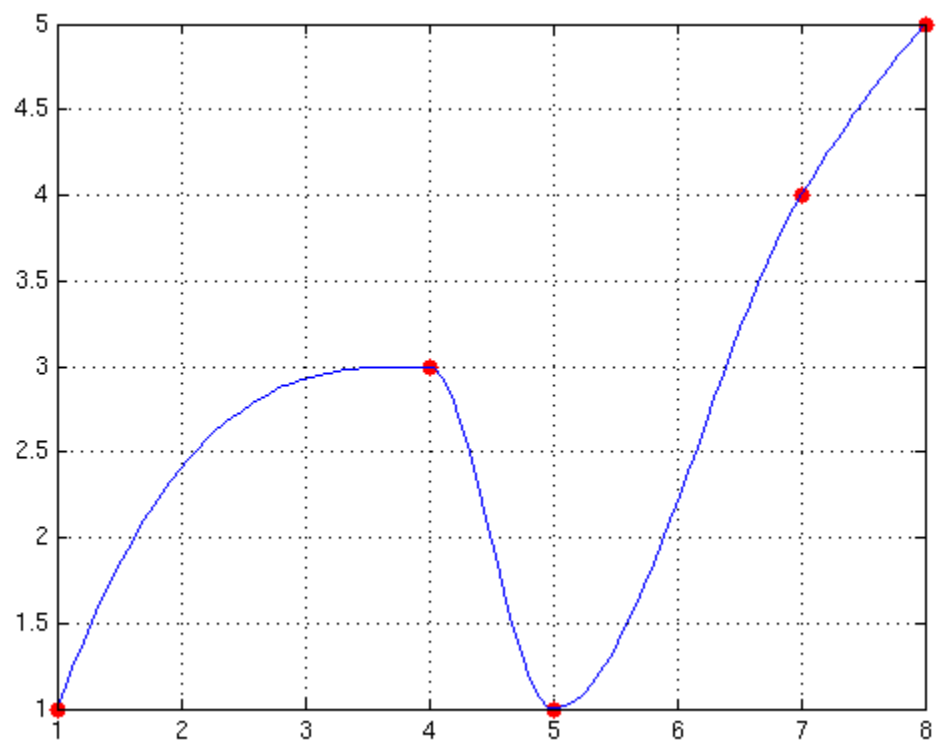
Class support for inputs *x*, *y*, *xx*:
float: double, single

See also *INTERP1*, *SPLINE*, *PPVAL*, *UNMKPP*.

Reference page in Help browser
doc pchip

```
c =  
    form: 'pp'  
breaks: [1 4 5 7 8]  
  coefs: [4x4 double]  
pieces: 4  
order: 4  
  dim: 1
```





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