

Statistics

▼ Linear Regression

All linear regression commands have been updated in Maple 2016 with a new option, **summarize**, that allows for the display of a summary for the given regression model.

`with(Statistics) :`

`X := Vector([1, 2, 3, 4, 5, 6], datatype=float) :`

`Y := Vector([2, 3, 4, 3.5, 5.8, 7], datatype=float) :`

By default, the **Fit** command returns the resulting regression model for the given model function:

`Fit(a + b * t + c * t^2, X, Y, t);`

$1.960000000000000 + 0.164999999999999 t + 0.110714285714286 t^2$

The **summarize** option includes a full summary for each of the regression coefficients, as well as values for the [r-squared](#) and [adjusted r-squared](#) for the model. Also, the [solution module](#) for regression commands has been extended with the ability to return values for r-squared, adjusted r-squared, and the value of the t-statistic for testing whether the corresponding regression coefficient is different than 0 and its corresponding probability.

`Fit(a + b * t + c * t^2, X, Y, t, summarize=true) :`

Summary:

Model: $1.9600000+.16500000*t+.11071429*t^2$

Coefficients:

	Estimate	Std. Error	t-value	P(> t)
a	1.9600	1.1720	1.6724	0.1930

b	0.1650	0.7667	0.2152	0.8434
c	0.1107	0.1072	1.0325	0.3778

R-squared: 0.9252, Adjusted R-squared: 0.8753

The **summarize** option can also be used to return an embedded table, which contains more details on the residuals:

*Fit(a + b * t + c * t^2, X, Y, t, **summarize** = embed) :*

Summary

Model: $1.9600000 + 0.16500000 t + 0.11071429 t^2$

Coefficients	Estimate	Standard Error	t-value	P(> t)
a	1.96000	1.17199	1.67237	0.193045
b	0.165000	0.766748	0.215194	0.843415
c	0.110714	0.107226	1.03253	0.377769

R-squared: 0.925169

Adjusted R-squared: 0.875282

▼ Residuals

Residual Sum of Squares	Residual Mean Square	Residual Standard Error	Degrees of Freedom
1.28771	0.429238	0.655163	3

Five Point Summary

Minimum	First Quartile	Median	Third Quartile	Maximum
-0.891429	-0.290357	0.155714	0.290595	0.548571

▼ Hypothesis Testing

The summarize option has also been added to all [hypothesis testing](#) commands. Previously, the `infolevel` command would have been required to print the results of a hypothesis test as a report.

`with(Statistics) :`

`X := Array([9, 10, 8, 4, 8, 3, 0, 10, 15, 9]) :`

`OneSampleChiSquareTest(X, 7, confidence = .95, summarize = embed) :`

Chi-Square Test on One Sample					
Null Hypothesis:		Sample drawn from population with standard deviation equal to 7			
Alternative Hypothesis:		Sample drawn from population with standard deviation not equal to 7			
Sample Size	Sample Standard Deviation	Distribution	Computed Statistic	Computed p-value	Confidence Interval
10.	4.24788	ChiSquare(9)	3.31429	0.0989571	2.92184 ..7.75496
Result:		Accepted: This statistical test does not provide enough evidence to conclude that the null hypothesis is false.			

▼ Summary and Tabulation

The [DataSummary](#), [FivePointSummary](#), and [FrequencyTable](#) commands can also accept a `summarize` option as well as be used to return summary statistics for [DataFrames](#):

`with(Statistics) :`

```
X := DataFrame((Sample(Uniform(0, 1), [50, 2])|LinearAlgebra:-RandomVector(50, generator = rand(0..3))))
```

	1	2	3
1	0.814723686393179	0.276025076998578	3
2	0.905791937075619	0.679702676853675	2
3	0.126986816293506	0.655098003973841	1
4	0.913375856139019	0.162611735194631	2
5	0.632359246225410	0.118997681558377	1
6	0.0975404049994095	0.498364051982143	0
7	0.278498218867048	0.959743958516081	1
8	0.546881519204984	0.340385726666133	0
...

`DataSummary(X, summarize = embed) :`

	1	2	3
mean	0.5661101110386353	0.48987882283825	1.22
standarddeviation	0.3125414653315035	0.2785520677947388\4	1.1830434635864795
skewness	-0.371651925679829\9	0.1158245251557512\3	0.3106598115832435
kurtosis	1.729650721370971	1.7837876411157099	1.5501229720154914
minimum	0.0318328463774206\76	0.0119020695012413\97	0.0
maximum	0.9705927817606157	0.9597439585160811	3.0
cumulativeweight	50.0	50.0	50.0

▼ Visualizations

There are many new visualizations in Maple 2016 for statistics and data analysis, including new options for creating [colorschemes](#) using point values:

with(Statistics) :

```
data := Matrix((Sample(Uniform(0, 1), [50, 2]) | LinearAlgebra:-RandomVector(50, generator = rand(0..3))))
```

[

50 x 3 Matrix

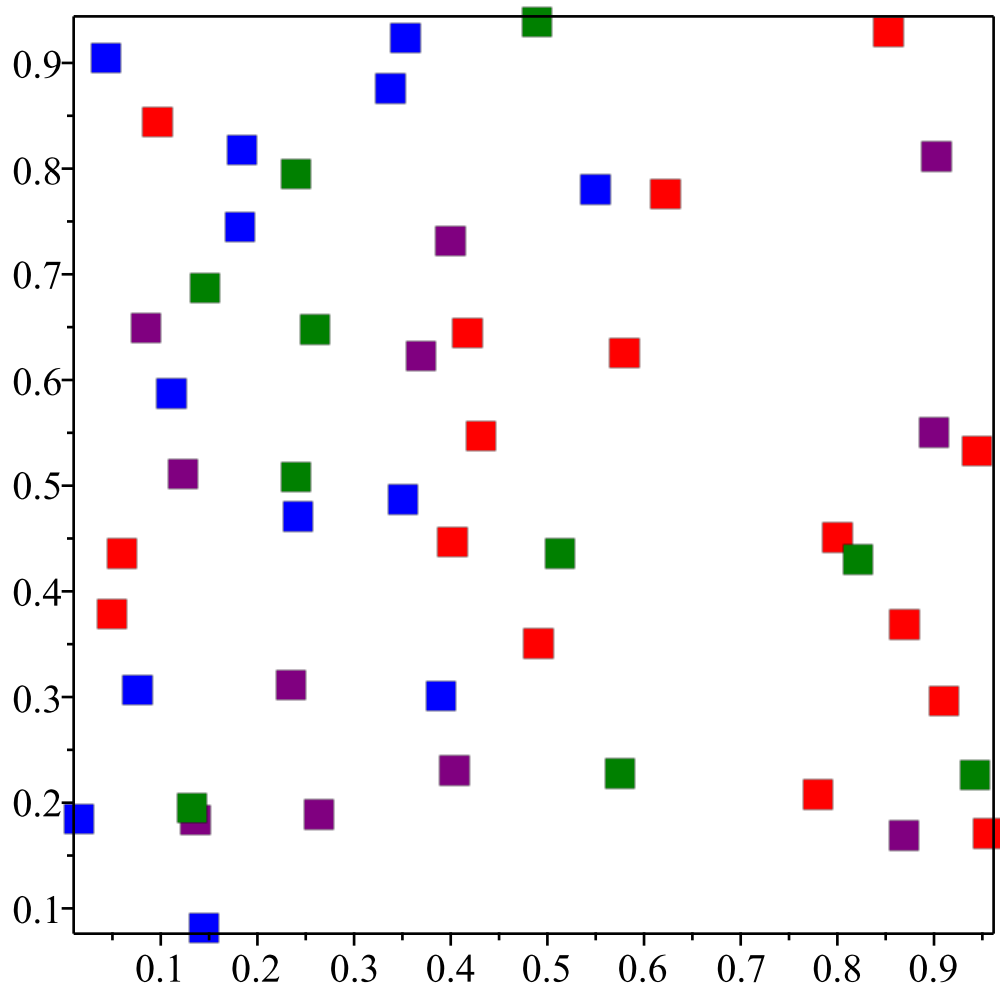
Data Type: anything

Storage: rectangular

Order: Fortran_order

]

```
ScatterPlot( data[ .., 1 ], data[ .., 2 ], symbolsize = 20, symbol = solidbox,
             colorscheme = [ "valuesplit", data[ .., 3 ], [ 0 = "Red", 1 = "Blue", 2 = "Green", 3 = "Purple" ] ] )
```



Maple 2016 also introduces a new visualization in Statistics for generating [heat maps](#). A heat map is a visualization method that represents the magnitude of the included data as a discrete density plot.

```
U := <seq(0..10)> :
```

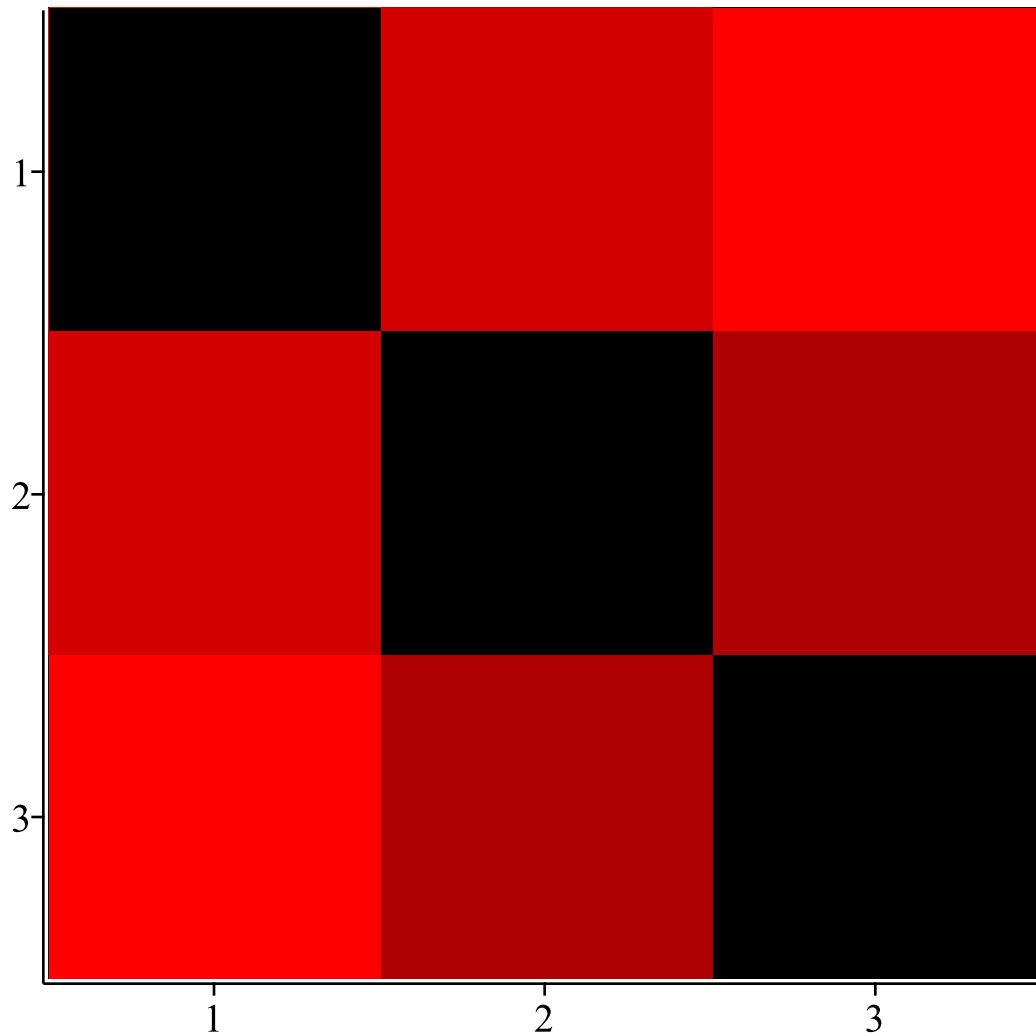
```
V := <seq(sin(i), i=0..10)> :
```

```
W := <seq(cos(i), i=0..10)> :
```

```
CM := CorrelationMatrix(Matrix([U, V, W]), ignore);
```

$$\begin{bmatrix} 1. & -0.116741765101327 & -0.354242435116850 \\ -0.116741765101327 & 1. & 0.0701662833954325 \\ -0.354242435116850 & 0.0701662833954325 & 1. \end{bmatrix}$$

```
HeatMap(CM, color = ["Red", "Black"]);
```



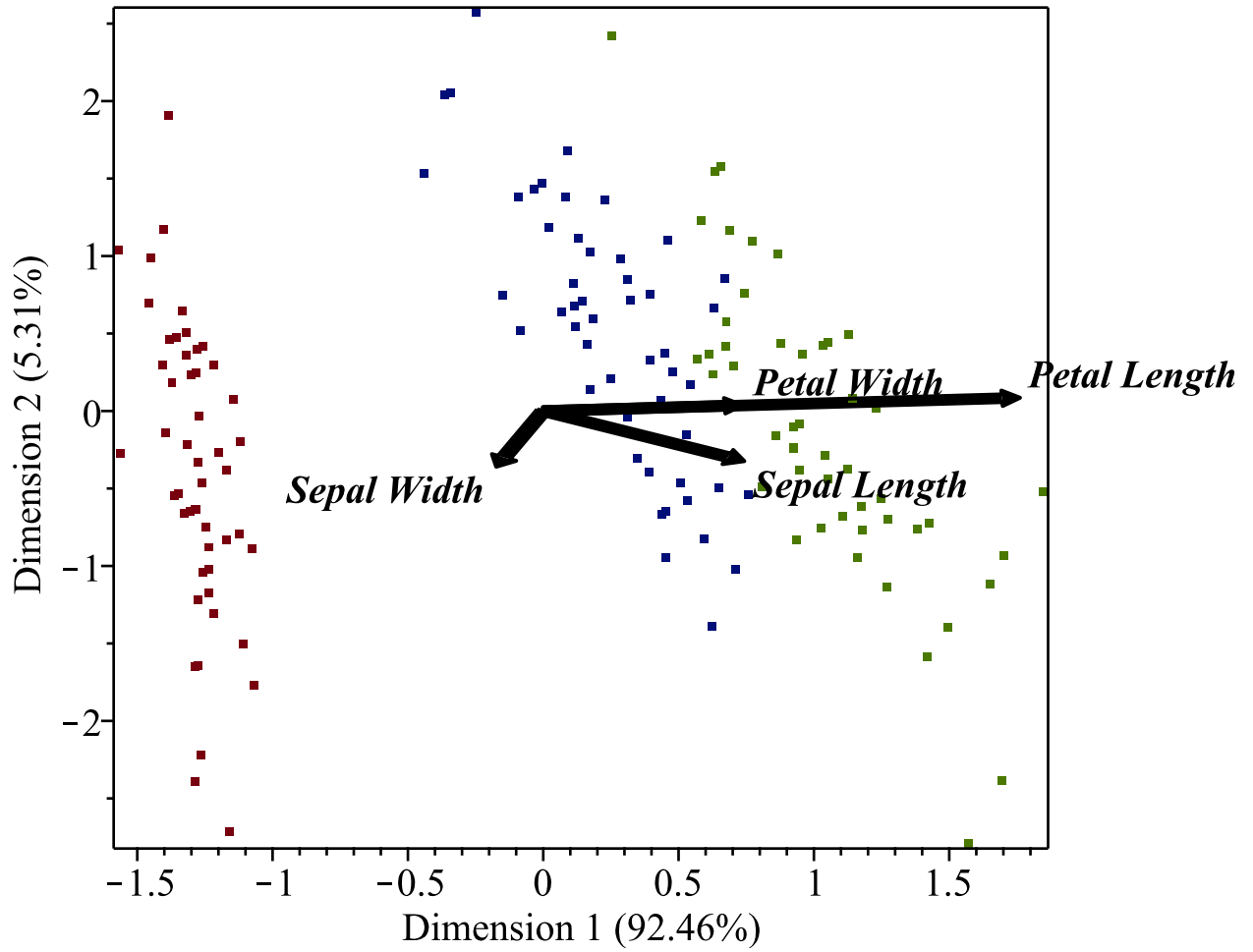
There are also two new visualizations related to [Principal Component Analysis](#): [Biplot](#), and [ScreePlot](#).

```
IrisDF := Import(FileTools:-JoinPath(["datasets", "iris.csv"], base = datadir) )
```

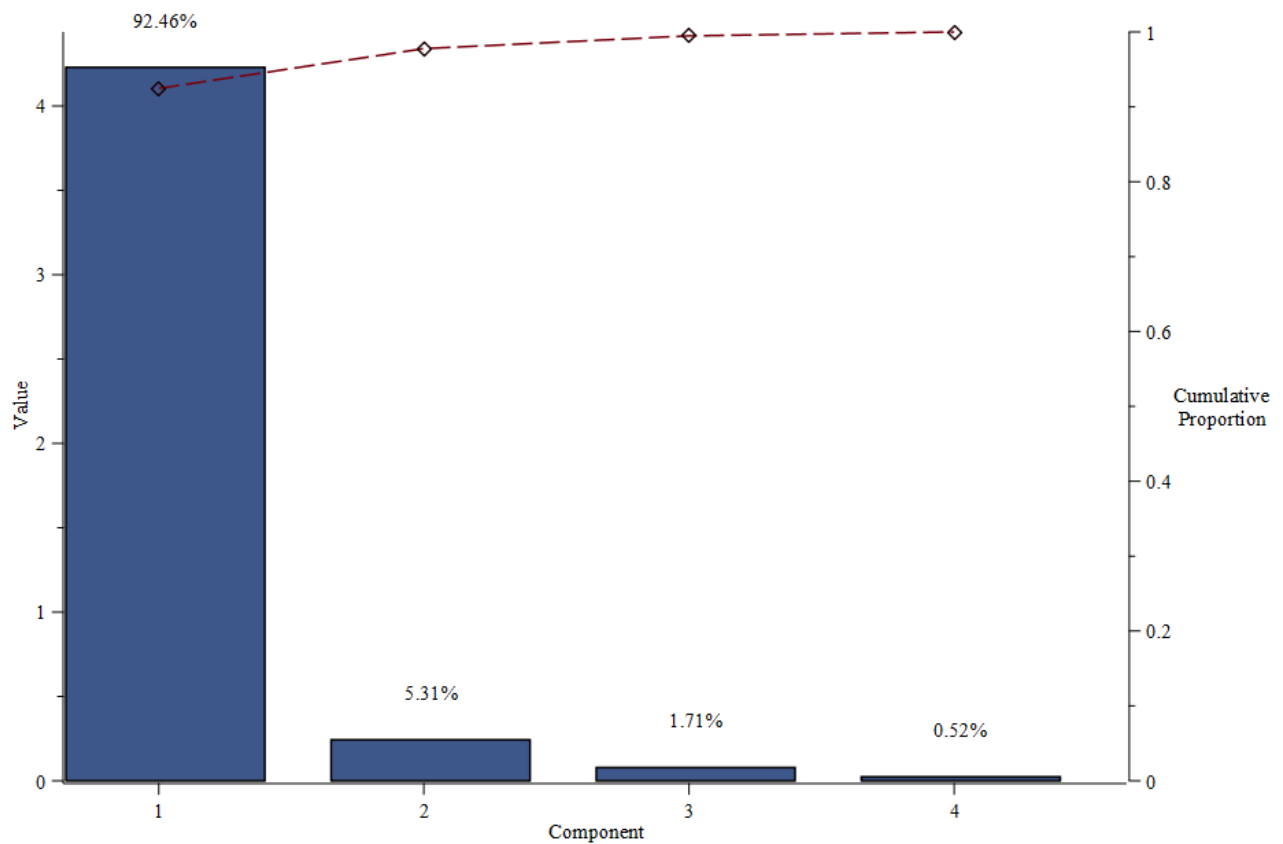
	<i>Sepal Length</i>	<i>Sepal Width</i>	<i>Petal Length</i>	<i>Petal Width</i>	<i>Species</i>
1	5.1	3.5	1.4	0.2	"setosa"
2	4.9	3	1.4	0.2	"setosa"
3	4.7	3.2	1.3	0.2	"setosa"
4	4.6	3.1	1.5	0.2	"setosa"
5	5	3.6	1.4	0.2	"setosa"
6	5.4	3.9	1.7	0.4	"setosa"
7	4.6	3.4	1.4	0.3	"setosa"
8	5	3.4	1.5	0.2	"setosa"
...

```
Biplot(IrisDF[['Sepal Length', 'Sepal Width', 'Petal Length', 'Petal Width']],
```

```
colorscheme = ["valuesplit", IrisDF[ `Species` ]], size = [600, "golden"])
```



```
ScreePlot(IrisDF[[ `Sepal Length`, `Sepal Width`, `Petal Length`, `Petal Width` ]])
```

The new [GridPlot](#) command is useful for visualizing multidimensional datasets. GridPlot generates a matrix of plots corresponding to the columns of a dataset.

```
GridPlot(IrisDF[ ['Sepal Length', 'Sepal Width', 'Petal Length', 'Petal Width']], upper
= ScatterPlot, lower = SunflowerPlot, width = 600, widthmode = pixels);
```

