CORRELATION ANALYSIS

Example 1:

```
proc contents data=sashelp._all_;
run;
proc print data=sashelp.class;
run;
```

	The SAS System							
Ob	s	Name	Sex	Age	Height	Weight		
	1	Alfred	М	14	69.0	112.5		
	2	Alice	F	13	56.5	84.0		
	3	Barbara	F	13	65.3	98.0		
	4	Carol	F	14	62.8	102.5		
	5	Henry	M	14	63.5	102.5		
	6	James	M	12	57.3	83.0		
	7	Jane	F	12	59.8	84.5		
	8	Janet	F	15	62.5	112.5		

Contains 19 observation

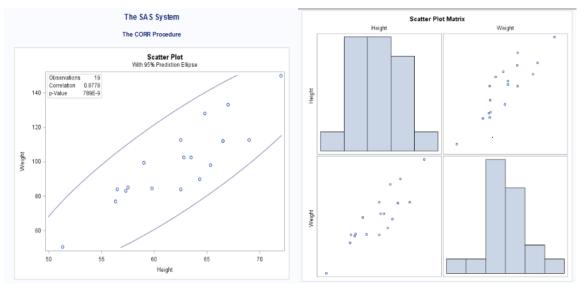
```
proc corr data=sashelp.class;
    var height weight;
run;
```

The CORR Procedure								
	2 Variables: Height Weight							
Simple Statistics								
Variable	N	Mean	Std I	Dev	Sum	Mini	mum	Maximum
Height	19	62.33684	5.12	708	1184	51.3	0000	72.00000
Weight	19	100.02632	22.77	393	1901	50.5	0000	150.00000
	Pe	earson Cor Prob	relatio > r und				N = 19	
		Height Weight					t	
	Н	eight	1	.000	00	-	87779 <.0001	
	W	eight	0.87779			1.	00000	1

Since the Pearson correlation coefficient is 0.87779, we can conclude that there is strong positive relationship between height and weight.

Since the probability value <.0001 we can conclude that correlation is highly significant.

proc corr data=sashelp.class plots=matrix (histogram) plot=scatter();
var height weight; run;



Scatter plot shows that the weight and height are linearly related as weight increase, height also increase.

EXAMPLE 2:

proc print data=sashelp.iris;
run;

Species	SepalLength	SepalWidth	PetalLength	PetalWidth
Setosa	50	33	14	2
Setosa	46	34	14	3
Setosa	46	36	10	2
Setosa	51	33	17	5
Setosa	55	35	13	2
	Setosa Setosa Setosa Setosa	Setosa 50 Setosa 46 Setosa 46 Setosa 51	Setosa 50 33 Setosa 46 34 Setosa 46 36 Setosa 51 33	Setosa 50 33 14 Setosa 46 34 14 Setosa 46 36 10 Setosa 51 33 17

NOTE: There were 150 observations read from the data set SASHELP.IRIS.

```
proc corr data=sashelp.iris;
run;
```

The CORR Procedure 4 Variables: SepalLength SepalWidth PetalLength PetalWidth

Simple Statistics								
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum	Label	
SepalLength	150	58.43333	8.28066	8765	43.00000	79.00000	Sepal Length (mm)	
SepalWidth	150	30.57333	4.35866	4586	20.00000	44.00000	Sepal Width (mm)	
PetalLength	150	37.58000	17.65298	5637	10.00000	69.00000	Petal Length (mm)	
PetalWidth	150	11.99333	7.62238	1799	1.00000	25.00000	Petal Width (mm)	

Pearson Correlation Coefficients, N = 150 Prob > r under H0: Rho=0							
	SepalLength	SepalWidth	PetalLength	PetalWidth			
SepalLength	1.00000	-0.11757	0.87175	0.81794			
Sepal Length (mm)		0.1519	<.0001	<.0001			
SepalWidth	-0.11757	1.00000	-0.42844	-0.36613			
Sepal Width (mm)	0.1519		<.0001	<.0001			
PetalLength	0.87175	-0.42844	1.00000	0.96287			
Petal Length (mm)	<.0001	<.0001		<.0001			
PetalWidth	0.81794	-0.36613	0.96287	1.00000			
Petal Width (mm)	<.0001	<.0001	<.0001				

proc corr data=sashelp.iris;
var sepallength petallength;
run;

The CORR Procedure

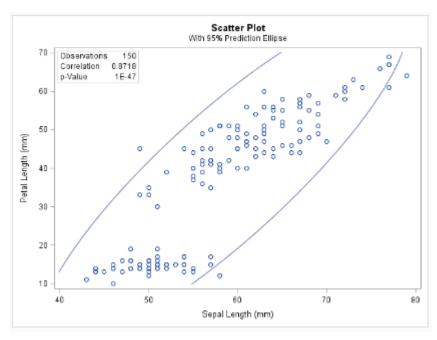
2 Variables: SepalLength PetalLength

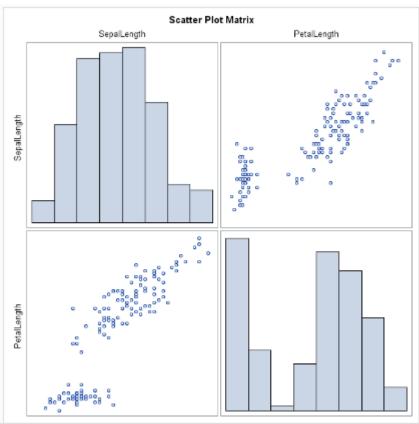
Simple Statistics							
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum	Label
SepalLength	150	58.43333	8.28066	8765	43.00000	79.00000	Sepal Length (mm)
PetalLength	150	37.58000	17.65298	5637	10.00000	69.00000	Petal Length (mm)

Pearson Correlation Coefficients, N = 150 Prob > r under H0: Rho=0						
SepalLength PetalLength						
SepalLength Sepal Length (mm)	1.00000	0.87175 <.0001				
PetalLength Petal Length (mm)	0.87175 <.0001	1.00000				

Based on the Pearson correlation coefficient, the value is 0.87175. Thus, we can conclude that there is strong positive relationship between sepal length and petal length.

proc corr data=sashelp.iris plots=matrix(histogram) plots=scatter();
var sepallength petallength;
run;





From the scatterplot of the variable sepal length and petal length, we can see that the points going from bottom left to the upper right, which is saying the correlation is positive.

EXAMPLE 3:

Use fish data from SAS help

title 'Fish Measurement Data';
proc corr data=sashelp.fish nomiss plots=matrix(histogram);
var Height Width Length3 Weight3;
run;

Fish Measurement Data

The CORR Procedure

4 Variables: Height Width Length3 Weight3

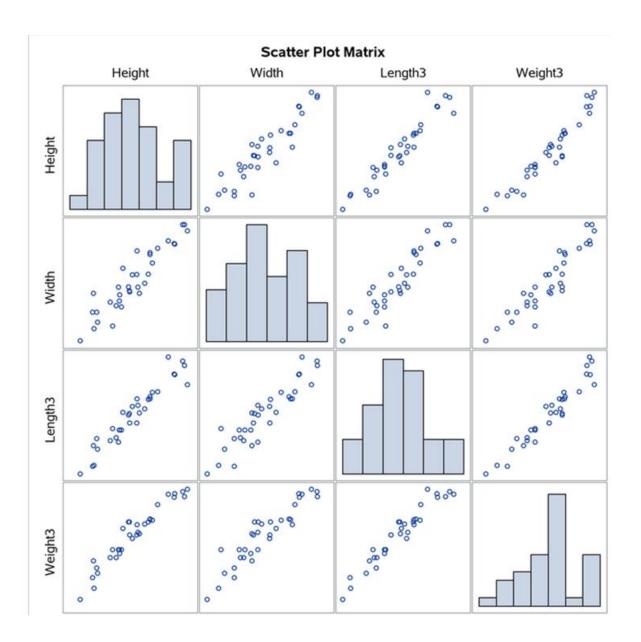
	Simple Statistics								
Variable	N	Mean	Std Dev	Sum	Minimum	Maximum			
Height	34	15.22057	1.98159	517.49950	11.52000	18.95700			
Width	34	5.43805	0.72967	184.89370	4.02000	6.74970			
Length3	34	38.38529	4.21628	1305	30.00000	46.50000			
Weight3	34	8.44751	0.97574	287.21524	6.23168	10.00000			

Output 2.8.2: Pearson Correlation Coefficients

Pearson Correlation Coefficients, N = 34 Prob > r under H0: Rho=0							
	Height	Width	Length3	Weight3			
Height	1.00000	0.92632 <.0001	0.95492 <.0001	0.96261 <.0001			
Width	0.92632 <.0001	1.00000	0.92171 <.0001	0.92789 <.0001			
Length3	0.95492 <.0001	0.92171 <.0001	1.00000	0.96523 <.0001			
Weight3	0.96261 <.0001	0.92789 <.0001	0.96523 <.0001	1.00000			

Since the Pearson correlation coefficient is more than 0.7, we can conclude that there is strong positive relationship between height, width and length.

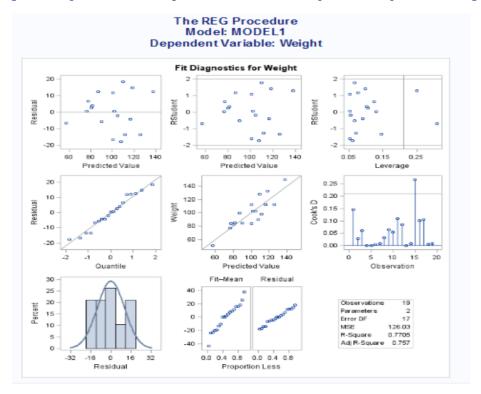
Since the Pearson correlation coefficient is 0.87779, we can conclude that there is strong positive relationship between height and weight.



REGRESSION ANALYSIS

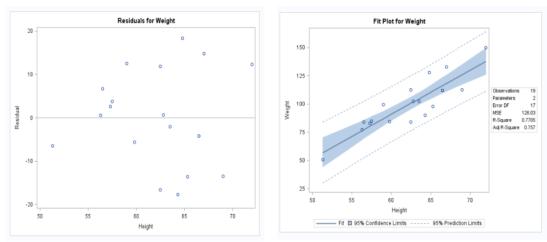
Example 1:

proc reg data=sashelp.class; model weight = height; run; quit;



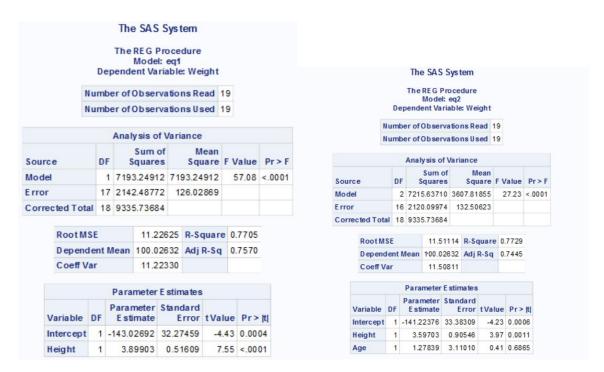
From residual versus predicted value plot, we can see that the data are randomly scattered which indicate that the regression is independent and has constant variance.

From histogram of percent versus residual plot, we can see that the data are the data is normally distributed.



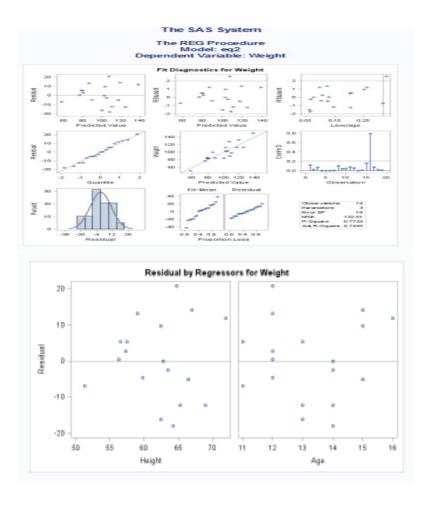
Plot of residual versus height show that data is linear.

```
proc reg data=sashelp.class rsquare;
  eq1: model weight=height;
  eq2: model weight=height age;
run;
```



Model for eq1 is significant as the p-value is less than 0.0001. Height is significant in estimating weight as the p-value is less than 0.0001. The coefficient parameter for intercept and height are -143.02692 and 3.89903 respectively.

Model for eq2 also significant as the p-value is less than 0.0001. Height is significant in estimating weight as the p-value is less than 0.0001While age is not significant in estimating weight as the p-value is 0.6865 greater than 0.05. The coefficient parameter for intercept, height and age are -141.122376, 3.59703and 1.27839 respectively.



Plot of residual versus height and residual versus age show random patterns which indicate that the models are independent and have constant variance.

REFERENCES

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