

Iris Data Set

TL

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Description

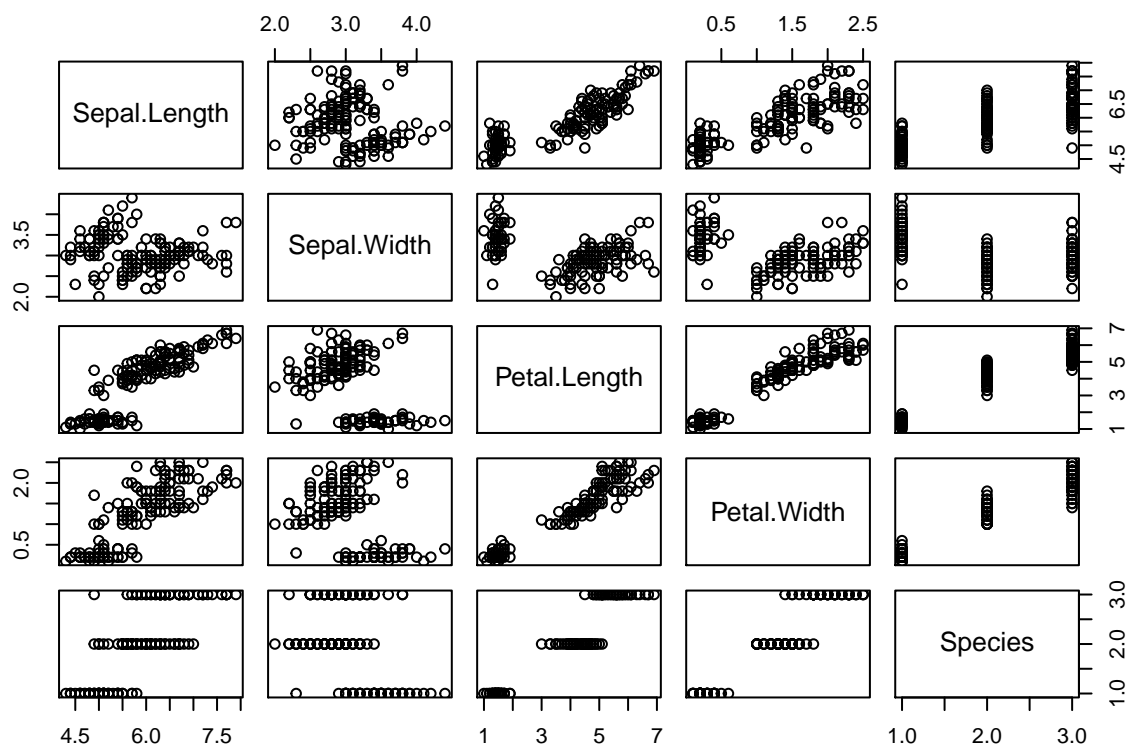
```
summary(iris)
```

```
##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
## Min.      :4.300    Min.      :2.000    Min.      :1.000    Min.      :0.100
## 1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
## Median :5.800    Median :3.000    Median :4.350    Median :1.300
## Mean   :5.843    Mean   :3.057    Mean   :3.758    Mean   :1.199
## 3rd Qu.:6.400    3rd Qu.:3.300    3rd Qu.:5.100    3rd Qu.:1.800
## Max.   :7.900    Max.   :4.400    Max.   :6.900    Max.   :2.500
##      Species
## setosa      :50
## versicolor:50
## virginica   :50
##
##
##
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100	setosa :50
2	1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300	versicolor:50
3	Median :5.800	Median :3.000	Median :4.350	Median :1.300	virginica :50
4	Mean :5.843	Mean :3.057	Mean :3.758	Mean :1.199	
5	3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800	
6	Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500	

Table 1: Summary of the data set

Scatterplot Matrix



```
iris[['Is.Versicolor']] <- as.numeric(iris[['Species']] == 'versicolor')
iris[['Is.Virginica']] <- as.numeric(iris[['Species']] == 'virginica')
fit.1 <- glm(Is.Versicolor ~ Petal.Length + Sepal.Length, data = iris)
fit.2 <- glm(Is.Virginica ~ Petal.Length + Sepal.Length, data = iris)
#summary(fit)
output <- capture.output(stargazer(fit.1, fit.2, title = 'Regression Results', summary=FALSE, header=F))
cat(output)
```

Table 2: Regression Results

	<i>Dependent variable:</i>	
	Is.Versicolor	Is.Virginica
	(1)	(2)
Petal.Length	0.148*** (0.043)	0.184*** (0.031)
Sepal.Length	-0.230** (0.092)	0.023 (0.067)
Constant	1.119*** (0.407)	-0.490* (0.294)
Observations	150	150
Log Likelihood	-94.823	-45.951
Akaike Inf. Crit.	195.646	97.902
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01	