

# Edgar Anderson's Iris Data

## Contents

<b>Description</b>	<b>1</b>
<b>Usage</b>	<b>1</b>
<b>Format</b>	<b>1</b>
<b>Source</b>	<b>2</b>
<b>Examples</b>	<b>2</b>

## Description

This famous (Fisher's or Anderson's) iris data set gives the measurements in centimeters of the variables sepal length and width and petal length and width, respectively, for 50 flowers from each of 3 species of iris. The species are *Iris setosa*, *versicolor*, and *virginica*.

## Usage

```
iris
```

## Format

iris is a data frame with 150 cases (rows) and 5 variables (columns) named:

- **Sepal.Length**
- **Sepal.Width**
- **Petal.Length**
- **Petal.Width**
- **Species**

## Source

Anderson, Edgar (1935). “The irises of the Gaspé Peninsula.” *Bulletin of the American Iris Society*, **59**: 2–5.

Fisher, Ronald A. (1936). “The use of multiple measurements in taxonomic problems.” *Annals of Eugenics*, **7** (Part **II**): 179–188.

## Examples

We investigate the Sepal and Petal leaves for the three species in the Iris data:

```
summary(iris)
```

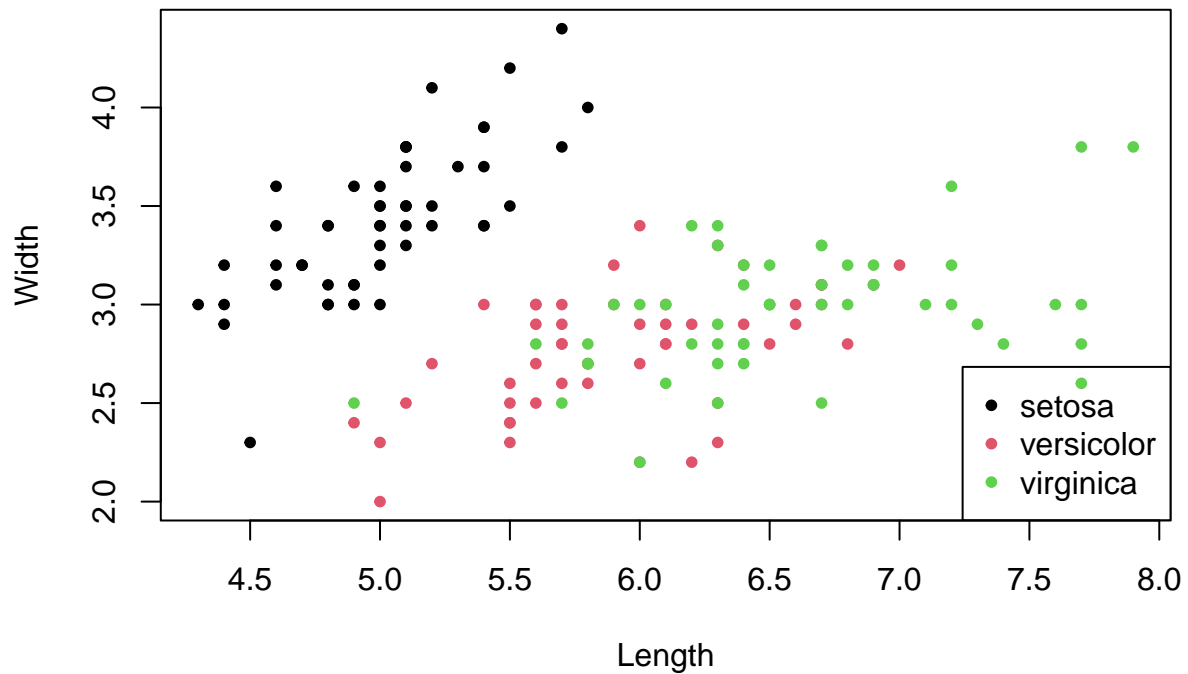
```
##      Sepal.Length      Sepal.Width      Petal.Length      Petal.Width
##  Min.       :4.300    Min.       :2.000    Min.       :1.000    Min.       :0.100
##  1st Qu.:5.100    1st Qu.:2.800    1st Qu.:1.600    1st Qu.:0.300
##  Median :5.800    Median :3.000    Median :4.350    Median :1.300
##  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
##
##
##
```

To examine the Sepal leaves, we select the length and the width:

```
llen <- iris$Sepal.Length
lwid <- iris$Sepal.Width
```

Then we plot the data:

```
plot(llen, lwid, xlab = "Length", ylab = "Width",
     pch = 20, col = as.numeric(iris$Species))
legend("bottomright", legend = levels(iris$Species), col = 1:3, pch = 20)
```



We can also select the Petal leaves:

```
llen <- iris$Petal.Length
lwid <- iris$Petal.Width
```

This gives us the following plot:

