A Test File

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September 17, 2005

A simple example that will run in any S engine: The integers from 1 to 10 are

```
[1] 1 2 3 4 5 6 7 8 9 10
```

We can also emulate a simple calculator:

```
> 1 + 1
[1] 2
> 1 + pi
[1] 4.141593
> sin(pi/2)
[1] 1
```

Now we look at Gaussian data:

One Sample t-test

```
data: x
t = 0.4896, df = 19, p-value = 0.63
alternative hypothesis: true mean is not equal to 0
95 percent confidence interval:
   -0.3163053   0.5094767
sample estimates:
   mean of x
0.09658574
```

Note that we can easily integrate some numbers into standard text: The third element of vector \mathbf{x} is -0.482273282274676, the p-value of the test is 0.63001.

Now we look at a summary of the famous iris data set, and we want to see the commands in the code chunks. Note that the summary needs to be print()ed explicitly, because eval would discard it otherwise. I consider this a feature, because it allows for much finer control on what gets into the final report.

```
> data(iris)
> print(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

> library(graphics)

> pairs(iris)

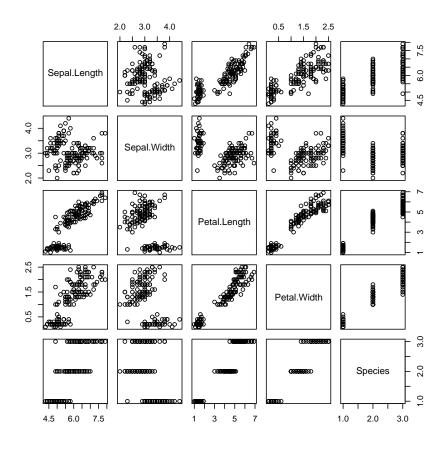


Figure 1: Pairs plot of the iris data.

> boxplot(Sepal.Length ~ Species, data = iris)

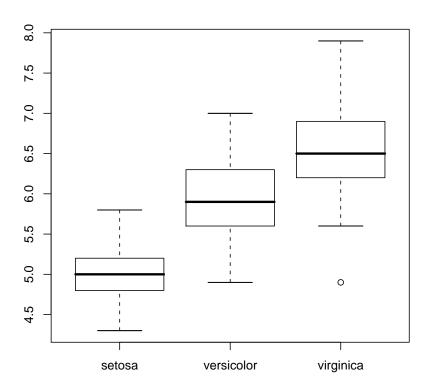


Figure 2: Boxplot of sepal length grouped by species.