

# Making a simple dataframe

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In this walkthrough we'll take a simple dataset that is presented as lists of separate numbers (eg 1,2,3,4) and put them into vectors. We'll then take these vectors and make simply dataframe. We'll then make a boxplot using ggplot2 and ggpubr.

## Working with data in 2 seperate vectors

- Data often come in 2 seperate sets of numbers, such as “the control group had lengths 4, 5, 6, 3, and 3, while the treatment group had length 11, 12, 6, 4 and 7”.
- One way to work with data like this in R is to load each set of numbers into a “vector” using the `c()` function

```
control <- c(4, 5,6,3,3)
treatment <- c(11,12,6,4,7)
```

## Summary Statistics

We can calculate the mean and other summary statistics of each vector on its own

```
mean(control)
```

```
## [1] 4.2
```

```
sd(control)
```

```
## [1] 1.30384
```

```
summary(control)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##       3.0     3.0     4.0     4.2     5.0     6.0
```

## Making a dataframe

To plot data its is almost always best to set it up in a dataframe.

Starting with the raw data we could make a dataframe like this

```
dat <- data.frame(length = c(4, 5,6,3,3,
                             11,12,6,4,7),
                  group = c("C","C","C","C","C",
                             "T","T","T","T","T"))
```

We can make a dataframe from the two vectors we've already typed up like this by surrounding them with a `c()` like this " `c(control, treatment)` "

```
dat <- data.frame(length = c(control,
                             treatment),
                  group = c("C","C","C","C","C",
                             "T","T","T","T","T"))
```

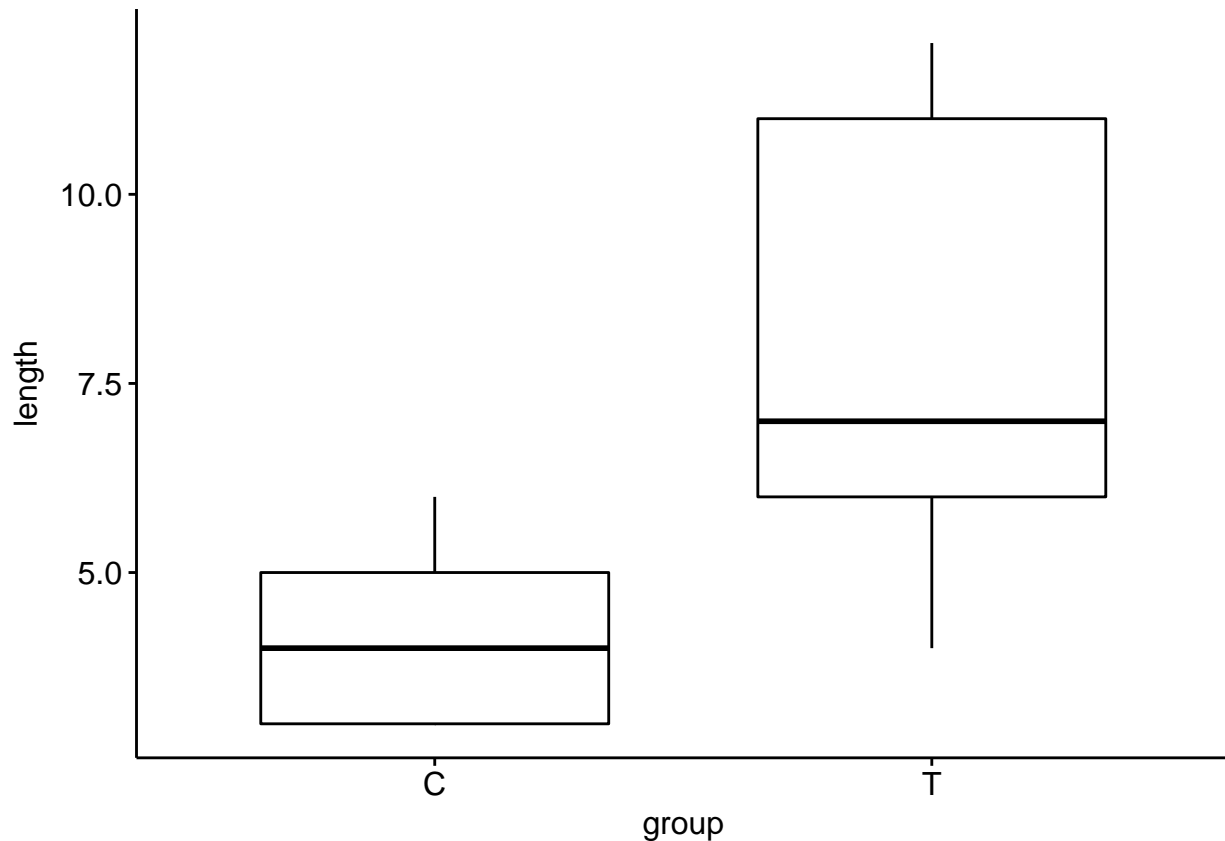
## Making a boxplot

We can make a boxplot from the data like this using ggpubr

```
library(ggplot2)
library(ggpubr)
```

```
## Loading required package: magrittr
```

```
ggboxplot(data = dat,
  y = "length",
  x = "group")
```

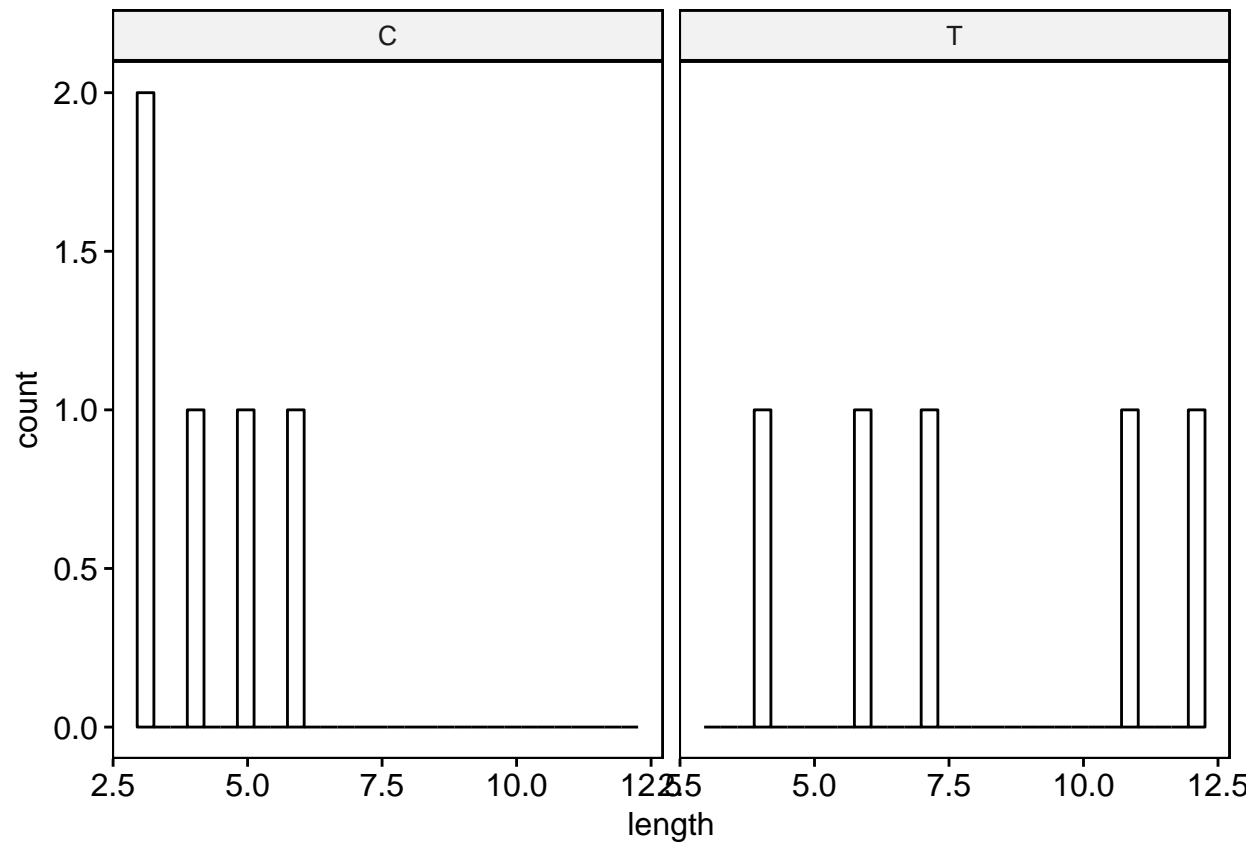


## Making a histogram

- An alternative to using a boxplot is a histogram.
- We can split up the data by treatment using facet.by = “group”
- Because my sample dataset has so few datapoints this doesn’t look very good.

```
gghistogram(data = dat,
  x = "length",
  facet.by = "group")
```

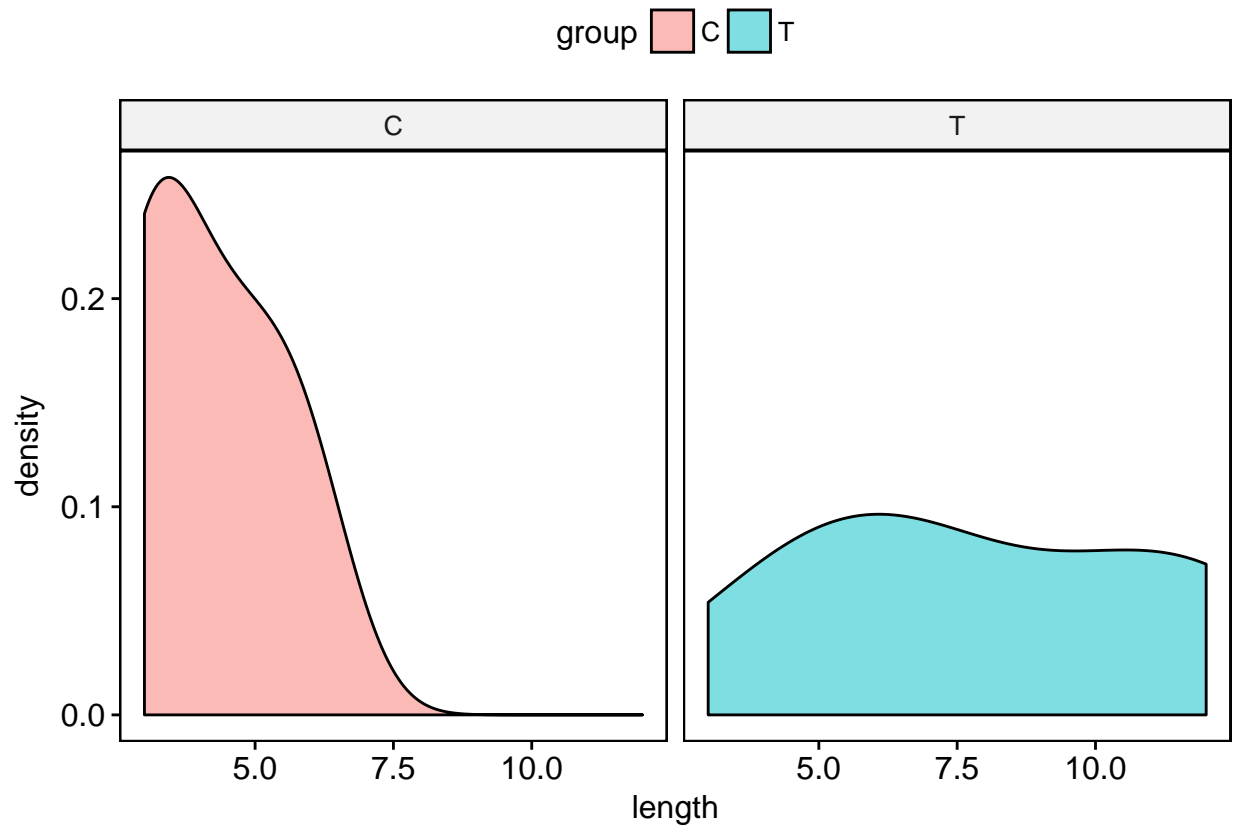
```
## Warning: Using `bins = 30` by default. Pick better value with the argument
## `bins`.
```



### Making a density plot

A plot that is similar to a histogram is a density plot.

```
ggdensity(data = dat,  
  x = "length",  
  facet.by = "group",  
  fill = "group")
```



### Advanced

You can save yourself some typing by using the `rep()` command when making your dataframe since the group codes “C” and “T” get repeated. IF you do this you need to make sure you keep track of all your parentheses.

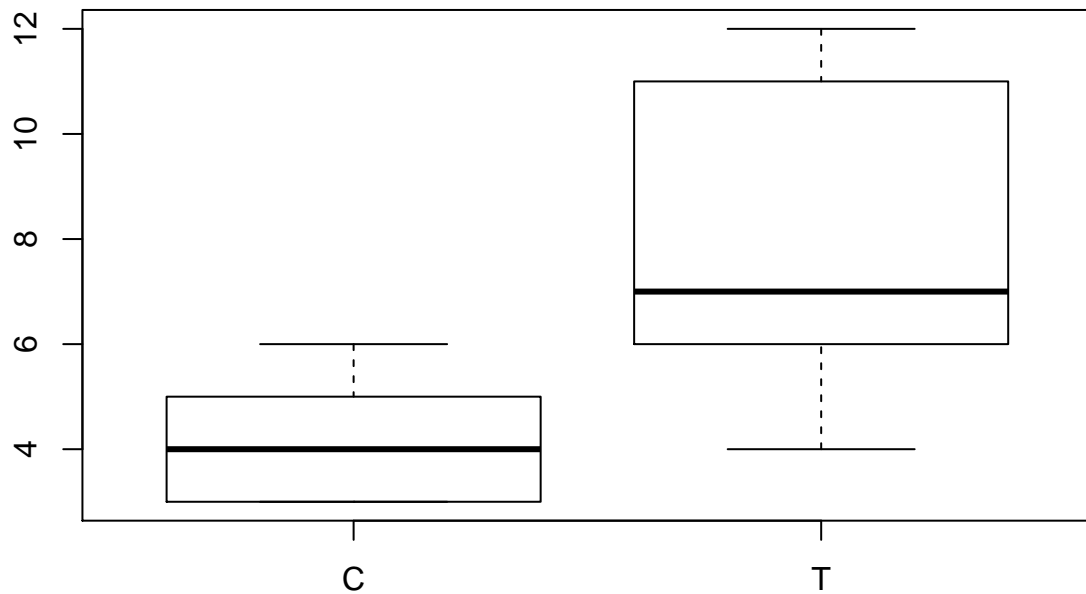
```
dat <- data.frame(length = c(control,
                             treatment),
                  group = c(rep("C",5),
                             rep("T",5)))
```

### Historical reference

#### Base R

In the old days (eg 2008...) we used to make our boxplot using the `boxplot` function.

```
boxplot(length ~ group, data = dat)
```



In the less old days (eg 2015) we used to make our boxplots using regular qplot or ggplot

#### ggplot's qplot

```
qplot(data = dat,  
      y = length,  
      x = group,  
      geom = "boxplot")
```

#### Standard ggplot

```
ggplot(data = dat,  
      aes(y = length,  
          x = group)) +  
  geom_boxplot()
```