

Cleaning__170523

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We often wish to tidy and reshape a dataset so that we can create certain plots. Here I introduce the two packages **tidyr** and **reshape2** to help the need and also to see how functions in **tidyr** and **reshape2** overlap and differ.

We first compare the functions `gather()`, `separate()` and `spread()`, from **tidyr**, with the functions `melt()`, `colsplit()` and `dcast()`, from **reshape2**.

The original dataset

```
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## 1         5.1         3.5         1.4         0.2   setosa
## 2         4.9         3.0         1.4         0.2   setosa
## 3         4.7         3.2         1.3         0.2   setosa
## 4         4.6         3.1         1.5         0.2   setosa
## 5         5.0         3.6         1.4         0.2   setosa
## 6         5.4         3.9         1.7         0.4   setosa
```

tidyr package

gather {tidyr}: takes multiple columns and collapses into key-value pairs, duplicating all other columns as needed. You use `gather()` when you notice that you have columns that are not variables.

Simply put, `gather()` takes wide-format data and turns it into long-format data

```
iris.tidyr <- iris %>%
  gather(key,value,-Species)
```

```
##   Species      key value
## 1  setosa Sepal.Length  5.1
## 2  setosa Sepal.Length  4.9
## 3  setosa Sepal.Length  4.7
## 4  setosa Sepal.Length  4.6
## 5  setosa Sepal.Length  5.0
## 6  setosa Sepal.Length  5.4
```

Our next step is to split the column `key` into two different columns: Part of a flower (Sepal or Petal) and Measure of that part (Length or Width), hence we use `separate()` function.

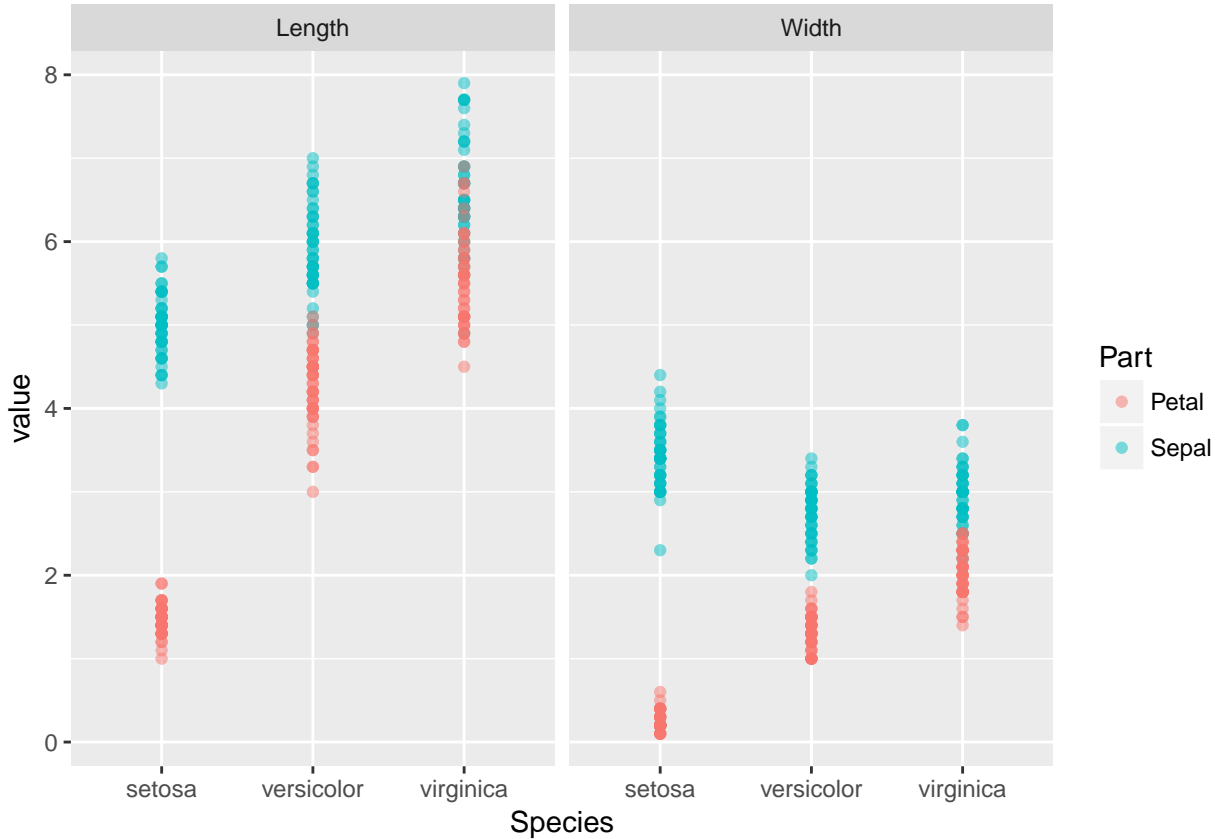
separate {tidyr}: turns a single character column into multiple columns.

```
iris.tidyr <- iris %>%
  gather(key,value,-Species) %>%
  separate(key,into=c("Part","Measure"),sep="\\.\\.\\.")
```

```
##   Species Part Measure value
## 1  setosa Sepal  Length  5.1
## 2  setosa Sepal  Length  4.9
## 3  setosa Sepal  Length  4.7
## 4  setosa Sepal  Length  4.6
## 5  setosa Sepal  Length  5.0
## 6  setosa Sepal  Length  5.4
```

With this dataset structure, we now can create a plot as shown below.

```
iris.tidyr %>%
  ggplot(aes(x = Species, y = value, col = Part)) +
  geom_point(alpha = 0.5) +
  facet_grid(. ~ Measure)
```



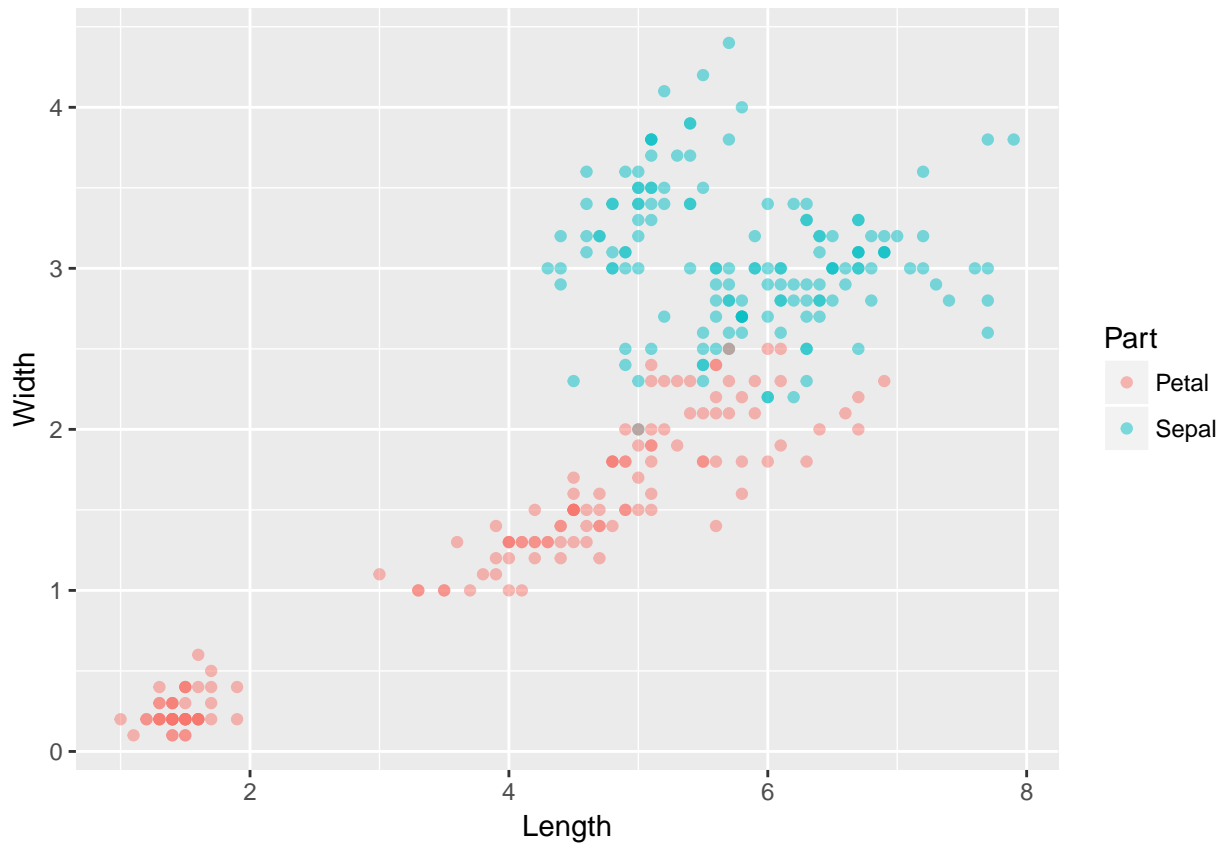
spread {tidyr}: spreads a key-value pair across multiple columns. In contrast to `gather()`, `spread()` takes long-format data and turns it into wide-format data.

```
iris$Flower <- 1:nrow(iris)
iris.tidyr <- iris %>%
  gather(key, value, - Species, - Flower) %>%
  separate(key, c("Part", "Measure"), "\\.") %>%
  spread(Measure, value)
```

```
##   Species Flower  Part Length Width
## 1  setosa      1  Petal   1.4   0.2
## 2  setosa      1  Sepal   5.1   3.5
## 3  setosa      2  Petal   1.4   0.2
## 4  setosa      2  Sepal   4.9   3.0
## 5  setosa      3  Petal   1.3   0.2
## 6  setosa      3  Sepal   4.7   3.2
```

With this dataset structure, we now can create a plot as shown below.

```
iris.tidyr %>%
  ggplot(aes(x=Length,y=Width,col=Part)) +
  geom_point(alpha=0.5)
```



reshape2 package

melt {reshape2}: converts an object into a molten data frame, giving same result with the gather() function from tidyr.

However, gather() cannot handle matrices or arrays, while melt() can!

```
iris.re <- iris %>%
  melt(id.vars="Species")
```

```
##   Species      variable value
## 1  setosa Sepal.Length   5.1
## 2  setosa Sepal.Length   4.9
## 3  setosa Sepal.Length   4.7
## 4  setosa Sepal.Length   4.6
## 5  setosa Sepal.Length   5.0
## 6  setosa Sepal.Length   5.4
```

colsplit {reshape2}: splits variable names that is a combination of multiple variables.

Again, we can achieve the same result with separate() function from tidyr, however, colsplit() operates only on a single column so we use cbind() to insert the new two columns in the data frame. While separate() performs all the operation at once.

```
iris$Flower <- 1:nrow(iris)
iris.re <- iris %>%
  melt(id.vars=c("Species", "Flower"))
iris.re = cbind(Species=iris.re[,1],
                Flower=iris.re[,2],
```

```
colsplit(iris.re[,3], "\\.", c("Part", "Measure")),
value=iris.re[,4])
```

```
##   Species Flower  Part Measure value
## 1  setosa      1 Sepal  Length  5.1
## 2  setosa      2 Sepal  Length  4.9
## 3  setosa      3 Sepal  Length  4.7
## 4  setosa      4 Sepal  Length  4.6
## 5  setosa      5 Sepal  Length  5.0
## 6  setosa      6 Sepal  Length  5.4
```

Again, the same result produced by `spread()` from `tidyr` can be obtained using `dcast()` from `reshape2` by specifying the correct formula.

cast {reshape2}: casts a molten data frame into an array or data frame.

```
iris.re = dcast(iris.re, formula=Flower+Species+Part ~Measure)
```

```
##   Flower Species  Part Length Width
## 1      1  setosa Petal    1.4    0.2
## 2      1  setosa Sepal    5.1    3.5
## 3      2  setosa Petal    1.4    0.2
## 4      2  setosa Sepal    4.9    3.0
## 5      3  setosa Petal    1.3    0.2
## 6      3  setosa Sepal    4.7    3.2
```

Sources:

<https://blog.rstudio.org/2014/07/22/introducing-tidyr/>

<http://www.milanor.net/blog/reshape-data-r-tidyr-vs-reshape2/>