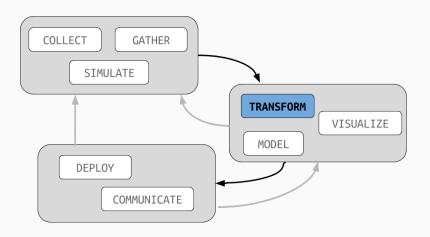
Lecture 25: Restructuring data

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In some cases we have datasets where multiple columns could be treated as individual observations. What does that mean? Think of the cancer dataset we used earlier in the semester, taking off just the cancer incidence rates:

cancer

```
## # A tibble: 1,961 x 5
##
   breast colorectal prostate lung melanoma
##
    <dbl>
             <dbl> <dbl> <dbl>
                                <dbl>
## 1
    127.8
             40.9 113.0 61.5
                                 14.5
## 2
    133.2 39.2 82.1 58.1
                                13.1
    101.6 36.9 117.9 35.1
                                13.7
## 3
## 4
    127.4 41.0 96.5 64.9
                                12.3
## 5 119.8
             37.5
                    121.4 69.7
                                15.0
## 6 150.8 46.2 110.6 74.9
                                25.4
## # ... with 1,955 more rows
```

A common question with this data is: how we could plot all of the cancer types on the same plot. The canonical way of doing this would be to make a new dataset where each row, instead of being a single county, is then just one incidence rate. In other words each county will have five rows associated with it. To do this we use the gather function. It is found in the package **tidyr**.

library(tidyr)

```
gather(cancer)
## # A tibble: 9,805 x 2
##
       key value
## <chr> <dbl>
## 1 breast 127.8
## 2 breast 133.2
## 3 breast 101.6
## 4 breast 127.4
## 5 breast 119.8
## 6 breast 150.8
## # ... with 9,799 more rows
```

If we want to give the name of the key' andvalue', those are given as the next two parameters to gather:

```
## # A tibble: 9,805 x 2
## type incidence
##
     <chr> <dbl>
## 1 breast 127.8
## 2 breast 133.2
## 3 breast 101.6
## 4 breast 127.4
## 5 breast 119.8
## 6 breast 150.8
## # ... with 9,799 more rows
```

gather(cancer, type, incidence)

In many cases, there are other variables that we want to be duplicated along with the other keys. For example, take the speed skating dataset, selecting off a few variables to make it a bit more tractable:

We can indicate the variables that should not be gathered by including them after the key and value terms with minus signs:

```
gather(speed, skater, value, -num_skater, -nationality)
```

```
## # A tibble: 23,520 x 4
##
    num_skater nationality skater value
                     <chr> <chr> <chr> <dbl>
##
         <int>
## 1
           175
                       RUS time_lap1 6.91
           100
## 2
                       CAN time_lap1 6.69
## 3
           138
                       ITA time lap1 6.87
           182
## 4
                      TUR time lap1 7.30
          92
## 5
                       AUS time_lap1 7.00
## 6
           190
                      USA time lap1
                                     6.69
## # ... with 2.351e+04 more rows
```

Or, we can not use the minus sign and include on those variables that should be gathered:

```
## # A tibble: 23,520 x 4
##
    num_skater nationality skater value
##
         <int>
                    <chr> <chr> <chr> <dbl>
## 1
           175
                      RUS time lap1 6.91
           100
                      CAN time_lap1 6.69
## 2
## 3
          138
                      ITA time lap1 6.87
## 4
        182
                      TUR time lap1 7.30
## 5
          92
                      AUS time_lap1 7.00
## 6
           190
                      USA time lap1 6.69
## # ... with 2.351e+04 more rows
```

The results are the same; in most cases one or the other will lead to less typing. There is also a complement to gathering called spreading, available through the function spread. You should not need that in this class because every dataset is maximally spread already. It is also a lot more difficult to use because you have to be careful about implicit missing values and what to do with them.