

# Data Science with R (Data Analytics)

**Basic Data Elements** 

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# **Outline For Today**

- Data Transformation
  - Reshape
  - Split
  - Combine
- Character Manipulation
- Dates and timestamps
- Accessing Database
- Getting Data from Web
  - API data sources
  - Connecting to an external database
- Other Data Resources



# **Data Transformation Overview**

#### **Subset**

Index operators [], subset(), and with() can be used to subset data according to various conditions.

```
#Three different ways to do the exact same thing!
data_sub = iris[iris$Species=='setosa', 3:5]
data_sub1 = subset(iris, Species=='setosa', 3:5)
data_sub2 = with(iris, iris[Species=='setosa', 3:5])
head(data_sub, 1); head(data_sub1, 1); head(data_sub2, 1)
```

1	Petal.Length 1.4	Petal.Width 0.2	Species setosa
1	Petal.Length 1.4	Petal.Width 0.2	Species setosa
1	Petal.Length 1.4	Petal.Width 0.2	Species setosa



#### **Transform**

The transform() function can be used to create a new column in an existing dataset.

```
iris_tr = transform(iris, v1=log(Sepal.Length))
head(iris_tr, 1)
```

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species v1
1 5.1 3.5 1.4 0.2 setosa 1.6292
```

```
#Equivalent:
iris_tr1 = iris
iris_tr1$v1 = log(iris$Sepal.Length)
head(iris_tr1, 1)
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species	v1
1	5.1	3.5	1.4	0.2	setosa	1.6292



#### **Discretize**

The **cut()** function can be used to transform a numeric variable into a categorical variable.

```
groupvec = quantile(iris\_tr$v1, c(0, 0.25, 0.50, 0.75, 1.0)) labels = c('A', 'B', 'C', 'D') iris\_tr$v2 = cut(iris\_tr$v1, breaks=groupvec, labels=labels, include.lowest=TRUE)
```

```
0% 25% 50% 75% 100%
1.458615 1.629241 1.757858 1.856298 2.066863
```

```
iris_tr[c(1,6), ]
```

groupvec

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species	v1	
v2							
1	5.1	3.5	1.4	0.2	setosa	1.629	Α
6	5.4	3.9	1.7	0.4	setosa	1.686	В



#### **Set Levels of a Factor**

```
vec = rep(c(0,1), c(4,6))
vec
```

[1] 0 0 0 0 1 1 1 1 1 1

```
#Converting to a factor and creating the labels/levels all at once.
vec_fac
vec_fac = factor(vec, labels=c('male','female'))
```

[1] male male male female female female female female female Levels: male female

#### **Set Levels of a Factor**

```
#First converting to a factor.

vec1 = factor(rep(c(0,1,3), c(4,6,2)))

vec1

[1] 0 0 0 0 1 1 1 1 1 1 3 3

Levels: 0 1 3
```

```
#Then creating the labels/levels.
levels(vec1) = c("male", "female", "male")
vec1
```

#### **Rename Levels of a Factor**

```
vec2 = factor(rep(c('b','a'), c(4,6)))
vec2
[1] b b b b a a a a a a
Levels: a b
levels(vec2)
[1] "a" "b"
relevel(vec2, ref='b') #Changing the reference level.
```

[1] b b b b a a a a a a Levels: b a



# **Data Reshape**

# What makes data wide or long?

Wide data has a column for each variable.

For example, this is **wide-format** data:

data = iris[, 1:4] #A wide dataset; columns are variables, rows are observations.

head(data, 5)

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
1	5.1	3.5	1.4	0.2
2	4.9	3.0	1.4	0.2
3	4.7	3.2	1.3	0.2
4	4.6	3.1	1.5	0.2
5	5.0	3.6	1.4	0.2

# What makes data wide or long?

Long-format data has a column for possible variable types and a column for the values of those variables.

For example, this is **Long-format** data:

	Species	variable	value
121	virginica	Sepal.Length	6.9
411	virginica	Petal.Length	5.1
549	versicolor	Petal.Width	1.1
170	setosa	Sepal.Width	3.8
63	versicolor	Sepal.Length	6.0
418	virginica	Petal.Length	6.7
314	setosa	Petal.Length	1.1
480	setosa	Petal.Width	0.2
567	virginica	Petal.Width	1.8
66	versicolor	Sepal.Length	6.7



#### Why we need to reshape the data?

Long-form data and wide-form data are used for different purposes in data analysis. The ultimate shape you want to get your data into will depend on what you are doing with it.

It turns out that you need wide-format data for some types of data analysis and long-format data for others. In reality, you need long-format data much more commonly than wide-format data. For example, ggplot2 requires long-format data (technically tidy data), plyr requires long-format data, and most modelling functions (such as lm(), glm(), and gam()) require long-format data. But people often find it easier to record their data in wide format.

So it's important to get comfortable converting back-and-forth between the two.

#### **Long and Wide-Form Data**

Use the **stack()** function to reshape to long form; use **unstack()** for wide form. Stacking vectors concatenates multiple vectors into a single vector along with a factor indicating where each observation originated. Unstacking reverses this operation.

```
data_I = stack(data) #A long dataset; treats all elements as pieces of data.
data w = unstack(data I) #A wide dataset; columns are variables, rows are
obs.
str(data_l); str(data_w)
'data.frame': 600 obs. of 2 variables:
$ values: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
$ ind : Factor w/ 4 levels "Petal.Length",..: 3 3 3 3 3 3 3 3 3 3 ...
'data.frame': 150 obs. of 4 variables:
$ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
$ Petal.Width: num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
$ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
$ Sepal.Width: num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
```



#### **Long and Wide-Form Data**

Let's take the last two columns of iris and reshape them to wide form.

```
subdata = iris[ ,4:5] #This subset of the overall data is in long form. subdata[c(1,51, 101), ]
```

```
Petal.Width Species

1 0.2 setosa

51 1.4 versicolor

101 2.5 virginica
```

```
subdata_w = unstack(subdata) #Each factor becomes a column in the wide
form.
```

head(subdata\_w, 1)

	setosa	versicolor	virginica
1	0.2	1.4	2.5



The reshape2 package lets you flexibly restructure and aggregate data using just two functions:

melt() (takes wide format data and "melts" it into long format data)cast() (takes long format data and "casts" it into wide format data).

- dcast() reshapes a molten data into a data frame
- acast() reshape a molten data into a vector/matrix/array

Think of working with metal: if you melt metal, it drips and becomes long. If you cast it into a mould, it becomes wide. Since you will most commonly work with data.frame objects, we'll explore the dcast function.

Let's take the long-format subset data and cast it into a wide-format data frame.

dcast uses a formula to describe the shape of the data. The arguments on the left refer to the ID variables and the arguments on the right refer to the measured variables.

```
#install.packages("reshape2")
library(reshape2)
                            #Cast long format to wide
format.
dcast(data=subdata,
                            #Specifying the data to manipulate.
   formula=Species ~ .,
                            #Species should be the main column; nothing else.
    value.var='Petal.Width',
                            #Values to fill in should come from Petal.Width.
    fun=mean)
                            #Aggregate the values by the mean.
    Species
    setosa
                  0.246
    versicolor 1.326
    virginica
                  2.026
```



To make use of the function we need to specify a data frame, the id variables (which will be left at their settings) and the measured variables (columns of data) to be stacked. The default assumption on measured variables is that it is all columns that are not specified as id variables.

```
iris_long = melt(data=iris, #Melt wide format to long format.
           id.vars = 'Species')
                                      #The main identification variable is
Species.
set.seed(5)
i = sample(nrow(iris_long), 5)
iris Iona[i, ]
                       variable
                                          value
         Species
121
         virginica
                       Sepal.Length
                                           6.9
411
                                           5.1
         virginica
                       Petal.Length
         versicolor
549
                       Petal.Width
                                           1.1
170
                       Sepal.Width
                                           3.8
         setosa
63
         versicolor
                        Sepal.Length
                                           6.0
```



Using dcast to get the mean (by species) of each variable:

```
dcast(data=iris_long,  #Specifying the data to manipulate.

formula=Species ~ variable,  #Species is main col.; swing variable col.

value.var = 'value',  #Values to fill in should come from value col.

fun=mean)  #Aggregate the values by the mean.
```

	Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
1	setosa	5.01	3.43	1.46	0.246
2	versicolor	5.94	2.77	4.26	1.326
3	virginica	6.59	2.97	5.55	2.026



# In-class exercise using Tips dataset

- 1. Come up one or two questions you want to answer from Tips dataset.
- 2. Present your code and results.



# **Example:** Is there a gender difference in the tipping habits?

Using the built-in tips dataset, get the mean tip amount by sex:

```
dcast(tips, sex ~ ., value.var='tip', fun=mean)
#Manipulate the tips data; sex is the main column; don't keep anything else; fill in
#values based on the tip column; aggregate the results by the mean.
```

```
sex .
Female 2.833448
```

2 Male 3.089618

Get means by sex and party size ("swing over" the size variable).

```
dcast(tips, sex ~ size, value.var='tip', fun=mean)
```

```
1
                         4
                                      6
sex
            2.53
                  3.25
                               5.14
Female 1.28
                         4.02
                                      4.60
Male
      1.92
            2.61
                   3.48
                         4.17
                               3.75
                                      5.85
```



# **Example:** Is there a gender difference in the tipping habits?

We want to aggregate the average total\_bill by sex. How can we do that?

```
dcast(tips, sex ~ . , value.var='total_bill', fun=mean)
```

```
sex .
```

- 1 Female 18.05690
- 2 Male 20.74408

# **Split and Combine Data**

#### **Merge Two Data Frames**

```
datax = data.frame(id=c(1,2,3), gender=c('M', 'F', 'M'))
datay = data.frame(id=c(3,1,2), name=c('tom','john','mary'))
datax; datay
        gender
    id
   1 M
2 2 F
3 3 M
    id
        name
   3 tom
2 1 john
   2
        mary
```

#### **Merge Two Data Frames**

Merging two data frames by common columns or row names (similar to *JOIN* in SQL):

```
merge(datax, datay, by='id')
```

```
id gender name
1 1 M john
2 2 F mary
3 3 M tom
```

# **Split Into Groups**

The **split()** function divides the data into groups defined by the factor specified in the second argument.

```
iris_split = split(iris, iris$Species)
class(iris_split)
```

```
[1] "list"
```

```
attributes(iris_split)
```

```
$names
[1] "setosa" "versicolor" "virginica"
```

```
str(iris_split)
```



#### **Split Into Groups**

```
List of 3
   $ setosa :'data.frame': 50 obs. of 5 variables:
    ..$ Sepal.Length: num [1:50] 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
    ..$ Sepal.Width: num [1:50] 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
    ..$ Petal.Length: num [1:50] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
    ..$ Petal.Width: num [1:50] 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
    ..$ Species : Factor w/ 3 levels "setosa", "versicolor", ..: 1 1 1 1 1 1 1 1 1 1 ...
   $ versicolor: 'data.frame': 50 obs. of 5 variables:
    ..$ Sepal.Length: num [1:50] 7 6.4 6.9 5.5 6.5 5.7 6.3 4.9 6.6 5.2 ...
    ..$ Sepal.Width: num [1:50] 3.2 3.2 3.1 2.3 2.8 2.8 3.3 2.4 2.9 2.7 ...
    ..$ Petal.Length: num [1:50] 4.7 4.5 4.9 4 4.6 4.5 4.7 3.3 4.6 3.9 ...
    ..$ Petal.Width: num [1:50] 1.4 1.5 1.5 1.3 1.5 1.3 1.6 1 1.3 1.4 ...
   ..$ Species : Factor w/ 3 levels "setosa", "versicolor", ..: 2 2 2 2 2 2 2 2 2 2 ...
   $ virginica: 'data.frame': 50 obs. of 5 variables:
    ..$ Sepal.Length: num [1:50] 6.3 5.8 7.1 6.3 6.5 7.6 4.9 7.3 6.7 7.2 ...
    ..$ Sepal.Width: num [1:50] 3.3 2.7 3 2.9 3 3 2.5 2.9 2.5 3.6 ...
    ..$ Petal.Length: num [1:50] 6 5.1 5.9 5.6 5.8 6.6 4.5 6.3 5.8 6.1 ...
    ..$ Petal.Width: num [1:50] 2.5 1.9 2.1 1.8 2.2 2.1 1.7 1.8 1.8 2.5 ...
                : Factor w/ 3 levels "setosa", "versicolor", ...: 3 3 3 3 3 3 3 3 3 3 ...
    ..$ Species
```



# 'Unsplit'

#### unsplit reverses the split operation:

```
iris_unsplit = unsplit(iris_split, iris$Species)
class(iris_unsplit)
```

```
[1] "data.frame"
```

```
iris_unsplit[c(1,51, 101), ]
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
51	7.0	3.2	4.7	1.4	versicolor
101	6.3	3.3	6.0	2.5	virginica



# **Split Example**

Using the tips dataset, let's split by sex and get some statistics from each subgroup.

```
library(reshape2) #Not using the reshape2 functions, just want the tips
dataset.
tips
tips_by_sex = split(tips, tips$sex)
head(tips_by_sex[[1]], 2)

total_bill tip sex smoker day time size
```

```
total_bill tip sex smoker day time size

1 17.0 1.01 Female No Sun Dinner 2

5 24.6 3.61 Female No Sun Dinner 4
```

```
ratio_fun = function(x) {
   sum(x$tip) / sum(x$total_bill)
}
```

# **Split Example**

Apply the <a href="ratio\_fun">ratio\_fun</a>() function to each member of the split list:

```
result = lapply(tips_by_sex, ratio_fun)
result
```

\$Female [1] 0.1569178

\$Male [1] 0.1489398

# **Character Manipulation**

#### **Character Manipulation**

Here are some basic functions for manipulating character data:

- nchar(): Count the number of characters
- strsplit(): Split the elements of a character vector
- paste(): Concatenate strings
- substr(): Substrings of a character vector
- gsub(): Replacement
- grep(): Pattern matching

#### **Character Functions:** *nchar*

#### Count the number of characters

fruit = 'apple orange grape banana'
nchar(fruit)

[1] 25



# **Character Functions:** *strsplit*

#### Split the elements of a character vector

```
split.string.list = strsplit(fruit, split=' ')
split.string.list
```

```
[[1]]
[1] "apple" "orange" "grape" "banana"
```

```
fruitvec = unlist(split.string.list)
fruitvec
```

```
[1] "apple" "orange" "grape" "banana"
```

#### **Character Functions:** *paste*

#### **Concatenate strings**

```
paste(fruitvec, collapse = ' ')
[1] "apple orange grape banana"

paste(fruitvec, collapse=',')
[1] "apple,orange,grape,banana"
```

```
paste(fruitvec, 'extra')
[1] "apple extra" "orange extra" "grape extra" "banana extra"

paste(fruitvec, 'extra', sep = '.')
[1] "apple.extra" "orange.extra" "grape.extra" "banana.extra"
```



#### **Example: Formulas Creation**

Paste is often useful when you want to create objects programmatically, such as generating a formula or creating variables on the fly.

```
n = 1:20
xvar = paste('x', n, sep = '')
right = paste(xvar, collapse = ' + ')
left = 'y ~'
my_formula = paste(left, right)
my_formula
```

```
[1] "y \sim x1 + x2 + x3 + x4 + x5 + x6 + x7 + x8 + x9 + x10 + x11 + x12 + x13 + x14 + x15 + x16 + x17 + x18 + x19 + x20"
```

#### **Character Functions:** *substr*

#### Substrings of a character vector

```
substr(fruit, 1, 5)
```

```
[1] "apple"
```

substr(fruit, 1, 7)

[1] "apple o"

substr(fruit, 9, 21)

[1] "ange grape ba"

## **Character Functions:** *gsub*

#### Replacement

gsub('apple', 'strawberry', fruit)

[1] "strawberry orange grape banana"

gsub('a', '?', fruit)

[1] "?pple or?nge gr?pe b?n?n?"

gsub('an', 'HA', fruit)

[1] "apple orHAge grape bHAHAa"

## **Character Functions:** *grep*

#### **Pattern Matching**

grep('grape', fruitvec)

[1] 3

grep('a', fruitvec)

[1] 1 2 3 4

grep('an', fruitvec)

[1] 2 4



We want to explore the information provided in packages on CRAN. The goal is to get a dataframe where one column contains the package name and another the author name.

```
#get packages table
#install.packages("XML")
library(XML)
web =
'http://cran.r-project.org/web/packages/available_packages_by_name.html'
packages = readHTMLTable(web, stringsAsFactors = FALSE)
```

Use str() to explore the packages object. The V1 column is what we need.

```
pnames = packages[[1]][ ,1]
length(pnames)
```

[1] 8586

```
pnames = pnames[2:11]
```

```
b = 'http://cran.r-project.org/web/packages/'
a = '/index.html'
urls = paste0(b, pnames, a)
```



Now urls is a character strings containing the ten web addresses. Let's explore one of them.

```
table = readHTMLTable(urls[1], stringsAsFactors=FALSE, header=FALSE)
info = table[[1]]
paste0(info$V1, info$V2)
```

- [1] "Version:2.0"
- [2] "Depends:R (≥ 2.10), nnet, quantreg, MASS, locfit"
- [3] "Published:2014-07-11"
- [4] "Author: Katalin Csillery, Louisiane Lemaire, Michael Blum and Olivier

Francois"

- [5] "Maintainer: Michael Blum < michael. blum at imag.fr>"
- [6] "License:GPL (≥ 3)"
- [7] "NeedsCompilation:no"
- [8] "In views:Bayesian"
- [9] "CRAN checks:abc results"



Get information on ten packages using lapply

```
tables = lapply(urls, readHTMLTable, stringsAsFactors=FALSE, header=FALSE) infos = lapply(tables, function(x) x[[1]]) infovec = lapply(infos, function(x) paste0(x$V1, x$V2))
```

Extract author names from infovec using grep and lapply. Finally, combine anamevec with pnames.

```
aname = lapply(infovec, function(x) x[grep('Author:', x)])
anamevec = lapply(aname, function(x) substr(x, 8, nchar(x)))
anamevec = unlist(anamevec)
name.df = data.frame(pnames, anamevec)
```

## **Manipulating Dates and Timestamps**

## **Two Types of Date-related Data**

- Date object: only contains date and time information
- Time Series object: normal data object with timestamp added

## **Date Objects**

- Date class: Only date; no time information
- POSIXct class: Time (with timezone) included with date

Use as.Date() to create date class object.

```
date1 = '1989-05-04'
date1 = as.Date(date1)
class(date1)
```

```
[1] "Date"
```

```
date1 = '05/04/1989'
date1 = as.Date(date1, format='%m/%d/%Y')
```

Manipulating a date class object:

date2 = date1 + 31

date2 - date1

Time difference of 31 days

date2 > date1

[1] TRUE

Count from 1970.01.01 in the date class object:

Sys.Date() - structure(0, class='Date')

Time difference of 16962 days

Create a vector of date class objects:

```
dates = seq(date1, length=4, by='day')
format(dates, '%w')
```

```
[1] "4" "5" "6" "0"
```

weekdays(dates)

```
[1] "Thursday" "Friday" "Saturday" "Sunday"
```



#### **POSIXct Class Object**

Use as.POSIXct to create POSIXct class objects.

```
time1 = '1989-05-04'

time1 = as.POSIXct(time1)

time1 = "2011-03-1 01:30:00"

time1 = as.POSIXct(time1, format="%Y-%m-%d %H:%M:%S")

time1 = as.POSIXct("2011-03-1 01:30:00", tz='GMT')

time2 = seq(from=time1, to=Sys.time(), by='month')
```

### **POSIXct Class Object**

Create a POSIXct class object based on a numeric object.

The ISOdatetime function can convert a numeric object into a POSIXct object.

```
time1 = ISOdatetime(2011,1,1,0,0,0)
rtimes = ISOdatetime(2013, rep(4:5,5), sample(30,10), 0, 0, 0)
```

The zoo and xts packages are recommended to deal with time series objects.

```
#install.packages("xts")
library(xts)
x = 1:4
y = seq(as.Date('2001-01-01'), length=4, by='day')
date1 = xts(x, y)
```

Extract or modify the content or time index of a time series object:

```
value = coredata(date1)
coredata(date1) = 2:5
time = index(date1)
```

While we are using the xts package, other data manipulation methods like subset and aggregate still work.

```
x = 5:2

y = seq(as.Date('2001-01-02'), length=4, by='day')

date2 = xts(x, y)

date3 = cbind(date1, date2)

names(date3) = c('v1', 'v2')

date4 = rbind(date1, date2)

names(date4) = 'value'
```

The window function can help us extract or modify a subset of a time series object (as observed between defined *start* and *end* times).

```
window(date4, start=as.Date("2001-01-04"))
```

```
value
2001-01-04 5
2001-01-04 3
2001-01-05 2
```

```
#lag() and diff() are still available
lag(date2)
diff(date2)
```

First load the quantmod package, download data from the Shanghai Stock Exchange composite index, and extract the 'close' column, which is the adjusted closing price.

```
#install.packages("quantmod")
library(quantmod)
options(getSymbols.auto.assign=FALSE)
library(xts)
SSEC = getSymbols('^SSEC', src='yahoo', from='2000-01-01')
head(SSEC, 3)
```

```
SSEC.Open SSEC.High SSEC.Low SSEC.Close

2000-01-04 1368.693 1407.518 1361.214 1406.371

2000-01-05 1407.829 1433.780 1398.323 1409.682

2000-01-06 1406.036 1463.955 1400.253 1463.942

SSEC.Volume SSEC.Adjusted

2000-01-04 0 1406.371

2000-01-05 0 1409.682

2000-01-06 0 1463.942
```



Compute the change rate, and find out the biggest change day:

```
data$ratio = with(data, diff(close)/close)
data.df = as.data.frame(data)
data.df[order(abs(data.df$ratio), decreasing=T), ][1:5, ]
```

	close	ratio
2007-02-27	2771.791	-0.09697993
2007-06-04	3670.401	-0.09000136
2001-10-23	1670.562	0.08972613
2008-09-19	2075.091	0.08638369
2008-04-24	3583.028	0.08503924



Is there a calendar effect? To check, we aggregate returns by month

```
#install.packages("lubridate")
library(lubridate)
data$mday = month(data)
res = aggregate(data$ratio, data$mday, mean, na.rm=TRUE)
cat(format(res*100, digits=2, scientific=F))
```



Tencent QQ is an instant messaging software service developed by Tencent. A QQ user can build a group to chat with friends. We'll explore QQ group chat data for this example. There are only two columns: the first column contains user IDs and the second column contains the chat times. So our questions are:

- Who chats the most in the group?
- When are people most likely to chat?

```
id time

1 8cha0 2011/7/8 12:11:13

2 2cha061 2011/7/8 12:11:49

3 6cha437 2011/7/8 12:13:36
```



```
time = as.POSIXct(data$time, tz='GMT')
id = as.factor(data$id)
data1 = data.frame(id, time)

user = as.data.frame(table(data1$id))
user = user[order(user$Freq, decreasing=T), ]
user[1:5, ] #getting the top five chat users
```

	Var1	Freq
91	7cha1	1511
86	6cha437	1238
66	4cha387	1100
98	8cha08	695
69	4cha69	533

```
data1$hour = hour(data1$time)
hours = as.data.frame(table(data1$hour))
hours = hours[order(hours$Freq, decreasing=T), ]
data1$wday = wday(data1$time)
wdays = as.data.frame(table(data1$wday))
wdays = wdays[order(wdays$Freq, decreasing=T), ]
```

# **Accessing Databases**

#### **Accessing Databases**

There are many database management systems (DBMS) for working with relational databases, and R can connect to all of the common ones.

The DBI package provides a unified syntax for accessing the following DBMS:

- SQLite
- MySQL
- MariaDB
- PostgreSQL
- Oracle

The RODBC package is an alternative that uses ODBC database connections

- This package is particularly useful for connecting to SQL Server/Access databases
- To use this package, you need to set up an ODBC data source on your machine

#### **Example: Reading Data from an SQLite Database**

```
#install.packages(c("DBI","RSQLite","learningr"))
library(DBI)
library(RSQLite)
driver = dbDriver("SQLite")
db_file = system.file("extdata", "crabtag.sqlite", package="learningr")
conn = dbConnect(driver, db_file)
query = "SELECT count(*) FROM Daylog"
(id_block = dbGetQuery(conn, query))
    count(*)
     405
dbDisconnect(conn)
[1] TRUE
```



# **Getting Data from Web**

#### The Different Routes of Data Retrieval

- If there is an API, use it.
- If there is no API but there is data in a table format, try readHTMLTable().
- If there is no API and the data is *unstructured*, you need to apply some form of web scraping.

#### **Tools for Web-data Retrieval**

- Packages for retrieval
  - RCurl
  - httr
- Packages for parsing

  - o rjson
  - RJSONIO
  - selectr
- Chrome Developer Tools

#### **Using an API**

Example: Get the weather data for your IP location from the Wunderground API.

- The RCurl package provides convenient functions to fetch URIs, get and post forms, etc.
- The RJSONIO package allows conversion to and from data in Javascript object notation (JSON) format

[1]"http://api.wunderground.com/api/a98d04ac43156c84/conditions/forecast/q/aut oip.json"



#### **Extracting Data Using Wunderground API**

```
fromurl = function(finalurl) {
  web = getURL(finalurl)
  raw = fromJSON(web)
  high = raw$forecast$simpleforecast$forecastday[[2]]$high['celsius']
  low = raw$forecast$simpleforecast$forecastday[[2]]$low['celsius']
  condition = raw$forecast$simpleforecast$forecastday[[2]]$conditions
  currenttemp = raw$current_observation$temp_c
  currentweather = raw$current_observation$weather
  city = as.character(raw$current_observation$display_location['full'])
  result = list(city=city, current=paste(currenttemp, '°C',
            currentweather, sep="),
  tomorrow=paste(low, '-', high,'°C', condition, sep="))
  names(result) = c('city', 'current', 'tomorrow')
  return(result)
}
```

#### **Local Weather Results**

```
fromurl(url)
```

```
$city
[1] "New York, NY"

$current
[1] "27°C Overcast"

$tomorrow
[1] "23-32°C Clear"
```

#### **Scraping Data From the Web**

The RCurl and XML packages provide useful parsing functions:

- getURL: Download a web page
- readHTMLTable: Read data from one or more HTML tables.
- htmlTreeParse/xmlTreeParse parses an XML or HTML file and generates an R structure representing the XML/HTML tree
- xmlApply applies a function to each of the children of an XMLNode
- getNodeSet: Find matching nodes in an internal XML tree/DOM



### **Example: Using readHTMLTable()**

Use readHTMLTable to get data on Available CRAN Packages:

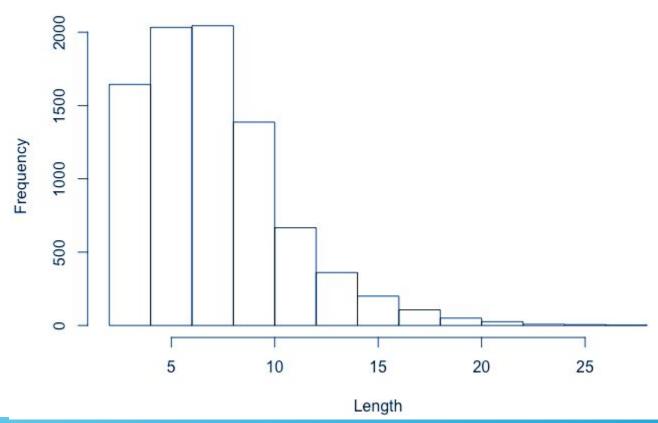
Date Package Title
2 2015-06-21 FAMILY A Convex Formulation for Modeling Interactions with
Strong\nHeredity



# **Example: Using readHTMLTable()**

```
res = nchar(data[,2])
hist(res, main="R Package Name Length", xlab="Length")
```

#### R Package Name Length





#### **Example: Getting XML Data**

```
library(XML)
url = "http://www.w3schools.com/xml/plant_catalog.xml"
xmlfile = xmlTreeParse(url) #download and parse XML
xmltop = xmlRoot(xmlfile) #get root node
xmlValue(xmltop[[10]][[1]]) #get leaf node data
```

```
[1] "Mayapple"
```

xmlValue(xmltop[['PLANT']][['COMMON']]) #get data from children of 'xmltop'

[1] "Bloodroot"



#### **Example: Getting XML Data**

xmlSApply(xmltop[[1]], xmlValue) #get data from each child of XML nodes

```
plantcat = xmlSApply(xmltop, function(x) xmlSApply(x, xmlValue))
plantcat_df = data.frame(t(plantcat),row.names=NULL)
plantcat_df[1:5,1:4]
```

	COMMON	BOTANICAL	ZONE	LIGHT
1	Bloodroot	Sanguinaria canadensis	4	Mostly Shady
2	Columbine	Aquilegia canadensis	3	Mostly Shady
3	Marsh Marigold	Caltha palustris	4	Mostly Sunny
4	Cowslip	Caltha palustris	4	Mostly Shady
5	Dutchman's-Breeches	Dicentra cucullaria	3	Mostly Shady



#### **Example: Getting HTML data**

In the following exercise, we extract critic reviews for *The Shawshank Redemption*.

```
library(RCurl)
library(XML)
url = 'http://www.imdb.com/title/tt0111161/criticreviews?ref_=tt_ov_rt'
raw = getURL(url)
data = htmlParse(raw)
xpath = '//tr[@itemprop="reviews"]/td[2]/div'
nodes = getNodeSet(data, xpath)
text = sapply(nodes, xmlValue)
```

# **Other Data Resources**

# **Government Open Data**

- Data.gov
- Socrata
- Some cities have their own open data websites, like San Francisco: sfgov.org
- UN open data
- WHO open data
- US Census
- USGS



#### **Economics and Finance**

- OECD
- UMD
- World Bank
- CBOE Futures Exchange
- Google Finance
- Google Trends
- NASDAQ
- OANDA
- Yahoo Finance

#### **Other Data Resources**

- Programmable Web
- InfoChimps
- Google Public Data Explorer
- Junar
- The New York Times API
- The Guardian Data Blog

