

# Workshop 2:

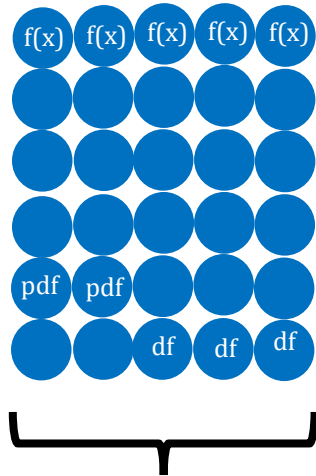
## The tidyverse and beyond

- we built this software on  
base R code



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# Recall



Base R:  
Comes  
pre-  
loaded

Functions

```
> base::|
```

max.col	{base}
mean	{base}
mean.Date	{base}
mean.default	{base}
mean.difftime	{base}
mean.POSIXct	{base}
mean.POSIXlt	{base}
mem.limits	{base}
memCompress	{base}

mean(x, ...)

Generic function for the (trimmed) arithmetic mean.

Press F1 for additional help

Data sets

```
> data()
```

faithful
freeny
infert
iris
iris3
islands
lh
longley

iris

This famous (Fisher's or Anderson's) iris data set gives the measurements in centimeters of the variables sepal length and width and petal length and width, respectively, for 50 flowers from each of 3 species of iris. The species are *Iris setosa*, *versicolor*, and *virginica*.

Press F1 for additional help

Conflicts

```
> filter|
```

filter	{dplyr}
filter_	{dplyr}
filter_all	{dplyr}
filter_at	{dplyr}
filter_if	{dplyr}
Filter	{base}
Filters	

filter(x, filter, method = c("convolution", "recursive"), sides = 2L, circular = FALSE, init = NULL)

Applies linear filtering to a univariate time series or to each series separately of a multivariate time series.

Press F1 for additional help

# Base R

## Cheat Sheet

### Getting Help

#### Accessing the help files

##### ?mean

Get help of a particular function.

**help.search('weighted mean')**

Search the help files for a word or phrase.

**help(package = 'dplyr')**

Find help for a package.

#### More about an object

##### str(iris)

Get a summary of an object's structure.

##### class(iris)

Find the class an object belongs to.

### Using Packages

#### install.packages('dplyr')

Download and install a package from CRAN.

#### library(dplyr)

Load the package into the session, making all its functions available to use.

#### dplyr::select

Use a particular function from a package.

#### data(iris)

Load a built-in dataset into the environment.

### Working Directory

#### getwd()

Find the current working directory (where inputs are found and outputs are sent).

#### setwd('C://file/path')

Change the current working directory.

Use projects in RStudio to set the working directory to the folder you are working in.

### Vectors

#### Creating Vectors

c(2, 4, 6)	2 4 6	Join elements into a vector
2:6	2 3 4 5 6	An integer sequence
seq(2, 3, by=0.5)	2.0 2.5 3.0	A complex sequence
rep(1:2, times=3)	1 2 1 2 1 2	Repeat a vector
rep(1:2, each=3)	1 1 1 2 2 2	Repeat elements of a vector

#### Vector Functions

<b>sort(x)</b>	<b>rev(x)</b>
Return x sorted.	Return x reversed.
<b>table(x)</b>	<b>unique(x)</b>
See counts of values.	See unique values.

#### Selecting Vector Elements

##### By Position

<b>x[4]</b>	The fourth element.
<b>x[-4]</b>	All but the fourth.
<b>x[2:4]</b>	Elements two to four.
<b>x[-(2:4)]</b>	All elements except two to four.
<b>x[c(1, 5)]</b>	Elements one and five.

##### By Value

<b>x[x == 10]</b>	Elements which are equal to 10.
<b>x[x &lt; 0]</b>	All elements less than zero.
<b>x[x %in% c(1, 2, 5)]</b>	Elements in the set 1, 2, 5.

##### Named Vectors

<b>x['apple']</b>	Element with name 'apple'.
-------------------	----------------------------

### Programming

#### For Loop

```
for (variable in sequence){  
  Do something  
}
```

##### Example

```
for (i in 1:4){  
  j <- i + 10  
  print(j)  
}
```

#### If Statements

```
if (condition){  
  Do something  
} else {  
  Do something different  
}
```

##### Example

```
if (i > 3){  
  print('Yes')  
} else {  
  print('No')  
}
```

#### While Loop

```
while (condition){  
  Do something  
}
```

##### Example

```
while (i < 5){  
  print(i)  
  i <- i + 1  
}
```

#### Functions

```
function_name <- function(var){  
  Do something  
  return(new_variable)  
}
```

##### Example

```
square <- function(x){  
  squared <- x*x  
  return(squared)  
}
```

### Reading and Writing Data

Also see the **readr** package.

Input	Output	Description
df <- read.table('file.txt')	write.table(df, 'file.txt')	Read and write a delimited text file.
df <- read.csv('file.csv')	write.csv(df, 'file.csv')	Read and write a comma separated value file. This is a special case of read.table/write.table.
load('file.RData')	save(df, file = 'file.Rdata')	Read and write an R data file, a file type special for R.

Conditions	a == b	Are equal	a > b	Greater than	a >= b	Greater than or equal to	is.na(a)	Is missing
	a != b	Not equal	a < b	Less than	a <= b	Less than or equal to	is.null(a)	Is null

## Types

Converting between common data types in R. Can always go from a higher value in the table to a lower value.

<code>as.logical</code>	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).
<code>as.numeric</code>	1, 0, 1	Integers or floating point numbers.
<code>as.character</code>	'1', '0', '1'	Character strings. Generally preferred to factors.
<code>as.factor</code>	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.

## Maths Functions

<code>log(x)</code>	Natural log.	<code>sum(x)</code>	Sum.
<code>exp(x)</code>	Exponential.	<code>mean(x)</code>	Mean.
<code>max(x)</code>	Largest element.	<code>median(x)</code>	Median.
<code>min(x)</code>	Smallest element.	<code>quantile(x)</code>	Percentage quantiles.
<code>round(x, n)</code>	Round to n decimal places.	<code>rank(x)</code>	Rank of elements.
<code>signif(x, n)</code>	Round to n significant figures.	<code>var(x)</code>	The variance.
<code>cor(x, y)</code>	Correlation.	<code>sd(x)</code>	The standard deviation.

## Variable Assignment

```
> a <- 'apple'
> a
[1] 'apple'
```




## The Environment

<code>ls()</code>	List all variables in the environment.
<code>rm(x)</code>	Remove x from the environment.
<code>rm(list = ls())</code>	Remove all variables from the environment.

You can use the environment panel in RStudio to browse variables in your environment.

## Matrices

```
m <- matrix(x, nrow = 3, ncol = 3)
Create a matrix from x.
```

 <code>m[2, ]</code>	- Select a row	<code>t(m)</code>	Transpose
 <code>m[, 1]</code>	- Select a column	<code>m %*% n</code>	Matrix Multiplication
 <code>m[2, 3]</code>	- Select an element	<code>solve(m, n)</code>	Find x in: $m * x = n$

## Lists

```
l <- list(x = 1:5, y = c('a', 'b'))
A list is a collection of elements which can be of different types.
```

<code>l[[2]]</code>	<code>l[1]</code>	<code>l\$x</code>	<code>l['y']</code>
Second element of l.	New list with only the first element.	Element named x.	New list with only element named y.




Also see the **dplyr** package.

## Data Frames



```
df <- data.frame(x = 1:3, y = c('a', 'b', 'c'))
A special case of a list where all elements are the same length.
```

x	y
1	a
2	b
3	c

### Matrix subsetting

<code>df[, 2]</code>	
<code>df[2, 1]</code>	
<code>df[2, 2]</code>	

### List subsetting

<code>df\$x</code>		<code>df[[2]]</code>	
<i>Understanding a data frame</i>			
<code>View(df)</code>	See the full data frame.		
<code>head(df)</code>	See the first 6 rows.		

`nrow(df)`  
Number of rows.

`ncol(df)`  
Number of columns.

`dim(df)`  
Number of columns and rows.

`cbind` - Bind columns.



`rbind` - Bind rows.



## Strings

Also see the **stringr** package.

<code>paste(x, y, sep = ' ')</code>	Join multiple vectors together.
<code>paste(x, collapse = ' ')</code>	Join elements of a vector together.
<code>grep(pattern, x)</code>	Find regular expression matches in x.
<code>gsub(pattern, replace, x)</code>	Replace matches in x with a string.
<code>toupper(x)</code>	Convert to uppercase.
<code>tolower(x)</code>	Convert to lowercase.
<code>nchar(x)</code>	Number of characters in a string.

## Factors

<code>factor(x)</code>	Turn a vector into a factor. Can set the levels of the factor and the order.
<code>cut(x, breaks = 4)</code>	Turn a numeric vector into a factor by 'cutting' into sections.

## Statistics

<code>lm(y ~ x, data=df)</code>	Linear model.	<code>t.test(x, y)</code>	Perform a t-test for difference between means.	<code>prop.test</code>	Test for a difference between proportions.
<code>glm(y ~ x, data=df)</code>	Generalised linear model.	<code>pairwise.t.test</code>	Perform a t-test for paired data.	<code>aov</code>	Analysis of variance.
<code>summary</code>	Get more detailed information out a model.				

## Distributions

	Random Variates	Density Function	Cumulative Distribution	Quantile
Normal	<code>rnorm</code>	<code>dnorm</code>	<code>pnorm</code>	<code>qnorm</code>
Poisson	<code>rpois</code>	<code>dpois</code>	<code>ppois</code>	<code>qpois</code>
Binomial	<code>rbinom</code>	<code>dbinom</code>	<code>pbinom</code>	<code>qbinom</code>
Uniform	<code>runif</code>	<code>dunif</code>	<code>punif</code>	<code>qunif</code>

## Plotting

Also see the **ggplot2** package.

 <code>plot(x)</code>	Values of x in order.	 <code>plot(x, y)</code>	Values of x against y.	 <code>hist(x)</code>	Histogram of x.
--	-----------------------	---	------------------------	--	-----------------

## Dates

See the **lubridate** package.

# Creating objects

For most of us, R is simply the creation of and manipulation of objects

```
new_object <- c(1, 2, 3)
```

- the objects are then fed into functions to create amazing new objects

```
amazing_new_object <- function(new_object)
```

Broadly speaking the following is true in R:

- information

- data frame <- function(information)

- plot <- function(data frame)

- model <- function(data frame)

# Data structures

Data structures encompass everything from our new object created in the last slide and beyond

Data can be stored in different forms

```
# double (for double precision floating point numbers)  
typeof(1)
```

```
# character  
typeof("string")
```

```
# logical  
typeof(FALSE)
```

```
# missing values are represented by NA  
example <- c(1, 2, NA, 4)
```

Other examples include integers and complex numbers

# Types of data structures I

# Vectors:

These come in two forms

- A: Atomic vectors contain exactly one type of data

```
all_numbers      <- c(1, 2, 0.5, -0.5, 3.4)
```

```
all_characters   <- c("One", "too", "3")
```

```
all_logical      <- c(TRUE, FALSE) # NOTE: Always type it out
```

- B: Lists allow combinations of different types of data

```
this_is_a_list   <- list(1, TRUE, "Three", "4")
```

```
typeof(this_is_a_list)  
[1] "list"
```

```
this_is_also_a_list <- list(all_numbers, all_characters, all_logical)
```

# Types of data structures I

# Matrices/Arrays:

- You can have a matrix of two or more dimensions  
`a_matrix <- matrix(1:9, 3, 3)`
- Vectors and matrices can only contain one type of data
- If you try to create a vector with more than one data type, then it will undergo coercion to the least common denominator
- The coercion rule goes:  
logical -> integer -> numeric -> complex -> character
- You can perform coercions yourself on vectors  
`nums_as_characters <- as.character(all_numbers)`  
`back_to_numbers <- as.numeric(nums_as_characters)`



# Worksheet

## ws2\_script1\_data\_structures\_I.R

# Types of data structures II

# Data frames:

- These are a special type of list
- Observations are in rows
- Variables are in columns
- Labels or other metadata may also be present
- ```
a_data_frame <- data.frame(number = 1:10,  
                             char   = sample(letters, 10),  
                             this_is_a_col_name = rep(c(TRUE, FALSE), 5))
```
- In the tidyverse data frames are called "tibbles"
  - Tibbles are one of the unifying features of the tidyverse
  - The two main differences are with;
    - (a) printing: tibbles have a defined print method
    - (b) subsetting: use of the `[` function
- Not all functions work with tibbles
- To convert back it's just a matter of typing:  

```
old_data_frame <- as.data.frame(your_tbl_df)
```

# Worksheet

## ws2\_script2\_data\_structures\_II

# Indexing

- Indexing can occur in one or two dimensions
- One dimension:

```
new_object <- c(1, 2, 3)
new_object[1]
[1] 1
```
- Two dimensions

```
a_data_frame[1, 1]

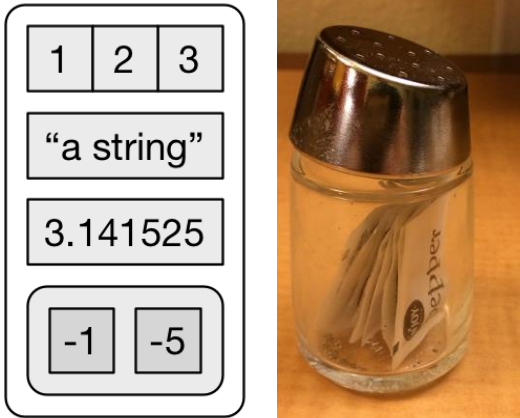
a_data_frame$number[1]
```
- In the tidyverse we don't use `[` much as `dplyr::filter()` and `dplyr::select()` allow you to solve the same problems
- However, given so much of the R has been written using these, it's worth recognising and understanding them

# Indexing

# Recall

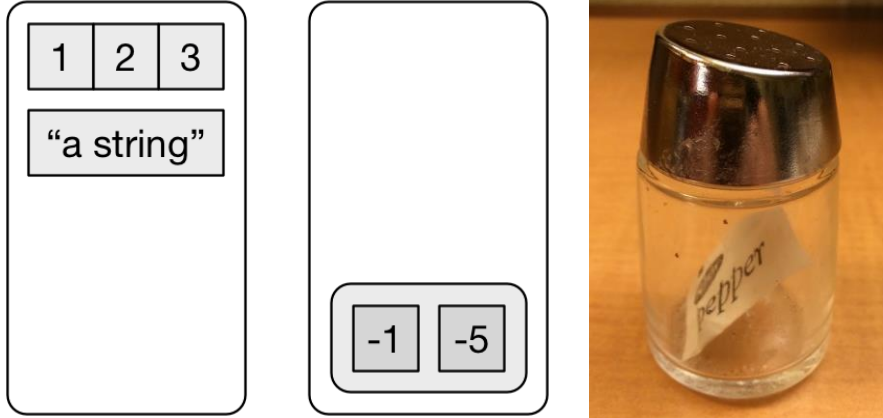
```
- this_is_also_a_list <- list(all_numbers, all_characters, all_logical)
```

a



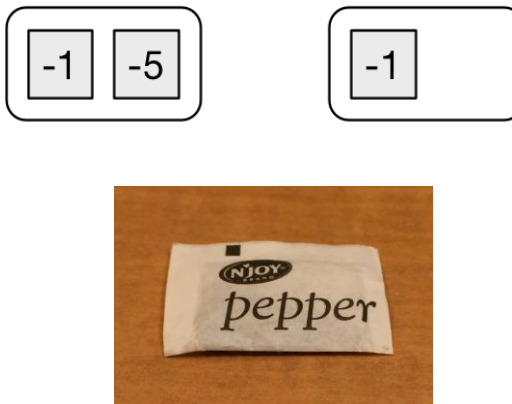
list

a[1:2]      a[4]



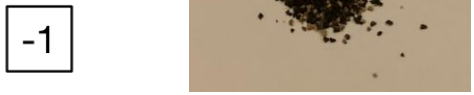
list

a[[4]]      a[[4]][1]



Vector

a[[4]][[1]]



Vector

# Important

- `[` extracts a sublist, results will be a list
- `[[` extracts a single component
- `$` is similar to `[[` for named elements of a list

# Worksheet

ws2\_script3\_indexing.R

# Types of data structures III

## # Factors

- In R, factors are used to work with categorical variables
- Historically they were easier to work with than characters, hence many baseR functions automatically convert characters to factors
- This does not happen in the tidyverse
- The forcats packages is designed to handle factors in the tidyverse
- One of the most important uses of factors is in statistical modeling; since categorical variables enter into statistical models differently than continuous variables, storing data as factors insures that the modeling functions will treat such data correctly

# Worksheet

`ws2_script4_factors.R`





# SPOOKY STORY TIME

Once upon a time.....

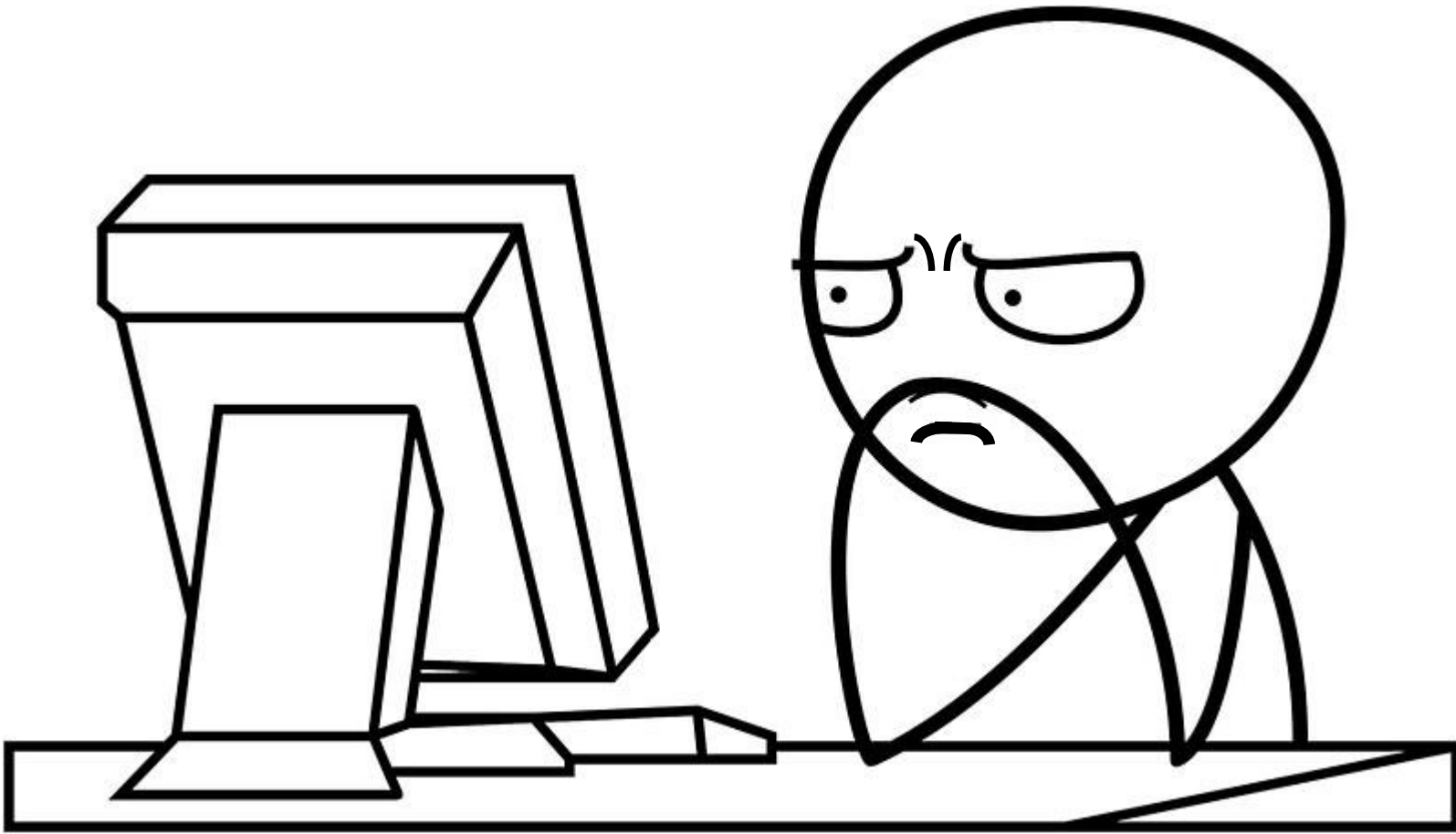
A Researcher collected some data

The Researcher was very good

The data was very good

But then.....

..... The Researcher inputted that data into Excel using default settings





# Then Excel converted some gene names to dates

Ziemann et al. *Genome Biology* (2016) 17:177  
DOI 10.1186/s13059-016-1044-7

Genome Biology

COMMENT

Open Access



## Gene name errors are widespread in the scientific literature

Mark Ziemann<sup>1</sup>, Yotam Eren<sup>1,2</sup> and Assam El-Osta<sup>1,3\*</sup>

### Abstract

The spreadsheet software Microsoft Excel, when used with default settings, is known to convert gene names to dates and floating-point numbers. A programmatic scan of leading genomics journals reveals that approximately one-fifth of papers with supplementary Excel gene lists contain erroneous gene name conversions.

frequently reused. Our aim here is to raise awareness of the problem.

We downloaded and screened supplementary files from 18 journals published between 2005 and 2015 using a suite of shell scripts. Excel files (.xls and .xlsx suffixes) were converted to tabular separated files (tsv) with ssconvert (v1.12.9). Each sheet within the Excel file was converted to a separate tsv file. Each column of data in the tsv file was screened for the presence of gene sym-

NCBI LocusLink

Search: LocusLink, Display: Brief, Organism: All, Query: Hs NEDD5, One of 1 Loci, Save All Loci

Click to Display: mRNA-Genomic Alignments (spanning 38716 bps)

PUB, OMIM, A REVIEW, UNIGENE, MAP, VAR, HOMOL, GDB, e!, UCSC

*Homo sapiens* Official Gene Symbol and Name (HGNC)

NEDD5: neural precursor cell expressed, developmentally down-regulated 5

LocusID: 4735

Overview, Submit GeneRIF

Locus Type: gene with protein product, function known or inferred

Product: neural precursor cell expressed, developmentally down-regulated 5

Alternate Symbols: DIFF6, SEPT2, hNedd5, KIAA0158

Relationships

Mouse Homology Maps:

| Map                    | Scale        | Gene           |
|------------------------|--------------|----------------|
| NCBI vs. MGD           | 1 cM         | 2-Sep Hs Mm    |
| UCSC vs. MGD           | 1 cM         | Sept2 Hs Mm    |
| UCSC vs. Hudson et al. | 1 1319.34 cR | AW208991 Hs Mm |

Map Information

# Excel also frequently got Clipboard amnesia



The answer, unfortunately, is **no**, you can't stop this from happening.

25

As described by [Joel Spolsky](#), developer and program manager for excel:



The official reason is that Excel doesn't really have cut and paste, it has move and copy. That's necessary because Excel automatically does reference fix up. For example, if cell A2 is defined as =A1, and you move cell A1 to A3, cell A2 will be updated to =A3.

If Excel actually cut things to the clipboard you would somehow need to have a reference pointing >into< the clipboard which is bizarre and for which there is no reasonable syntax. In other words, Excel doesn't want to leave you with dangling references during a move operation and isn't confident that it would be able to fix them up correctly when you completed the move by selecting "Paste."

Joel Spolsky 3/9/2004

[source](#)

What this means is that because of the difficulty inherent in the way excel maintains *references*, at the time of development there was no good way to store these references outside of excel and have them remain dynamic to be re-inserted. Once you change *focus* excel's ability to retain your original references is lost.

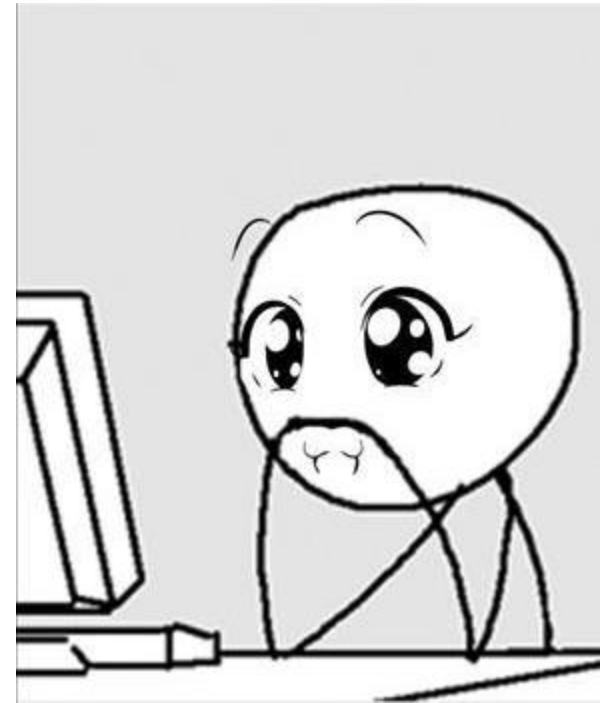
Unfortunately, MS does not consider this a bug.

Then one day.....

The Researcher accidentally used sort  
without selecting all the columns  
and inadvertently randomised  
the data.....

Help is at hand: R for Excel Users

<http://rex-analytics.com/things-live-r-r-excel-users/>



# Importing data with readr

- Used to read plain text rectangular files into R (e.g. csv)
- `read_csv` is the equivalent of `read.csv` in base R
- readr has a number of advantages over base R import function
  - ~10X faster
  - produces tibbles
  - doesn't convert character vectors to factors
  - more reproducible (readr code on your computer is likely to work on another computer)

# Data Import

with readr, tibble, and tidyr

## Cheat Sheet



R's **tidyverse** is built around **tidy data** stored in **tibbles**, an enhanced version of a data frame.



The front side of this sheet shows how to read text files into R with **readr**.



The reverse side shows how to create tibbles with **tibble** and to layout tidy data with **tidyr**.

### Other types of data

Try one of the following packages to import other types of files

- **haven** - SPSS, Stata, and SAS files
- **readxl** - excel files (.xls and .xlsx)
- **DBI** - databases
- **jsonlite** - json
- **xml2** - XML
- **httr** - Web APIs
- **rvest** - HTML (Web Scraping)

## Write functions

Save **x**, an R object, to **path**, a file path, with:

**write\_csv**(x, path, na = "NA", append = FALSE, col\_names = !append)

Tibble/df to comma delimited file.

**write\_delim**(x, path, delim = " ", na = "NA", append = FALSE, col\_names = !append)

Tibble/df to file with any delimiter.

**write\_excel\_csv**(x, path, na = "NA", append = FALSE, col\_names = !append)

Tibble/df to a CSV for excel

**write\_file**(x, path, append = FALSE)

String to file.

**write\_lines**(x, path, na = "NA", append = FALSE)

String vector to file, one element per line.

**write\_rds**(x, path, compress = c("none", "gz", "bz2", "xz", ...))

Object to RDS file.

**write\_tsv**(x, path, na = "NA", append = FALSE, col\_names = !append)

Tibble/df to tab delimited files.

## Read functions

### Read tabular data to tibbles

These functions share the common arguments:

**read\_\***(file, col\_names = TRUE, col\_types = NULL, locale = default\_locale(), na = c("", "NA"), quoted\_na = TRUE, comment = "", trim\_ws = TRUE, skip = 0, n\_max = Inf, guess\_max = min(1000, n\_max), progress = interactive())

a,b,c  
1,2,3  
4,5,NA

| A | B | C  |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**read\_csv()**

Reads comma delimited files.

**read\_csv**("file.csv")

a;b;c  
1;2;3  
4;5;NA

| A | B | C  |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**read\_csv2()**

Reads Semi-colon delimited files.

**read\_csv2**("file2.csv")

a|b|c  
1|2|3  
4|5|NA

| A | B | C  |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**read\_delim**(delim, quote = "\"", escape\_backslash = FALSE, escape\_double = TRUE) Reads files with any delimiter.

**read\_delim**("file.txt", delim = "|")

a b c  
1 2 3  
4 5 NA

| A | B | C  |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**read\_fwf**(col\_positions)

Reads fixed width files.

**read\_fwf**("file.fwf", col\_positions = c(1, 3, 5))

**read\_tsv()**

Reads tab delimited files. Also **read\_table()**.

**read\_tsv**("file.tsv")

### Useful arguments

a,b,c  
1,2,3  
4,5,NA

**Example file**

**write\_csv**(path = "file.csv",  
x = **read\_csv**("a,b,c|n1,2,3|n4,5,NA"))

|   |   |    |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**Skip lines**

**read\_csv**("file.csv",  
skip = 1)

| A | B | C  |
|---|---|----|
| 1 | 2 | 3  |
| 4 | 5 | NA |

**No header**

**read\_csv**("file.csv",  
col\_names = FALSE)

| A | B | C |
|---|---|---|
| 1 | 2 | 3 |

**Read in a subset**

**read\_csv**("file.csv",  
n\_max = 1)

| x | y | z  |
|---|---|----|
| A | B | C  |
| 1 | 2 | 3  |
| 4 | 5 | NA |

**Provide header**

**read\_csv**("file.csv",  
col\_names = c("x", "y", "z"))

| A  | B  | C  |
|----|----|----|
| 1  | 2  | 3  |
| NA | NA | NA |

**Missing Values**

**read\_csv**("file.csv",  
na = c("4", "5", "9"))

### Read non-tabular data

**read\_file**(file, locale = default\_locale())

Read a file into a single string.

**read\_file\_raw**(file)

Read a file into a raw vector.

**read\_lines**(file, skip = 0, n\_max = -1L, locale = default\_locale(), na = character(), progress = interactive())

Read each line into its own string.

**read\_lines\_raw**(file, skip = 0, n\_max = -1L, progress = interactive())

Read each line into a raw vector.

**read\_log**(file, col\_names = FALSE, col\_types = NULL, skip = 0, n\_max = -1, progress = interactive())

Apache style log files.

## Parsing data types

readr functions guess the types of each column and convert types when appropriate (but will NOT convert strings to factors automatically).

A message shows the type of each column in the result.

```
## Parsed with column specification:
## cols(
##   age = col_integer(),
##   sex = col_character(),
##   earn = col_double()
## )
#> # A tibble: 1 x 3
#>   age sex   earn
#>   <int> <chr> <dbl>
#> 1   23 male  45678
```

age is an integer  
sex is a character  
earn is a double (numeric)

1. Use **problems()** to diagnose problems

**x** <- **read\_csv**("file.csv"); **problems(x)**

2. Use a **col\_** function to guide parsing

- **col\_guess()** - the default
- **col\_character()**
- **col\_double()**
- **col\_euro\_double()**
- **col\_datetime**(format = "") Also **col\_date**(format = "") and **col\_time**(format = "")
- **col\_factor**(levels, ordered = FALSE)
- **col\_integer()**
- **col\_logical()**
- **col\_number()**
- **col\_numeric()**
- **col\_skip()**

```
x <- read_csv("file.csv", col_types = cols(
  A = col_double(),
  B = col_logical(),
  C = col_factor()
))
```

3. Else, read in as character vectors then parse with a **parse\_** function.

- **parse\_guess**(x, na = c("", "NA"), locale = default\_locale())
- **parse\_character**(x, na = c("", "NA"), locale = default\_locale())
- **parse\_datetime**(x, format = "", na = c("", "NA"), locale = default\_locale()) Also **parse\_date()** and **parse\_time()**
- **parse\_double**(x, na = c("", "NA"), locale = default\_locale())
- **parse\_factor**(x, levels, ordered = FALSE, na = c("", "NA"), locale = default\_locale())
- **parse\_integer**(x, na = c("", "NA"), locale = default\_locale())
- **parse\_logical**(x, na = c("", "NA"), locale = default\_locale())
- **parse\_number**(x, na = c("", "NA"), locale = default\_locale())

**x\$A** <- **parse\_number**(x\$A)



## Tibbles - an enhanced data frame

The **tibble** package provides a new S3 class for storing tabular data, the tibble. Tibbles inherit the data frame class, but improve two behaviors:

- **Display** - When you print a tibble, R provides a concise view of the data that fits on one screen.
- **Subsetting** - [ always returns a new tibble, [[ and \$ always return a vector.
- **No partial matching** - You must use full column names when subsetting

```
# A tibble: 234 x 6
  manufacturer model displ
<chr> <chr> <dbl>
1 audi a4 1.8
2 audi a4 1.8
3 audi a4 2.0
4 audi a4 2.0
5 audi a4 2.0
6 audi a4 2.8
7 audi a4 3.1
8 audi a4 quattro 1.8
9 audi a4 quattro 1.8
10 audi a4 quattro 2.0
# ... with 224 more rows, and 3
# more variables: year <int>,
# cyl <int>, trans <chr>
```

tibble display

```
156 1999 6 auto(14)
157 1999 6 auto(14)
158 2000 6 auto(14)
159 2000 6 auto(s4)
160 1999 4 manual(m5)
161 1999 4 auto(14)
162 2000 4 manual(m5)
163 2000 4 manual(m5)
164 2000 4 auto(14)
165 2000 4 auto(14)
166 1999 4 auto(14)
[ reached getOption("max.print")
  -- omitted 58 rows --]
```

data frame display

A large table to display

- Control the default appearance with options:
 

```
options(tibble.print_max = n,
        tibble.print_min = m, tibble.width = Inf)
```
- View entire data set with **View(x, title)** or **glimpse(x, width = NULL, ...)**
- Revert to data frame with **as.data.frame()** (required for some older packages)

### Construct a tibble in two ways

**tibble(...)**  
Construct by columns.  
*tibble(x = 1:3,*  
*y = c("a", "b", "c"))*

Both make this tibble

**tribble(...)**  
Construct by rows.  
*tribble(*  
 *~x, ~y,*  
 *1, "a",*  
 *2, "b",*  
 *3, "c")*

```
A tibble: 3 x 2
  x     y
<int> <dbl>
1     1  a
2     2  b
3     3  c
```

**as\_tibble(x, ...)** Convert data frame to tibble.

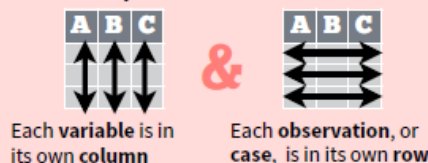
**enframe(x, name = "name", value = "value")**  
Converts named vector to a tibble with a names column and a values column.

**is\_tibble(x)** Test whether x is a tibble.

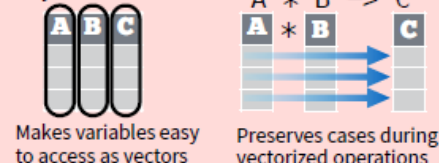
## Tidy Data with tidyr

**Tidy data** is a way to organize tabular data. It provides a consistent data structure across packages.

A table is tidy if:



Tidy data:



## Reshape Data - change the layout of values in a table

Use **gather()** and **spread()** to reorganize the values of a table into a new layout. Each uses the idea of a key column: value column pair.

**gather(data, key, value, ..., na.rm = FALSE, convert = FALSE, factor\_key = FALSE)**

Gather moves column names into a key column, gathering the column values into a single value column.

table4a

| country | 1990 | 2000 |
|---------|------|------|
| A       | 0.7K | 2K   |
| B       | 37K  | 80K  |
| C       | 212K | 213K |

→

| country | year | cases |
|---------|------|-------|
| A       | 1990 | 0.7K  |
| B       | 1990 | 37K   |
| C       | 1990 | 212K  |
| A       | 2000 | 2K    |
| B       | 2000 | 80K   |
| C       | 2000 | 213K  |

key value

*gather(table4a, `1990`, `2000`,  
key = "year", value = "cases")*

**spread(data, key, value, fill = NA, convert = FALSE, drop = TRUE, sep = NULL)**

Spread moves the unique values of a key column into the column names, spreading the values of a value column across the new columns that result.

table2

| country | year | type  | count |
|---------|------|-------|-------|
| A       | 1999 | cases | 0.7K  |
| A       | 1999 | pop   | 19M   |
| A       | 2000 | cases | 2K    |
| A       | 2000 | pop   | 20M   |
| B       | 1999 | cases | 37K   |
| B       | 1999 | pop   | 172M  |
| B       | 2000 | cases | 80K   |
| B       | 2000 | pop   | 174M  |
| C       | 1999 | cases | 212K  |
| C       | 1999 | pop   | 1T    |
| C       | 2000 | cases | 213K  |
| C       | 2000 | pop   | 1T    |

key value

*spread(table2, type, count)*

## Handle Missing Values

**drop\_na(data, ...)**

Drop rows containing NA's in ... columns.

x

| x1 | x2 |
|----|----|
| A  | 1  |
| B  | NA |
| C  | NA |
| D  | 3  |
| E  | NA |

→

| x1 | x2 |
|----|----|
| A  | 1  |
| D  | 3  |

*drop\_na(x, x2)*

**fill(data, ..., .direction = c("down", "up"))**

Fill in NA's in ... columns with most recent non-NA values.

x

| x1 | x2 |
|----|----|
| A  | 1  |
| B  | NA |
| C  | NA |
| D  | 3  |
| E  | NA |

→

| x1 | x2 |
|----|----|
| A  | 1  |
| B  | 1  |
| C  | 1  |
| D  | 3  |
| E  | 3  |

*fill(x, x2)*

**replace\_na(data, replace = list(), ...)**

Replace NA's by column.

x

| x1 | x2 |
|----|----|
| A  | 1  |
| B  | NA |
| C  | NA |
| D  | 3  |
| E  | NA |

→

| x1 | x2 |
|----|----|
| A  | 1  |
| B  | 2  |
| C  | 2  |
| D  | 3  |
| E  | 2  |

*replace\_na(x, list(x2 = 2), x2)*

## Expand Tables - quickly create tables with combinations of values

**complete(data, ..., fill = list())**

Adds to the data missing combinations of the values of the variables listed in ...  
*complete(mtcars, cyl, gear, carb)*

**expand(data, ...)**

Create new tibble with all possible combinations of the values of the variables listed in ...  
*expand(mtcars, cyl, gear, carb)*

## Split and Combine Cells

Use these functions to split or combine cells into individual, isolated values.

**separate(data, col, into, sep = "[^:alnum:]]+", remove = TRUE, convert = FALSE, extra = "warn", fill = "warn", ...)**

Separate each cell in a column to make several columns.

table3

| country | year | rate     |
|---------|------|----------|
| A       | 1999 | 0.7K/19M |
| A       | 2000 | 2K/20M   |
| B       | 1999 | 37K/172M |
| B       | 2000 | 80K/174M |
| C       | 1999 | 212K/1T  |
| C       | 2000 | 213K/1T  |

→

| country | year | cases | pop |
|---------|------|-------|-----|
| A       | 1999 | 0.7K  | 19M |
| A       | 2000 | 2K    | 20M |
| B       | 1999 | 37K   | 172 |
| B       | 2000 | 80K   | 174 |
| C       | 1999 | 212K  | 1T  |
| C       | 2000 | 213K  | 1T  |

*separate(table3, rate,  
into = c("cases", "pop"))*

**separate\_rows(data, ..., sep = "[^:alnum:]."]+, convert = FALSE)**

Separate each cell in a column to make several rows. Also **separate\_rows()**.

table3

| country | year | rate     |
|---------|------|----------|
| A       | 1999 | 0.7K/19M |
| A       | 2000 | 2K/20M   |
| B       | 1999 | 37K/172M |
| B       | 2000 | 80K/174M |
| C       | 1999 | 212K/1T  |
| C       | 2000 | 213K/1T  |

→

| country | year | rate |
|---------|------|------|
| A       | 1999 | 0.7K |
| A       | 1999 | 19M  |
| A       | 2000 | 2K   |
| A       | 2000 | 20M  |
| B       | 1999 | 37K  |
| B       | 1999 | 172M |
| B       | 2000 | 80K  |
| B       | 2000 | 174M |
| C       | 1999 | 212K |
| C       | 1999 | 1T   |
| C       | 2000 | 213K |
| C       | 2000 | 1T   |

*separate\_rows(table3, rate)*

**unite(data, col, ..., sep = "\_", remove = TRUE)**

Collapse cells across several columns to make a single column.

table5

| country | century | year |
|---------|---------|------|
| Afghan  | 19      | 99   |
| Afghan  | 20      | 0    |
| Brazil  | 19      | 99   |
| Brazil  | 20      | 0    |
| China   | 19      | 99   |
| China   | 20      | 0    |

→

| country | year |
|---------|------|
| Afghan  | 1999 |
| Afghan  | 2000 |
| Brazil  | 1999 |
| Brazil  | 2000 |
| China   | 1999 |
| China   | 2000 |

*unite(table5, century, year,  
col = "year", sep = "")*

# Importing data from stata and other stats packages

- Stata, SPSS and SAS file formats can be imported and exported using the tidyverse package "haven"

## Importing data with other packages

- Very often the data you handle may not be your own
  - collaborators
  - government
  - online databases
  - publications
  - webpages

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# Worksheet

ws2\_script5\_data\_import.R

# Introductory R Workshops

~~Week 1 (13<sup>th</sup> February):~~

~~Take a parachute and jump (into the tidyverse)~~

- ~~— tidying and visualisation of NGS data~~
- ~~— using sample R scripts~~

~~Week 2 (20<sup>th</sup> February):~~

~~We built this software on base R code~~

- ~~— overview and structure of R syntax~~

**Week 3 (27<sup>th</sup> February):**

**Sending an SOS to the world**

- how to identify with errors in your code and get help

