

# Basics in R

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# Workshop overview

## Part 1: Getting to know R

- Basic operations in R

## Part 2: Hands-on practice

- Work on sample data
- Plot pretty graphs

## Part 3: Learning to be independent

- Seek help from the web

# Recap

# apply function

```
d1[,c(10:14)]=apply(d1[,c(10:14)], MARGIN=2, FUN=as.numeric)
```

↑  
Store the  
output in an  
object

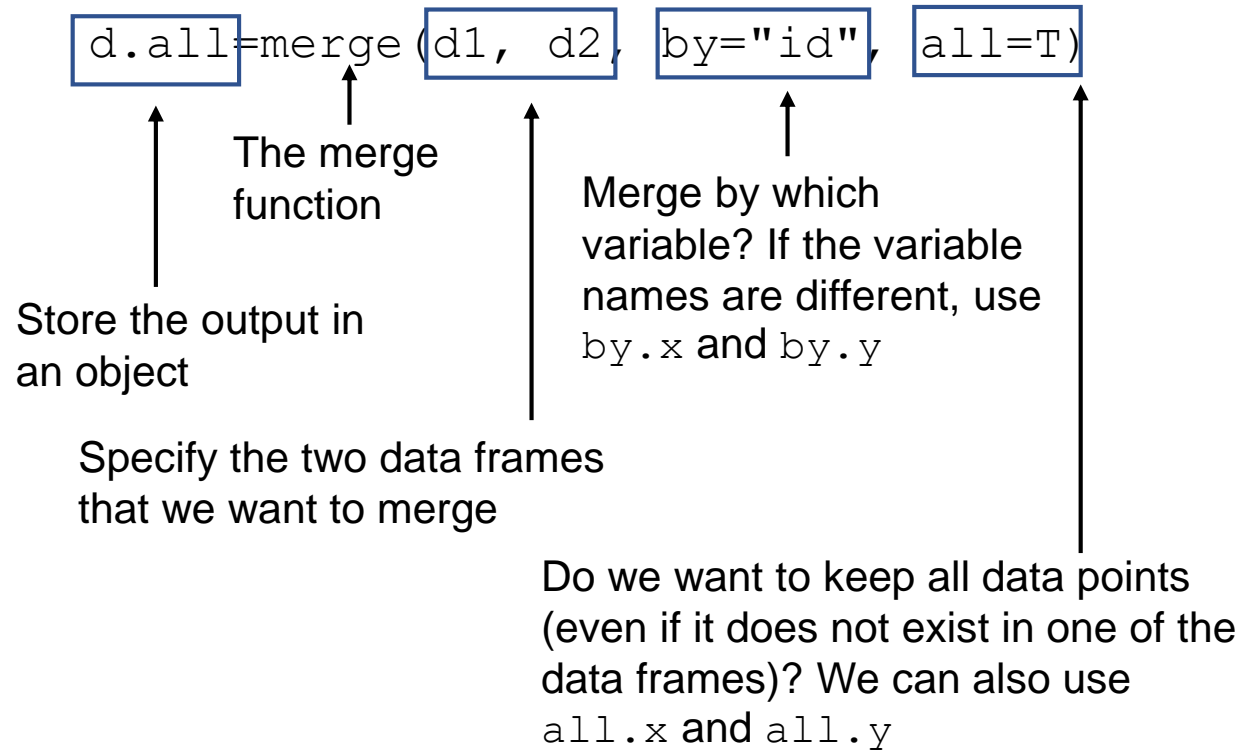
↑  
The apply  
function

↑  
Specify the data frame  
- Here, we only want  
columns 10 to 14

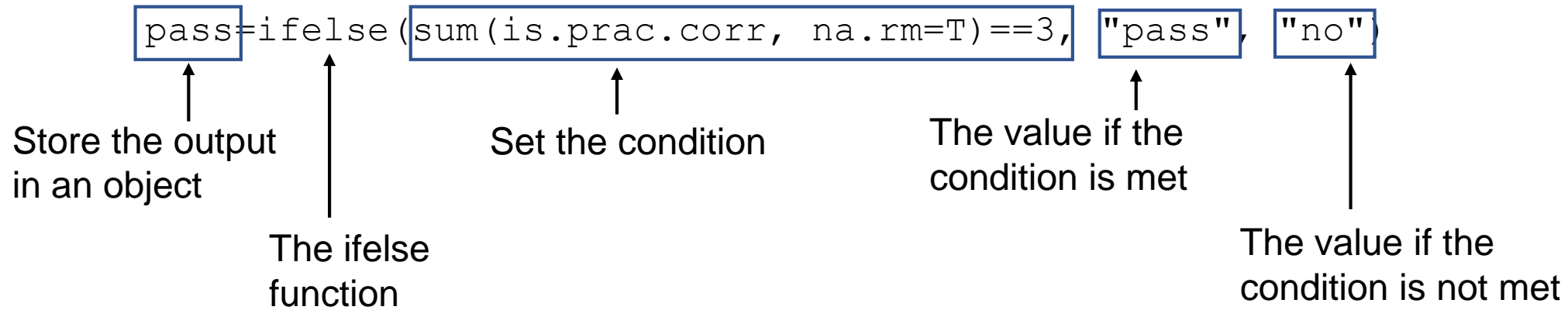
↑  
Margin 1: per row  
Margin 2: per column

↑  
Which function to apply?  
- We could also use  
*mean*, *sum*, etc., or even  
write our own function

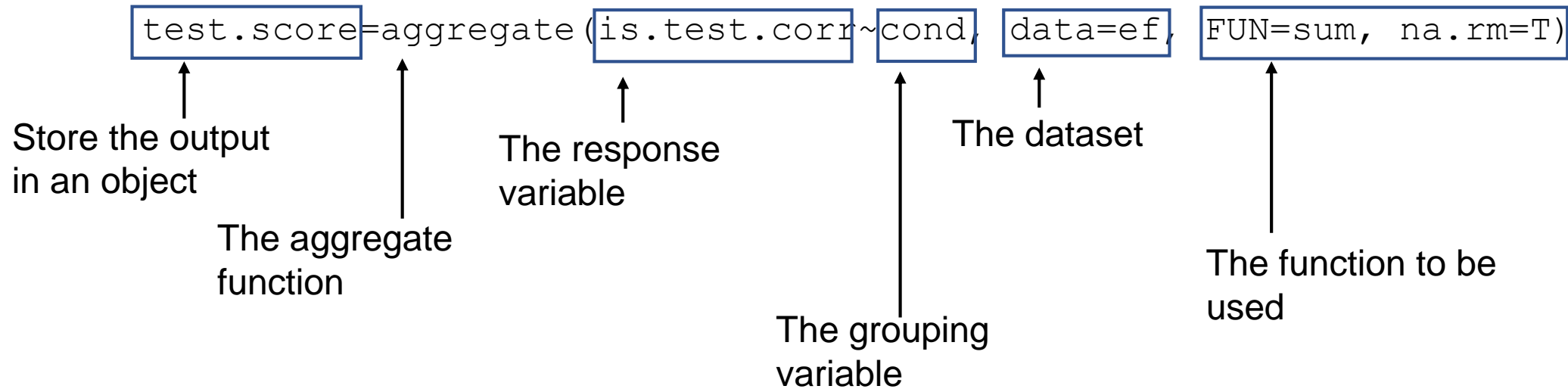
# merge function



# ifelse function



# aggregate **function**



# for loop

- This is how a for loop works in general:

```
for (i in 1:n){  
  ...  
}
```

What we want to do with the items

You can read this as “for  $i$  in 1 to  $n$ ”, where

- $n$  is the total number of items
- $i$  is the index of each of these items

\* Remember  
to indent the  
codes

```
for (i in 1:length(eflist)){  
  ef=read.csv(paste0(path="./mock_ef/", eflist[i])) ##use this instead of the file's name  
  ef$id=as.character(ef$id)  
  ef$cond=factor(ef$cond, level=c("neut", "cong", "inco"))  
  ef[ef==""] = NA  
  ##calculate scores  
  is.prac.corr=ifelse(ef$prac.resp==ef$prac.corr, 1, 0)  
  pass=ifelse(sum(is.prac.corr, na.rm=T)==3, "pass", "no")  
  ef$is.test.corr=ifelse(ef$test.resp==ef$test.corr, 1, 0)  
  test.score=aggregate(is.test.corr~cond, data=ef, FUN=sum, na.rm=T)  
  ##extract data  
  xx=c() ##create an empty holder  
  xx$id=ef$id[1]  
  xx$pass=pass  
  xx$neut=test.score$is.test.corr[1]  
  xx$cong=test.score$is.test.corr[2]  
  xx$inco=test.score$is.test.corr[3]  
  ##add data to an empty dataframe  
  d.ef=rbind(d.ef, xx)  
}
```

\* Change 1

\* Change 2: empty data frame should be  
placed before the for loop



# Plot

```
##plot 1: scatterplot
par(mar=c(3.4, 3.4, 0.5, 0.5), mgp=c(1.8, 0.5, 0), tcl=-0.3, las=1)
plot(x=x.pass$age, y=x.pass$read.dur, #ylim=c(0, 10),
     xlab="Age (year)", ylab="Reading duration (minutes)",
     pch=8, cex=1.3, lwd=2, col="blue",
     cex.axis=1.3, cex.lab=1.8)
##add line
age.read=lm(x.pass$read.dur~x.pass$age)
abline(age.read, lty=2, lwd=3)
```

To save/export the plot, you can either

- Click plots -> export
- Or use functions like `jpeg()` or `pdf()`

```
jpeg("plot1.jpg", width=500)
... (the plotting script)
dev.off()
```

# Plotting parameters

- To place both plots side-by-side:

```
##both plots together
par(mfrow=c(1, 2), mar=c(3.4, 3.6, 2, 2), mgp=c(2, 0.5, 0), tcl=-0.3, las=1)
##plot reading duration
plot(x.pass$grp, x.pass$read.dur, main="Reading duration",
     xlab="Group", ylab="Reading duration (minutes)", xaxt="n",
     cex.axis=1.3, cex.lab=1.6, lwd=2, col="deepskyblue")
mtext(text=c("interested", "not interested"), at=c(1, 2), side=1,
      line=0.4, cex=1.3)
##plot brain activation
plot(x.pass$grp, x.pass$avg.hbo, main="Brain activation",
     xlab="Group", ylab="Brain activation (HbO)", xaxt="n",
     cex.axis=1.3, cex.lab=1.6, lwd=2, col="deepskyblue")
mtext(text=c("interested", "not interested"), at=c(1, 2), side=1,
      line=0.4, cex=1.3)
```

\*Added more space

\*Added title

# **Part 3**

## **Learning to be independent**

# Some useful websites for further reading

- <https://learningstatisticswithr.com/book/introR.html>
- <https://www.learnbyexample.org/r-introduction/>
- Or, you could just google what you need, like what I always do.
- You don't need to know the codes by heart. You just need to know what tools are available, and then google the codes whenever you need to use them.
- You can also have a snippets file to save codes that you reuse frequently.
- Practice makes perfect: it will become easier after several rounds of coding

# Widely used packages

- Many R users (whom I know) use *tidyverse* to clean their data and *ggplot2* for plotting.
- Some R users find *tidyverse* to be more intuitive, but I think it is because they are more familiar with *tidyverse*.
- I prefer to use base R but this is just a personal preference.
- I would say to go for whatever that you are most comfortable with.
- Most websites provide solutions using base R and *tidyverse*:
  - Google: How to calculate means by group?
  - <https://www.statology.org/r-mean-by-group/>
- I like this website when it comes to plotting:  
[www.r-graph-gallery.com](http://www.r-graph-gallery.com)

<https://www.tidyverse.org/>  
<https://dplyr.tidyverse.org/>  
<https://ggplot2.tidyverse.org/>

## Method 1: Use base R.

```
aggregate(df$col_to_aggregate, list(df$col_to_group_by), FUN=mean)
```

## Method 2: Use the dplyr() package.

```
library(dplyr)

df %>%
  group_by(col_to_group_by) %>%
  summarise_at(vars(col_to_aggregate), list(name = mean))
```

# Data mining & PCA

- <https://www.geeksforgeeks.org/data-mining-in-r/>
- <https://www.tutorialspoint.com/exploring-data-mining-with-r>
- <https://rpubs.com/uriel623/670548>
- <https://www.geeksforgeeks.org/confirmatory-factor-analysis-in-r/>
- I won't go into details for two main reasons:
  - They're not something I usually do, so I'm not familiar with these topics (I'd be happy to talk about mixed effect models though)
  - My aim is to cover the basics in R, so they're beyond the workshop's scope
- The links above give good explanation and examples and they build upon what I presented today (except that they use *dplyr* while I use base R to achieve the same thing)

# Thank you!

## That's the end of the workshop.

Please fill in the feedback form:

<https://forms.gle/3eaJEAUYUD39auZ9A>

