

Getting started, getting the data in

Function	Source Package	Description	Example
Getting activated			
<code>install.packages()</code>	Base	Installs packages from CRAN	<code>install.packages("ggplot2")</code>
<code>library()</code>	Base	Packages must be loaded at the beginning of each R session.	<code>library(ggplot2)</code>
<code>file.path()</code>	Base	Codes a file path that can be called in subsequent functions.	<code>src <- file.path("/users/filepath")</code>
Generating R objects			
<code>c()</code>	Base	Combines elements into a vector	<pre>x <- c(1,2,3) y <- c("A","B","C") x <- factor(x=vector, levels=c(1,2,3), labels=c("A","B","C"))</pre>
<code>factor()</code>	Base	Generates a categorical factor variables	
<code>quantcut()</code>	gtools	Categorizes a continuous variable quantiles	<pre>x <- quantcut(x=vector, q=4) x <- sample(x=vector, size=n, replace=TRUE)</pre>
<code>sample()</code>	Base	Randomly samples a set of numbers	
<code>rnorm()</code>	Base	Generates a random normal distribution	<code>x <- rnorm(n=n, mean=10, sd=5)</code>
<code>list()</code>	Base	Generates a list	<code>mylist <- list(a, b, c)</code>
<code>data.frame()</code>	Base	Generates a data frame	<code>df <- data.frame(a,b,c, stringsAsFactors=F)</code>
Explore the data			
<code>summary()</code>	Base	Summarizes a continuous variable	<code>summary(example_object)</code>
<code>table()</code>	Base	Produces frequencies for a variable	<code>table(example_object, useNA="always")</code>
<code>prop.table()</code>	Base	Produces proportions for a table() object	<code>prop.table(table(example_object, useNA="always"))</code>
<code>mean()</code> / <code>median()</code>	Base	Calculates the mean / median of a continuous variable	<pre>mean(x, na.rm=TRUE) median(x, na.rm=TRUE)</pre>
<code>min()</code> / <code>max()</code>	Base	Calculates the minimum / maximum of a continuous variable	<pre>min(x, na.rm=TRUE) max(x, na.rm=TRUE)</pre>
<code>is.na()</code>	Base	TRUE/FALSE - is a value NA	<code>is.na(x)</code>
<code>str()</code>	Base	Returns the structure of an object	<code>str(data.frame.example_object)</code>
<code>nrow()</code> / <code>ncol()</code>	Base	Returns the number of rows/cols in a data frame	<code>nrow(data.frame.example_object)</code>
<code>head()</code> / <code>tail()</code>	Base	Prints the first/last 6 observations of a vector or data frame	<pre>head(data.frame.example_object) tail(data.frame.example_object)</pre>
<code>identical()</code>	Base	TRUE/FALSE - are two objects exactly identical	<code>identical(example_object1, example_object2)</code>
Working with external data			
<code>read.table()</code>	Base	Read a flat delimited file into R	<code>df <- read.table(file.path, stringsAsFactors=F)</code>
<code>read_sas()</code>	Haven	Read a .sas7bdat file into R	<code>df <- read_sas(file.path)</code>
<code>read.xlsx()</code>	openxls	Read an excel spreadsheet into R	<code>df <- read.xlsx(spreadsheet.path,</code>

	x		sheet=1)
<code>save()</code>	Base	Saves objects as an RDATA file	<code>save(object, file="Object name.Rdata")</code>
<code>saveRDS()</code>	Base	Save an object as an RDS file	<code>saveRDS(data.frame, file="Object name.RDS")</code>
<code>load()</code>	Base	Load's an RDATA file into R	<code>load(file="Object name.Rdata")</code>
<code>readRDS()</code>	Base	Reads an RDS file into R	<code>df <- readRDS("Object name.RDS")</code>

Subsetting and filtering functions

Function	Source Package	Description	Example
Variable selection			
<code>x[i]</code>	Base	Select the <i>i</i> -value of x	<code>x[5]</code>
<code>df[row_i,col_i]</code>	Base	select the i-row and i-column of df-data frame	<code>df[5, 10]</code>
<code>df[, "var_name"]</code>	Base	select the "var_name" variable in a data frame	<code>summary(df[, "BMI"])</code>
<code>df\$variable</code>	Base	select the "var_name" variable in a data frame	<code>summary(df\$BMI)</code>
<code>names()</code>	Base	Print the variable names in a data frame	<code>names(df)</code>
<code>names() <- c()</code>	Base	Rename the variable names in a data frame	<code>names(df) <- c("SUBJID", "DOB", "BMI", "AGE")</code>
Operators			
<code><, <=</code>	Base	Less than / less than or equal to	<code>var1 < var2</code> <code>var1 <= var2</code>
<code>>, >=</code>	Base	Greater than/ greater than or equal to	<code>var1 > var2</code> <code>var1 >= var2</code>
<code>%in%</code>	Base	Includes	<code>var1 %in% var2</code>
<code>==</code>	Base	Exactly equal to	<code>var1 == var2</code>
<code>&</code>	Base	And	<code>df[df\$x==1 & df\$y==2,]</code>
<code>!</code>	Base	Not	<code>df[df\$x != 1,]</code>
<code> </code>	Base	Or	<code>df[df\$x ==1 df\$x==2,]</code>
Manipulating datasets			
<code>rbind()</code>	Base	Concatenates rows of 2 data frames	<code>df3 <- rbind(df1, df2)</code>
<code>cbind()</code>	Base	Combines columns of 2 data frames	<code>df3 <- cbind(df1, df2)</code>
<code>merge()</code>	Base	Merges to data frames based on a merging variable	<code>df3 <- merge(x=df1, y=df2, by= "mergevar", all.x=T, all.y=T)</code>
<code>order()</code>	Base	Sorts / orders an object	<code>df3 <- df4[order(df4\$mergevar),]</code>

Helpful cleaning functions

Function	Description	Example
Variable transformations		
<code>ifelse(x, yes, no)</code>	Basic if-else-then conditional function for binary logic	<code>ifelse(x=="A", yes=1, no=0)</code>

ifelse(x, yes, ifelse(x, yes, no))	Nested if-else-then function for multiple conditional logic expressions	ifelse(x=="A", 1, ifelse(x=="B", 2, 3))
toupper() / tolower()	Converts a character string to all upper- or lower-case	toupper("my name is dave") [1] "MY NAME IS DAVE"
quantile()	Generates quantile cutpoints	quantile(x, c(0.25, 0.5, 0.75), na.rm=TRUE)
gtools::quantcut()	Categorizes a continuous variable into quantiles	x <- quantcut(x=vector, q=4)
Arithmetic operations		
+	Addition	x + y
-	Subtraction	x - y
/	Division	x / y
*	Multiplication	x * y
** or ^	Exponentiation	x**2 or x^2
log() / exp()	Natural log and exponentiation	log(x); exp(x)
sqrt() / nthroot	Square-root or Nth-root	sqrt(100); nthroot(100, 3)
floor() / ceiling()	Round down / up to the nearest integer	floor(10.5) round(pi)
round(x, digits=0)	Round to some decimal place	[1] 3.141593
Dates		
as.Date()	Creates a date object	as.Date("2019-06-25")
Sys.Date()	Today's date	as.Date(x, origin="1970-01-01")
Sys.time()	System date and time	R.date <- as.Date(SAS.date, origin="1960-01-01") Sys.Date() Sys.time()

Renaming all types of things

Function	Description	Example
Renaming variables		
names()	Rename all the variables in a data frame	names(df) <- c("Var1", "Var2", "Var3")
dplyr::Rename(dat, newname=old)	Renames individual variables and dropping the old ones	Rename(df, Newvar1 = OldVar1, Newvar2 = Oldvar2)
Factor variables		
factor(x, levels, labels)	Create a factor variable. Order of the variable is set by levels , formats are set with labels	factor(x, levels=c(1,2,3), labels="Never", "Current", "Former")
levels()	Renames the formats in a factor variable	levels(x, c("Never smoker", "Current smoker", "Former smoker"))
relevel(x, ref=)	Changes the reference group of a factor variable	relevel(x, "Current smoker")
Apply() functions		

<code>apply(x, margin, fun)</code>	Apply a function (fun) across variable X. Margin=1 applies the function across each observation Margin=2 applies the function for each column	# Sums 3 variables for each observation <code>apply(df[,c("var1", "var2", "var3")], 1, sum)</code> # Sums each of the three variables <code>apply(df[,c("var1", "var2", "var3")], 2, sum)</code> # Creates a frequency table for three variables <code>lapply(df[,c("var1", "var2", "var3")], table)</code>
<code>lapply(x, fun)</code>	List-apply. Applies a function across heterogenous data	# Converts all variables to numeric <code>lapply(df[,names(df)], as.numeric)</code>
<code>tapply(continuous, categorical, function)</code>	Useful for getting summary statistics for each level of a categorical variable.	# Get range of BMI for each level of category BMI <code>tapply(df\$BMI, df\$BMICAT, summary)</code>

Handling duplicates

Function	Description	Example
<code>uplicated()</code>	TRUE/FALSE - is a value a duplicate of another	<code>uplicated(x)</code>
<code>!uplicated()</code>	TRUE/FALSE - is a value NOT duplicated of another	<code>!uplicated(x)</code>
<code>table(duplicated(x))</code>	Frequency table - how many duplicates?	<code>table(duplicated(x))</code>
<code>df[!duplicated(df\$ID),]</code>	Remove the duplicate ID observations from data frame "df"	<code>nodup <- df[!duplicated(df\$ID),]</code> <code>wide <- reshape(data = long, varying=c("var1", "var2"), vnames = "newvar", idvar = "SubjectID", direction = "wide")</code>
<code>reshape(data, varying, v.names, idvar, direction)</code>	Reshape a dataset from long-form to wide form, or vice versa	<code>long <- reshape(data = long, varying=c("var1", "var2"), vnames = "newvar", idvar = "SubjectID", direction = "long")</code>

Stay functional!