

BaseR Programming and Data Manipulation

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The following was completed to complete Homework 2 for ST558 and practice basic programming and data manipulation in R.

Task One: Basic Vector Practice

Question 1

```
#Creating two vectors called pre and post using provided data

pre <- c(130, 128, 116, 124, 133, 134, 118, 126, 114, 127,
        141, 138, 128, 140, 137, 131, 120, 128, 139, 135)

post <- c(114, 98, 113, 99, 107, 116, 113, 111, 119, 117,
        101, 119, 130, 122, 106, 106, 124, 102, 117, 113)
```

Question 2

```
#Assigning patient IDs to the vectors
patient <- paste("Subject", 1:20, sep = "_")
names(pre) <- patient
names(post) <- patient
```

Question 3

```
#Creating a vector of the difference in blood pressure pre vs post treatment
diff_bp <- pre-post
diff_bp
```

Subject_1	Subject_2	Subject_3	Subject_4	Subject_5	Subject_6	Subject_7
16	30	3	25	26	18	5
Subject_8	Subject_9	Subject_10	Subject_11	Subject_12	Subject_13	Subject_14
15	-5	10	40	19	-2	18
Subject_15	Subject_16	Subject_17	Subject_18	Subject_19	Subject_20	
31	25	-4	26	22	22	

Question 4

```
#Finding the mean difference in blood pressure
mean(diff_bp)
```

```
[1] 17
```

Question 5

```
#Identifying patients that experienced a decrease in blood pressure after treatment
which(diff_bp > 0)
```

Subject_1	Subject_2	Subject_3	Subject_4	Subject_5	Subject_6	Subject_7
1	2	3	4	5	6	7
Subject_8	Subject_10	Subject_11	Subject_12	Subject_14	Subject_15	Subject_16
8	10	11	12	14	15	16
Subject_18	Subject_19	Subject_20				
18	19	20				

Question 6

```
#Subsetting the difference vector to only return patients with a positive change
diff_bp[which(diff_bp > 0)]
```

Subject_1	Subject_2	Subject_3	Subject_4	Subject_5	Subject_6	Subject_7
16	30	3	25	26	18	5
Subject_8	Subject_10	Subject_11	Subject_12	Subject_14	Subject_15	Subject_16
15	10	40	19	18	31	25
Subject_18	Subject_19	Subject_20				
26	22	22				

Question 7

```
#Calculating the mean blood pressure difference of patients with a positive change
mean(diff_bp[which(diff_bp > 0)])
```

```
[1] 20.64706
```

Task Two

Question 1

```
#Combining vectors into a data frame
bp_df_trt <- data.frame(patient, pre, post, diff_bp)
colnames(bp_df_trt) <- c("patient", "pre_bp", "post_bp", "diff_bp")
```

Question 2

```
#Subsetting the data frame to only include patients with a positive change
bp_df_trt[(bp_df_trt$diff_bp < 0),]
```

	patient	pre_bp	post_bp	diff_bp
Subject_9	Subject_9	114	119	-5
Subject_13	Subject_13	128	130	-2
Subject_17	Subject_17	120	124	-4

Question 3

```
#Adding a data column identifying if the patient's post bp is normal (less than 120)
bp_df_trt$normal <- bp_df_trt$post_bp < 120
```

Question 4

```
#Displaying the final treatment blood pressure data frame
knitr::kable(bp_df_trt)
```

```
Warning: 'xfun::attr()' is deprecated.
Use 'xfun::attr2()' instead.
See help("Deprecated")
Warning: 'xfun::attr()' is deprecated.
Use 'xfun::attr2()' instead.
See help("Deprecated")
```

	patient	pre_bp	post_bp	diff_bp	normal
Subject_1	Subject_1	130	114	16	TRUE
Subject_2	Subject_2	128	98	30	TRUE
Subject_3	Subject_3	116	113	3	TRUE
Subject_4	Subject_4	124	99	25	TRUE
Subject_5	Subject_5	133	107	26	TRUE
Subject_6	Subject_6	134	116	18	TRUE
Subject_7	Subject_7	118	113	5	TRUE
Subject_8	Subject_8	126	111	15	TRUE
Subject_9	Subject_9	114	119	-5	TRUE
Subject_10	Subject_10	127	117	10	TRUE
Subject_11	Subject_11	141	101	40	TRUE
Subject_12	Subject_12	138	119	19	TRUE
Subject_13	Subject_13	128	130	-2	FALSE
Subject_14	Subject_14	140	122	18	FALSE
Subject_15	Subject_15	137	106	31	TRUE
Subject_16	Subject_16	131	106	25	TRUE
Subject_17	Subject_17	120	124	-4	FALSE
Subject_18	Subject_18	128	102	26	TRUE
Subject_19	Subject_19	139	117	22	TRUE
Subject_20	Subject_20	135	113	22	TRUE

Task Three

Question 1

```
#Creating a data frame for patients who received a placebo
#Only permanently saving the final dataset
bp_df_placebo <- data.frame(names_plac = paste("Subject", 1:10, sep = "_"),
                             pre_bp = c(138, 135, 147, 117, 152, 134, 114, 121, 131, 130),
                             post_bp = c(105, 136, 123, 130, 134, 143, 135, 139, 120, 124))
bp_df_placebo$diff_bp <- bp_df_placebo$pre_bp - bp_df_placebo$post_bp
bp_df_placebo$normal <- bp_df_placebo$post_bp < 120

bp_df_placebo
```

	names_plac	pre_bp	post_bp	diff_bp	normal
1	Subject_1	138	105	33	TRUE
2	Subject_2	135	136	-1	FALSE
3	Subject_3	147	123	24	FALSE
4	Subject_4	117	130	-13	FALSE
5	Subject_5	152	134	18	FALSE
6	Subject_6	134	143	-9	FALSE
7	Subject_7	114	135	-21	FALSE
8	Subject_8	121	139	-18	FALSE
9	Subject_9	131	120	11	FALSE
10	Subject_10	130	124	6	FALSE

Question 2

```
#Creating a list with 2 elements, the treatment dataset and the placebo dataset
bp_list <- list(treatment=bp_df_trt, placebo=bp_df_placebo)
```

Question 3

```
#Accessing the treatment dataset from the list in 3 different ways
bp_list[1]
```

\$treatment

	patient	pre_bp	post_bp	diff_bp	normal
Subject_1	Subject_1	130	114	16	TRUE
Subject_2	Subject_2	128	98	30	TRUE
Subject_3	Subject_3	116	113	3	TRUE
Subject_4	Subject_4	124	99	25	TRUE
Subject_5	Subject_5	133	107	26	TRUE
Subject_6	Subject_6	134	116	18	TRUE
Subject_7	Subject_7	118	113	5	TRUE
Subject_8	Subject_8	126	111	15	TRUE
Subject_9	Subject_9	114	119	-5	TRUE
Subject_10	Subject_10	127	117	10	TRUE
Subject_11	Subject_11	141	101	40	TRUE
Subject_12	Subject_12	138	119	19	TRUE
Subject_13	Subject_13	128	130	-2	FALSE
Subject_14	Subject_14	140	122	18	FALSE
Subject_15	Subject_15	137	106	31	TRUE
Subject_16	Subject_16	131	106	25	TRUE
Subject_17	Subject_17	120	124	-4	FALSE
Subject_18	Subject_18	128	102	26	TRUE
Subject_19	Subject_19	139	117	22	TRUE
Subject_20	Subject_20	135	113	22	TRUE

bp_list[[1]]

	patient	pre_bp	post_bp	diff_bp	normal
Subject_1	Subject_1	130	114	16	TRUE
Subject_2	Subject_2	128	98	30	TRUE
Subject_3	Subject_3	116	113	3	TRUE
Subject_4	Subject_4	124	99	25	TRUE
Subject_5	Subject_5	133	107	26	TRUE
Subject_6	Subject_6	134	116	18	TRUE
Subject_7	Subject_7	118	113	5	TRUE
Subject_8	Subject_8	126	111	15	TRUE
Subject_9	Subject_9	114	119	-5	TRUE
Subject_10	Subject_10	127	117	10	TRUE
Subject_11	Subject_11	141	101	40	TRUE
Subject_12	Subject_12	138	119	19	TRUE
Subject_13	Subject_13	128	130	-2	FALSE
Subject_14	Subject_14	140	122	18	FALSE
Subject_15	Subject_15	137	106	31	TRUE
Subject_16	Subject_16	131	106	25	TRUE

Subject_17	Subject_17	120	124	-4	FALSE
Subject_18	Subject_18	128	102	26	TRUE
Subject_19	Subject_19	139	117	22	TRUE
Subject_20	Subject_20	135	113	22	TRUE

```
bp_list$treatment
```

	patient	pre_bp	post_bp	diff_bp	normal
Subject_1	Subject_1	130	114	16	TRUE
Subject_2	Subject_2	128	98	30	TRUE
Subject_3	Subject_3	116	113	3	TRUE
Subject_4	Subject_4	124	99	25	TRUE
Subject_5	Subject_5	133	107	26	TRUE
Subject_6	Subject_6	134	116	18	TRUE
Subject_7	Subject_7	118	113	5	TRUE
Subject_8	Subject_8	126	111	15	TRUE
Subject_9	Subject_9	114	119	-5	TRUE
Subject_10	Subject_10	127	117	10	TRUE
Subject_11	Subject_11	141	101	40	TRUE
Subject_12	Subject_12	138	119	19	TRUE
Subject_13	Subject_13	128	130	-2	FALSE
Subject_14	Subject_14	140	122	18	FALSE
Subject_15	Subject_15	137	106	31	TRUE
Subject_16	Subject_16	131	106	25	TRUE
Subject_17	Subject_17	120	124	-4	FALSE
Subject_18	Subject_18	128	102	26	TRUE
Subject_19	Subject_19	139	117	22	TRUE
Subject_20	Subject_20	135	113	22	TRUE

Question 4

```
#Displaying the pre_bp column from the placebo dataset
bp_list[[2]][2]
```

	pre_bp
1	138
2	135
3	147
4	117
5	152

6	134
7	114
8	121
9	131
10	130

Task Four

Question 1

```
#Initializing status columns in both treatment and placebo datasets
bp_list$treatment$status <- character(20)
bp_list$placebo$status <- character(10)
```

Question 2

```
#Using a for loop to iterate through the 20 treatment patients
#and classify the post-treatment bp as high, borderline, or optimal
for(i in 1:20){
  if(bp_list$treatment$post_bp[i] > 130) {
    bp_list$treatment$status[i] <- "high"
  } else if(bp_list$treatment$post_bp[i] > 120){
    bp_list$treatment$status[i] <- "borderline"
  } else {
    bp_list$treatment$status[i] <- "optimal"
  }
}

bp_list$treatment
```

	patient	pre_bp	post_bp	diff_bp	normal	status
Subject_1	Subject_1	130	114	16	TRUE	optimal
Subject_2	Subject_2	128	98	30	TRUE	optimal
Subject_3	Subject_3	116	113	3	TRUE	optimal
Subject_4	Subject_4	124	99	25	TRUE	optimal
Subject_5	Subject_5	133	107	26	TRUE	optimal
Subject_6	Subject_6	134	116	18	TRUE	optimal
Subject_7	Subject_7	118	113	5	TRUE	optimal
Subject_8	Subject_8	126	111	15	TRUE	optimal
Subject_9	Subject_9	114	119	-5	TRUE	optimal
Subject_10	Subject_10	127	117	10	TRUE	optimal
Subject_11	Subject_11	141	101	40	TRUE	optimal
Subject_12	Subject_12	138	119	19	TRUE	optimal
Subject_13	Subject_13	128	130	-2	FALSE	borderline
Subject_14	Subject_14	140	122	18	FALSE	borderline

Subject_15	Subject_15	137	106	31	TRUE	optimal
Subject_16	Subject_16	131	106	25	TRUE	optimal
Subject_17	Subject_17	120	124	-4	FALSE	borderline
Subject_18	Subject_18	128	102	26	TRUE	optimal
Subject_19	Subject_19	139	117	22	TRUE	optimal
Subject_20	Subject_20	135	113	22	TRUE	optimal

Question 3

```
#Using a for loop to iterate through the 10 placebo patients
#and classify the post-placebo bp as high, borderline, or optimal
for(i in 1:10){
  if(bp_list$placebo$post_bp[i] > 130) {
    bp_list$placebo$status[i] <- "high"
  } else if(bp_list$placebo$post_bp[i] > 120){
    bp_list$placebo$status[i] <- "borderline"
  } else {
    bp_list$placebo$status[i] <- "optimal"
  }
}

bp_list$placebo
```

	names_plac	pre_bp	post_bp	diff_bp	normal	status
1	Subject_1	138	105	33	TRUE	optimal
2	Subject_2	135	136	-1	FALSE	high
3	Subject_3	147	123	24	FALSE	borderline
4	Subject_4	117	130	-13	FALSE	borderline
5	Subject_5	152	134	18	FALSE	high
6	Subject_6	134	143	-9	FALSE	high
7	Subject_7	114	135	-21	FALSE	high
8	Subject_8	121	139	-18	FALSE	high
9	Subject_9	131	120	11	FALSE	optimal
10	Subject_10	130	124	6	FALSE	borderline

Task Five

```
#Writing a function that takes in a list of two dataframes, and calculates a
#certain statistic for the pre, post, and difference bp columns of both dataframes
task5_func <- function(list, stat = "mean") {
  my_fun <- get(stat)
  vec_values <- c(my_fun(list[[1]]$pre_bp),
                  my_fun(list[[1]]$post_bp),
                  my_fun(list[[1]]$diff_bp),
                  my_fun(list[[2]]$pre_bp),
                  my_fun(list[[2]]$post_bp),
                  my_fun(list[[2]]$diff_bp)
                )
  vec_names <- c(paste(attributes(list)$names[[1]], "pre_bp", stat, sep = " "),
                 paste(attributes(list)$names[[1]], "post_bp", stat, sep = " "),
                 paste(attributes(list)$names[[1]], "diff_bp", stat, sep = " "),
                 paste(attributes(list)$names[[2]], "pre_bp", stat, sep = " "),
                 paste(attributes(list)$names[[2]], "post_bp", stat, sep = " "),
                 paste(attributes(list)$names[[2]], "diff_bp", stat, sep = " ")
                )
  names(vec_values) <- vec_names
  return(list(vec_values))
}
```

```
#Using the function to calculate mean, var, sd, min, and max
#from the bp treatment and placebo datasets
task5_func(bp_list)
```

```
[[1]]
treatment pre_bp mean treatment post_bp mean treatment diff_bp mean
          129.35          112.35          17.00
placebo pre_bp mean placebo post_bp mean placebo diff_bp mean
          131.90          128.90           3.00
```

```
task5_func(bp_list, "var")
```

```
[[1]]
treatment pre_bp var treatment post_bp var treatment diff_bp var
          64.55526          74.76579          153.68421
placebo pre_bp var placebo post_bp var placebo diff_bp var
          149.87778          124.98889          341.33333
```

```
task5_func(bp_list, "sd")
```

```
[[1]]
```

treatment	pre_bp	sd	treatment	post_bp	sd	treatment	diff_bp	sd
		8.034629			8.646721			12.396944
placebo	pre_bp	sd	placebo	post_bp	sd	placebo	diff_bp	sd
		12.242458			11.179843			18.475209

```
task5_func(bp_list, "min")
```

```
[[1]]
```

treatment	pre_bp	min	treatment	post_bp	min	treatment	diff_bp	min
		114			98			-5
placebo	pre_bp	min	placebo	post_bp	min	placebo	diff_bp	min
		114			105			-21

```
task5_func(bp_list, "max")
```

```
[[1]]
```

treatment	pre_bp	max	treatment	post_bp	max	treatment	diff_bp	max
		141			130			40
placebo	pre_bp	max	placebo	post_bp	max	placebo	diff_bp	max
		152			143			33