# R 프로그래밍 #3

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# Top-programming-languages (2018)

Language Rank	Types	Spectrum Ranking
1. Python	● 🖵 🛢	100.0
2. C++		99.7
3. Java		97.5
4. C		96.7
<b>5.</b> C#		89.4
6. PHP		84.9
<b>7.</b> R	<b>-</b>	82.9
8. JavaScript		82.6
<b>9.</b> Go	₩ 🖵	76.4
10. Assembly		74.1

https://spectrum.ieee.org/at-work/innovation/the-2018-top-programming-languages

#### In the last class

- Object vector, factor, matrix, ...
- vector numeric, character, logical
- matrix Two dimension numeric, char, logical...

```
> seq
> rep
> length
> which
> paste
> sample
> matrix
```

matrix indexing

#### In the last class

• Define a function

```
my_sine <- function(x){
     y <- sin(x)
     return(y)
}</pre>
```

- Load once (Ctrl + Enter)
- Use

```
> my_sine(pi)
```

- This returns the sine of pi
  - one parameter: x
  - one argument: pi

# Exercise 3-1) matrix

# make directory C:\Rprog\03
setwd(" C:/Rprog/03 ")

- 1. Build a matrix, 'mymat' consisting of 100 rows and 3 columns with initial value of 0
- Generate 1 to 100 sequential values and save them to the first column of mymat
- 3. Generate 100 odd numbers ranging from 1 to 200 and save them to the second column of mymat
- 4. Generate 100 even numbers ranging from 1 to 200 and save them to the third column of mymat
- 5. Show values at 2, 3, 4, 5 rows in the second column
- 6. Substrate 1 from all elements of mymat
- 7. Substrate values in the second column from the values in the first column, and save them to 'mysub'
- 8. Show the mean/sum/squared sum of mysub

#### Object – <del>vector</del>, <del>factor</del>, <del>matrix</del>, data.frame, list

- Basic data structure in R
  - Numeric vector
  - Logical vector
  - Character vector
- Use 'class' function

```
> x <- c(10.4, 6.4, 1.7)

> class(x)

> x <- c(10.4, 6.4, 1.7, "test")

> matrix(c(1,2,3,4), 2, 2)

> matrix(c(1,2,3,"4"), 2, 2)

> matrix(c(1,2,3,TRUE), 2, 2)
```

# **Object - Data frames**

A preferred way to store data in R

```
ids <- 1:10
idnames <- paste("Name", ids, sep="")
students <- data.frame(ids, idnames)
students
class(ids)
class(idnames)
class(students)
class(students[,1])
class(students[,2])
students <- data.frame(ids, idnames, stringsAsFactors = F)
class(students[,2])
students[1,]</pre>
```

# variables ids idnames 1 Name1 2 Name2 3 Name3 4 Name4 5 Name5 6 Name5 6 Name6 7 Name7 8 Name8 9 Name9 10 Name10

# Data frames indexing

Collection of variables → use \$

```
students$ids
students[,1]
students[,"ids"]
```

## **Object - Lists**

A collection of objects
It could consist of vectors, matrix, array, functions, and so on

```
parent_names <- c("Fred", "Mary")
number_of_children <- 2
child_ages <- c(4, 7, 9)
data.frame(parent_names, number_of_children, child_ages)
lst <- list(parent_names, number_of_children, child_ages)
lst[1]
lst[[1]]
class(lst[1])
class(lst[1])
lst[[1]][1]
lst[[1]][c(1,2)]</pre>
```

# Data I/O

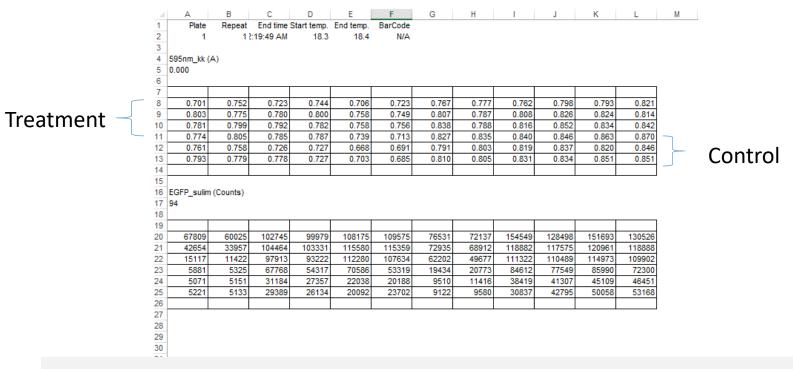
# Text file write / read

```
x <- c(1,2,3,4)
y <- c(5,6,7,8)
xy<-data.frame(x=x, y=y)

write.table(xy, file="table_write.txt")
write.table(xy, file="table_write.txt", quote=F)
write.table(xy, file="table_write.txt", quote=F, row.names=F)
write.table(xy, file="table_write.txt", quote=F, row.names=F, sep=",")
write.table(xy, file="table_write.csv", quote=F, row.names=F, sep=",")</pre>
```

```
mydata<-read.table(file="table_write.csv")
mydata<-read.table(file="table_write.csv", sep=",")
mydata<-read.table(file="table_write.csv", sep=",", header=T)
plot(mydata$x, mydata$z)
```

#### Data read from excel files



Mean difference between Treatment vs. Control in each sample



# **Excel**

6												
7			Т		Т							
8	0.701	0.752	0.723	0.744	0.706	0.723	0.767	0.777	0.762	0.798	0.793	0.82
9	0.803	0.775	0.780	0.800	0.758	0.749	0.807	0.787	0.808	0.826	0.824	0.81
10	0.781	0.799	0.792	0.782	0.758	0.756	0.838	0.788	0.816	0.852	0.834	0.84
11	0.774	0.805	0.785	0.787	0.739	0.713	0.827	0.835	0.840	0.846	0.863	0.87
12	0.761	0.758	0.726	0.727	0.668	0.691	0.791	0.803	0.819	0.837	0.820	0.84
13	0.793	0.779	0.778	0.727	0.703	0.685	0.810	0.805	0.831	0.834	0.851	0.85
14												
15												
16	EGFP_sulim	(Counts)										
17	94											
18												
19												
20	67809	60025	102745	99979	108175	109575	76531	72137	154549	128498	151693	13052
21	42654	33957	104464	103331	115580	115359	72935	68912	118882	117575	120961	11888
22	15117	11422	97913	93222	112280	107634	62202	49677	111322	110489	114973	10990
23	5881	5325	67768	54317	70586	53319	19434	20773	84612	77549	85990	7230
24	5071	5151	31184	27357	22038	20188	9510	11416	38419	41307	45109	4645
25	5221	5133	29389	26134	20092	23702	9122	9580	30837	42795	50058	5316
26												
27												
28												
29												
30												
31	41860								128251		_	11977
32	5391	5203	42780	35936	37572	32403	12689	13923	51289	53884	60386	5730
33												
34												
35												



# readxl package

Download Rprog04-fl.xls from <a href="https://github.com/greendaygh/Rprog2019">https://github.com/greendaygh/Rprog2019</a> Link: <a href="https://github.com/greendaygh/Rprog2019/raw/master/Rprog04-fl.xls">https://github.com/greendaygh/Rprog2019/raw/master/Rprog04-fl.xls</a> copy the file to working directory

```
install.packages(readx1)
library(readx1)
dir()

mydata <- read_excel("Rprog04-fl.xls", sheet=2, skip = 6, col_names=F)

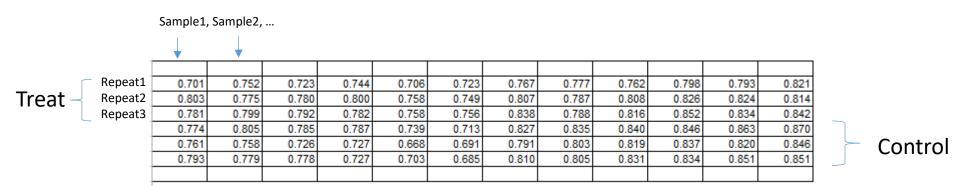
myod <- as.data.frame(mydata[1:9, ])
mygfp <- as.data.frame(mydata[12:21, ])

myod[,1] <- as.numeric(myod[,1])
mygfp[,1] <- as.numeric(mygfp[,1])</pre>
```

- \*\* tibble vs. data.frame
- never changes type of the inputs
- never changes name of variables
- never create rownames

모든 경우 그렇지는 않으며 해당 excel file 특이적으로 첫번째 컬럼이 character임

# **Data manipulation**



```
# OD
myod_treat <- myod[2:4,]
myod_control <- myod[5:7,]

sample_names <- paste("Sample", c(1:12), sep="")
replicate_labels <- paste("Rep", c(1:3), sep="")

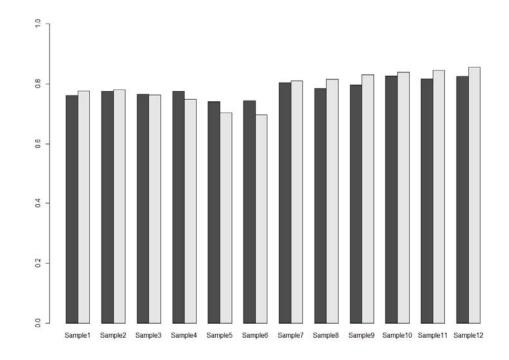
rownames(myod_treat) <- replicate_labels
colnames(myod_treat) <- sample_names
rownames(myod_control) <- replicate_labels
colnames(myod_control) <- sample_names</pre>
```

# Mean comparison

```
mean_treat <- colMeans(myod_treat)
mean_control <- colMeans(myod_control)

plot(mean_treat, type="h")
barplot(mean_treat, ylim=c(0,1))

mean_test <- data.frame(mean_treat, mean_control)
barplot(t(mean_test), ylim=c(0,1), beside=T)</pre>
```



## Exercise 3-2) standard deviation

- Make a function for computing standard deviations
  - Function name: mysd
  - Input parameter: inx (numeric vector, length(inx)>1)
  - Return: standard deviation of inx
- Generate a variable x with 10 random numbers using 'sample' from 1 to 100
- Use mysd to compute the standard deviation of x

sample mean = 
$$\bar{x} = \frac{1}{n}(x_1 + x_2 + \dots + x_n) = \frac{1}{n}\sum_{i} x_i$$
  
sample standard deviation =  $\sqrt{s^2} = \sqrt{\frac{1}{n-1}\sum_{i}(x_i - \bar{x})^2}$ 

# **Apply**

```
Sample1 Sample2 Sample3 Sample4 Sample5 Sample6 Sample7 Sample8 Sample9 Sample10 Sample11 Sample12 Rep1 0.7738588 0.8049214 0.7846458 0.7871608 0.7393147 0.7132604 0.8267264 0.8352386 0.8397562 0.8459013 0.8631678 0.8699542 Rep2 0.7607952 0.7582134 0.7259247 0.7272937 0.6677032 0.6911640 0.7911676 0.8031119 0.8193607 0.8374564 0.8198136 0.8460365 Rep3 0.7925900 0.7791847 0.7780503 0.7274179 0.7033402 0.6846401 0.8104981 0.8053088 0.8314057 0.8338089 0.8511754 0.8506644
```

```
apply(myod_control, 1, mean)
apply(myod_control, 2, mean)

1: row (가로)
2: column (세로)
```

```
apply(myod_control, 2, function(x){
   xmean <- mean(x)
   return(xmean)
})
apply(myod_control, 2, sd)</pre>
```

# The apply function

For matrices, vectorized functions are applied to each element

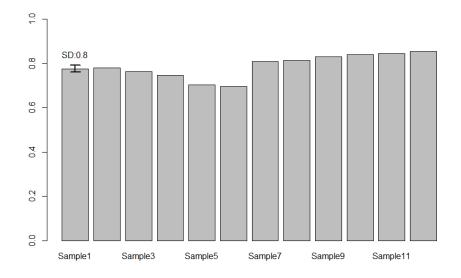
```
# a common pattern to generate matrix
rowSums(m)
colSums(m)
apply(m, 1, mean)
apply(m, 2, mean)
sapply(m, sum)

?sweep
sweep(m, 1, 10)
sweep(m, 1, 10, "+")
sweep(m, 1, 10, "/")
```

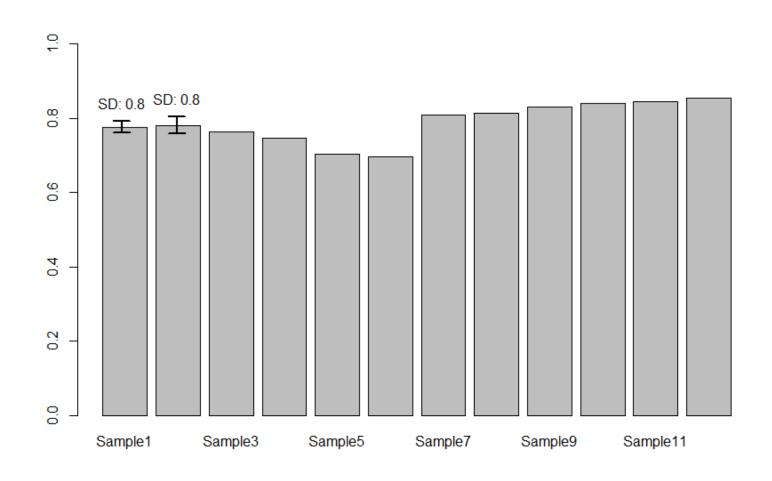
# barplot with sd

```
control_mean <- apply(myod_control, 2, function(x){mean(x)})
control_sd <- apply(myod_control, 2, mysd)

barplot(control_mean, width=0.8, space=0.2, col="gray", ylim=c(0,1))
arrows(0.5, control_mean[1], 0.5, control_mean[1]+control_sd[1], length=0.1, angle=90)
arrows(0.5, control_mean[1], 0.5, control_mean[1]-control_sd[1], length=0.1, angle=90)
lab <- paste("SD:", round(control_mean[1]+control_sd[1],1))
text(0.5, control_mean[1]+control_sd[1]+0.05, labels = lab)</pre>
```



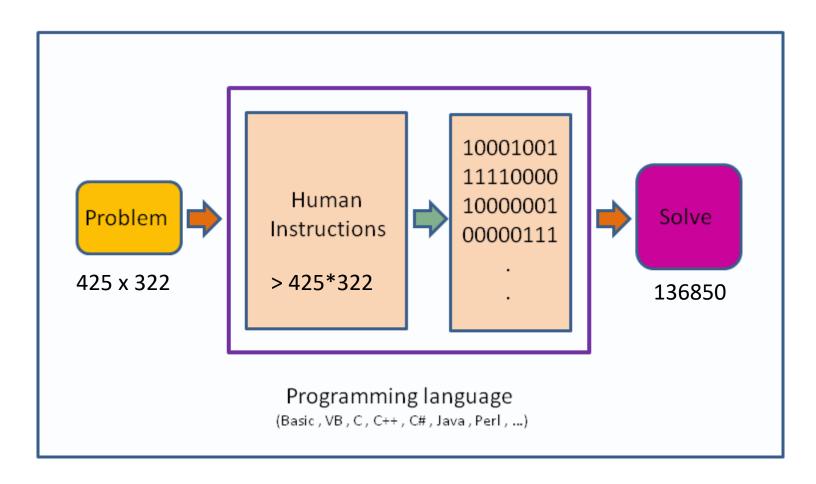
# Exercise 3-3) barplot with sd



# Programming

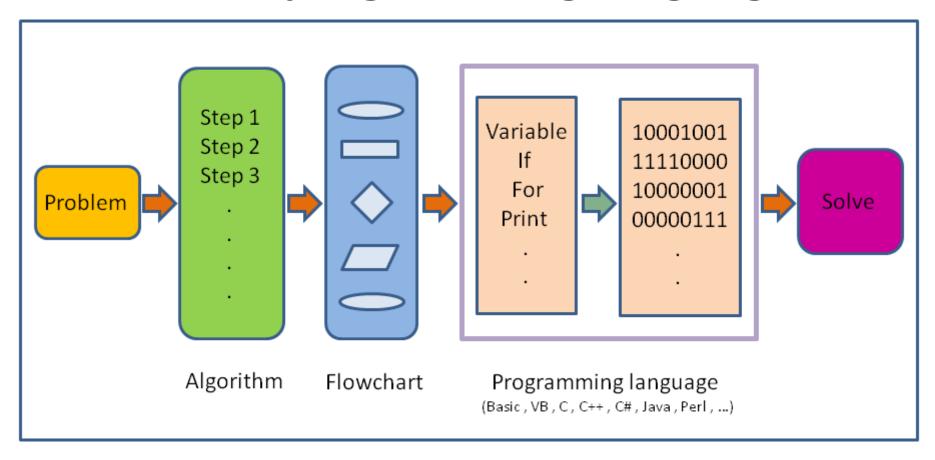
if, if else, for, while, break

# What is a programming language?



https://en.wikiversity.org/wiki/Introduction\_to\_Programming/About\_Programming

# What is a programming language?



# if / if else

```
if(condition) {
       expression
if(condition) {
       expression1
} else {
       expression2
if(condition) {
       expression1
} else if (condition2) {
       expression2
} else {
       expression3
```

All of the binary operators in R are vectorized, operating element by element on their arguments, recycling values as needed. These operators include:

+	addition	-	subtraction	*	${\it multiplication}$
/	division	^	Exponentiation	%%	Modulus
		%/%	Integer Division		

Comparison operators will return the logical values TRUE or FALSE, or NA if any elements involved in a comparison are missing.

<	less than	>	greater than	<=	l.t. or equal
>=	g.t. or equal		equality	!	non-equality

Logical operators come in elementwise and pairwise forms.

&	elementwise and	&&	pairwise and	!	negation
-	elementwise or	П	pairwise or	xor()	exclusive or

The %in% operator can be used to test if a value is present in a vector or array.

# if / if else

```
x <- 5
if(x > 0){
   print("Positive number")
if(x > 0) print("Positive number")
x <- -5
if(x > 0){
       print("Non-negative number")
} else {
       print("Negative number")
if(x > 0)
       print("Non-negative number")
else
       print("Negative number")
```

# for loop

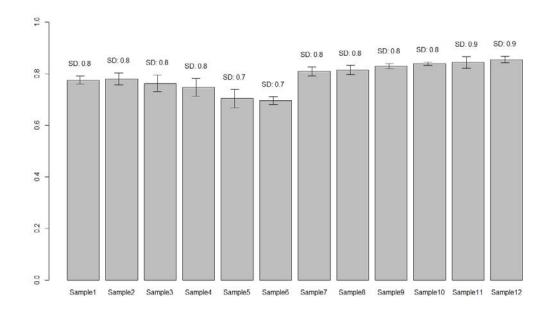
```
for(val in sequence){
                statement
i ← value
          x \leftarrow c(2,5,3,9,8,11,6)
           count <- 0
           for(i in x) {
                if(i %% 2 == 0) count <- count+1
           print(count)
          x \leftarrow c(2,5,3,9,8,11,6)
i ← index
           count <- 0
           for(i in 1:length(x)) {
                  val <- x[i]
               if(val > mean(x)) count <- count+1</pre>
           print(count)
           cat("count:", count, "\n")
```

# Exercise 3-4) for

```
barplot(control_mean, width=0.83, space=0.2, col="gray", ylim=c(0,1))

arrows(0.5, control_mean[1], 0.5, control_mean[1]+control_sd[1], length=0.1, angle=90)
arrows(0.5, control_mean[1], 0.5, control_mean[1]-control_sd[1], length=0.1, angle=90)
lab <- paste("SD:", round(control_mean[1]+control_sd[1],1))
text(0.5, control_mean[1]+control_sd[1]+0.05, labels = lab)

arrows(1.5, control_mean[2], 1.5, control_mean[2]+control_sd[2], length=0.1, angle=90)
arrows(1.5, control_mean[2], 1.5, control_mean[2]-control_sd[2], length=0.1, angle=90)
lab <- paste("SD:", round(control_mean[2]+control_sd[2],1))
text(1.5, control_mean[2]+control_sd[2]+0.05, labels = lab)</pre>
```



# Exercise 3-5) Writing a function for...

- read excel file (case, control)
- compare means
  - compute difference in each sample
  - plot bar graph
- return mean differences as a numeric vector

```
mean_comparison <- function(filename, plot_flag){
    # read excel file
    # get case control matrix (OD values only)
    # compute mean, sd
    # barplot
    # return mean differences
}</pre>
```

# What if...

595nm_kk (	(A)								
0.000									
0.701	0.752	0.723	0.744	0.706		0.777	0.762		
0.803	0.775	0.780	0.800	0.758		0.787	0.808		
0.781	0.799	0.792	0.782	0.758		0.788	0.816		
0.774	0.805	0.785	0.787	0.739		0.835	0.840		
0.761	0.758	0.726	0.727	0.668					

#### **Next**

- R programming
  - maze
- Data manipulation
  - dplyr