

# Geospatial Analysis with R

Class 10



# Today

- A little R/Rmarkdown syntax
- Control structures (\*apply)

# Create your own data

- Create three matrices, `m1`, `m2`, `m3`
  - `m1`:
    - Random number seed: `set.seed(100)`
    - `V1` = 1:20
    - `V2` 20 random numbers from 1:100
    - `V3` 20 numbers from `rnorm`, mean = 500, sd = 100)
  - `m2`: Same variables, but `set.seed(200)`
  - `m3`: Same variables, but `set.seed(300)`
- Create vector `v`: 20 random draws from `LETTERS[1:5]`; `set.seed(1)`
- Create 2 `data.frames`: `dat1` (`m1` & `v`), `dat2` (`m2` & `v`)
  - Name the column holding `v` "GRP"
- Create list `l` combining `m1`, `m2`, `m3`, `dat1`, `dat2`, `v`
  - Name the list elements the same as the object name.

```
set.seed(100)
m1 <- cbind(V1 = 1:20, V2 = sample(1:100, size = 20, replace = TRUE),
            V3 = rnorm(n = 20, mean = 500, sd = 100))
set.seed(200)
m2 <- cbind(V1 = 1:20, V2 = sample(1:100, size = 20, replace = TRUE),
            V3 = rnorm(n = 20, mean = 500, sd = 100))
set.seed(300)
m3 <- cbind(V1 = 1:20, V2 = sample(1:100, size = 20, replace = TRUE),
            V3 = rnorm(n = 20, mean = 500, sd = 100))
set.seed(1)
v <- sample(LETTERS[1:5], size = 20, replace = TRUE)
dat1 <- data.frame(m1, "GRP" = v)
dat2 <- data.frame(m2, "GRP" = v)
l <- list(m1, m2, m3, dat1, dat2, v)
names(l) <- c("m1", "m2", "m3", "dat1", "dat2", "v")
```

# Control structures

- Branching
  - Not usually vectorized
  - `ifelse` for vectorized branching
  - often done within looping
- Looping
  - `for`, `while`
  - `*apply`
    - A special form of looping
    - Intended for *applying* a function to data
    - Returns results of loop directly into object

# lapply

- Apply function to vector, return list

```
l2 <- l[c("m1", "m2", "m3")]  
lapply(l2, mean)
```

```
## $m1  
## [1] 187.5092  
##  
## $m2  
## [1] 183.919  
##  
## $m3  
## [1] 197.5562
```

# sapply

- Apply function to vector, return simplest possible output

```
l2 <- l[c("m1", "m2", "m3")]  
sapply(l2, mean)
```

```
##           m1           m2           m3  
## 187.5092 183.9190 197.5562
```

# apply

- Apply function to margins of array or matrix, return vector, array, or list

```
apply(l2[[1]], MARGIN = 1, FUN = mean)
```

```
## [1] 180.3295 179.2091 179.6122 194.6613 188.1127 184.0228 183.3715  
## [8] 199.0285 157.5395 253.0099 176.7303 225.8020 189.3987 210.4468  
## [15] 170.1874 179.7183 155.3259 192.3648 146.4090 204.9025
```

```
apply(l2[[1]], MARGIN = 2, FUN = mean)
```

```
##      V1      V2      V3  
## 10.5000 46.8000 505.2275
```



## \*apply with anonymous functions

- Use anonymous functions to pass an iterator
- For more complex operations inside `{ }`

```
# Simple  
o <- lapply(1:2, function(x) l2[[x]])  
o
```

```
## [[1]]  
##      V1 V2      V3  
## [1,]  1 31 508.9886  
## [2,]  2 26 509.6274  
## [3,]  3 56 479.8366  
## [4,]  4  6 573.9840  
## [5,]  5 47 512.3380  
## [6,]  6 49 497.0683  
## [7,]  7 82 461.1146  
## [8,]  8 38 551.0856  
## [9,]  9 55 408.6186  
## [10,] 10 18 731.0297  
## [11,] 11 63 456.1910  
## [12,] 12 89 576.4061  
## [13,] 13 29 526.1961  
## [14,] 14 40 577.3405  
## [15,] 15 77 418.5621  
## [16,] 16 67 456.1549  
## [17,] 17 21 427.9778  
## [18,] 18 36 523.0945
```

```
# More complex
o2 <- lapply(1:5, function(x) {
  12[[1]][x] - 12[[2]][x]
})
o2
```

```
## [[1]]
## [1] 0
##
## [[2]]
## [1] 0
##
## [[3]]
## [1] 0
##
## [[4]]
## [1] 0
##
## [[5]]
## [1] 0
```

## Create data using `lapply`

- Let's recreate our matrix examples

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```
seeds <- c(100, 200, 300) # or 100 * 1:3
l3 <- lapply(seeds, function(x) {
  set.seed(x)
  m <- cbind(V1 = 1:20,
             V2 = sample(1:100, size = 20, replace = TRUE),
             V3 = rnorm(n = 20, mean = 500, sd = 100))
})
names(l3) <- paste0("m", 1:3)
```

# Check the values

- Let's check them now against original values

```
m1 == l3$m1  
m2 == l3$m2  
all(m1 == l3$m1)  
all(m2 == l3$m2)
```

- Etc, but we could do this check with a looping function!

## Check using `lapply`

- We know that `l[1:3]` contains `m1`, `m2`, `m3`, and so does `l3`, so

```
lapply(1:3, function(x) all(l[[x]] == l3[[x]]))
```

- More compact

```
sapply(1:3, function(x) all(l[[x]] == l3[[x]]))
```

## Looping practice

- Write a `for` loop that iterates through the vector `1:10` and prints the iterator `i` multiplied by 10

```
for(i in 1:10) print(i * 10)
```



# Looping practice

- Write a `for` loop that iterates through the vector `1:10` and prints the iterator `i` multiplied by 10
- Do the same, but instead of print `i * 10`, catch the result in a predefined empty list `o`

```
for(i in 1:10) print(i * 10)  
  
o <- list()  
for(i in 1:10) o[[i]] <- i * 10
```

# Looping practice

- Write a `for` loop that iterates through the vector `1:10` and prints the iterator `i` multiplied by 10
- Do the same, but instead of print `i * 10`, catch the result in a predefined empty list `o`
- Do the same as above, but use an `lapply` that assigns output to `o`

```
for(i in 1:10) print(i * 10)

o <- list()
for(i in 1:10) o[[i]] <- i * 10

o <- lapply(1:10, function(x) x * 10)
```

# Looping practice

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- Do the same, but instead of print `i * 10`, catch the result in a predefined empty list `o`
- Do the same as above, but use an `lapply` that assigns output to `o`
- Do the same as above, but use `sapply` instead of `lapply`

```
for(i in 1:10) print(i * 10)

o <- list()
for(i in 1:10) o[[i]] <- i * 10

o <- lapply(1:10, function(x) x * 10)

o <- sapply(1:10, function(x) x * 10)
```

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- Do the same as above, but use an `lapply` that assigns output to `o`
- Do the same as above, but use `sapply` instead of `lapply`
- Let's use `sapply` to find which elements of `l` are `matrix`

```
for(i in 1:10) print(i * 10)

o <- list()
for(i in 1:10) o[[i]] <- i * 10

o <- lapply(1:10, function(x) x * 10)

o <- sapply(1:10, function(x) x * 10)

sapply(l, is.matrix)
```



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- Do the same as above, but use an `lapply` that assigns output to `o`
- Do the same as above, but use `sapply` instead of `lapply`
- Let's use `sapply` to find which elements of `l` are `matrix`
- Let's use `lapply` to calculate the `colMeans` of matrices and `data.frames` in `l`
- Use `lapply` to calculate and output `mean` and `sd` of each matrix in `l`

```
for(i in 1:10) print(i * 10)

o <- list()
for(i in 1:10) o[[i]] <- i * 10

o <- lapply(1:10, function(x) x * 10)

o <- sapply(1:10, function(x) x * 10)

sapply(l, is.matrix)

lapply(l[1:5], function(x) colMeans(x[, 1:3]))
lapply(1:5, function(x) colMeans(l[[x]][, 1:3]))
```

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- Do the same as above, but use an `lapply` that assigns output to `o`
- Do the same as above, but use `sapply` instead of `lapply`
- Let's use `sapply` to find which elements of `l` are `matrix`
- Let's use `lapply` to calculate the `colMeans` of matrices and `data.frames` in `l`
- Use `lapply` to calculate and output `mean` and `sd` of each matrix in `l`
- Do the same, but use conditional to test whether each element of `l` is `matrix`, then calculate `mean`, `sd`.

```
lapply(l, function(x) {  
  if(is.matrix(x)) c("mu" = mean(x), "sd" = sd(x))  
})  
  
sapply(l, function(x) {  
  if(is.matrix(x)) {  
    c("mu" = mean(x), "sd" = sd(x))  
  } else if(is.data.frame(x)) {  
    c("mu" = mean(unlist(x[, 1:3])), "sd" = sd(unlist(x[, 1:3])))  
  } else {  
    c("mu" = NA, "sd" = NA)  
  }  
})
```