

apply_functions

2023-04-04

```
?lapply
?mapply
```

The `apply` functions allow us to apply a function to a vector or list of values iteratively. This helps minimize errors in code and makes the analyses more efficient. With `lapply()` and `sapply` functions, we can only provide one argument to iterate on

With `mapply()`, we can provide multiple arguments to iterate. It probably also returns a vector or simplified data structure as result.

`sapply()` function simplifies the output to a vector (or the simplest data structure possible), while `lapply()` returns an output in the form of a list.

```
get_mass_from_length_theropoda <- function(length){
  mass <- 0.73 * length ^ 3.63
  return(mass)
}
```

```
theropoda_lengths <- c(17.8013631070471, 20.3764452071665, 14.0743486294308, 25.65782386974, 26.0952008)
```

```
get_mass_from_length_theropoda(length = theropoda_lengths)
```

```
## [1] 25262.027 41253.332 10767.568 95233.732 101260.017 40775.516
## [7] 24072.130 4785.145 39129.521 29666.193 26830.297 64700.869
## [13] 42768.180 94697.262 79013.471 103955.226 92798.465 41901.983
## [19] 17439.569 41055.045 37544.201 25198.303 12928.490 36388.290
## [25] 34962.862 80307.929 8854.525 50183.194 28846.165 35735.369
## [31] 115908.187 31765.368 58958.713 5561.862 28349.410 15418.314
## [37] 9218.648 1197.666 94407.873 19552.500
```

```
theropoda_masses <- get_mass_from_length_theropoda(length = theropoda_lengths)
```

```
mylist <- (theropoda_masses)
second_list <- c(mylist, list(c("Luna", "Avi", "Anita")))
second_list[[1]]
```

```
## [1] 25262.03
```

```
data.frame(theropoda_masses, c("Anita", "Avi", "Luna", "Maria"))
```

```
##      theropoda_masses c..Anita....Avi....Luna....Maria..
## 1      25262.027                                Anita
```

## 2	41253.332	Avi
## 3	10767.568	Luna
## 4	95233.732	Maria
## 5	101260.017	Anita
## 6	40775.516	Avi
## 7	24072.130	Luna
## 8	4785.145	Maria
## 9	39129.521	Anita
## 10	29666.193	Avi
## 11	26830.297	Luna
## 12	64700.869	Maria
## 13	42768.180	Anita
## 14	94697.262	Avi
## 15	79013.471	Luna
## 16	103955.226	Maria
## 17	92798.465	Anita
## 18	41901.983	Avi
## 19	17439.569	Luna
## 20	41055.045	Maria
## 21	37544.201	Anita
## 22	25198.303	Avi
## 23	12928.490	Luna
## 24	36388.290	Maria
## 25	34962.862	Anita
## 26	80307.929	Avi
## 27	8854.525	Luna
## 28	50183.194	Maria
## 29	28846.165	Anita
## 30	35735.369	Avi
## 31	115908.187	Luna
## 32	31765.368	Maria
## 33	58958.713	Anita
## 34	5561.862	Avi
## 35	28349.410	Luna
## 36	15418.314	Maria
## 37	9218.648	Anita
## 38	1197.666	Avi
## 39	94407.873	Luna
## 40	19552.500	Maria

```

mass_from_length <- function(length, a,b){
  mass <- 0.73 * length ^ 3.63
  return(mass)
}
new_masses <- mass_from_length(length = theropoda_lengths)
#rm(new_lengths) # The rm function allows to remove objects from the R environment.

theropoda_masses == new_masses

```

```

## [1] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
## [16] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
## [31] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE

```

```
all(theropoda_masses == new_masses) # tests that all values in a logical vector are equal to TRUE
```

```
## [1] TRUE
```

```
all.equal(theropoda_masses, new_masses)
```

```
## [1] TRUE
```

```
a_values <- c(0.759, 0.751, 0.74, 0.746, 0.759, 0.751, 0.749, 0.751, 0.738, 0.768, 0.736, 0.749, 0.746,
```

```
b_values <- c(3.627, 3.633, 3.626, 3.633, 3.627, 3.629, 3.632, 3.628, 3.633, 3.627, 3.621, 3.63, 3.631,
```

```
mass_from_length <- function(length = theropoda_lengths, a = a_values, b = b_values){  
  mass <- a * length ^ b  
  return(mass)  
}
```

<- the scope operator or double arrow, allows creating and modifying objects in parent variables

```
dino_data <- data.frame(theropoda_lengths, a_values, b_values) %>% mutate(massses = get_mass_from_length(theropoda_lengths, a_values, b_values))  
print(dino_data)
```

##	theropoda_lengths	a_values	b_values	massses
## 1	17.801363	0.759	3.627	25262.027
## 2	20.376445	0.751	3.633	41253.332
## 3	14.074349	0.740	3.626	10767.568
## 4	25.657824	0.746	3.633	95233.732
## 5	26.095201	0.759	3.627	101260.017
## 6	20.311154	0.751	3.629	40775.516
## 7	17.566324	0.749	3.632	24072.130
## 8	11.256343	0.751	3.628	4785.145
## 9	20.081903	0.738	3.633	39129.521
## 10	18.607163	0.768	3.627	29666.193
## 11	18.099189	0.736	3.621	26830.297
## 12	23.065969	0.749	3.630	64700.869
## 13	20.579885	0.746	3.631	42768.180
## 14	25.617925	0.744	3.632	94697.262
## 15	24.371433	0.749	3.628	79013.471
## 16	26.284725	0.751	3.626	103955.226
## 17	25.475378	0.744	3.639	92798.465
## 18	20.464209	0.754	3.626	41901.983
## 19	16.073826	0.774	3.635	17439.569
## 20	20.349417	0.751	3.629	41055.045
## 21	19.854399	0.763	3.642	37544.201
## 22	17.788981	0.749	3.632	25198.303
## 23	14.801642	0.741	3.633	12928.490
## 24	19.684091	0.754	3.629	36388.290
## 25	19.468589	0.746	3.620	34962.862
## 26	24.480778	0.755	3.619	80307.929
## 27	13.335996	0.764	3.638	8854.525

```
## 28      21.506599    0.758    3.627  50183.194
## 29      18.464030    0.760    3.621  28846.165
## 30      19.586153    0.748    3.628  35735.369
## 31      27.084752    0.745    3.628 115908.187
## 32      18.960937    0.756    3.635  31765.368
## 33      22.482917    0.739    3.624  58958.713
## 34      11.732572    0.733    3.621   5561.862
## 35      18.375885    0.757    3.621  28349.410
## 36      15.537505    0.747    3.632  15418.314
## 37      13.484875    0.741    3.627   9218.648
## 38       7.685612    0.752    3.624   1197.666
## 39      25.596335    0.752    3.634  94407.873
## 40      16.588285    0.748    3.621  19552.500
```

```
theropoda_lengths < 20
```

```
## [1] TRUE FALSE TRUE FALSE FALSE FALSE TRUE TRUE FALSE TRUE TRUE FALSE
## [13] FALSE FALSE FALSE FALSE FALSE FALSE TRUE FALSE TRUE TRUE TRUE TRUE
## [25] TRUE FALSE TRUE FALSE TRUE TRUE FALSE TRUE FALSE TRUE TRUE TRUE
## [37] TRUE TRUE FALSE TRUE
```

```
mass_from_length_max <- function(length) {
  if(length < 20) {
    mass <- 0.73 * length ^ 3.63
    return(mass)
  } else {
    return(NA)
  }
}
```

```
mass_from_length_max <- function(length) {
  if(length < 20) {
    mass <- 0.73 * length ^ 3.63
    mass <- NA
  }
}
```

```
mass_from_length_max <- function(length) {
  if(length < 20) {
    mass <- 0.73 * length ^ 3.63
    return(mass)
  }
}
```

```
get_mass_from_length_theropoda(length = theropoda_lengths)
```

```
## [1] 25262.027 41253.332 10767.568 95233.732 101260.017 40775.516
## [7] 24072.130 4785.145 39129.521 29666.193 26830.297 64700.869
## [13] 42768.180 94697.262 79013.471 103955.226 92798.465 41901.983
## [19] 17439.569 41055.045 37544.201 25198.303 12928.490 36388.290
```

```
## [25] 34962.862 80307.929 8854.525 50183.194 28846.165 35735.369
## [31] 115908.187 31765.368 58958.713 5561.862 28349.410 15418.314
## [37] 9218.648 1197.666 94407.873 19552.500
```

```
sapply(theropoda_lengths, mass_from_length_max)
```

```
## [[1]]
## [1] 25262.03
##
## [[2]]
## NULL
##
## [[3]]
## [1] 10767.57
##
## [[4]]
## NULL
##
## [[5]]
## NULL
##
## [[6]]
## NULL
##
## [[7]]
## [1] 24072.13
##
## [[8]]
## [1] 4785.145
##
## [[9]]
## NULL
##
## [[10]]
## [1] 29666.19
##
## [[11]]
## [1] 26830.3
##
## [[12]]
## NULL
##
## [[13]]
## NULL
##
## [[14]]
## NULL
##
## [[15]]
## NULL
##
## [[16]]
## NULL
##
```

```

## [[17]]
## NULL
##
## [[18]]
## NULL
##
## [[19]]
## [1] 17439.57
##
## [[20]]
## NULL
##
## [[21]]
## [1] 37544.2
##
## [[22]]
## [1] 25198.3
##
## [[23]]
## [1] 12928.49
##
## [[24]]
## [1] 36388.29
##
## [[25]]
## [1] 34962.86
##
## [[26]]
## NULL
##
## [[27]]
## [1] 8854.525
##
## [[28]]
## NULL
##
## [[29]]
## [1] 28846.17
##
## [[30]]
## [1] 35735.37
##
## [[31]]
## NULL
##
## [[32]]
## [1] 31765.37
##
## [[33]]
## NULL
##
## [[34]]
## [1] 5561.862
##

```

```
## [[35]]
## [1] 28349.41
##
## [[36]]
## [1] 15418.31
##
## [[37]]
## [1] 9218.648
##
## [[38]]
## [1] 1197.666
##
## [[39]]
## NULL
##
## [[40]]
## [1] 19552.5
```

#Homework

```
dinosaur_lengths <- read.csv(file = "../data raw/dinosaur_lengths.csv")
```

```
get_mass_from_length_by_name <- function(length, dinosaur_name) {
  if (dinosaur_name == "Stegosauria") {
    a <- 10.95
    b <- 2.64
  } else if (dinosaur_name == "Theropoda") {
    a <- 0.73
    b <- 3.63
  } else if (dinosaur_name == "Sauropoda") {
    a <- 214.44
    b <- 1.46
  } else {
    a = NA
    b = NA
  }
  mass <- a * length^b
  return(mass)
}
```

```
mapply(get_mass_from_length_by_name, length = dinosaur_lengths$lengths, dinosaur_name = dinosaur_lengths$dinosaur_name)
```

```
## [1] 24341.681 NA NA 22114.190 NA NA
## [7] 57349.470 14160.494 49677.749 42105.917 10221.747 15339.988
## [13] 70624.102 23883.825 28552.864 18801.370 19438.673 NA
## [19] 19607.970 16032.845 NA 50350.112 15969.078 29582.848
## [25] 15201.456 12980.541 9937.867 9599.415 49245.963 23846.751
## [31] 53805.661 53326.467 NA 15554.977 18544.119 NA
## [37] NA 82492.318 17909.041 38694.503 80303.181 19592.802
## [43] 10614.785 29560.809 71658.477 NA 83961.661 NA
## [49] 26284.040 21766.002 63571.873 5480.255 33917.314 22778.032
## [55] 13819.165 21154.149 17635.099 14577.594 NA 14032.340
## [61] 30231.694 NA 11293.886 72743.800 23679.901 64258.574
```

##	[67]	14931.085	16323.818	NA	NA	NA	7599.703
##	[73]	NA	NA	NA	NA	46920.035	70529.031
##	[79]	9484.528	NA	68340.494	44959.626	NA	48249.486
##	[85]	11730.174	NA	52295.177	NA	NA	NA
##	[91]	40358.292	38891.137	30878.439	19125.425	NA	NA
##	[97]	8697.216	19627.357	NA	NA	13411.390	33157.499
##	[103]	10874.733	24554.930	16819.494	18421.449	NA	19645.723
##	[109]	38206.241	53196.019	22346.109	NA	22685.103	NA
##	[115]	13613.983	34685.790	NA	18654.525	NA	101482.428
##	[121]	89149.257	NA	20820.837	NA	22232.852	59702.598
##	[127]	NA	16321.774	22748.880	NA	NA	NA
##	[133]	NA	25987.768	49818.253	13106.766	NA	32112.443
##	[139]	NA	16984.463	10859.926	93973.020	52342.265	19151.788
##	[145]	NA	13954.186	NA	15021.820	35933.327	140435.607
##	[151]	20467.332	23869.639	NA	NA	15211.979	57098.945
##	[157]	23588.700	27381.008	85932.513	NA	9331.295	NA
##	[163]	NA	32005.502	16613.444	7904.857	NA	26352.263
##	[169]	19880.480	15543.679	15493.654	13546.034	NA	36095.081
##	[175]	42437.608	NA	NA	51637.913	NA	44120.181
##	[181]	9535.583	59840.348	NA	NA	NA	44822.176
##	[187]	14232.684	34751.496	11292.437	NA	NA	NA
##	[193]	22002.082	19554.166	13223.770	NA	NA	68935.505
##	[199]	9172.206	90096.476	25796.762	50594.426	61952.966	20132.528
##	[205]	NA	13979.439	15481.074	12104.000	21789.436	54009.090
##	[211]	13812.364	8071.939	21144.506	44097.848	16250.303	70065.996
##	[217]	11170.349	22826.560	40885.088	17292.043	18394.391	50267.629
##	[223]	70791.032	28464.276	41431.346	NA	14242.918	NA
##	[229]	NA	52014.366	32865.058	NA	11906.150	17964.362
##	[235]	14844.497	13079.836	76048.107	18843.875	NA	30737.511
##	[241]	37983.026	18711.957	22636.970	29868.755	42799.606	NA
##	[247]	43632.463	103600.943	NA	NA	10330.761	23659.805
##	[253]	19126.024	17175.845	28017.230	54437.041	NA	20657.057
##	[259]	13275.051	NA	8222.362	NA	108964.075	NA
##	[265]	5845.741	26356.588	NA	59636.239	14857.582	45043.701
##	[271]	47427.024	NA	NA	11807.182	27575.709	18177.367
##	[277]	NA	22108.648	33908.940	NA	NA	NA
##	[283]	NA	45862.941	23366.240	16165.694	10263.470	NA
##	[289]	24026.928	33497.651	NA	15770.110	48190.121	33107.401
##	[295]	20523.437	21387.730	15771.706	12632.938	28352.199	10401.651
##	[301]	41162.369	16740.472	29576.590	28831.907	21622.906	NA
##	[307]	26736.709	18663.882	10872.689	13072.222	35308.681	17145.703
##	[313]	19620.530	1550.370	NA	11509.202	16574.358	94984.150
##	[319]	9448.048	56370.430	NA	47899.078	27521.456	24907.229
##	[325]	12800.024	34456.895	NA	19137.794	9084.302	NA
##	[331]	20396.019	7636.822	15452.482	NA	11482.576	NA
##	[337]	21323.042	17062.973	24482.018	19394.529	61929.256	NA
##	[343]	29113.203	53044.431	17891.216	21665.733	21611.857	13917.623
##	[349]	21715.000	NA	10525.601	31777.548	45932.499	16396.801
##	[355]	NA	21020.829	9499.589	NA	11886.269	13597.168
##	[361]	NA	32610.060	50496.496	23180.857	20838.975	27426.143
##	[367]	51655.501	52241.022	27527.983	40947.425	26691.614	23152.573
##	[373]	43419.737	44236.593	60396.602	15878.961	70561.697	17374.235
##	[379]	10332.362	34844.884	NA	43839.492	NA	10259.928
##	[385]	24344.124	NA	23490.643	15151.289	40052.674	31011.453


```
## [391]      NA 36300.595 28716.671 21434.730      NA 27977.292
## [397] 13912.492      NA      NA 45387.391 21638.866 12782.316
## [403]      NA      NA      NA 74279.377 19250.194 19647.872
## [409] 39022.265      NA      NA 9446.876 33097.292      NA
## [415] 23694.389 15501.027 13490.363 7311.070 63156.403 40543.550
## [421] 19942.976      NA      NA 26888.995      NA 18102.809
## [427] 125939.133      NA      NA 14393.863      NA 62045.506
## [433] 60194.052 36753.957      NA      NA 32061.537      NA
## [439] 67466.670 17627.746 24171.682 25917.752 67098.902      NA
## [445] 17699.295 18903.752 13127.745 17295.450 42209.926 23426.667
## [451] 118937.988      NA 18165.832      NA 46816.660      NA
## [457] 53237.908 23121.375 25937.746      NA 47637.068      NA
## [463] 127540.554      NA 12313.099 24276.516 15500.675 16109.794
## [469] 15965.471 54296.492      NA      NA 14365.977 153749.934
## [475] 59143.016 18524.301 6227.675 13606.978      NA      NA
## [481] 49146.996 103896.484 38059.728 41076.716      NA 30013.153
## [487] 41805.513 20113.277 24071.440      NA      NA 8489.727
## [493] 24349.181      NA      NA 44921.367 26262.993 16883.382
## [499] 14444.693      NA
```

```
dinosaur_lengths %>%
rowwise %>%
mutate(masses = get_mass_from_length_by_name(lengths, species))
```

```
## # A tibble: 500 x 3
## # Rowwise:
##   species      lengths masses
##   <chr>         <dbl>   <dbl>
## 1 Stegosauria    18.5 24342.
## 2 Ankylosauria   16.4    NA
## 3 Ankylosauria   23.7    NA
## 4 Sauropoda      23.9 22114.
## 5 Ankylosauria   21.7    NA
## 6 Ankylosauria   21.4    NA
## 7 Theropoda      22.3 57349.
## 8 Theropoda      15.2 14160.
## 9 Theropoda      21.4 49678.
## 10 Stegosauria   22.8 42106.
## # ... with 490 more rows
```

```
library("ggplot2")
ggplot(data = dinosaur_lengths, mapping = aes(x = lengths)) +
geom_histogram() +
facet_wrap(~species)
```

```
## 'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.
```

