

Apply functions

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11/15/2022

Exercise 1

Write a function named `mass_from_length_theropoda()` that takes `length` as an argument to get an estimate of mass for Theropoda dinosaurs. Use the equation $\text{mass} <- 0.73 * \text{length}^{3.63}$. Copy and run the code below to generate the object `theropoda_lengths` in your R environment. Pass the entire vector to your function (by giving it as value for the `length` argument); this calculates the mass for each length value in the vector `theropoda_lengths`.

```
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)
```

Calculate the mass for each length values in the vector 'get_mass_from_length_theropoda'

```
get_mass_from_length_theropoda(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
```

Create a new version of the function named `mass_from_length()` that uses the equation $\text{mass} <- a * \text{length}^b$ and takes `length`, `a` and `b` as arguments. In the function arguments, set the default values for `a` to 0.73 and `b` to 3.63. If you run this function with just the `length` data from Part 1, you should get the same result as Part 1. Copy the data below into R and call your function using the vector of lengths from Part 1 (above) and these vectors of `a` and `b` values to estimate the mass for the dinosaurs using different values of `a` and `b`.

```
mass_from_length <- function(length, a = 0.73, b = 3.63) {  
  mass <- a * length^b  
  return(mass)  
}  
  
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(length = theropoda_lengths, a = a_values, b = b_values)
```

```
## [1] 26039.686 42825.603 10800.224 98273.049 104257.481 41822.386
## [7] 24840.644 4899.022 39915.948 30937.922 26354.908 66384.865
## [13] 43837.944 97141.451 80553.856 105556.405 97374.660 42760.136
## [19] 18749.274 42109.012 40674.182 26003.425 13229.824 37472.789
## [25] 34684.033 80187.272 9460.977 51630.571 29253.772 36399.306
## [31] 117511.962 33384.288 58581.226 5462.316 28637.745 15864.172
## [37] 9284.810 1218.755 98522.609 19534.524
```

```
mapply(mass_from_length, theropoda_lengths, a_values, b_values )
```

```
## [1] 26039.686 42825.603 10800.224 98273.049 104257.481 41822.386
## [7] 24840.644 4899.022 39915.948 30937.922 26354.908 66384.865
## [13] 43837.944 97141.451 80553.856 105556.405 97374.660 42760.136
## [19] 18749.274 42109.012 40674.182 26003.425 13229.824 37472.789
## [25] 34684.033 80187.272 9460.977 51630.571 29253.772 36399.306
## [31] 117511.962 33384.288 58581.226 5462.316 28637.745 15864.172
## [37] 9284.810 1218.755 98522.609 19534.524
```

Create a data frame for this data using the code `dino_data <- data.frame(theropoda_lengths, a_values, b_values)`. Use `dplyr` to add a new masses column to this data frame (using `mutate()` and your function) and print the result to the console.

```
dino_data <- data.frame(length = theropoda_lengths, as = a_values, bs = b_values)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
## filter, lag

## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

```
dino_data %>%
  mutate(masses = mass_from_length(length, as, bs))
```

```
##      length    as    bs    masses
## 1 17.801363 0.759 3.627 26039.686
## 2 20.376445 0.751 3.633 42825.603
## 3 14.074349 0.740 3.626 10800.224
## 4 25.657824 0.746 3.633 98273.049
## 5 26.095201 0.759 3.627 104257.481
## 6 20.311154 0.751 3.629 41822.386
## 7 17.566324 0.749 3.632 24840.644
## 8 11.256343 0.751 3.628 4899.022
## 9 20.081903 0.738 3.633 39915.948
## 10 18.607163 0.768 3.627 30937.922
## 11 18.099189 0.736 3.621 26354.908
```

```
## 12 23.065969 0.749 3.630 66384.865
## 13 20.579885 0.746 3.631 43837.944
## 14 25.617925 0.744 3.632 97141.451
## 15 24.371433 0.749 3.628 80553.856
## 16 26.284725 0.751 3.626 105556.405
## 17 25.475378 0.744 3.639 97374.660
## 18 20.464209 0.754 3.626 42760.136
## 19 16.073826 0.774 3.635 18749.274
## 20 20.349417 0.751 3.629 42109.012
## 21 19.854399 0.763 3.642 40674.182
## 22 17.788981 0.749 3.632 26003.425
## 23 14.801642 0.741 3.633 13229.824
## 24 19.684091 0.754 3.629 37472.789
## 25 19.468589 0.746 3.620 34684.033
## 26 24.480778 0.755 3.619 80187.272
## 27 13.335996 0.764 3.638 9460.977
## 28 21.506599 0.758 3.627 51630.571
## 29 18.464030 0.760 3.621 29253.772
## 30 19.586153 0.748 3.628 36399.306
## 31 27.084752 0.745 3.628 117511.962
## 32 18.960937 0.756 3.635 33384.288
## 33 22.482917 0.739 3.624 58581.226
## 34 11.732572 0.733 3.621 5462.316
## 35 18.375885 0.757 3.621 28637.745
## 36 15.537505 0.747 3.632 15864.172
## 37 13.484875 0.741 3.627 9284.810
## 38 7.685612 0.752 3.624 1218.755
## 39 25.596335 0.752 3.634 98522.609
## 40 16.588285 0.748 3.621 19534.524
```

Create a new version of your `mass_from_length_theropoda()` function from Part 1 of Exercise 1 called `mass_from_length_max()`. This function should only calculate a mass if the value of length passed to the function is less than 20. If length is greater than 20, return NA instead.

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

Use `sapply()` and this new function to estimate the mass for the `theropoda_lengths` data from Exercise 1.

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

```
## [1] 25262.027      NA 10767.568      NA      NA      NA 24072.130
## [8] 4785.145       NA 29666.193 26830.297      NA      NA      NA
## [15]      NA      NA      NA      NA 17439.569      NA 37544.201
## [22] 25198.303 12928.490 36388.290 34962.862      NA 8854.525      NA
## [29] 28846.165 35735.369      NA 31765.368      NA 5561.862 28349.410
## [36] 15418.314 9218.648 1197.666      NA 19552.500
```

A Data Set of Dinosaur Lengths

Download the CSV file of data on dinosaur lengths with species names into your data folder and import it using `read.csv()`

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

Write a function `get_mass_from_length_by_name()` that uses the equation $\text{mass} <- a * \text{length}^b$ to estimate the size of a dinosaur from its length. This function should take two arguments, the length and the name of the dinosaur group. Inside this function use if/else if/else statements to check to see if the name is one of the following values and if so set `a` and `b` to the appropriate values.

Stegosauria: $a = 10.95$ and $b = 2.64$ (Seebacher 2001). Theropoda: $a = 0.73$ and $b = 3.63$ (Seebacher 2001). Sauropoda: $a = 214.44$ and $b = 1.46$ (Seebacher 2001). If the name is not any of these values set `a = NA` and `b = NA`.

```
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)  
}
```

Use this function and `mapply()` to calculate the estimated mass for each dinosaur. You'll need to pass the data to `mapply()` as single vectors or columns, not the whole data frame.

```
mapply(get_mass_from_length_by_name, length = dino_lengths$lengths, dinosaur_name = dino_lengths$species)
```

```
##      [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
```

Using dplyr, add a new masses column to the data frame (using rowwise(), mutate() and your function) and print the result to the console.

```
library(dplyr)
dino_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
```

Using ggplot2, make a histogram of dinosaur masses with one subplot for each species (remember facet_wrap()).

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

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

