loops-prep

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#Basic For Loops in R

```
for (item in list_of_items){
    do_something(item)
}
volumes <- c(1.6,3,8)
for (volume in volumes){
    mass <- 2.65 * volume^0.9
    print(mass)
}

volume <- volumes[1]
mass <- 2.65 * volume^0.9
print(mass)
volume <- volumes[2]
mass <- 2.65 * volume^0.9
print(mass)
volume <- volumes[3]
mass <- 2.65 * volume^0.9
print(mass)
volume <- volumes[3]
mass <- 2.65 * volume^0.9
print(mass)</pre>
```

#Looping By Index

```
volumes <- c(1.6,3,8) # i is short for index
masses <- vector(length = length(volumes), mode = "numeric")
for (i in 1:length(volumes)){
  mass <- 2.65 * volumes[i]^0.9
  masses[i] <- mass
}</pre>
```

#Looping Over Multiple Objects

```
volumes <- c(1.6,3,8)
b0 <- c(2.65, 1.28, 3.29)
b1 <- c(0.9, 1.1, 1.2)
masses <- vector(length = length(volumes), mode = "numeric")
for (i in 1:length(volumes)){
  mass <- 2.65 * volumes[i]^b1[i]
  masses[i] <- mass
}</pre>
```

Looping Using Functions

```
est_mass <- function (volume){</pre>
  if (volume > 5) {
    mass \leftarrow 2.65 * volume ^ 0.9
    }else {
    mass <- NA
    }
    return(mass)
}
volumes = c(1.6, 3, 8)
masses <- vector(mode = "numeric", length = length(volumes))</pre>
for (i in 1:length(volumes)) {
  mass <- est_mass(volumes[i])</pre>
  masses[i] <- mass</pre>
}
masses_apply <- sapply(volumes, est_mass)</pre>
#Looping Over Files
download.file("https://www.datacarpentry.org/semester-biology/data/locations.zip", "locations.zip")
unzip("locations.zip")
data_files<- list.files(pattern = "locations-")</pre>
results<- vector(mode = "integer", length(data_files))</pre>
for(i in 1:length(data_files)){
  data <- read.csv(data_files[i])</pre>
  count <- nrow(data)</pre>
  results[i] <- count</pre>
}
```