

# Homework 5

Nathaniel Haulk

10/28/2021

```
library("devtools")

#devtools::install_github("daijiang/neonDivData")

library("neonDivData")

library("tidyverse")
```

## Question 1:

```
#sort(unique(data_plant$taxon_name))

## Pulls the genus for each plant
data_plant$genus = sub(" .*", "", data_plant$taxon_name)

## Selects a random number 100 times between 1 and the length of data_plant
i = sample(1:nrow(data_plant), 100, replace=FALSE)

## Records the genus at the random
genus_samp = data_plant$genus[i]

## Prints out 100 genus
print(genus_samp)
```

```
##      [1] "Ambrosia"      "Aristida"      "Panicum"
##      [4] "Rhus"          "Vulpia"        "Bromus"
##      [7] "Viola"         "Solanum"       "Carex"
##     [10] "Juncus"        "Chamaesyce"    "Lindera"
##     [13] "Vulpia"        "Uvularia"      "Ledum"
##     [16] "Pinus"         "Acer"          "Potentilla"
##     [19] "Tragopogon"    "Gaultheria"    "Rubus"
##     [22] "Salix"         "Ambrosia"      "Prunus"
##     [25] "Anemone"       "Sabal"         "Setaria"
##     [28] "Lycopodiella" "Leersia"       "Pinus"
##     [31] "Malacothrix"   "Stylosanthes"  "Carex"
##     [34] "Convallaria"   "Pinus"         "Eriophorum"
##     [37] "Bouteloua"     "Arnica"        "Empetrum"
##     [40] "Rubus"         "Galium"        "Toxicodendron"
```

```
## [43] "Dichantherium"      "Hesperostipa"      "Hibiscus"
## [46] "Dryas"              "Descurainia"       "Schizachyrium"
## [49] "Vaccinium"          "Polygala"          "Dasyochloa"
## [52] "Onoclea"            "Fraseria"          "Vernonia"
## [55] "Cephalanthus"      "Clarkia"           "Hesperostipa"
## [58] "Quercus"            "Lonicera"          "Gymnanthes"
## [61] "Xerophyllum"        "Lyonia"            "Trifolium"
## [64] "Euphorbia"          "Elymus"            "Solidago"
## [67] "Acer"               "Zizia"             "Krascheninnikovia"
## [70] "Cylindropuntia"     "Rumex"             "Liatris"
## [73] "Hibiscus"           "Helianthus"        "Liatris"
## [76] "Rhus"               "Osmunda"           "Carex"
## [79] "Anemone"            "Apodanthera"       "Polystichum"
## [82] "Dasyochloa"         "Fraxinus"          "Echinacea"
## [85] "Bromus"             "Juglans"           "Acer"
## [88] "Vaccinium"          "Machaeranthera"    "Prosopis"
## [91] "Anthoxanthum"       "Blechnum"          "Phlox"
## [94] "Andropogon"         "Cryptantha"        "Onoclea"
## [97] "Aralia"             "Artemisia"         "Carya"
## [100] "Sanicula"
```

## Question 2:

```
data_plant$taxon_name2 = sub("(\\w\\s\\w+).*", "\\1", data_plant$taxon_name)

## Selects a random number 100 times between 1 and the length of data_plant
i = sample(1:nrow(data_plant), 100, replace=FALSE)

## Records the genus and species name at the random
taxon2_samp = data_plant$taxon_name2[i]

## Prints out data
print(taxon2_samp)
```

```
## [1] "Arctagrostis latifolia"      "Polygonatum biflorum"
## [3] "Chamaesyce arizonica"       "Aristida adscensionis"
## [5] "Lotus unifoliolatus"        "Desmanthus illinoensis"
## [7] "Clitoria mariana"           "Festuca brachyphylla"
## [9] "Tridens flavus"             "Vaccinium uliginosum"
## [11] "Heuchera richardsonii"      "Grammitis hookeri"
## [13] "Grindelia squarrosa"        "Aira caryophyllea"
## [15] "Rosa arkansana"             "Mahonia nervosa"
## [17] "Vernonia angustifolia"      "Echinocereus reichenbachii"
## [19] "Vaccinium myrsinites"       "Bothriochloa ischaemum"
## [21] "Sibbaldia procumbens"       "Symphoricarpos orbiculatus"
## [23] "Poa pratensis"              "Sorghum bicolor"
## [25] "Panicum dichotomiflorum"    "Polygonum perfoliatum"
## [27] "Boehmeria cylindrica"       "Monotropa uniflora"
## [29] "Acer negundo"               "Acer saccharum"
## [31] "Viola sp"                   "Acalypha sp"
## [33] "Dirca palustris"            "Pedicularis lapponica"
```

## [35]	"Betula alleghaniensis"	"Betula occidentalis"
## [37]	"Equisetum hyemale"	"Metrosideros polymorpha"
## [39]	"Rosa arkansana"	"Pinus palustris"
## [41]	"Tragopogon dubius"	"Vitis rotundifolia"
## [43]	"Populus tremuloides"	"Pteridium aquilinum"
## [45]	"Gymnocarpium sp"	"Elymus elymoides"
## [47]	"Eriogonum effusum"	"Lechea torreyi"
## [49]	"Lepidium densiflorum"	"Echinacea angustifolia"
## [51]	"Gymnanthes lucida"	"Euphorbia davidii"
## [53]	"Aristida purpurea"	"Ceanothus sp"
## [55]	"Trifolium microcephalum"	"Acalypha gracilens"
## [57]	"Rubus chamaemorus"	"Geranium columbinum"
## [59]	"Chamaesyce nutans"	"Achyranthes aspera"
## [61]	"Galactia volubilis"	"Ilex decidua"
## [63]	"Melochia pyramidata"	"Lactuca biennis"
## [65]	"Polygonum virginianum"	"Acer negundo"
## [67]	"Fragaria virginiana"	"Eriophorum russeolum"
## [69]	"Delphinium hansenii"	"Pteridium aquilinum"
## [71]	"Mycelis muralis"	"Athyrium filix"
## [73]	"Amianthium muscitoxicum"	"Carex spp"
## [75]	"Yucca elata"	"Eriophorum russeolum"
## [77]	"Pteridium aquilinum"	"Eriophorum vaginatum"
## [79]	"Lindera benzoin"	"Eriogonum effusum"
## [81]	"Uvularia perfoliata"	"Cornus florida"
## [83]	"Hibiscus laevis"	"Sedum sp"
## [85]	"Rubus trivialis"	"Galax urceolata"
## [87]	"Galium sp"	"Eriophorum vaginatum"
## [89]	"Chamaecrista fasciculata"	"Lactuca tatarica"
## [91]	"Oxalis stricta"	"Veratrum sp"
## [93]	"Equisetum pratense"	"Ulmus sp"
## [95]	"Circaea lutetiana"	"Teucrium canadense"
## [97]	"Celastrus orbiculatus"	"Monarda fistulosa"
## [99]	"Serenoa repens"	"Amaranthus palmeri"

### Question 3

```
## Creates data frame n_1 that counts the number of sample_areas with a size of 1 m2 in data_plant
n_1 = data_plant %>%
  group_by(siteID) %>%
  summarise(richness_1m2 = sum(sample_area_m2 == "1"))

## Creates data frame n_10 that counts the number of sample_areas with a size of 1 and 10 m2 in data_plant
n_10 = data_plant %>%
  group_by(siteID) %>%
  summarise(richness_10m2 = sum(sample_area_m2 == "1" | sample_area_m2 == "10"))

## Creates data frame n_100 that counts the number of sample_areas with a size of 1, 10, and 100 m2 in data_plant
n_100 = data_plant %>%
  group_by(siteID) %>%
  summarise(richness_100m2 = sum(sample_area_m2 == "1" | sample_area_m2 == "10" | sample_area_m2 == "100"))
```

```
## Joins n_1, n_10, n_100
n_all = left_join(n_1, n_10) %>%
  left_join(n_100)
```

```
## Joining, by = "siteID"
## Joining, by = "siteID"
```

```
## prints n_all
print(n_all)
```

```
## # A tibble: 47 x 4
##   siteID richness_1m2 richness_10m2 richness_100m2
##   <chr>         <int>         <int>         <int>
## 1 ABBY           7587          13603          16946
## 2 BARR           4615           6041           6616
## 3 BART           5875          10659          13354
## 4 BLAN           8803          15270          19798
## 5 BONA           4091           5925           6983
## 6 CLBJ           10226          15656          18328
## 7 CPER           23663          37876          44284
## 8 DCFS           12067          16649          19622
## 9 DEJU           8673          13000          14918
## 10 DELA          12012          22521          29326
## # ... with 37 more rows
```

## Question 4

```
## creates n_all_longer that records site richness based off spatial scale.
n_all_longer = n_all %>%
  pivot_longer(!siteID, names_to = "spatial_scale", values_to = "richness")

## Prints dataframe
print(n_all_longer)
```

```
## # A tibble: 141 x 3
##   siteID spatial_scale richness
##   <chr>   <chr>         <int>
## 1 ABBY   richness_1m2      7587
## 2 ABBY   richness_10m2     13603
## 3 ABBY   richness_100m2    16946
## 4 BARR   richness_1m2      4615
## 5 BARR   richness_10m2     6041
## 6 BARR   richness_100m2    6616
## 7 BART   richness_1m2      5875
## 8 BART   richness_10m2     10659
## 9 BART   richness_100m2    13354
## 10 BLAN  richness_1m2      8803
## # ... with 131 more rows
```

## Question 5