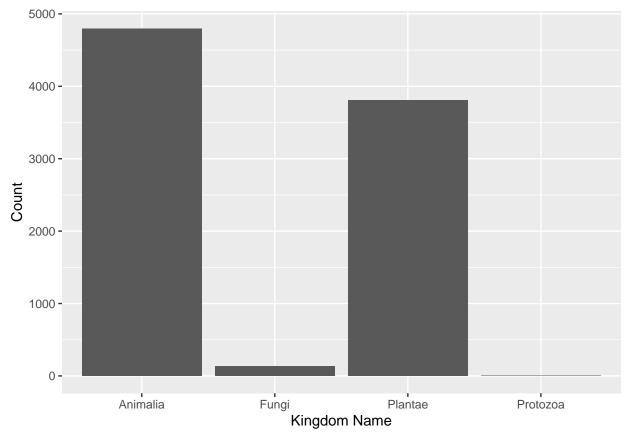
bassinat

Lindsey Weyant

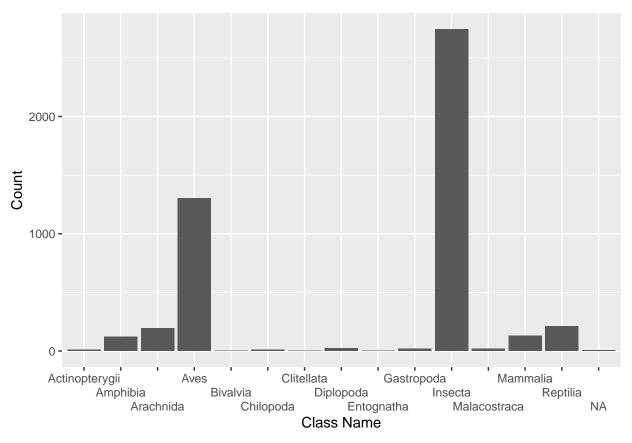
12/2/2021

```
inat%>%
  ggplot(aes(x = taxon_kingdom_name)) +
  geom_bar() +
  labs(x = "Kingdom Name", y = "Count", Title = "Kingdoms of the ECWA Watershed")
```



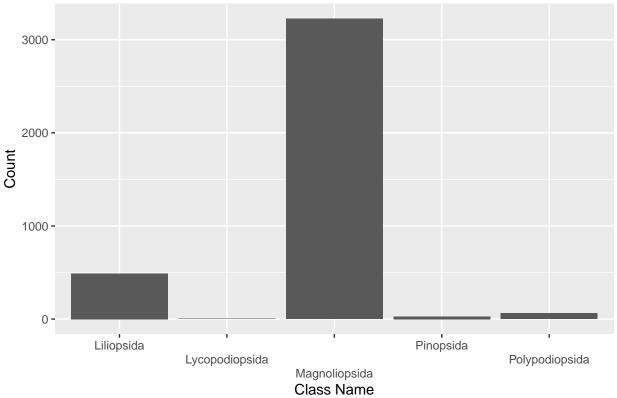
```
inatfungi <- inat %>%
  filter(taxon_kingdom_name == "Fungi")
inatfungi %>%
  count(scientific_name)
```

```
## 2 Amanita flavorubens
## 3 Amanita parcivolvata
## 4 Amanita persicina
## 5 Amanita rubescens
                                   1
## 6 Apioperdon pyriforme
                                   2
## 7 Aureoboletus betula
                                   3
## 8 Auricularia americana
## 9 Baeospora myosura
                                   1
## 10 Cantharellus cinnabarinus
## # ... with 58 more rows
inatprotozoa <- inat %>%
 filter(taxon_kingdom_name == "Protozoa")
inatprotozoa %>%
 count(scientific_name)
## # A tibble: 3 x 2
## scientific name
    <chr>
                        <int>
## 1 Diachea leucopodia
## 2 Fuligo septica
## 3 Lycogala epidendrum
inatanimalia <- inat %>%
 filter(taxon_kingdom_name == "Animalia")
inatanimalia %>%
   count(scientific_name, sort=TRUE)
## # A tibble: 974 x 2
##
     scientific_name
                                  n
##
     <chr>
                              <int>
## 1 Cardinalis cardinalis
                                 75
## 2 Turdus migratorius
                                 58
## 3 Buteo lineatus
## 4 Thryothorus ludovicianus
                                 50
## 5 Papilio polyxenes
                                 45
## 6 Danaus plexippus
                                 43
## 7 Sciurus carolinensis
                                 43
## 8 Storeria dekayi
                                 43
## 9 Xylocopa virginica
                                 43
## 10 Argiope aurantia
                                 39
## # ... with 964 more rows
inatanimalia %>%
 ggplot(aes(x=taxon_class_name)) +
 geom_bar() +
 scale_x_discrete(guide = guide_axis(n.dodge=3)) +
 labs(x= "Class Name", y = "Count", Title = "Animalia Observations by Class")
```



```
inatplantae <- inat %>%
 filter(taxon_kingdom_name == "Plantae")
inatplantae %>%
  count(scientific_name)
## # A tibble: 607 x 2
##
      scientific_name
##
      <chr>
  1 Abelmoschus esculentus
## 2 Acalypha hispida
## 3 Acalypha rhomboidea
## 4 Acer buergerianum
                                 1
## 5 Acer floridanum
                                 9
                                24
  6 Acer negundo
  7 Acer palmatum
                                 2
## 8 Acer rubrum
                                12
   9 Acer saccharinum
                                 2
## 10 Acer saccharum
                                 1
## # ... with 597 more rows
inatplantae %>%
  ggplot(aes(x=taxon_class_name)) +
  geom_bar() +
  scale_x_discrete(guide = guide_axis(n.dodge=3)) +
  labs(x= "Class Name", y = "Count", title = "Plantae Observations by Class")
```

Plantae Observations by Class



```
inatanimalia %>%
 filter(taxon_class_name == "Insecta") %>%
count(scientific_name, sort = TRUE)
## # A tibble: 719 x 2
##
     scientific_name
                             n
##
      <chr>
                         <int>
## 1 Papilio polyxenes
## 2 Danaus plexippus
                            43
## 3 Xylocopa virginica
## 4 Harmonia axyridis
                            39
## 5 Bombus impatiens
                            34
## 6 Papilio glaucus
                            34
## 7 Plathemis lydia
                            33
## 8 Junonia coenia
                            28
## 9 Phyciodes tharos
                            28
## 10 Alaus oculatus
                            26
## # ... with 709 more rows
inatanimalia %>%
  filter(taxon_class_name == "Aves") %>%
count(scientific_name, sort = TRUE)
## # A tibble: 120 x 2
##
      scientific_name
                                   n
```

<int>

75

##

<chr>

1 Cardinalis cardinalis

```
## 2 Turdus migratorius
                                  58
## 3 Buteo lineatus
                                  56
## 4 Thryothorus ludovicianus
                                  50
## 5 Setophaga coronata
                                  39
## 6 Spinus tristis
                                  39
## 7 Ardea herodias
                                  36
## 8 Sayornis phoebe
                                  35
## 9 Dryobates pubescens
                                  32
## 10 Mimus polyglottos
## # ... with 110 more rows
inatplantae %>%
  filter(taxon_class_name == "Magnoliopsida") %>%
count(scientific_name, sort = TRUE)
## # A tibble: 489 x 2
##
      scientific name
##
      <chr>
                              <int>
## 1 Alliaria petiolata
                                111
## 2 Ficaria verna
                                107
## 3 Impatiens capensis
                                 91
## 4 Phytolacca americana
                                 63
## 5 Liriodendron tulipifera
## 6 Hedera helix
                                 61
## 7 Ligustrum sinense
                                 61
                                 54
## 8 Toxicodendron radicans
## 9 Liquidambar styraciflua
                                 52
## 10 Sassafras albidum
                                 51
## # ... with 479 more rows
inatplantae %>%
  filter(taxon_class_name == "Liliopsida") %>%
count(scientific_name, sort = TRUE)
## # A tibble: 95 x 2
##
      scientific_name
                                  n
##
      <chr>
                              <int>
## 1 Microstegium vimineum
                                139
## 2 Arisaema triphyllum
                                 28
## 3 Smilax rotundifolia
                                 23
## 4 Commelina communis
                                 18
## 5 Allium vineale
                                 16
## 6 Dioscorea polystachya
## 7 Chasmanthium latifolium
                                 14
## 8 Tipularia discolor
## 9 Lycoris radiata
                                 10
## 10 Iris pseudacorus
                                  9
## # ... with 85 more rows
inatanimalia %>%
  filter(taxon_class_name == "Amphibia") %>%
  count(scientific_name, sort = TRUE)
## # A tibble: 14 x 2
##
      scientific_name
                                    n
##
      <chr>>
                                <int>
```

```
## 1 Lithobates palustris
                                  23
## 2 Hyla chrysoscelis
                                   20
## 3 Acris crepitans
                                  17
## 4 Hyla cinerea
                                  16
## 5 Lithobates catesbeianus
                                   13
## 6 Lithobates clamitans
                                   9
## 7 Anaxyrus americanus
## 8 Ambystoma opacum
                                   4
## 9 Eurycea cirrigera
                                   3
## 10 Anaxyrus fowleri
                                    2
## 11 Lithobates sphenocephalus
## 12 Notophthalmus viridescens
                                    2
## 13 Pseudacris crucifer
                                    2
## 14 Pseudacris feriarum
                                    1
inatanimalia %>%
 filter(taxon_class_name == "Reptilia") %>%
count(scientific_name, sort = TRUE)
```

```
## # A tibble: 29 x 2
     scientific_name
##
                                     n
##
      <chr>
                                 <int>
## 1 Storeria dekayi
## 2 Anolis carolinensis
                                    25
## 3 Haldea striatula
                                    22
## 4 Pantherophis alleghaniensis
                                    18
## 5 Agkistrodon contortrix
                                    14
## 6 Scincella lateralis
                                    13
## 7 Trachemys scripta
                                    13
## 8 Chelydra serpentina
                                     8
## 9 Thamnophis sirtalis
                                     7
## 10 Carphophis amoenus
## # ... with 19 more rows
```