

# BBS Species Updates Notebook

## Check differences from years of BBS species

to develop taxonomic harmonization workflow

```
# optional  
# rm(ls())  
require(here)
```

Loading required package: here

here() starts at /Users/billspat/Code/SpaCELab/atx/Avian-Interaction-Database

```
require(readr)
```

Loading required package: readr

```
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

```
R_L0 <- here::here('R/v2')
source(file.path(R_L0, 'bbs_specieslist_L0_psb.R'))
```

Loading required package: curl

Using libcurl 8.7.1 with LibreSSL/3.3.6

Attaching package: 'curl'

The following object is masked from 'package:readr':

```
parse_date

print(L0_dir)
```

```
[1] "/Users/billspat/Code/SpaCELab/atx/Avian-Interaction-Database/L0"
```

we've already downloaded the 2023 file and don't have the URL handy for re-download, so read it in directly using the guessed encoding

```
bbs.splist.2023.file <- file.path(L0_dir, "bbs_splist_2022_L0.csv")
encoding.2023 <- readr::guess_encoding(bbs.splist.2023.file)
encoding.2023$encoding[1]
```

```
[1] "ISO-8859-1"
```

not sure why this isn't UTF8 unless this file was downloaded on a windows computer and saved here.

read into a data frame and check it

```
bbs.splist.2023<-read.csv(bbs.splist.2023.file ,fileEncoding=encoding.2023$encoding[1])
print(nrow(bbs.splist.2023))
```

```
[1] 760
```

```
head(bbs.splist.2023)
```

Seq	AOU	English_Common_Name	French_Common_Name
1	6 1770	Black-bellied Whistling-Duck	Dendrocygne à ventre noir
2	7 1780	Fulvous Whistling-Duck	Dendrocygne fauve
3	8 1760	Emperor Goose	Oie empereur
4	9 1690	Snow Goose	Oie des neiges
5	10 1691	(Blue Goose) Snow Goose	Oie des neiges (forme bleue)
6	11 1700	Ross's Goose	Oie de Ross
		Spanish_Common_Name	ORDER Family Genus
1		Dendrocygna autumnalis	Anseriformes Anatidae Dendrocygna
2		Dendrocygna bicolor	Anseriformes Anatidae Dendrocygna
3		Anser canagicus	Anseriformes Anatidae Anser
4		Anser caerulescens	Anseriformes Anatidae Anser
5	Anser caerulescens (blue form)	Anseriformes Anatidae	Anser
6		Anser rossii	Anseriformes Anatidae Anser
		Species	genus_species
1		autumnalis	Dendrocygna autumnalis
2		bicolor	Dendrocygna bicolor
3		canagicus	Anser canagicus
4		caerulescens	Anser caerulescens
5	caerulescens (blue form)	Anser caerulescens (blue form)	Anser caerulescens (blue form)
6		rossii	Anser rossii

Read using re-useable function that also

```
bbs.splist.2023 <- read_bbs_2023(bbs.splist.2023.file)
head(bbs.splist.2023)
```

Seq	AOU	English_Common_Name	French_Common_Name	
1	6 1770	Black-bellied Whistling-Duck	Dendrocygne à ventre noir	
2	7 1780	Fulvous Whistling-Duck	Dendrocygne fauve	
3	8 1760	Emperor Goose	Oie empereur	
4	9 1690	Snow Goose	Oie des neiges	
5	10 1691	(Blue Goose) Snow Goose	Oie des neiges (forme bleue)	
6	11 1700	Ross's Goose	Oie de Ross	
	Order	Family	Genus	Species
1	Anseriformes	Anatidae	Dendrocygna	autumnalis
2	Anseriformes	Anatidae	Dendrocygna	bicolor
3	Anseriformes	Anatidae	Anser	canagicus
4	Anseriformes	Anatidae	Anser	caerulescens

```

5 Anseriformes Anatidae      Anser caerulescens (blue form)
6 Anseriformes Anatidae      Anser                  rossii
                           genus_species
1      Dendrocygna autumnalis
2      Dendrocygna bicolor
3      Anser canagicus
4      Anser caerulescens
5 Anser caerulescens (blue form)
6      Anser rossii

```

download 2024 file (with 2023 data) if it's not downloaded already, read and alter columns for comparison

```

bbs.splist.2024.file <- file.path(L0_dir,"BBS2024_SpeciesListL0.csv")
# this only downloads and modifies if we don't have one already
bbs.splist.2024 <- download_current_bbs_csv(bbs_csv_file_path = bbs.splist.2024.file)

```

Warning in download\_current\_bbs\_csv(bbs\_csv\_file\_path = bbs.splist.2024.file):  
 BBS file exists  
 /Users/billspat/Code/SpaCELab/atx/Avian-Interaction-Database/L0/BBS2024\_SpeciesListL0.csv  
 not overwriting. set overwrite = TRUE to write new file

```
head(bbs.splist.2024)
```

	Seq	AOU	English_Common_Name	French_Common_Name
1	6	1770	Black-bellied Whistling-Duck	Dendrocygne à ventre noir
2	7	1780	Fulvous Whistling-Duck	Dendrocygne fauve
3	8	1760	Emperor Goose	Oie empereur
4	9	1690	Snow Goose	Oie des neiges
5	10	1691	(Blue Goose) Snow Goose	Oie des neiges (forme bleue)
6	11	1700	Ross's Goose	Oie de Ross
	Order	Family	Genus	Species
1	Anseriformes	Anatidae	Dendrocygna	autumnalis
2	Anseriformes	Anatidae	Dendrocygna	bicolor
3	Anseriformes	Anatidae	Anser	canagicus
4	Anseriformes	Anatidae	Anser	caerulescens
5	Anseriformes	Anatidae	Anser caerulescens	(blue form)
6	Anseriformes	Anatidae	Anser	rossii
			genus_species	
1			Dendrocygna autumnalis	
2			Dendrocygna bicolor	

```
3           Anser canagicus
4           Anser caerulescens
5 Anser caerulescens (blue form)
6           Anser rossii
```

### Compare sp list to find changed species

check names

```
print('2023')
```

```
[1] "2023"
```

```
print(names(bbs.splist.2023))
```

```
[1] "Seq"           "AOU"           "English_Common_Name"
[4] "French_Common_Name" "Order"        "Family"
[7] "Genus"          "Species"       "genus_species"
```

```
print('2024')
```

```
[1] "2024"
```

```
print(names(bbs.splist.2024))
```

```
[1] "Seq"           "AOU"           "English_Common_Name"
[4] "French_Common_Name" "Order"        "Family"
[7] "Genus"          "Species"       "genus_species"
```

Compare

```
diffs <- difference_bbs_species_list_2023_24(bbs.splist.2023,bbs.splist.2024 )
diffs
```

```

# A tibble: 8 x 7
  AOU differing_values_diffEng~1 differing_values_dif~2 differing_values_dif~3
  <int> <chr>                  <chr>                  <chr>
1 3340 "Northern Goshawk -> Amer~ "Autour des palombes ~ ""
2 4123 "(unid. Red/Yellow Shafte~ ""
3 4641 "Pacific-slope Flycatcher~ "Moucherolle côtier -- ""
4 4640 "Cordilleran Flycatcher -- "Moucherolle des ravi~ ""
5 4642 "unid. Cordilleran / Paci~ "unid Moucherolle côt~ ""
6 7421 "Swinhoe\u0092s White-eye~ ""
7 5212 "Unid. Cassia Crossbill /~ ""
8 6556 "(unid. Myrtle/Audubon's)~ ""

# i abbreviated names: 1: differing_values_diffEnglish_Common_Name,
#   2: differing_values_diffFrench_Common_Name, 3: differing_values_diffOrder
# i 3 more variables: differing_values_diffFamily <chr>,
#   differing_values_diffGenus <chr>, differing_values_diffSpecies <chr>

```

### **Species differences & anomalies notes:**

The current comparison lists these species as changing since last time: differing\_values\_diffEnglish\_Common\_N

- Northern Goshawk -> American Goshawk
- (unid. Red/Yellow Shafted) Northern Flicker -> (unid. Red / Yellow Shafted) Northern Flicker
- Pacific-slope Flycatcher -> (Pacific-slope Flycatcher) Western Flycatcher
- Cordilleran Flycatcher -> (Cordilleran Flycatcher) Western Flycatcher
- unid. Cordilleran / Pacific-slope Flycatcher -> (unid. Cordilleran / Pac-slope) Western Flycatcher
- Swinhoe-ís White-eye -> Swinhoe,Ã³s White-eye
- Unid. Cassia Crossbill / Red Crossbill -> unid. Cassia Crossbill / Red Crossbill
- (unid. Myrtle/Audubon's) Yellow-rumped Warbler -> (unid. Myrtle / Audubon's) Yellow-rumped Warbler

The species re-assignment checking will occur in the next script: bbs\_specieslist\_L1.R

Then, if combining with bbs\_obs data: AvianInteractionData\_L1.R