

Filter Data

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Preliminaries

```
> library(fishWiDNR) # for setDBClasses(), expandCounts()
> library(dplyr)      # for select(), filter()
> library(FSA)        # for Summarize()

> setwd("C:/aaaWork/Web/fishR/Courses/WiDNR_Statewide_2015/Day1_IntroR_FMDData")
> d <- read.csv("FMDB_Sawyer.csv", stringsAsFactors=FALSE)
> d <- setDBClasses(d, type="RDNR")
> d <- expandCounts(d, ~Number.of.Fish, ~Length.or.Lower.Length.IN+Length.Upper.IN, new.name="Len")
> names(d)
```

[1] "County"	"Waterbody.Name"	"WBIC"
[4] "Survey.Year"	"Station.Name"	"Swims.Station.Id"
[7] "Site.Seq.No"	"Srvy.Seq.No"	"Survey.Begin.Date"
[10] "Survey.End.Date"	"Survey.Status"	"Data.Entry.Name"
[13] "Entry.Date"	"Visit.Fish.Seq.No"	"Visit.Type"
[16] "Gear"	"Sample.Date"	"Substation.Name"
[19] "Target.Species"	"Fish.Data.Seq.No"	"Net.Number"
[22] "Species.Code"	"Species"	"Length.or.Lower.Length.IN"
[25] "Length.Upper.IN"	"Length.or.Lower.Length.MM"	"Length.Upper.MM"
[28] "Weight.Pounds"	"Weight.Grams"	"Gender"
[31] "Disease"	"Injury.Type"	"Age..observed.annuli."
[34] "Edge.Counted.Desc"	"Age.Structure"	"Mark.Given"
[37] "Mark.Found"	"Second.Mark.Found"	"Tag.Number.Given"
[40] "Second.Tag.Number.Given"	"Tag.Number.Found"	"Second.Tag.Number.Found"
[43] "YOY"	"Entry.Date.1"	"Last.Update.Date"
[46] "Data.Ent.Name"	"Last.Update.Name"	"Invalid.Species"
[49] "Non.Standard.Bin"	"Length.Unit.Error"	"Length.Outside.Range"
[52] "Count.Outside.Range"	"Status.Code"	"Len"
[55] "lennote"		

Selecting Variables – select()

```
> d1 <- select(d, Waterbody.Name, Gear, Survey.Year, Species, Len, Weight.Pounds, Gender, Mark.Given)
> head(d1)
```

	Waterbody.Name	Gear	Survey.Year	Species	Len	Weight.Pounds	Gender	Mark.Given
11	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	5.0		NA	
12	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	9.8		NA	
13	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	10.2		NA	
14	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	10.3		NA	
15	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	12.0		NA	
16	ASHEGON LAKE	FYKE NET	2010	BLACK CRAPPIE	12.2		NA	

```
> tail(d1)
```

	Waterbody.Name	Gear	Survey.Year	Species	Len	Weight.Pounds	Gender	Mark.Given
52724	TEAL LAKE	FYKE NET	2010	WALLEYE	15.0	NA	M	
52725	TEAL LAKE	FYKE NET	2010	WALLEYE	15.3	NA	M	
52726	TEAL LAKE	FYKE NET	2010	WALLEYE	17.1	NA	M	
52727	TEAL LAKE	FYKE NET	2010	WALLEYE	17.0	NA	M	
52728	TEAL LAKE	FYKE NET	2010	WALLEYE	17.8	NA	M	
52729	TEAL LAKE	FYKE NET	2010	WALLEYE	17.8	NA	M	

```
> tmp <- select(d, County:Swims.Station.Id)
> head(tmp)
```

	County	Waterbody.Name	WBIC	Survey.Year		Station.Name	Swims.Station.Id
11	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674
12	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674
13	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674
14	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674
15	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674
16	SAWYER	ASHEGON LAKE	2448800	2010	ASHEGON LAKE_GENERAL	LAKE STATION	10005674

```
> tmp <- select(d, -(Station.Name:Status.Code))
> head(tmp)
```

	County	Waterbody.Name	WBIC	Survey.Year	Len	lennote
11	SAWYER	ASHEGON LAKE	2448800	2010	5.0	Observed length
12	SAWYER	ASHEGON LAKE	2448800	2010	9.8	Observed length
13	SAWYER	ASHEGON LAKE	2448800	2010	10.2	Observed length
14	SAWYER	ASHEGON LAKE	2448800	2010	10.3	Observed length
15	SAWYER	ASHEGON LAKE	2448800	2010	12.0	Observed length
16	SAWYER	ASHEGON LAKE	2448800	2010	12.2	Observed length

```
> tmp <- select(d, starts_with("Length")) # there is also an ends_with
> names(tmp)
```

```
[1] "Length.or.Lower.Length.IN" "Length.Upper.IN" "Length.or.Lower.Length.MM"
[4] "Length.Upper.MM" "Length.Unit.Error" "Length.Outside.Range"
```

```
> tmp <- select(d, Srvy.Seq.No, Species, Len, contains("Mark"))
> head(tmp)
```

	Srvy.Seq.No	Species	Len	Mark.Given	Mark.Found	Second.Mark.Found
11	56064296	BLACK CRAPPIE	5.0			
12	56064296	BLACK CRAPPIE	9.8			
13	56064296	BLACK CRAPPIE	10.2			
14	56064296	BLACK CRAPPIE	10.3			
15	56064296	BLACK CRAPPIE	12.0			
16	56064296	BLACK CRAPPIE	12.2			

Selecting Individuals – filter()

```
> levels(d1$Gear)
```

```
[1] "BACKPACK SHOCKER" "BOOM SHOCKER"      "BOTTOM GILL NET"   "FYKE NET"
[5] "MINI BOOM SHOCKER" "STREAM SHOCKER"
```

```
> xtabs(~Gear,data=d1)
```

Gear	BACKPACK SHOCKER	BOOM SHOCKER	BOTTOM GILL NET	FYKE NET	MINI BOOM SHOCKER
	1049	18944	182	25177	386
STREAM SHOCKER	7850				

```
> xtabs(~Waterbody.Name+Gear,data=d1)      # only partial results shown
```

Waterbody.Name	Gear			
	BACKPACK SHOCKER	BOOM SHOCKER	BOTTOM GILL NET	FYKE NET
ASHEGON LAKE	0	0	0	98
BADGER CREEK	105	0	0	0
BARBER LAKE	0	661	0	2179
BARKER LAKE	0	0	19	0
BILLY BOY FLOWAGE	0	0	0	104
BLACK DAN LAKE	0	554	0	953
BLAISDELL LAKE	0	0	7	0
BLUEBERRY LAKE	0	61	0	0
BRUNET RIVER	0	0	0	0
CHIPPANAZIE CREEK	69	0	0	0
CHIPPEWA RIVER	0	0	140	0
CONNORS LAKE	0	735	0	2126
DURPHEE LAKE	0	693	0	386
EAST FORK CHIPPEWA RIVER	0	0	0	0
EDDY CREEK	0	0	0	0

```
> tmp <- filter(d1,Waterbody.Name=="BARBER LAKE")
```

```
> xtabs(~Waterbody.Name,tmp)      # only partial results shown
```

Waterbody.Name	ASHEGON LAKE	BADGER CREEK	BARBER LAKE	BARKER LAKE
	0	0	2840	0
BILLY BOY FLOWAGE		BLACK DAN LAKE	BLAISDELL LAKE	BLUEBERRY LAKE
	0	0	0	0
BRUNET RIVER		CHIPPANAZIE CREEK	CHIPPEWA RIVER	CONNORS LAKE
	0	0	0	0
DURPHEE LAKE	EAST FORK CHIPPEWA RIVER		EDDY CREEK	FLAMBEAU RIVER
0	0		0	0

```
> tmp <- droplevels(tmp)
```

```
> xtabs(~Waterbody.Name,tmp)
```

Waterbody.Name
BARBER LAKE
2840

```
> tmp <- filter(d1, Waterbody.Name %in% c("BARBER LAKE", "LAKE CHETAC"))
> tmp <- droplevels(tmp)
> xtabs(~Waterbody.Name, tmp)
```

```
Waterbody.Name
BARBER LAKE LAKE CHETAC
      2840      6946
```

```
> LCblg <- filter(d1, Waterbody.Name=="LAKE CHETAC", Species=="BLUEGILL")
> xtabs(~Gear, LCblg)
```

```
Gear
BACKPACK SHOCKER      BOOM SHOCKER  BOTTOM GILL NET      FYKE NET MINI BOOM SHOCKER
      0              398              0              191              0
STREAM SHOCKER
      0
```

```
> LCblg <- filter(LCblg, Gear=="BOOM SHOCKER")
> Summarize(~Len, data=LCblg)
```

```
      n      mean      sd      min      Q1      median      Q3      max percZero
398.000  5.984    1.163    3.000    5.000    6.200    6.900    8.900    0.000
```

```
> LCblgPREF <- filter(LCblg, Len>=7)
> Summarize(~Len, data=LCblgPREF)
```

```
      n      mean      sd      min      Q1      median      Q3      max percZero
95.0000  7.3632    0.3461    7.0000    7.1000    7.3000    7.5000    8.9000    0.0000
```

```
> sturgWts <- filter(d1, Species=="LAKE STURGEON", !is.na(Weight.Pounds))
> head(sturgWts)
```

```
Waterbody.Name      Gear Survey.Year      Species  Len Weight.Pounds Gender Mark.Given
1  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 58.0      43.9      U      PIT
2  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 61.5      70.5      U      PIT
3  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 59.7      55.6      U      PIT
4  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 62.5      66.5      U
5  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 55.7      38.8      U
6  BARKER LAKE BOTTOM GILL NET      2010 LAKE STURGEON 56.4      45.7      U      PIT
```

Application Assignment

Create a script that performs the following tasks:

1. Load and prepare (set classes, expand counts, examine structure) your FM data in R (**HINT:** use all or some of your script from the first application assignment). Call this the *original data.frame*.
2. Create a data.frame that removes all variables related to the database (e.g., when datum was entered, who entered it, error flags, etc.).
3. Examine the sample size per water body and gear combination in the original data.frame.

4. Isolate (from the original data.frame) a water body of your choice and show the number of each species captured (in all gears).
5. Isolate (from the original data.frame) three water bodies of your choice and make one table that shows the number of each species captured in each water body (regardless of gear).
6. Isolate (from the original data.frame) one species of fish from one gear used in one waterbody.
 - Construct a table of frequency of each sex.
 - Summarize the length variable.
7. (*Time Permitting*) Suppose the waterbody and species you chose above has a minimum length limit (make up the minimum length). Isolate those fish that would be legal. Show that your filtering was successful.
8. (*Time Permitting*) Repeat the previous question but for a protected slot.
9. (*Time Permitting*) Repeat the previous question but for a harvest slot.
10. (*Time Permitting*) List all water bodies and species for which a weight in pounds was recorded (begin with the original data.frame).

Save your script!