

# Filter Data

*Derek H. Ogle, Northland College*

*4-Mar-2015*

## Preliminaries

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

> setwd("C:/aaaWork/Web/fishR/Courses/WiDNR_Statewide_2015/Day1_IntroR_FMDData")
> d <- read.csv("FMDB_Sawyer_MultiYr_APEX.csv", stringsAsFactors=FALSE, na.strings=c("-", "NA", ""))
> 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"	"Survey.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)
> headtail(d1)
```

	Waterbody.Name	Gear	Survey.Year	Species	Len	Weight.Pounds	Gender
1	GRINDSTONE LAKE	FYKE NET	2003	CREEK CHUB	NA	NA	<NA>
2	GRINDSTONE LAKE	FYKE NET	2003	WALLEYE	NA	NA	<NA>
3	GRINDSTONE LAKE	FYKE NET	2003	NORTHERN PIKE	NA	NA	<NA>
448038	LAKE CHIPPEWA BOOM	SHOCKER	2002	WALLEYE	13.0	NA	<NA>
448039	ISLAND LAKE BOOM	SHOCKER	2007	LARGEMOUTH BASS	7.3	NA	<NA>
448043	BLAISDELL LAKE BOOM	SHOCKER	2006	MUSKELLUNGE	13.7	NA	<NA>

  

	Mark.Given
1	<NA>
2	<NA>
3	<NA>
448038	<NA>
448039	<NA>
448043	<NA>

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

	County	Waterbody.Name	WBIC	Survey.Year	Station.Name
1	SAWYER	GRINDSTONE LAKE	2391200	2003	GRINDSTONE LAKE_GENERAL LAKE STATION
2	SAWYER	GRINDSTONE LAKE	2391200	2003	GRINDSTONE LAKE_GENERAL LAKE STATION
3	SAWYER	GRINDSTONE LAKE	2391200	2003	GRINDSTONE LAKE_GENERAL LAKE STATION
448038	SAWYER	LAKE CHIPPEWA	2399700	2002	LAKE CHIPPEWA_GENERAL LAKE STATION
448039	SAWYER	ISLAND LAKE	2381800	2007	ISLAND LAKE_GENERAL LAKE STATION
448043	SAWYER	BLAISDELL LAKE	2402200	2006	BLAISDELL LAKE_GENERAL LAKE STATION

  

	Swims.Station.Id
1	10005586
2	10005586
3	10005586
448038	10005605
448039	10005570
448043	10005611

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

	County	Waterbody.Name	WBIC	Survey.Year	Len	lennote
1	SAWYER	GRINDSTONE LAKE	2391200	2003	NA	Observed length
2	SAWYER	GRINDSTONE LAKE	2391200	2003	NA	Observed length
3	SAWYER	GRINDSTONE LAKE	2391200	2003	NA	Observed length
448038	SAWYER	LAKE CHIPPEWA	2399700	2002	13.0	Expanded length
448039	SAWYER	ISLAND LAKE	2381800	2007	7.3	Expanded length
448043	SAWYER	BLAISDELL LAKE	2402200	2006	13.7	Expanded 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,Survey.Seq.No,Species,Len,contains("Mark"))
> headtail(tmp)
```

	Survey.Seq.No	Species	Len	Mark.Given	Mark.Found	Second.Mark.Found
1	51723	CREEK CHUB	NA	<NA>	<NA>	<NA>
2	51726	WALLEYE	NA	<NA>	<NA>	<NA>
3	51726	NORTHERN PIKE	NA	<NA>	<NA>	<NA>
448038	51356	WALLEYE	13.0	<NA>	<NA>	<NA>
448039	97739	LARGEMOUTH BASS	7.3	<NA>	<NA>	<NA>
448043	94228	MUSKELLUNGE	13.7	<NA>	<NA>	<NA>

## Selecting Individuals – filter()

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

[1] "BACKPACK SHOCKER"	"BOOM SHOCKER"
[3] "BOTTOM GILL NET"	"DIP NET"
[5] "FLOATING GILL NET"	"FYKE NET"
[7] "HOOK AND LINE"	"LONG LINE SHOCKER"
[9] "MINI BOOM SHOCKER"	"MINI FYKE NET"
[11] "MINI FYKE NET WITH TURTLE EXCLUSION"	"MINI FYKE NET WITHOUT TURTLE EXCLUSION"
[13] "SEINE"	"STREAM SHOCKER"

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

Gear

BACKPACK SHOCKER	BOOM SHOCKER
9467	131432
BOTTOM GILL NET	DIP NET
342	189
FLOATING GILL NET	FYKE NET
2883	193217
HOOK AND LINE	LONG LINE SHOCKER
1688	72
MINI BOOM SHOCKER	MINI FYKE NET
4479	15525
MINI FYKE NET WITH TURTLE EXCLUSION	MINI FYKE NET WITHOUT TURTLE EXCLUSION
13873	24856
SEINE	STREAM SHOCKER
2458	47565

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

Waterbody.Name	BACKPACK SHOCKER	BOOM SHOCKER	BOTTOM GILL NET	DIP NET
ALDER CREEK	182	0	0	0
ASHEGON LAKE	0	58	0	0
BADGER CREEK	105	0	0	0
BARBER CREEK	90	0	0	0
BARBER LAKE	0	979	0	0
BARKER LAKE	0	381	25	0
BEAVER CREEK	0	0	0	0
BENSON CREEK	74	0	0	0
BILLY BOY FLOWAGE	0	92	0	0
BLACK DAN LAKE	0	1732	0	0
BLACK LAKE	0	213	0	0

BLAISDELL LAKE	0	404	41	0
BLUEBERRY CREEK	52	0	0	0
BLUEBERRY LAKE	0	979	0	0
BRUNET RIVER	133	0	0	0

```
> tmp <- filter(d1,Waterbody.Name=="BARBER LAKE")
> xtabs(~Waterbody.Name,data=tmp) # only partial results shown
```

Waterbody.Name				
ALDER CREEK	ASHEGON LAKE	BADGER CREEK	BARBER CREEK	BARBER LAKE
0	0	0	0	3727
BARKER LAKE	BEAVER CREEK	BENSON CREEK	BILLY BOY FLOWAGE	BLACK DAN LAKE
0	0	0	0	0
BLACK LAKE	BLAISDELL LAKE	BLUEBERRY CREEK	BLUEBERRY LAKE	BRUNET RIVER
0	0	0	0	0
CALLAHAN LAKE				
0				

```
> tmp <- droplevels(tmp)
> xtabs(~Waterbody.Name,data=tmp)
```

Waterbody.Name	
BARBER LAKE	
3727	

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

Waterbody.Name	
BARBER LAKE	LAKE CHETAC
3727	14827

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

Gear		
BOOM SHOCKER	FYKE NET	MINI FYKE NET
1005	191	327

```
> LCblg <- filter(LCblg,Gear=="BOOM SHOCKER")
> Summarize(~Len,data=LCblg,digits=2)
```

n	mean	sd	min	Q1	median	Q3	max	percZero
1005.00	6.16	1.08	2.60	5.50	6.20	7.00	9.80	0.00

```
> LCblgPREF <- filter(LCblg,Len>=7)
> Summarize(~Len,data=LCblgPREF,digits=2)
```

n	mean	sd	min	Q1	median	Q3	max	percZero
259.00	7.45	0.43	7.00	7.20	7.30	7.65	9.80	0.00

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

	Waterbody.Name	Gear	Survey.Year	Species	Len	Weight.Pounds	Gender	Mark.Given
1	CHIPPEWA RIVER	DIP NET	2006 LAKE	STURGEON	54.3	32.0	M	PIT
2	CHIPPEWA RIVER	DIP NET	2006 LAKE	STURGEON	59.7	47.0	F	PIT
3	CHIPPEWA RIVER	DIP NET	2006 LAKE	STURGEON	54.8	37.0	M	PIT
415	BARKER LAKE BOTTOM	GILL NET	2012 LAKE	STURGEON	58.3	34.2	<NA>	PIT
416	BARKER LAKE BOTTOM	GILL NET	2012 LAKE	STURGEON	60.9	50.6	<NA>	PIT
417	BARKER LAKE BOTTOM	GILL NET	2012 LAKE	STURGEON	60.9	50.6	<NA>	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!**