# Analysis of a GRTS Survey Design for a Finite Resource

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#### 1 Preliminaries

This document presents analysis of a GRTS survey design for a finite resource. The finite resource used in the analysis is small lakes in Florida. The analysis will include calculation of three types of population estimates: (1) estimation of proportion and size (number of lakes) for site evaluation status categorical variables; (2) estimation of proportion and size for lake condition categorical variables; and (3) estimation of the cumulative distribution function (CDF) and percentiles for quantitative variables. Testing for difference between CDFs from subpopulations also will be presented.

The initial step is to use the library function to load the spsurvey package. After the package is loaded, a message is printed to the R console indicating that the spsurvey package was loaded successfully.

Load the spsurvey package

```
> # Load the spsurvey package
> library(spsurvey)
>
```

Version 2.5 of the spsurvey package was loaded successfully.

## 2 Read the survey design and analytical variables data file

The original Florida small lakes data file contains more than 3,800 records and 29 basins. To produce a more manageable number of records, only six basins were retained in the data that will be analyzed, which produced a file containing 930 records.

The next step is to read the data file, which includes both survey design variables and analytical variables. The read delim function is used to read the tab-delimited file and assign it to a data frame named FL\_lakes. The nrow function is used to determine the number of rows in the FL\_lakes data frame, and the resulting value is assigned to an object named nr. Finally, the initial six lines and the final six lines in the FL\_lakes data frame are printed using the head and tail functions, respectively.

Read the survey design and analytical variables data file

```
> # Read the data file and determine the number of rows in the file
> FL_lakes <- read.delim("FL_lakes.tab")
> nr <- nrow(FL_lakes)
>
```

Display the initial six lines in the data file.

```
> # Display the initial six lines in the data file
> head(FL_lakes)
```

```
siteID xcoord
                          ycoord
                                              basin
                                                              status
                                                                            TNT
                                       wgt
1 FLW03414-0014 8635535 12860896 5.369048 NWFWMD-1
                                                             Sampled
                                                                         Target
2 FLW03414-0046 8636136 12886783 5.369048 NWFWMD-1 Physical Barrier
                                                                         Target
3 FLW03414-0062 8617834 12869126 5.369048 NWFWMD-1
                                                           NonTarget NonTarget
4 FLW03414-0078 8673500 12883071 5.369048 NWFWMD-1 Physical Barrier
                                                                         Target
5 FLW03414-0086 8631884 12816428 5.369048 NWFWMD-1
                                                           NonTarget NonTarget
6 FLW03414-0118 8607699 12856644 5.369048 NWFWMD-1
                                                           NonTarget NonTarget
 pH_cat coliform_cat oxygen turbidity
   (0,6]
                (0,5]
                         9.9
2
    <NA>
                 <NA>
                                     NA
                          NA
```

3	<na></na>	<na></na>	NA	NA
4	<na></na>	<na></na>	NA	NA
5	<na></na>	<na></na>	NA	NA
6	<na></na>	<na></na>	NA	NA

>

Display the final six lines in the data file.

```
> # Display the final six lines in the data file
> tail(FL_lakes)
```

```
xcoord
                                                                                TNT
            siteID
                              ycoord
                                          wgt
                                                 basin
                                                                  status
925 FLW03414-3878 8880656 12694963 4.80791 SWFWMD-4
                                                                     Dry
                                                                             Target
926 FLW03414-3886 8892406 12732977 4.80791 SWFWMD-4
                                                                 Sampled
                                                                             Target
927 FLW03414-3894 8836528 12723056 4.80791 SWFWMD-4
                                                                     Dry
                                                                             Target
928 FLW03414-3918 8923107 12725502 4.80791 SWFWMD-4 Landowner Denial
                                                                             Target
929 FLW03414-3926 8861298 12715824 4.80791 SWFWMD-4
                                                                     Dry
                                                                             Target
930 FLW03414-3950 8888601 12715641 4.80791 SWFWMD-4
                                                               NonTarget NonTarget
    pH_cat coliform_cat oxygen turbidity
925
      <NA>
                    <NA>
                              NA
                                         NA
926
     (6,8]
                  (5,50]
                            1.98
                                       8.2
927
      <NA>
                    <NA>
                              NA
                                         NA
928
      < NA >
                    <NA>
                              NΑ
                                        NA
929
      < NA >
                    <NA>
                              NA
                                         NA
930
      < NA >
                    <NA>
                              NA
                                         NA
```

>

The sample of small lakes in Florida is displayed in Figure 1. The sample sites for each basin are displayed using a unique color. First, the levels function is used to extract the set of basin names, and the result is assigned to object basin. Next, the rainbow function is called to select a set of six colors, and the result is assigned to object cols. The plot function is then used to produce the basic figure, but plotting of sample points is suppressed. The for function is used to loop through the set of six basins and plot color-coded points for each basin using the points function. Finally, the legend function is used to add a legend to the figure, and the title function is used to create a figure title.

```
+ ind <- FL_lakes$basin == basins[i]
+ points(FL_lakes$xcoord[ind], FL_lakes$ycoord[ind], pch=20, cex=0.4,
+ col=cols[i])
+ }
> legend(x="topright", inset=0.05, legend=basins, pch=20, cex=1, col=cols)
> title("Plot of Florida Small Lake Sites Color-Coded by Basin")
```

## 3 Analysis of site status evaluation variables

The first analysis that will be examined is calculation of extent estimates for site status evaluation variables. Extent is measured both by the proportion of the resource in status evaluation categories and by size of the resource in each category. For a finite resource like lakes, size refers to the number of lakes in a category. For calculating extent estimates (and for all of the analyses we will consider), the survey design weights are incorporated into the calculation process. Two site status variables will be examined: (1) status, which classifies lakes into six evaluation categories and (2) TNT, which classifies lakes as either "Target" or "NonTarget". The table and addmargins functions are used to create tables displaying the count for each code (level) of the two status variables.

```
> addmargins(table(FL_lakes$status))
```

A table displaying the number of values for each level of the status variable follows:

Dry	Landowner Denial	${\tt NonTarget}$
223	119	317
Otherwise Unsampleable	Physical Barrier	Sampled
1	99	171
Sum		
930		

> addmargins(table(FL\_lakes\$TNT))

A table displaying the number of values for each level of the TNT variable follows:

NonTarget Target Sum 317 613 930

## Plot of Florida Small Lake Sites Color-Coded by Basin

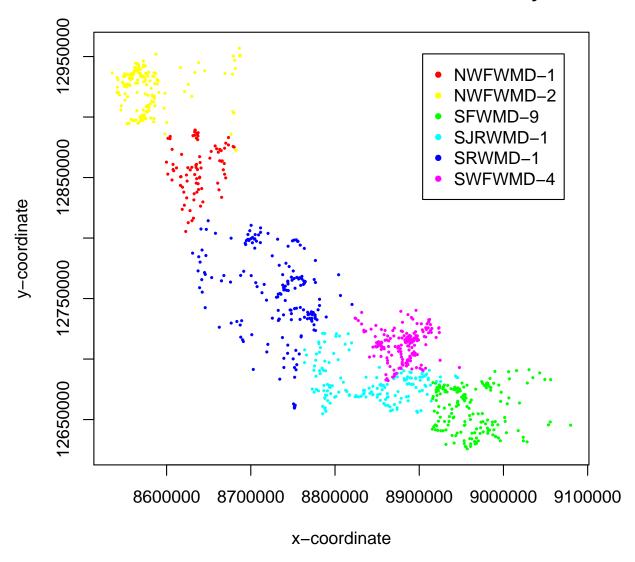


Figure 1: Florida Small Lake Sample Sites.

The cat analysis function in the spsurvey package will be used to calculate extent estimates. Four data frames constitute the primary input to the cat. analysis function. The first column (variable) in the four data frames provides the unique identifier (site ID) for each sample site and is used to connect records among the data frames. The siteID variable in the FL\_lakes data frame is assigned to the siteID variable in the data frames. The four data frames that will be created are named as follows: sites, subpop, design, and data.cat. The sites data frame identifies sites to use in the analysis and contains two variables: (1) siteID - site ID values and (2) Use - a logical vector indicating which sites to use in the analysis. The rep (repeat) function is used to assign the value TRUE to each element of the Use variable. Recall that nr is an object containing the number of rows in the FL\_lakes data frame. The subpop data frame defines populations and, optionally, subpopulations for which estimates are desired. Unlike the sites and design data frames, the subpop data frame can contain an arbitrary number of columns. The first variable in the subpopt data frame identifies site ID values and each subsequent variable identifies a type of population, where the variable name is used to identify type. A type variable identifies each site with a character value. If the number of unique values for a type variable is greater than one, then the set of values represent subpopulations of that type. When a type variable consists of a single unique value, then the type does not contain subpopulations. For this analysis, the subpop data frame contains three variables: (1) siteID - site ID values, (2) CombinedBasins - which will be used to calculate estimates for all of the basins combined, and (3) Basin - which will be used to calculate estimates for each basin individually. The basin variable in the FL\_lakes data frame is assigned to the Basin variable in the subpop data frame. The design data frame consists of survey design variables. For the analysis under consideration, the design data frame contains the following variables: (1) siteID - site ID values; (2) wgt - final, adjusted, survey design weights; (3) xcoord - x-coordinates for location; and (4) ycoord - y-coordinates for location. The wgt, xcoord, and ycoord variables in the design data frame are assigned values using variables with the same names in the FL\_lakes data frame. Like the subpop data frame, the data cat data frame can contain an arbitrary number of columns. The first variable in the data.cat data frame identifies site ID values and each subsequent variable identifies a response variable. The two response variables are Status and Target\_NonTarget, which are assigned the status and TNT variables, respectively, in the FL\_lakes data frame. Missing data (NA) is allowed for the response variables, which are the only variables in the input data frames for which NA values are allowed.

Create the sites data frame.

Create the design data frame.

Create the data.cat data frame.

Use the cat.analysis function to calculate extent estimates for the site status evaluation variables.

```
> # Calculate extent estimates for the site status evaluation variables
> Extent_Estimates <- cat.analysis(sites, subpop, design, data.cat)
>
```

The extent estimates for all basins combined are displayed using the print function. The object produced by cat. analysis is a data frame containing thirteen columns. The first five columns identify the population (Type), subpopulation (Subpopulation), response variable (Indicator), levels of the response variable (Category), and number of values in a category (NResp). A category labeled "Total" is included for each combination of population, subpopulation, and response variable. The next four columns in the data frame provide results for the proportion estimates: the proportion estimate (Estimate.P), standard error of the estimate (StdError.P), lower confidence bound (LCB95Pct.P), and upper confidence bound (UCB95Pct.P). Argument conf for cat. analysis allows control of the confidence bound level. The default value for conf is 95, hence the column names for confidence bounds contain the value 95. Supplying a different value to the conf argument will be reflected in the confidence bound names. Confidence bounds are obtained using the standard error and the Normal distribution multiplier corresponding to the confidence level. The final four columns in the data frame provide results for the size (units) estimates: the units estimate (Estimate.U), standard error of the estimate (StdError.U), lower confidence bound (LCB95Pct.U), and upper confidence bound (UCB95Pct.U).

```
> # Print the extent estimates for all basins combined
> print(Extent_Estimates[c(1:7, 45:47),])
```

	Type	Subpopulation	Indicator	Category	NResp
1	${\tt CombinedBasins}$	All Basins	Status	Dry	223
2	${\tt CombinedBasins}$	All Basins	Status	Landowner Denial	119
3	${\tt CombinedBasins}$	All Basins	Status	${\tt NonTarget}$	317
4	CombinedBasins	All Basins	Status	Otherwise Unsampleable	1

```
5
  CombinedBasins
                     All Basins
                                                         Physical Barrier
                                                                              99
                                            Status
  CombinedBasins
                      All Basins
                                            Status
                                                                   Sampled
                                                                             171
7
  CombinedBasins
                     All Basins
                                                                     Total
                                                                             930
                                            Status
45 CombinedBasins
                     All Basins Target_NonTarget
                                                                NonTarget
                                                                             317
46 CombinedBasins
                     All Basins Target_NonTarget
                                                                    Target
                                                                             613
47 CombinedBasins
                     All Basins Target_NonTarget
                                                                     Total
                                                                             930
     Estimate.P StdError.P LCB95Pct.P
                                        UCB95Pct.P
                                                     Estimate.U StdError.U
1
    23.01117939 0.97789814
                             21.094534
                                        24.9278245 1184.155291
                                                                 50.188531
2
    13.32737468 0.99049216
                             11.386046
                                        15.2687037
                                                     685.826701
                                                                 50.967147
3
    36.91250997 1.15995817
                             34.639034
                                        39.1859862 1899.517763
                                                                 60.260564
4
     0.09422536 0.08497475
                              0.000000
                                         0.2607728
                                                       4.848837
                                                                   4.372792
5
     8.47917794 0.71507723
                              7.077652
                                         9.8807036
                                                     436.338497
                                                                 36.766620
6
    18.17553265 1.03356643
                             16.149780
                                        20.2012856
                                                     935.312910
                                                                 53.169549
7
   100.0000000 0.00000000 100.000000 100.0000000 5146.000000
                                                                  9.275053
45
    36.91250997 1.15995817
                             34.639034
                                        39.1859862 1899.517763
                                                                 60.260564
46
    63.08749003 1.15995817
                             60.814014
                                        65.3609663 3246.482237
                                                                 59.166424
47 100.0000000 0.00000000 100.000000 100.000000 5146.000000
                                                                   9.275053
  LCB95Pct.U UCB95Pct.U
    1085.7876 1282.52300
1
2
               785.72047
     585.9329
3
    1781.4092 2017.62630
4
       0.0000
                13.41935
5
               508.39975
     364.2772
6
     831.1025 1039.52331
7
    5127.8212 5164.17877
45
    1781.4092 2017.62630
46
    3130.5182 3362.44630
47
    5127.8212 5164.17877
>
```

The write.csv function is used to store the extent estimates as a comma-separated value (csv) file. Files in csv format can be read by programs such as Microsoft Excel.

```
> write.csv(Extent_Estimates, file="Extent_Estimates.csv")
```

### 4 Analysis of lake condition variables

The second analysis that will be examined is estimating resource proportion and size for lake condition variables. Two lake condition variables will be examined: (1) pH\_cat, which classifies lakes by categories of pH value and (2) coliform\_cat, which classifies lakes by categories of fecal coliform count. The table and addmargins functions are used to create tables displaying the count for each level of the two lake condition variables.

> addmargins(table(FL\_lakes\$pH\_cat))

A table displaying the number of values for each level of the pH category variable follows:

> addmargins(table(FL\_lakes\$coliform\_cat))

A table displaying the number of values for each level of the fecal coliform category variable follows:

As for extent estimates, the cat.analysis function will be used to calculate condition estimates. The sites data frame for this analysis differs from the one used to calculate extent estimates. The Use logical variables in sites is set equal to the value "Sampled", so that only sampled sites are used in the analysis. The subpop and design data frames created in the prior analysis can be reused for this analysis. The data.cat data frame contains the two lake condition variables: pHCat and ColiformCat. Variables pH\_cat and coliform\_cat in the FL\_lakes data frame are assigned to pHCat and ColiformCat, respectively.

Create the sites data frame.

Create the data.cat data frame.

Use the cat.analysis function to calculate estimates for the lake condition variables.

```
> # Calculate estimates for the categorical variables
> Condition_Estimates <- cat.analysis(sites, subpop, design, data.cat)
>
```

Print the lake condition estimates for all basins combined.

- > # Print the condition estimates for all basins combined
- > print(Condition\_Estimates[c(1:4, 28:32),])

	Туре	Subpopu	ılation	Indicator	Category N	Resp Estima	te.P
1	CombinedBasins	All	Basins	${ t pHCat}$	(0,6]	78 42.91	5056
2	CombinedBasins	All	Basins	${ t pHCat}$	(6,8]	82 50.39	6558
3	CombinedBasins	All	Basins	${ t pHCat}$	(8,14]	11 6.68	8386
4	CombinedBasins	All	Basins	pHCat	Total	171 100.00	0000
28	CombinedBasins	All	Basins Co	liformCat	(0,5]	97 55.98	6933
29	CombinedBasins	All	Basins Co	liformCat	(5,50]	40 24.10	8155
30	CombinedBasins	All	Basins Co	liformCat	(50,500]	31 18.52	1502
31	CombinedBasins	All	Basins Co	liformCat (5	00,5e+03]	2 1.38	3410
32	CombinedBasins	All	Basins Co	liformCat	Total	170 100.00	0000
	StdError.P LCB	95Pct.P	UCB95Pct.	P Estimate.U	StdError.U	LCB95Pct.U	UCB95Pct.U
1	2.8530505 37	.323179	48.50693	2 401.39005	26.886965	348.69257	454.08754
2	3.0180108 44	.481366	56.31175	1 471.36552	28.637754	415.23655	527.49448
3	1.5603961 3	.630066	9.74670	6 62.55734	14.557867	34.02444	91.09023
4	0.0000000 100	.000000	100.00000	0 935.31291	7.447521	920.71604	949.90978
28	2.8761564 50	.349770	61.62409	6 519.19950	26.470305	467.31866	571.08035
29	3.0417644 18	.146407	30.06990	4 223.56900	28.568114	167.57652	279.56147
30	2.4596628 13	.700651	23.34235	2 171.76069	22.738993	127.19309	216.32830
31	0.8268103 0	.000000	3.00392	9 12.82917	7.673900	0.00000	27.86974
32	0.0000000 100	.000000	100.00000	0 927.35836	7.435967	912.78414	941.93259

>

Use the write.csv function to write the condition estimates as a csv file.

> write.csv(Condition\_Estimates, file="Condition\_Estimates.csv")

# 5 Analysis of lake condition variables correcting for population size

The frame is a data structure containing spatial location data in addition to other attributes regarding a resource of interest and is used to create a survey design. A frame often takes the form of a shapefile. The frame can be used to obtain size values (e.g., number of lakes) for the populations and subpopulations examined in an analysis. Examination of the Estimates.U column in the Condition\_Estimates data frame produced by cat.analysis reveals that the estimated Total value for both condition variables and each combination of population value and subpopulation value does not sum to the corresponding frame size value. For example, the Total entry in the Estimate.U column for the pHcat variable, population "CombinedBasins" and subpopulation "All Basins" is 935 (rounded to a whole number). The corresponding

frame size value is 5,146. The popsize (population size) argument to cat.analysis provides a mechanism for forcing the Total category to equal a desired value. First, the c (combine) function is used to create a named vector of frame size values for each basin. Output from the c function is assigned to an object named framesize. The popsize argument is a list, which is a particular type of R object. The popsize list must include an entry for each population type included in the subpop data frame, i.e., CombinedBasins and Basin for this analysis. The sum function applied to framesize is assigned to the CombinedBasins entry in the popsize list. Recall that the basin population type contains subpopulations, i.e., basins. When a population type contains subpopulations, the entry in the popsize list also is a list. The as list function is applied to framesize, and the result is assigned to the Basin entry in the popsize list.

Assign frame size values.

Use the cat.analysis function to calculate estimates for the lake condition variables.

```
> Condition_Estimates_popsize <- cat.analysis(sites, subpop, design, data.cat,
+ popsize=list(CombinedBasins=sum(framesize),</pre>
```

+ Basin=as.list(framesize)))

Print the lake condition estimates for all basins combined.

- > # Print the lake condition estimates for all basins combined
- > print(Condition\_Estimates\_popsize[c(1:4, 28:32),])

	Туре	Subpopu	lation	Ιı	ndicator	Category 1	NResp	Estimat	e.P
1	${\tt CombinedBasins}$	All	Basins		${ t pHCat}$	(0,6]	78	42.915	5056
2	${\tt CombinedBasins}$	All	Basins		${ t pHCat}$	(6,8]	82	50.396	3558
3	${\tt CombinedBasins}$	All	Basins		${ t pHCat}$	(8,14]	11	6.688	386
4	${\tt CombinedBasins}$	All	Basins		${ t pHCat}$	Total	171	100.000	0000
28	${\tt CombinedBasins}$	All	Basins	Col	iformCat	(0,5]	97	55.986	933
29	${\tt CombinedBasins}$	All	Basins	Col	iformCat	(5,50]	40	24.108	3155
30	${\tt CombinedBasins}$	All	Basins	Col	iformCat	(50,500]	31	18.521	.502
31	${\tt CombinedBasins}$	All	Basins	Col	iformCat (5	00,5e+03]	2	1.383	3410
32	${\tt CombinedBasins}$	All	Basins	Col	iformCat	Total	170	100.000	0000
	StdError.P LCBS	95Pct.P	UCB95Pc	t.P	Estimate.U	StdError.	J LCB9	95Pct.U	UCB95Pct.U
1	2.8530505 37	.323179	48.506	932	2208.40876	146.81798	3 192	20.6508	2496.1667
2	3.0180108 44	.481366	56.311	751	2593.40689	155.3068	1 228	39.0111	2897.8027
3	1.5603961 3	.630066	9.746	706	344.18435	80.29798	3 18	36.8032	501.5655
4	NA	NA		NA	5146.00000	N.	A	NA	NA
28	2.8761564 50	. 349770	61.624	1096	2881.08756	148.0070	1 259	90.9992	3171.1760

```
29
    3.0417644
                18.146407
                            30.069904 1240.60567
                                                    156.52920
                                                                 933.8141
                                                                            1547.3973
                                                                            1201.1974
30
    2.4596628
                13.700651
                            23.342352
                                        953.11648
                                                    126.57425
                                                                 705.0355
31
    0.8268103
                 0.000000
                             3.003929
                                         71.19029
                                                     42.54766
                                                                   0.0000
                                                                             154.5822
32
           NA
                                   NA 5146.00000
                                                           NA
                                                                       NA
                                                                                   NA
                       NA
```

>

Use the write.csv function to write the condition estimates as a csv file.

```
> write.csv(Condition_Estimates_popsize, file="Condition_Estimates_popsize.csv")
```

## 6 Analysis of quantitative variables

The third analysis that will be examined is estimating the CDF and percentiles for quantitative variables. Two quantitative variables will be examined: (1) oxygen - dissolved oxygen value and (2) turbidity - turbidity value. The summary function is used to summarize the data structure of the two quantitative variables.

```
> summary(FL_lakes$oxygen)
```

Summarize the data structure of the dissolved oxygen variable:

```
Min. 1st Qu. Median Mean 3rd Qu. Max. NA's 0.830 4.880 6.870 6.468 8.310 12.480 759
```

> summary(FL\_lakes\$turbidity)

Summarize the data structure of the turbidity variable:

```
Min. 1st Qu. Median Mean 3rd Qu. Max. NA's 0.150 1.100 1.700 8.055 3.800 400.000 759
```

The cont.analysis function will be used to calculate estimates for quantitative variables. Input to the cont.analysis function is the same as input for the cat.analysis function except that the data frame containing response variables is named cont.data rather than cat.data. The sites, subpop, and design data frames created in the analysis of lake condition variables can be reused for this analysis. The data.cont data frame contains the two quantitative variables: DissolvedOxygen and Turbidity. Variables oxygen and turbidity in the FL\_lakes data frame are assigned to DissolvedOxygen and Turbidity, respectively. The popsize argument is included in the call to cont.analysis.

Create the data.cont data frame.

Use the cont.analysis function to calculate CDF and percentile estimates for the quantitative variables.

```
> CDF_Estimates <- cont.analysis(sites, subpop, design, data.cont,
+ popsize=list(CombinedBasins=sum(framesize),
+ Basin=as.list(framesize)))</pre>
```

The object produced by cont.analysis is a list containing two objects: (1) CDF, a data frame containing the CDF estimates and (2) Pct, a data frame containing percentile estimates plus estimates of population values for mean, variance, and standard deviation. Format for the CDF data frame is analogous to the data frame produced by cat.analysis. For the CDF data frame, however, the fourth column is labeled Value and contains the value at which the CDF was evaluated. Unlike the data frames produced by the other analysis functions we have examined, the Pct data frame contains only nine columns since there is a single set of estimates rather than two sets of estimates. In addition, the fourth column is labeled Statistic and identifies either a percentile or the mean, variance, or standard deviation. Finally, since percentile estimates are obtained by inverting the CDF estimate, the percentile estimates do not have a standard error value associated with them.

Use the write.csv function to write the CDF estimates as a csv file.

```
> write.csv(CDF_Estimates$CDF, file="CDF_Estimates.csv")
```

The cont.cdfplot function in spsurvey can be used to produce a PDF file containing plots of the CDF estimates. The primary arguments to cont.cdfplot are a character string containing a name for the PDF file and the CDF data frame in the CDF\_Estimates object. In addition, we make use of the logx argument to cont.cdfplot, which controls whether the CDF estimate is displayed using a logarithmic scale for the x-axis. The logx argument accepts two values: (1) "", do not use a logarithmic scale and (2) "x" - use a logarithmic scale. For this analysis, dissolved oxygen is displayed using the original response scale and turbidity is displayed using a logarithmic scale.

Produce a PDF file containing plots of the CDF estimates.

```
> cont.cdfplot("CDF_Estimates.pdf", CDF_Estimates$CDF, logx=c("","x"))
>
```

Print the percentile estimates for dissolved oxygen for all basins combined.

```
> # Print the percentile estimates for dissolved oxygen for all basins combined
> print(CDF_Estimates$Pct[1:10,])
```

```
Indicator
             Type Subpopulation
                                                       Statistic NResp Estimate
  CombinedBasins
                     All Basins DissolvedOxygen
                                                            5Pct
                                                                     8 1.578342
2
  CombinedBasins
                     All Basins DissolvedOxygen
                                                           10Pct
                                                                    17 2.285793
3
  CombinedBasins
                     All Basins DissolvedOxygen
                                                           25Pct
                                                                    42 4.624982
  CombinedBasins
                     All Basins DissolvedOxygen
                                                           50Pct
                                                                    83 6.809475
5
  CombinedBasins
                     All Basins DissolvedOxygen
                                                           75Pct
                                                                   129 8.333775
6
  CombinedBasins
                     All Basins DissolvedOxygen
                                                                   153 9.428672
                                                           90Pct
7
  CombinedBasins
                     All Basins DissolvedOxygen
                                                           95Pct
                                                                   163 9.996570
8
  CombinedBasins
                     All Basins DissolvedOxygen
                                                            Mean
                                                                   171 6.477253
9
  CombinedBasins
                     All Basins DissolvedOxygen
                                                                   171 6.442747
                                                        Variance
10 CombinedBasins
                     All Basins DissolvedOxygen Std. Deviation
                                                                   171 2.538257
            StdError
                      LCB95Pct
                                 UCB95Pct
                     0.9546438
1
                                 2.003976
2
                     1.7532592
                                 3.384501
3
                     4.1087503
                                 5.506396
4
                     6.5621691
                                 7.142007
5
                     7.9711324
                                 8.553456
6
                     9.0237184
                                 9.884125
7
                     9.7570792 10.457057
8
  0.148905597115604 6.1854029
                                 6.769102
  0.561664353995088 5.3419051
                                 7.543589
10 0.110639786234289 2.3214067
                                 2.755107
```

Use the write.csv function to write the percentile estimates as a csv file.

>

```
> write.csv(CDF_Estimates$Pct, file="Percentile_Estimates.csv")
```

The cont.cdftest function in spsurvey can be used to test for statistical difference between the CDFs from subpopulations. For this analysis we will test for statistical difference between the CDFs from the six basins. The cont.cdftest function will test all possible pairs of basins. Arguments to cont.cdftest are the same as arguments to cont.analysis. Since we are interested only in testing among basins, the subpop data frame is subsetted to include only the siteID and Basin variables. Note that the popsize argument was modified from prior examples to include only the entry for Basin.

```
> CDF_Tests <- cont.cdftest(sites, subpop[,c(1,3)], design, data.cont,
+ popsize=list(Basin=as.list(framesize)))</pre>
```

The print function is used to display results for dissolved oxygen of the statistical tests for difference between CDFs for basins. The object produced by cont.cdftest is a data frame containing eight columns. The first column (Type) identifies the population. The second

and third columns (Subpopulation\_1 and Subpopulation\_2) identify the subpopulations. The fourth column (Indicator) identifies the response variable. Column five contains values of the test statistic. Six test statistics are available, and the default statistic is an F-distribution version of the Wald statistic, which is identified in the data frame as "Wald-F". The default statistic is used in this analysis. For further information about the test statistics see the help file for the cdf.test function in spsurvey, which includes a reference for the test for differences in CDFs. Columns six and seven (Degrees\_of\_Freedom\_1 and Degrees\_of\_Freedom\_2) provide the numerator and denominator degrees of freedom for the Wald test. The final column (p\_Value) provides the p-value for the test.

- > # Print results of the statistical tests for difference between CDFs from
- > # basins for dissolved oxygen
- > print(CDF\_Tests, digits=3)

	Туре	Subpopulation_1	Subpopulation_2	Indicator	Wald_F
1	Basin	NWFWMD-1	NWFWMD-2	DissolvedOxygen	3.1442
2	Basin	NWFWMD-1	SFWMD-9	DissolvedOxygen	4.4795
3	Basin	NWFWMD-1	SJRWMD-1	DissolvedOxygen	20.2917
4	Basin	NWFWMD-1	SRWMD-1	DissolvedOxygen	0.3048
5	Basin	NWFWMD-1	SWFWMD-4	DissolvedOxygen	10.6685
6	${\tt Basin}$	NWFWMD-2	SFWMD-9	DissolvedOxygen	2.6095
7	${\tt Basin}$	NWFWMD-2	SJRWMD-1	DissolvedOxygen	6.1606
8	${\tt Basin}$	NWFWMD-2	SRWMD-1	DissolvedOxygen	2.8194
9	${\tt Basin}$	NWFWMD-2	SWFWMD-4	DissolvedOxygen	3.8223
10	${\tt Basin}$	SFWMD-9	SJRWMD-1	DissolvedOxygen	12.7598
11	${\tt Basin}$	SFWMD-9	SRWMD-1	DissolvedOxygen	6.0877
12	${\tt Basin}$	SFWMD-9	SWFWMD-4	DissolvedOxygen	14.1179
13	${\tt Basin}$	SJRWMD-1	SRWMD-1	${\tt DissolvedOxygen}$	16.9733
14	${\tt Basin}$	SJRWMD-1	SWFWMD-4	${\tt DissolvedOxygen}$	5.2374
15	${\tt Basin}$	SRWMD-1	SWFWMD-4	${\tt DissolvedOxygen}$	6.4086
16	${\tt Basin}$	NWFWMD-1	NWFWMD-2	Turbidity	0.5751
17	${\tt Basin}$	NWFWMD-1	SFWMD-9	Turbidity	1.5886
18	${\tt Basin}$	NWFWMD-1	SJRWMD-1	Turbidity	1.1966
19	${\tt Basin}$	NWFWMD-1	SRWMD-1	Turbidity	1.8996
20	${\tt Basin}$	NWFWMD-1	SWFWMD-4	Turbidity	11.3469
21	${\tt Basin}$	NWFWMD-2	SFWMD-9	Turbidity	0.2456
22	${\tt Basin}$	NWFWMD-2	SJRWMD-1	Turbidity	0.2944
23	${\tt Basin}$	NWFWMD-2	SRWMD-1	Turbidity	0.4627
24	${\tt Basin}$	NWFWMD-2	SWFWMD-4	Turbidity	11.0052
25	${\tt Basin}$	SFWMD-9	SJRWMD-1	Turbidity	0.3688
26	${\tt Basin}$	SFWMD-9	SRWMD-1	Turbidity	0.0753
27	${\tt Basin}$	SFWMD-9	SWFWMD-4	Turbidity	13.5140
28	${\tt Basin}$	SJRWMD-1	SRWMD-1	Turbidity	0.6625
29	${\tt Basin}$	SJRWMD-1	SWFWMD-4	Turbidity	17.2017

30	Basin	SRWMD-1	SWFWMD-4	Turbidity	9.7487
	Degrees_of_	Freedom_1 Degree	es_of_Freedom_2	p_Value	
1		2	55	5.09e-02	
2		2	57	1.56e-02	
3		2	57	2.21e-07	
4		2	54	7.39e-01	
5		2	51	1.35e-04	
6		2	55	8.27e-02	
7		2	55	3.85e-03	
8		2	52	6.88e-02	
9		2	49	2.87e-02	
10		2	57	2.63e-05	
11		2	54	4.13e-03	
12		2	51	1.32e-05	
13		2	54	1.91e-06	
14		2	51	8.54e-03	
15		2	48	3.41e-03	
16		2	55	5.66e-01	
17		2	57	2.13e-01	
18		2	57	3.10e-01	
19		2	54	1.59e-01	
20		2	51	8.39e-05	
21		2	55	7.83e-01	
22		2	55	7.46e-01	
23		2	52	6.32e-01	
24		2	49	1.13e-04	
25		2	57	6.93e-01	
26		2	54	9.28e-01	
27		2	51	1.95e-05	
28		2	54	5.20e-01	
29		2	51	1.95e-06	
30		2	48	2.80e-04	

Use the write.csv function to write CDF test results as a csv file.

```
> # Write CDF test results as a csv file
> write.csv(CDF_Tests, file="CDF_Tests.csv")
>
```

>