

Analysis of a GRTS Survey Design for an Area Resource

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1 Preliminaries

This document presents analysis of a GRTS survey design for an area resource. The area resource used in the analysis is estuaries in South Carolina. Although a stratified survey design was used to sample estuaries, analyses will be conducted as if the design was unstratified. Instead, strata will be used to define subpopulations for analysis. The strata employed in the survey were: (1) open water and (2) tidal creeks. The analysis will include calculation of three types of population estimates: (1) estimation of proportion and size (area of estuaries) for site evaluation status categorical variables; (2) estimation of proportion and size for estuary 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 2.2 of the spsurvey package was loaded successfully.

2 Read the survey design and analytical variables data file

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 SC_estuaries. The nrow function is used to determine the number of rows in the SC_estuaries 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 SC_estuaries 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
> SC_estuaries <- read.delim("SC_estuaries.tab")
> nr <- nrow(SC_estuaries)
>
```

Display the initial six lines in the data file.

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

	siteID	xcoord	ycoord	wgt	stratum	status	IBI_status	IBI_score
1	EEOW00-001	1549320	1263020	10.47515	Open Water	Sampled	Good	3.5
2	EEOW00-002	1487548	1226750	10.47515	Open Water	Sampled	Good	4.0
3	EEOW00-003	1442831	1159768	10.47515	Open Water	Sampled	Good	4.0
4	EEOW00-004	1425151	1148860	10.47515	Open Water	Sampled	Good	4.5
5	EEOW00-005	1432172	1140588	10.47515	Open Water	Sampled	Good	4.5
6	EEOW00-006	1540551	1280557	10.47515	Open Water	Sampled	Marginal	2.5
	WQ_status	WQ_score						
1	Good	4.3						
2	Good	4.6						


```

3      Good      5.0
4      Good      5.0
5      Good      5.0
6      Good      4.2

```

```
>
```

Display the final six lines in the data file.

```

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

```

	siteID	xcoord	ycoord	wgt	stratum	status	IBI_status
130	EETC99-035	1441835	1151395	1.41106	Tidal Creek	NonTarget	<NA>
131	EETC99-036	1535449	1247775	1.41106	Tidal Creek	Sampled	Good
132	EETC99-037	1500880	1225190	1.41106	Tidal Creek	Sampled	Marginal
133	EETC99-038	1440733	1147398	1.41106	Tidal Creek	Sampled	Good
134	EETC99-039	1468504	1179279	1.41106	Tidal Creek	Sampled	Good
135	EETC99-040	1430671	1151686	1.41106	Tidal Creek	Sampled	Marginal

	IBI_score	WQ_status	WQ_score
130	NA	<NA>	NA
131	3.5	Good	4.3
132	2.5	Poor	3.0
133	3.0	Marginal	3.7
134	4.0	Good	4.3
135	2.5	Marginal	3.7

```
>
```

The sample of estuaries in South Carolina is displayed in Figure 1. The sample sites for each stratum are displayed using a unique color. First, the levels function is used to extract the set of unique strata values, and the result is assigned to object stratum. Next, the rainbow function is called to select a set of two 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 two unique strata values and plot the color-coded points for each stratum 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.

```

> stratum <- levels(SC_estuaries$stratum)
> cols <- rainbow(2)
> plot(SC_estuaries$xcoord, SC_estuaries$ycoord, type="n", xlab="x-coordinate",
+       ylab="y-coordinate")

```



```

> for(i in 1:2) {
+   ind <- SC_estuaries$stratum == stratum[i]
+   points(SC_estuaries$xcoord[ind], SC_estuaries$ycoord[ind], pch=20,
+         col=cols[i])
+ }
> legend(x="topright", inset=0.05, legend=stratum, pch=20, cex=0.8, col=cols)
> title("Plot of South Carolina Estuary Sites Color-Coded by Stratum")

```

3 Analysis of site status evaluation variables

The first analysis that will be examined is calculation of extent estimates for a site status evaluation variable. Extent is measured both by the proportion of the resource in status evaluation categories and by size of the resource in each category. For an area resource like estuaries, size refers to the area of estuaries 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. The site status variable named status will be examined, which classifies estuaries into two evaluation categories: "Sampled" and "NonTarget". The table and addmargins functions are used to create tables displaying the count for each code (level) of the status variable.

```

> addmargins(table(SC_estuaries$status))

```

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

NonTarget	Sampled	Sum
19	116	135

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 SC_estuaries 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 SC_estuaries 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 subpop data

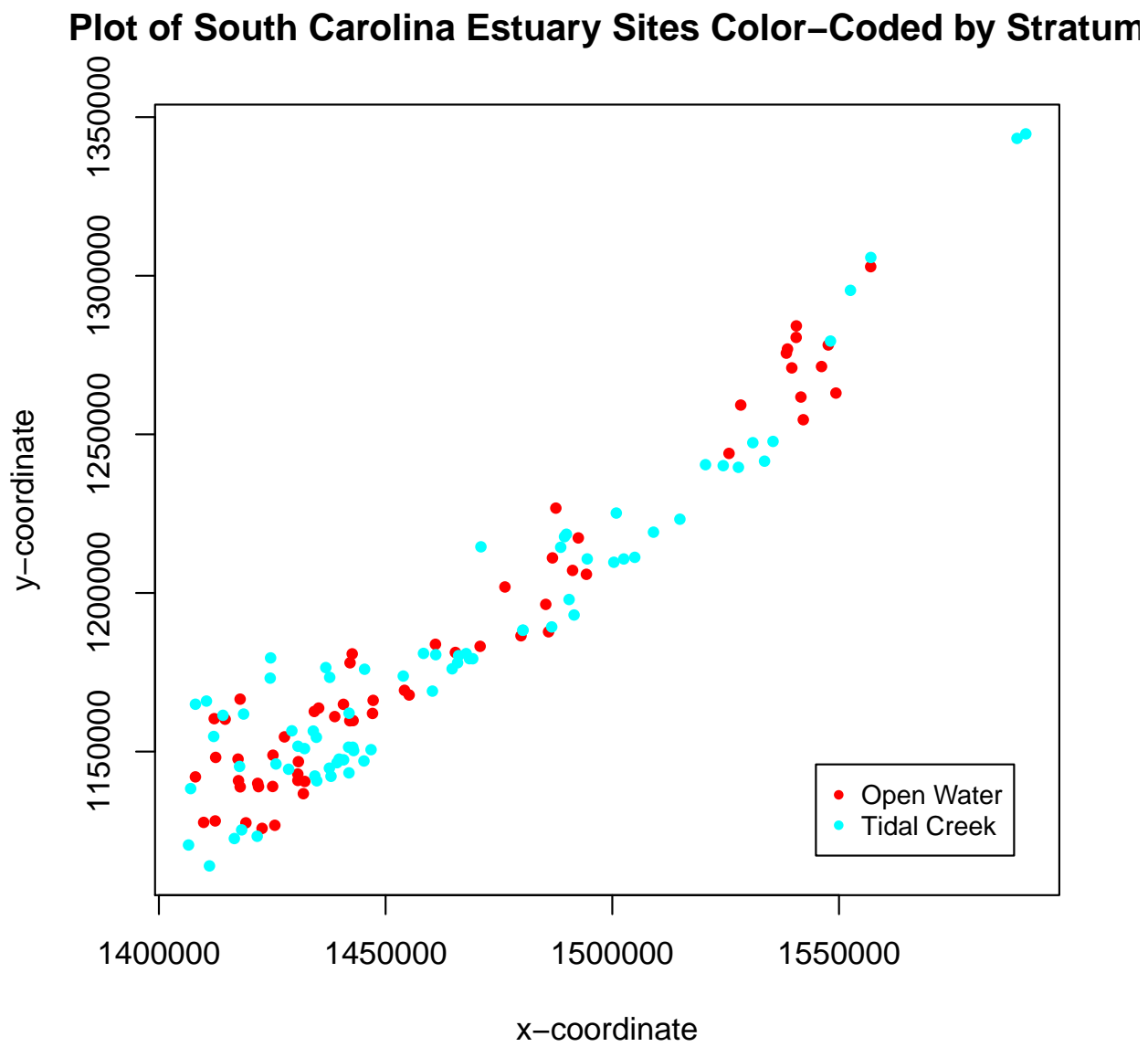


Figure 1: South Carolina Estuary Sample Sites.

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) All_Estuaries - which will be used to calculate estimates for all of the sample sites combined, and (3) Estuary_Type - which will be used to calculate estimates for each stratum individually. The stratum variable in the SC_estuaries data frame is assigned to the Estuary_Type 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 SC_estuaries 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 response variable is Status, which is assigned the status variable in the SC_estuaries 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.

```
> sites <- data.frame(siteID=SC_estuaries$siteID,
+                      Use=rep(TRUE, nr))
```

Create the subpop data frame.

```
> subpop <- data.frame(siteID=SC_estuaries$siteID,
+                      All_Estuaries=rep("All Estuaries", nr),
+                      Estuary_Type=SC_estuaries$stratum)
```

Create the design data frame.

```
> design <- data.frame(siteID=SC_estuaries$siteID,
+                      wgt=SC_estuaries$wgt,
+                      xcoord=SC_estuaries$xcoord,
+                      ycoord=SC_estuaries$ycoord)
```

Create the data.cat data frame.

```
> data.cat <- data.frame(siteID=SC_estuaries$siteID,
+                      Status=SC_estuaries$status)
```

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 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
> print(Extent_Estimates)
```

	Type	Subpopulation	Indicator	Category	NResp	Estimate.P	StdError.P
1	All_Estuaries	All Estuaries	Status	NonTarget	19	4.885244	1.335908
2	All_Estuaries	All Estuaries	Status	Sampled	116	95.114756	1.335908
3	All_Estuaries	All Estuaries	Status	Total	135	100.000000	0.000000
4	Estuary_Type	Open Water	Status	NonTarget	1	1.666667	1.443963
5	Estuary_Type	Open Water	Status	Sampled	59	98.333333	1.443963
6	Estuary_Type	Open Water	Status	Total	60	100.000000	0.000000
7	Estuary_Type	Tidal Creek	Status	NonTarget	18	24.000000	3.904808
8	Estuary_Type	Tidal Creek	Status	Sampled	57	76.000000	3.904808
9	Estuary_Type	Tidal Creek	Status	Total	75	100.000000	0.000000
	LCB95Pct.P	UCB95Pct.P	Estimate.U	StdError.U	LCB95Pct.U	UCB95Pct.U	
1	2.266912	7.503575	35.87424	9.445802e+00	17.36081	54.38767	
2	92.496425	97.733088	698.46458	4.380900e+01	612.60052	784.32864	
3	100.000000	100.000000	734.33882	4.210738e+01	651.80986	816.86778	
4	0.000000	4.496782	10.47515	9.075442e+00	0.000000	28.26270	
5	95.503218	100.000000	618.03414	9.075442e+00	600.24660	635.82168	
6	100.000000	100.000000	628.50930	3.476399e-14	628.50930	628.50930	
7	16.346718	31.653282	25.39909	4.132439e+00	17.29965	33.49852	
8	68.346718	83.653282	80.43044	4.132439e+00	72.33100	88.52987	
9	100.000000	100.000000	105.82952	4.435338e-15	105.82952	105.82952	

>

The `write.table` 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.table(Extent_Estimates, file="Extent_Estimates.csv", sep=",",
+             row.names=FALSE)
```

4 Analysis of estuary condition variables

The second analysis that will be examined is estimating resource proportion and size for estuary condition variables. Two estuary condition variables will be examined: (1) `IBI_Status`, which classifies estuaries by benthic IBI (index of biotic integrity) status categories and (2) `WQ_Status`, which classifies estuaries by WQ (water quality) status categories. The `table` and `addmargins` functions are used to create tables displaying the count for each level of the two estuary condition variables.

```
> addmargins(table(SC_estuaries$IBI_status))
```

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

Good	Marginal	Poor	Sum
99	14	3	116

```
> addmargins(table(SC_estuaries$WQ_status))
```

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

Good	Marginal	Poor	Sum
83	29	4	116

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 estuary condition variables: `IBI_Status` and `WQ_Status`. Variables `IBI_status` and `WQ_status` in the `SC_estuaries` data frame are assigned to `IBI_Status` and `WQ_Status`, respectively.

Create the sites data frame.


```
> sites <- data.frame(siteID=SC_estuaries$siteID,
+                      Use=SC_estuaries$status == "Sampled")
```

Create the data.cat data frame.

```
> data.cat <- data.frame(siteID=SC_estuaries$siteID,
+                        IBI_Status=SC_estuaries$IBI_status,
+                        WQ_Status=SC_estuaries$WQ_status)
```

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

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

Print the estuary condition estimates for all sites combined.

```
> # Print the condition estimates for all basins combined
> print(Condition_Estimates[c(1:3, 16:19),])
```

	Type	Subpopulation	Indicator	Category	NResp	Estimate.P	StdError.P
1	All_Estuaries	All Estuaries	IBI_Status	Good	99	86.183869	3.063318
2	All_Estuaries	All Estuaries	IBI_Status	Marginal	14	11.912344	3.111665
3	All_Estuaries	All Estuaries	IBI_Status	Poor	3	1.903787	1.349921
16	All_Estuaries	All Estuaries	WQ_Status	Total	116	100.000000	0.000000
17	Estuary_Type	Open Water	WQ_Status	Good	51	86.440678	4.011639
18	Estuary_Type	Open Water	WQ_Status	Marginal	8	13.559322	4.011639
19	Estuary_Type	Open Water	WQ_Status	Total	59	100.000000	0.000000
	LCB95Pct.P	UCB95Pct.P	Estimate.U	StdError.U	LCB95Pct.U	UCB95Pct.U	
1	80.179876	92.187861	601.96380	4.147219e+01	520.67980	683.24780	
2	5.813592	18.011097	83.20351	2.217377e+01	39.74371	126.66330	
3	0.000000	4.549583	13.29728	9.383494e+00	0.00000	31.68858	
16	100.000000	100.000000	698.46458	3.977348e+01	620.50999	776.41917	
17	78.578010	94.303346	534.23290	2.479330e+01	485.63893	582.82688	
18	5.696654	21.421990	83.80124	2.479330e+01	35.20726	132.39521	
19	100.000000	100.000000	618.03414	3.061318e-14	618.03414	618.03414	

```
>
```

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

```
> write.table(Condition_Estimates, file="Condition_Estimates.csv", sep="," ,
+             row.names=FALSE)
```


5 Analysis of estuary 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., area of estuaries) 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 `IBI_status` variable, population "All_Estuaries" and subpopulation "All Estuaries" is 698 square kilometers (rounded to a whole number). The corresponding frame size value is 734 square kilometers. The `popsiz` (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 `popsiz` argument is a list, which is a particular type of R object. The `popsiz` list must include an entry for each population type included in the subpop data frame, i.e., `All_Estuaries` and `Estuary_Type` for this analysis. The `sum` function applied to `framesize` is assigned to the `All_Estuaries` entry in the `popsiz` list. Recall that the `Estuary_Type` population contains subpopulations, i.e., stratum categories. When a population type contains subpopulations, the entry in the `popsiz` list also is a list. The `as.list` function is applied to `framesize`, and the result is assigned to the `Estuary_Type` entry in the `popsiz` list.

Assign frame size values.

```
> framesize <- c("Open Water"=628.509298, "Tidal Creek"=105.829522)
```

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

```
> Condition_Estimates_popsiz <- cat.analysis(sites, subpop, design, data.cat,
+   popsiz=list(All_Estuaries=sum(framesize),
+               Estuary_Type=as.list(framesize)))
```

Print the estuary condition estimates for IBI status.

```
> # Print the estuary condition estimates for all sites combined
> print(Condition_Estimates_popsiz[1:12,])
```

	Type	Subpopulation	Indicator	Category	NResp	Estimate.P	StdError.P
1	All_Estuaries	All Estuaries	IBI_Status	Good	99	86.183869	3.063318

2	All_Estuaries	All	Estuaries	IBI_Status	Marginal	14	11.912344	3.111665
3	All_Estuaries	All	Estuaries	IBI_Status	Poor	3	1.903787	1.349921
4	All_Estuaries	All	Estuaries	IBI_Status	Total	116	100.000000	NA
5	Estuary_Type	Open	Water	IBI_Status	Good	51	86.440678	3.461229
6	Estuary_Type	Open	Water	IBI_Status	Marginal	7	11.864407	3.526629
7	Estuary_Type	Open	Water	IBI_Status	Poor	1	1.694915	1.528238
8	Estuary_Type	Open	Water	IBI_Status	Total	59	100.000000	NA
9	Estuary_Type	Tidal	Creek	IBI_Status	Good	48	84.210526	4.344105
10	Estuary_Type	Tidal	Creek	IBI_Status	Marginal	7	12.280702	3.959455
11	Estuary_Type	Tidal	Creek	IBI_Status	Poor	2	3.508772	2.090309
12	Estuary_Type	Tidal	Creek	IBI_Status	Total	57	100.000000	NA
	LCB95Pct.P	UCB95Pct.P	Estimate.U	StdError.U	LCB95Pct.U	UCB95Pct.U		
1	80.179876	92.187861	632.881606	22.495132	588.791958	676.971253		
2	5.813592	18.011097	87.476970	22.850167	42.691465	132.262475		
3	0.000000	4.549583	13.980245	9.912994	0.000000	33.409355		
4	NA	NA	734.338820	NA	NA	NA		
5	79.656794	93.224562	543.287698	21.754147	500.650354	585.925042		
6	4.952341	18.776473	74.568900	22.165192	31.125922	118.011878		
7	0.000000	4.690206	10.652700	9.605117	0.000000	29.478383		
8	NA	NA	628.509298	NA	NA	NA		
9	75.696237	92.724816	89.119597	4.597346	80.108966	98.130229		
10	4.520312	20.041092	12.996608	4.190273	4.783824	21.209392		
11	0.000000	7.605702	3.713317	2.212164	0.000000	8.049078		
12	NA	NA	105.829522	NA	NA	NA		

>

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

```
> write.table(Condition_Estimates_popsizes, file="Condition_Estimates_popsizes.csv",
+             sep=";", row.names=FALSE)
```

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) IBI_score - IBI score and (2) WQ_score - WQ score. The summary function is used to summarize the data structure of the two quantitative variables.

```
> summary(SC_estuaries$IBI_score)
```

Summarize the data structure of the IBI score variable:

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
1.000	3.000	3.500	3.612	4.125	5.000	19

```
> summary(SC_estuaries$WQ_score)
```

Summarize the data structure of the WQ score variable:

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
2.700	4.000	4.600	4.407	5.000	5.000	19

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 estuary condition variables can be reused for this analysis. The `data.cont` data frame contains the two quantitative variables: `IBLScore` and `WQ_Score`, which contain the numeric scores for the IBI and WQ variables, respectively. Variables `IBI_score` and `WQ_score` in the `SC_estuaries` data frame are assigned to `IBLScore` and `WQ_Score`, respectively. The `popsiz` argument is included in the call to `cont.analysis`.

Create the `data.cont` data frame.

```
> data.cont <- data.frame(siteID=SC_estuaries$siteID,
+                          IBI_Score=SC_estuaries$IBI_score,
+                          WQ_Score=SC_estuaries$WQ_score)
```

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

```
> CDF_Estimates <- cont.analysis(sites, subpop, design, data.cont,
+   popsiz=list(All_Estuaries=sum(framesize),
+               Estuary_Type=as.list(framesize)))
```

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.table` function to write the CDF estimates as a csv file.


```
> write.table(CDF_Estimates$CDF, file="CDF_Estimates.csv", sep="," ,
+             row.names=FALSE)
```

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.

Produce a PDF file containing plots of the CDF estimates.

```
> cont.cdfplot("CDF_Estimates.pdf", CDF_Estimates$CDF)
>
```

Print the percentile estimates for IBI score for all sites combined.

```
> # Print the percentile estimates for IBI score for all sites combined
> print(CDF_Estimates$Pct[1:10,])
```

	Type	Subpopulation	Indicator	Statistic	NResp	Estimate
1	All_Estuaries	All Estuaries	IBI_Score	5Pct	3	1.9835561
2	All_Estuaries	All Estuaries	IBI_Score	10Pct	6	2.2809551
3	All_Estuaries	All Estuaries	IBI_Score	25Pct	17	2.8823748
4	All_Estuaries	All Estuaries	IBI_Score	50Pct	60	3.5875846
5	All_Estuaries	All Estuaries	IBI_Score	75Pct	87	4.1208723
6	All_Estuaries	All Estuaries	IBI_Score	90Pct	87	4.4496095
7	All_Estuaries	All Estuaries	IBI_Score	95Pct	110	4.6753552
8	All_Estuaries	All Estuaries	IBI_Score	Mean	116	3.7144320
9	All_Estuaries	All Estuaries	IBI_Score	Variance	116	0.6908874
10	All_Estuaries	All Estuaries	IBI_Score	Std. Deviation	116	0.8311964
		StdError	LCB95Pct	UCB95Pct		
1			1.5615314	2.1490635		
2			2.0393930	2.5134122		
3			2.6702423	3.0793319		
4			3.3695953	3.7713425		
5			3.9325709	4.3115620		
6			4.2517257	4.9369713		
7			4.4424379	5.0000000		
8	0.0721515473688744	3.5730176	3.8558464			
9	0.0866221699439855	0.5211111	0.8606637			
10	0.0521069224449442	0.7290687	0.9333241			

```
>
```

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


```
> write.table(CDF_Estimates$Pct, file="Percentile_Estimates.csv", sep=",",
+             row.names=FALSE)
```

The `cont.cdfctest` function in `spsurvey` can be used to test for statistical difference between the CDFs from subpopulations. For this analysis we will use the `cont.cdfctest` function to test for statistical difference between the CDFs from the two strata. Arguments to `cont.cdfctest` are the same as arguments to `cont.analysis`. Since we are interested only in testing among strata, the subpop data frame is subsetting to include only the `siteID` and `Estuary_Type` variables. Note that the `popsiz` argument was modified from prior examples to include only the entry for `Estuary_Type`.

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

The `print` function is used to display results for IBI score of the statistical tests for difference between CDFs for strata. The object produced by `cont.cdfctest` 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 strata CDFs for
> # IBI score and WQ score
> print(CDF_Tests)
```

	Type	Subpopulation_1	Subpopulation_2	Indicator	Wald_F
1	Estuary_Type	Open Water	Tidal Creek	IBI_Score	2.981072
2	Estuary_Type	Open Water	Tidal Creek	WQ_Score	14.585353
		Degrees_of_Freedom_1	Degrees_of_Freedom_2	p_Value	
1		2	109	5.489145e-02	
2		2	109	2.438223e-06	

```
>
```

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

```
> # Write CDF test results as a csv file
> write.table(CDF_Tests, file="CDF_Tests.csv", sep=",", row.names=FALSE)
>
```