

Calculating estimates for stratified random samples

Here is some R code that finds the stratified estimator of the population mean under simple random sampling within each stratum and an estimate of its standard deviation.

I begin by calling a very simple stratified population that I have constructed.

```
> library("RCurl")
> x<-getURL("http://users.stat.umn.edu/~gmeeden/classes/5201/moredata/stratpop1.txt")
> X<-read.table(textConnection(x), header=T)
> names(X)
```

```
[1] "y"      "strat"
```

```
> X
```

```
      y strat
1 10.21     b
2  6.05     a
3  8.80     b
4  3.35     a
5  5.94     a
6 11.22     b
7  7.87     b
8  5.57     a
9 13.02     b
10 9.69     b
11 6.60     a
12 9.23     b
```

```
> y<-X[,1]
> strat<-X[,2]
> foo<-split(y,strat)
```

You see there are two strata *a* and *b*, the first has 5 units and the second has 7 units. To make it easier to work with the strata I created *foo* using the `split` command. In R terminology *foo* is a list where each element of the list is a stratum. The `sapply` command lets me find information about each stratum in the population.

```
> foo
```

```
$a
[1] 6.05 3.35 5.94 5.57 6.60
```

```
$b
[1] 10.21 8.80 11.22 7.87 13.02 9.69 9.23
```

```

> foo[[1]]

[1] 6.05 3.35 5.94 5.57 6.60

> stratsizes<-sapply(foo,length)
> stratsizes

a b
5 7

> popsize<-length(X[,1])
> popsize

[1] 12

> WW<-stratsizes/popsize
> WW

      a      b
0.4166667 0.5833333

> stratmns<-sapply(foo,mean)
> stratmns

      a      b
5.50200 10.00571

```

Next I find a random sample of size 2 from stratum a and size 3 from stratum b. I then use the sapply command to find the strata means for the sample. Then I get the point estimate of the population mean. Next I find the estimated variance of our estimate and its standard error. Finally I find an approximate 95% confidence interval for the mean. Note this makes no sense here but it is included for completeness.

```

> smpsizes<-c(2,3)
> foo.smp<-list()
> set.seed(1962)
> for(i in seq(along=foo)){
+   foo.smp[[i]]<-sample(foo[[i]],smpsizes[i])
+ }
> names(foo.smp)<-names(foo)
> foo.smp

$a
[1] 6.05 5.94

$b
[1] 7.87 9.23 9.69

```

```

> smpmns<-sapply(foo.smp,mean)
> smpmns

      a      b
5.995 8.930

> est<-sum(WW*smpmns)
> est

[1] 7.707083

> smpvar<-sapply(foo.smp,var)
> smpvar

      a      b
0.00605 0.89560

> ratiosmptopop<-smpsizes/stratsizes
> ratiosmptopop

      a      b
0.4000000 0.4285714

> estvarofest<-sum((WW^2)*(1-ratiosmptopop)*(1/smpsizes)*smpvar)
> SEofest<-sqrt(estvarofest)
> SEofest

[1] 0.2415849

> lwbd<-est - 1.96*SEofest
> upbd<-est + 1.96*SEofest
> c(lwbd,upbd)

[1] 7.233577 8.180590

```

These commands should help you do HW assignment 4. Note for some of the bigger problems you will not want to look at all of X.