

Chapter 3, stratified random sampling

```
#####Chapter 3: Stratified random sampling#####
```

```
library(survey)
```

```
## Loading required package: grid
```

```
## Loading required package: Matrix
```

```
## Loading required package: survival
```

```
##  
## Attaching package: 'survey'
```

```
## The following object is masked from 'package:graphics':  
##  
## dotchart
```

```
library(SDaA)  
data(agstrat)  
head(agstrat) #take a look of the first 6 obsns
```

```
##           county state acres92 acres87 acres82 farms92 farms87 farms82  
## 1    PIERCE COUNTY   NE  297326  332862  319619    725    857    865  
## 2   JENNINGS COUNTY   IN  124694  131481  139111    658    671    751  
## 3    WAYNE COUNTY    OH  246938  263457  268434   1582   1734   1866  
## 4  VAN BUREN COUNTY   MI  206781  190251  197055   1164   1278   1464  
## 5   OZAUKEE COUNTY   WI   78772   85201   89331    448    483    527  
## 6 CLEARWATER COUNTY   MN  210897  229537  213105    583    699    693  
##  largef92 largef87 largef82 smallf92 smallf87 smallf82 region    rn  
## 1      54      54      42      58      67      48    NC   805  
## 2      14      13      14      42      36      38    NC   241  
## 3      20      19      16     175     186     184    NC   913  
## 4      23      17      9      56      66      55    NC   478  
## 5       6       5       5      56      49      48    NC  1028  
## 6      34      32      23       8      19      13    NC   496  
##      weight  
## 1 10.23301  
## 2 10.23301  
## 3 10.23301  
## 4 10.23301  
## 5 10.23301  
## 6 10.23301
```

```
nrow(agstrat) #number of rows
```

```
## [1] 300
```

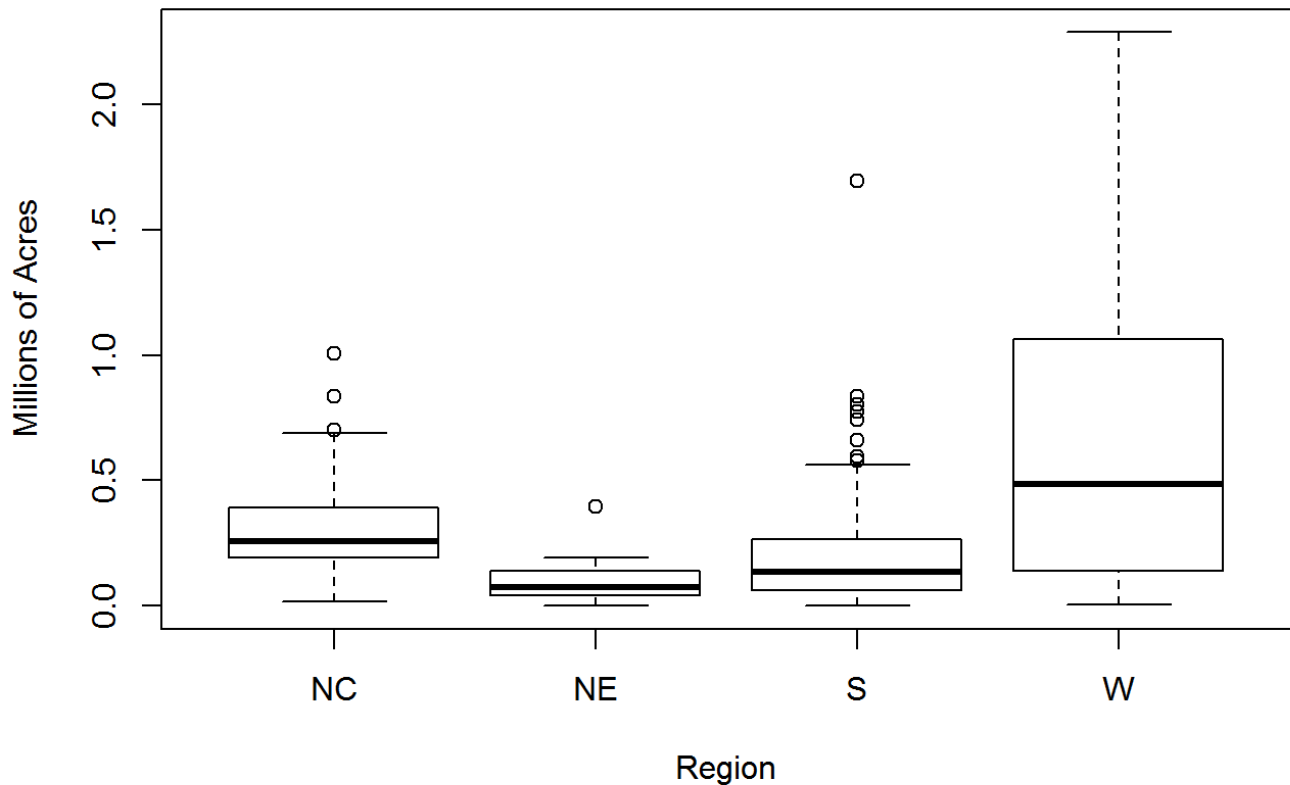
```
agstrat$region #take a look of the four regions, NC, NE, S, W
```

```
##      [1] NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC
##     [24] NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC
##     [47] NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC
##     [70] NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC
##     [93] NC NC NC NC NC NC NC NC NC NC NC NC NC NC NC NE NE NE NE NE NE NE NE NE NE NE NE NE
##    [116] NE NE NE NE NE NE NE NE NE NE S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [139] S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [162] S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [185] S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [208] S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [231] S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S  S
##    [254] S  S  S  S  S  S  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W
##    [277] W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W  W
##   [300] W
## Levels: NC NE S W
```

```
table(agstrat$region)
```

```
##
##  NC  NE   S   W
## 103  21 135  41
```

```
boxplot(acres92/10^6 ~ region, xlab = "Region", ylab = "Millions of Acres", data =
agstrat)
```



#notice the large variability in western region

```
dstr <- svydesign(id = ~1, strata = ~region, fpc = c(rep(1054,103),rep(220,21),rep(1382,135),rep(422,41)), weight = ~weight, data = agstrat)
dstr2 <- svydesign(id = ~1, strata = ~region, fpc = c(rep(1054,103),rep(220,21),rep(1382,135),rep(422,41)), data = agstrat)
dstr #
```

```
## Stratified Independent Sampling design
## svydesign(id = ~1, strata = ~region, fpc = c(rep(1054, 103),
##      rep(220, 21), rep(1382, 135), rep(422, 41)), weight = ~weight,
##      data = agstrat)
```

dstr2 #without weight info, it should give the same results

```
## Stratified Independent Sampling design
## svydesign(id = ~1, strata = ~region, fpc = c(rep(1054, 103),
##      rep(220, 21), rep(1382, 135), rep(422, 41)), data = agstrat)
```

```
##calculate mean, SE and confidence interval
smean<-svymean(~acres92, dstr)
smean
```

```
##          mean      SE
## acres92 295561 16380
```

```
smean2<-svymean(~acres92, dstr2)
smean2
```

```
##           mean      SE
## acres92 295561 16380
```

```
confint(smean, level=.95,df=296) #note df =n-H = 300-4
```

```
##           2.5 %    97.5 %
## acres92 263325 327796.5
```

```
##calculate total, SE and CI
stotal<-svyttotal(~acres92, dstr)
stotal
```

```
##           total      SE
## acres92 909736033 50417248
```

```
confint(stotal, level=.95,df=296)
```

```
##           2.5 %    97.5 %
## acres92 810514348 1008957719
```

```
#calculate S^2 for stratum NC
newdata1<-agstrat[which(agstrat$region=='NC'),]
var(newdata1$acres92)
```

```
## [1] 29618183543
```

```
##you can calculate the quantities stratum by stratum together
tapply(agstrat$acres92,agstrat$region,mean) #calculate mean for each region
```

```
##           NC           NE           S           W
## 300504.16  97629.81 211315.04 662295.51
```

```
tapply(agstrat$acres92,agstrat$region,var) #calculate variance for each region
```

```
##           NC           NE           S           W
## 29618183543  7647472708  53587487856 396185950266
```

```
tapply(agstrat$acres92,agstrat$region,sd) #calculate sd for each region
```

```
##           NC           NE           S           W
## 172099.34  87449.83 231489.71 629433.04
```

```
##calculate var(\bar y_h) for the four regions
svyby(~acres92, ~region, dstr, svymean, keep.var = TRUE)
```

```
##      region  acres92      se
## NC      NC 300504.16 16107.59
## NE      NE  97629.81 18149.49
## S       S 211315.04 18925.35
## W       W 662295.51 93403.65
```

```
#calculate var(\bar y_NC) by direct formula
sqrt(1-103/1054)*172099.34/sqrt(103)
```

```
## [1] 16107.59
```

```
##calculate var(\bar t_h) for the four regions
svyby(~acres92, ~region, dstr, svyttotal, keep.var = TRUE)
```

```
##      region  acres92      se
## NC      NC 316731379 16977399
## NE      NE  21478558  3992889
## S       S 292037390 26154840
## W       W 279488705 39416342
```

```
#####Select a stratified random #sampling#####
#install.packages("sampling")
library(sampling)
```

```
##
## Attaching package: 'sampling'
```

```
## The following objects are masked from 'package:survival':
##
##      cluster, strata
```

```
data(agpop)
agpop2<-agpop[order(agpop$region),]
idselected<-strata(agpop2,stratanames=c("region"),size=c(3,2,4,1), method="srswor"
)
idselected
```

```
##      region ID_unit      Prob Stratum
## 30      NC      30 0.002846300      1
## 537     NC     537 0.002846300      1
## 856     NC     856 0.002846300      1
## 1149    NE    1149 0.009090909      2
## 1160    NE    1160 0.009090909      2
## 1463     S    1463 0.002894356      3
## 1619     S    1619 0.002894356      3
## 1877     S    1877 0.002894356      3
## 1912     S    1912 0.002894356      3
## 2908     W    2908 0.002369668      4
```

```
table(idselected$region)
```

```
##
## NC NE  S  W
##  3  2  4  1
```

```
getdata(agpop2,idselected)
```

```
##      county state acres92 acres87 acres82 farms92 farms87
## 554  DICKINSON COUNTY  IA  202249  211002  227557    554    593
## 1342 OTTER TAIL COUNTY  MN  821073  876319  924670   2509   2925
## 2042  GEAUGA COUNTY   OH   65266   72766   74347    622    702
## 1975  JEFFERSON COUNTY NY  300559  338401  368352    894   1058
## 1986  ONONDAGA COUNTY NY  145329  158276  179015    636    772
## 341  OKEECHOBEE COUNTY FL  351885  384169  381895    418    400
## 497   THOMAS COUNTY   GA  174020  217103  215350    465    507
## 1515  ISSAQUENA COUNTY MS  113734  121019  111115    109    107
## 1550  SHARKEY COUNTY  MS  181946  177963  210485    125    127
## 1909  DE BACA COUNTY  NM 1343237 1296777 1165799    191    191
##      farms82 largef92 largef87 largef82 smallf92 smallf87 smallf82 region
## 554      652      30      34      31      38      47      43      NC
## 1342    3245     116     107     96     83     120     101     NC
## 2042     753       1       2       2     50     56     54     NC
## 1975    1245     31     25     22     25     32     40     NE
## 1986     835     24     21     22     56     68     77     NE
## 341     326     75     79     80     75     58     31     S
## 497     540     40     39     45     35     35     41     S
## 1515     134     32     36     31      0      4      4     S
## 1550     162     61     56     67      1      5      0     S
## 1909     204     91     86     88      9     22     23     W
##      ID_unit      Prob Stratum
## 554      30 0.002846300      1
## 1342     537 0.002846300      1
## 2042     856 0.002846300      1
## 1975    1149 0.009090909      2
## 1986    1160 0.009090909      2
## 341     1463 0.002894356      3
## 497     1619 0.002894356      3
## 1515    1877 0.002894356      3
## 1550    1912 0.002894356      3
## 1909    2908 0.002369668      4
```

##method could be srswr, systematic etc

```
insample<-c(sample(1:1054,103,replace=FALSE),  
sample(1055:1274,21,replace=FALSE),  
sample(1275:2656,135,replace=FALSE),sample(2657:3078,41,replace=FALSE))  
strsample<-agpop2[insample,]
```