# FINAL EXAM STAT 5201 Fall 2018 

Due on the class Canvas web site or in Room 313 Ford Hall on Tuesday, December 18 at 11:00 am In the second case please deliver to the office staff of the School of Statistics

## READ BEFORE STARTING

You must work alone and may only discuss these questions with the TA or Professor Meeden. You may use the class notes, the text and any other sources of material you have access to.

Start each answer on a new page and make sure that you name is on each page
If I discover a misprint or error in a question I will post a correction on the class web page. In case you think you have found an error you should check the class home page before contacting us.

Forty-seven persons took the exam. the scores ranged from 100 to 5 . There were 7 in the 90 's, 10 in the 80 's, 10 in the 70 's, 7 in the 60 's, 7 in the 50 's, 3 in the 40 's and 2 in the 30 's.

1. Find a recent survey reported in a newspaper, magazine or on the web. Briefly describe the survey. What are the target population and sampled population? What conclusions are drawn from the survey in the article. Do you think these conclusions are justified? What are the possible sources of bias in the survey? Please be specific but brief.
2. Information for a population with four strata is given just below.

| stratum | size | Est var | cost |
| :--- | :---: | :---: | :---: |
| 1 | 2000 | 450 | 10 |
| 2 | 1500 | 300 | 15 |
| 3 | 1000 | 500 | 20 |
| 4 | 1200 | 200 | 10 |

The second column of the table gives the sizes of the four strata, the next is our guess for the variance for each strata and the last, the cost per sampling an observation in each strata.
i) Using this information, find the optimal allocation of a sample of size 100.
ii) Suppose we want an estimate of the population mean with an variance of approximately 10. Based on the information in the table what would be the optimal allocation.

ANS i) For stratum $h$ let $N_{h}$ be its size and $w_{h}=N_{h} / N, s d_{h}$ be our guess for the standard deviation and $c_{h}$ the square root of the cost. Then in R I did the following

```
W<-c(20/57,15/57,10/57, 12/57)
> sd<-c(sqrt(450),sqrt(300),sqrt(500),sqrt (200))
> c<-c(sqrt(10),sqrt(15),sqrt(20),sqrt(10))
> n<-W*Sd/c
[1] 2.3537558 1.1768779 0.8771930 0.9415023
> nn<-100*(n/sum(n))
> nn
[1] 44.00095 22.00048 16.39819 17.60038
```

ii) From classwork or the text the solution is given by $n_{h}=\lambda \frac{w_{h} s d_{h}}{c_{h}}$ where $\lambda$ is the solution to the equation

$$
\lambda=\frac{\sum_{h} w_{h} s d_{h} c_{h}}{10+\sum_{h} w_{h}^{2} s d_{h}^{2} / N_{h}}
$$

This gives $\lambda=6.77$ with the corresponding allocation of $16,8,6$ and 6 .
3. From a population of 20 clusters a simple random sample of 4 clusters was taken. Within in each sampled cluster a further random sample was taken. The data are in a matrix denoted by mxclus. The following two commands will let you see the data in $R$.

```
library("RCurl")
```

load(url("http://users.stat.umn.edu/~gmeeden/classes/5201/moredata/clusf18.RData")) mxclus

The cluster sizes are only known for the clusters in the sample. Find the value of the ratio estimator for estimating the population mean and give an estimate of its variance.

ANS Using the handout on the class web page entitled working with cluster samples I found the estimate to be 25.21 and with an estimated variance of 19.51.
4. Consider the following table of sums of weights from a sample; each entry in the table is the sum of the sampling weights for persons in the sample falling in that classification (for example, the sum of the sampling weights for the number of women between the ages of 20 and 29 is 98 .)

|  | Age |  |  |  | Sum of |
| :--- | :---: | :---: | :---: | :---: | :---: |
|  | $20-29$ | $30-39$ | $40-49$ | $50-59$ | weights |
| Male | 183 | 447 | 522 | 416 | 1568 |
| Female | 98 | 377 | 395 | 467 | 1337 |
| Sum of | 281 | 824 | 917 | 883 |  |
| weights |  |  |  |  |  |

Assume it is known the that the population contains 1800 men and 1200 women and 300 persons between the ages of $20-29,900$ between 30-39, 1000 between $40-49$ and 800 between $50-59$. Readjust the cell weights so that in the new table the marginal weights agree with the known marginal population weights.

ANS
To do my raking I used the function altrake which can be found on the class web page in the handout called calibration using quadprog. Here is my code.

```
>mx<-rbind(c(183,447,522,416),c(98,377, 395,467))
>out<-altrake(mx,c(1800, 1200),c(300,900,1000, 800))
>out
> out
    [,1] [,2] [,3] [,4]
[1,] 210.6091 537.0491 622.9238 429.418
[2,] 89.3909 362.9509 377.0762 370.582
> apply(out,1,sum)
[1] 1800 1200
> apply(out,2,sum)
[1] 300 900 1000 800
```

5. Let $p=\left(p_{1}, \ldots, p_{n}\right)$ be a probability vector. That is each $p_{i}>0$ and they sum to one. Let $\Delta$ be the set of all such possible probability vectors. Let $\alpha=\left(\alpha_{1}, \ldots, \alpha_{n}\right)$ where each $\alpha_{i}>0$. We say that $p$ has the Dirichlet distribution with parameter $\alpha$ if its probability density function over $\Delta$ is given by

$$
f\left(p_{1}, \ldots, p_{n}\right)=\frac{\Gamma\left(\sum_{i=1}^{n} \alpha_{i}\right)}{\prod_{i=1}^{n} \Gamma\left(\alpha_{i}\right)} \prod_{i=1}^{n} p_{i}^{\alpha_{i}-1}
$$

Let $\alpha_{0}=\sum_{i=1}^{n} \alpha_{i}$. In class it was stated that

$$
E\left(p_{i}\right)=\alpha_{i} / \alpha_{0} \quad \text { and } \quad \operatorname{Var}\left(p_{i}\right)=\frac{\alpha_{i}\left(\alpha_{0}-\alpha_{i}\right)}{\alpha_{0}^{2}\left(\alpha_{0}+1\right)}
$$

i) Show that

$$
\operatorname{cov}\left(p_{i}, p_{j}\right)=-\frac{\alpha_{i}, \alpha_{j}}{\alpha_{0}^{2}\left(\alpha_{0}+1\right)}
$$

Remember that for any $b>0$ we have $\Gamma(b+1)=b \Gamma(b)$
ii) Let $a_{1}, \ldots, a_{n}$ be postive real numbers and assume that $p$ has the Dirichlet distribution with parameter $\alpha$. Find an expression for $\operatorname{Var}\left(\sum_{i=1}^{n} a_{i} p_{i}\right)$. You may answer the second part even if you can not do the first part.

ANS i) to keep the notation simple we will find the answer when $i=1$ and $j=2$.

$$
\begin{aligned}
E\left(p_{1} p_{2}\right) & =\frac{\Gamma\left(\alpha_{0}\right)}{\prod_{i=1}^{n} \Gamma\left(\alpha_{i}\right)} \int \cdots \int p_{1}^{\alpha_{1}+1-} p_{2}^{\alpha_{2}+1-1} \prod_{i=3}^{n} p_{i}^{\alpha_{i}-1} d p_{1} \cdots d p_{n} \\
& =\frac{\Gamma\left(\alpha_{0}\right)}{\prod_{i=1}^{n} \Gamma\left(\alpha_{i}\right)} \frac{\Gamma\left(\alpha_{1}+1\right) \Gamma\left(\alpha_{2}+1\right)}{\Gamma\left(\alpha_{0}+2\right)} \prod_{i=3}^{n} \Gamma\left(\alpha_{i}\right) \\
& =\frac{\alpha_{1} \alpha_{2}}{\alpha_{0}\left(\alpha_{0}+1\right)}
\end{aligned}
$$

Next we note that

$$
\begin{aligned}
\operatorname{cov}\left(p_{1}, p_{2}\right) & =E\left(p_{1} p_{2}\right)-E\left(p_{1}\right) E\left(p_{2}\right) \\
& =\frac{\alpha_{1} \alpha_{2}}{\alpha_{0}\left(\alpha_{0}+1\right)}-\frac{\alpha_{1} \alpha_{2}}{\alpha_{0}^{2}} \\
& =-\frac{\alpha_{i}, \alpha_{j}}{\alpha_{0}^{2}\left(\alpha_{0}+1\right)}
\end{aligned}
$$

ii) Recalling the well know formula for the variance of a sum we will have the answer by substitution.

$$
\operatorname{Var}\left(\sum_{i=1}^{n} a_{i} p_{i}\right)=\sum_{i=1}^{n} a_{i}^{2} \operatorname{Var}\left(p_{i}\right)+2 \sum_{i<j} a_{i} a_{j} \operatorname{cov}\left(p_{i}, p_{j}\right)
$$

6. In this problem you must preform a small simulation study to compare the ratio estimator and the regression estimator when the model generating the population is the correct one for the regression estimator. Use the following code to generate two different populations; the first with $s d=10$ and the second with $s d=5$. For each of the two populations take 400 simple random samples of size 10 and then of size 50. Compare both the point and interval estimators for the population total for the two estimators in the four cases. Discuss your results. Finally, construct a population such that when the sample size is 10 the average absolute error of the ratio estimator is at least four times larger that of the regression estimator.
```
set.seed(22334455)
```

popx<-rgamma $(1000,4)+40$
popy<-rnorm(1000,100 + 2*popx,sd)

ANS In each row of the following we have the average values of the estimator, its absolute error, the lower bound of the $95 \%$ confidence interval, the length of the interval and the proportion of times the interval contained the true total. Note that for the first population there is not much difference between the two estimators even for the larger sample size. It is only for the second population where the regression estimator does significantly better.

Results when $s d=10$, true total $=187783.3$ and $\operatorname{cov}(y, x)=0.41$
n=10
ansrt 187782.82681 .436181359 .712846 .120 .9325
ansrg 188019.3 2626.122182132 .211774 .150 .9100
n=50
[,1] [,2] [,3] [,4] [,5]
ansrt 187806.61103 .800184962 .65688 .1170 .97
ansrg 187647.61069 .034185020 .95253 .3440 .94
Results when $s d=5$, true total= 187936.8 and $\operatorname{cov}(y, x)=0.66$ n=10

```
    [,1] [,2] [,3] [,4] [,5]
ansrt 187880.7 1687.242 183877.0 8007.345 0.925
ansrg 188054.8 1313.061 185111.3 5887.077 0.910
n=50
```

    [,1] [,2] [,3] [,4] [,5]
    ansrt 188031.7678 .1392186283 .03497 .4450 .96
ansrg 187868.9534 .5169186555 .62626 .6720 .94
For the last part take popy<-rnorm(1000,200 + 2*popx,2)
keeping popx the same. In this case $\operatorname{cor}(\mathrm{y}, \mathrm{x})=0.91$ and
I got for $\mathrm{n}=10$.
[,1] [,2] [,3] [,4] [,5]
ansrt 287827.82400 .8158282058 .811538 .0820 .905
ansrg $288076.1 \quad 525.2245286898 .7 \quad 2354.8310 .910$

Here is the code I used to get the ans.

```
set.seed(22334455)
```

$\mathrm{x}<-\mathrm{rgamma}(1000,4)+40$

```
y<-rnorm(1000,100 + 4*popx,5)
ansrtreg<-function(y,x,n,R)
    {
        N<-length(y)
        ans<-matrix(0,2,5)
        for(i in 1:R){
            smp<-sort(sample(1:N,n))
            ansrt<-ratiotot(smp,y,x)
            ansrg<-regtot(smp,y,x)
            ans<-ans+ rbind(ansrt,ansrg)
        }
        return(ans/R)
    }
```

Here ratiotot and regtot are just the functions found in the $R$ handouts on the class web page dealing with the ratio estimator and the regression estimator.
7. Using the same popx as in problem 6 we now get popy by doing the following
popy<-rnorm(1000,100 + 4*popx,5)
For this population $\operatorname{cor}(p o p y, p o p x)=0.865$. Here you need to do a simulation study where you are estimating this correlation. You will take 300 samples of size $n=60$ doing pps sampling proportional to popx.

After you get the units in a sample and their design weights renormalize these weights so that they sum to 60 , the sample size. The next step is to simulate 500 complete copies of the population using the weighted polya posterior. You can do this using the function wtpolyap that is described in the $R$ handout using polyapost on the class web page. For each simulated complete copy of the population compute the correlation between the two variables. Using these 500 simulate correlations, find your estimate, its relative bias, its absolute error, the length of the approximated 0.95 credible interval and whether or not this interval contains the true correlation. Finally present the average of these quantities over the 300 samples.

Note if est is an estimate of the true median, $m d$, then the relative bias is equal to (est-md)/md.
Here are my results. The point estimate is working fine, the interval estimate is probably a bit short.

```
rel bias abs err len of intv freq of cov
-0.005989091 0.029182719 0.128871517 0.910000000
```

Here is the code that I used to get the answer.

```
anscor<-function(y,x,n,Rsim,R)
    {
        N<-length(y)
        wts<-sum(x)/(n*x)
        ans<-rep(0,4)
        trucor<-cor(y,x)
        for(i in 1:R){
        smp<-sort(sample(1:N,n,prob=x))
        swts<-wts[smp]
        swts<-n*swts/sum(swts)
        simcor<-rep(0,Rsim)
        for(j in 1:Rsim){
            out<-wtpolyap(smp,swts,N-n)
            simcor[j]<-cor(y[out],x[out])
        }
        est<-mean(simcor)
        bds<-as.numeric(quantile(simcor,c(0.025,0.975)))
        if(bds[1]<=trucor & trucor<=bds[2]) cov<-1
        else cov<-0
        ans<-ans+c((est-trucor)/trucor,abs(est-trucor),bds [2]-bds [1], cov)
    }
    cat("true cor", trucor,"\n")
    return(ans/R)
    }
```

