Finite Population Sampling

Fernando TUSELL

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Outline

Introduction

Sampling of independent observations

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Sampling without replacement

Stratified sampling

Taking samples

• We have been assuming samples

$$X_1, X_2, \ldots, X_n$$

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 - When we sample an infinite population: seeing one value does not affect the probability of seeing the same or another value.

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When we sample with replacement.

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- This makes sense:
 - When we sample an infinite population: seeing one value does not affect the probability of seeing the same or another value.
 - When we sample with replacement.
- With finite populations without replacement, what we see affects the probability of what is yet to be seen.

 With infinite populations, precision depends only on sample size.

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- With infinite populations, precision depends only on sample size.
- Usually, standard error of estimation is ^σ/_n where n is sample size and σ² the population variance.
- If estimator is consistent we approach (but never quite hit with certainty) the true value of the parameter.

If population is finite of size N, we could inspect all units and estimate anything with certainty:

$$\hat{m} = \frac{X_1 + X_2 + \ldots + X_n}{n}$$

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would verify $m = \hat{m}$ if n = N.

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 - If $n/N \approx 0$, independent sampling good approximation.

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- All parameters can, in principle, be known with certainty!
- With $n \neq N$,
 - If $n/N \approx 0$, independent sampling good approximation.
 - If $n/N \gg 0$, we have to take into account that we are looking at a substantial portion of the population.

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An overview of things to come

We will see:

What makes sampling without replacement more complex.

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 What relationship there is among independent and non-independent sampling.

An overview of things to come

We will see:

What makes sampling without replacement more complex.

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- What relationship there is among independent and non-independent sampling.
- What other types of sampling exist.

The central approximation

▶ Requirement: replacement of "large" population size *N*.

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- ▶ If *n* is "large" and X_1, \ldots, X_n "near" independent,

$$\overline{X} = \frac{X_1 + \ldots + X_n}{n} \sim N(m, \sigma^2/n)$$

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Then,

$$\operatorname{Prob}\left(\overline{X} - z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}} \le m \le \overline{X} + z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}}\right) = 1 - \alpha$$

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Estimation of the population total

Since T = Nm, we just have multiply by N the extremes of the interval for m.

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Estimation of the population total

- Since T = Nm, we just have multiply by N the extremes of the interval for m.
- ► Hence,

$$\operatorname{Prob}\left(N\overline{X}-Nz_{\alpha/2}\sqrt{\frac{\sigma^2}{n}} \leq T \leq N\overline{X}+Nz_{\alpha/2}\sqrt{\frac{\sigma^2}{n}}\right) = 1-\alpha$$

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• If X_i is a binary variable, \overline{X} is the sample proportion.

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• Usual estimate of variance is $\hat{p}(1-\hat{p})/n$.

- If X_i is a binary variable, \overline{X} is the sample proportion.
- We have $\overline{X} \sim N(p, pq/n)$
- Usual estimate of variance is $\hat{p}(1-\hat{p})/n$.
- Sometimes we use a (conservative) estimate: pq ≤ 0.5, hence a bound for σ² is 0.5/n.

Sampling error with confidence $1 - \alpha$.

From

$$\operatorname{Prob}\left(\overline{X} - z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}} \le m \le \overline{X} + z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}}\right) = 1 - \alpha$$

we see that we will be off the true value *m* by less than $z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}}$ with probability $1 - \alpha$.

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• This is called the " $1 - \alpha$ (sampling) error".

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we see that we will be off the true value *m* by less than $z_{\alpha/2}\sqrt{\frac{\sigma^2}{n}}$ with probability $1 - \alpha$.

- This is called the " 1α (sampling) error".
- "Sampling error" also used to mean standard deviation of the estimate.

Example:

What n do we need so that with confidence 0.95 the error in the estimation of a proportion is less than 0.03?

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• Confidence 0.95 means $z_{\alpha/2} = 1.96$

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- Confidence 0.95 means $z_{\alpha/2} = 1.96$
- Want $0.03 > 1.96\sqrt{\frac{\sigma^2}{n}}$. Worst case scenario is $\sigma^2 = 0.25$.

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► Therefore, $n > \frac{(1.96)^2 \times 0.25}{0.03^2} = 1067.11$ will do. Will take n = 1068.

Interesting facts (I)

 Under independent sampling (infite population or sampling with replacement), required sample size depends only on variance and precsion required.
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- Questions like: "Is a sample of 4% enough?" are badly posed.
- ► n = 4% of a population with N = 10000 insufficient to give a precision of 0.03 with confidence 0.95.

Interesting facts (I)

- Under independent sampling (infite population or sampling with replacement), required sample size depends only on variance and precsion required.
- Questions like: "Is a sample of 4% enough?" are badly posed.
- ► n = 4% of a population with N = 10000 insufficient to give a precision of 0.03 with confidence 0.95.
- ► ... but n = 0.3% of a population with N = 1000000 will be more than enough!

Finite Population Sampling — Sampling of independent observations

Interesting facts (II)

As long as populations are large detail is expensive!

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- As long as populations are large detail is expensive!
- To estimate a proportion in the CAPV with the precision stated requires about n = 1068.

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- To estimate the same proportion for each of the three Territories with the same precision, requires three times as large a sample!

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Interesting facts (II)

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- ► To estimate a proportion in the CAPV with the precision stated requires about n = 1068.
- To estimate the same proportion for each of the three Territories with the same precision, requires three times as large a sample!
- Subpopulation estimates have much lower precision than those for the whole population.

In independent sampling,

$$E[\overline{x}] = E\left[\frac{X_1 + \ldots + X_n}{n}\right]$$
$$= \frac{m + m + \ldots + m}{n} = \frac{nm}{n} = m$$

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- Without replacement, distribution of X_i depends on what other values are already present in the sample.

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- ► E[X_i] = m irrespective of what other values are in the sample.
- ▶ Without replacement, distribution of X_i depends on what other values are already present in the sample.
- The same result as for independent sampling is true!

Theorem 1

In a finite population of size N with $m = \sum_{i=1}^{N} y_i/N$, for samples Y_1, \ldots, Y_n without replacement of size n < N we have:

 $E[\overline{Y}] = m$

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Proof

- Y_1, Y_2, \ldots, Y_n are the elements of the sample.
- y_1, y_2, \ldots, y_N are the elements of the population.

• There are
$$\binom{N}{n} = \frac{N!}{(N-n)!n!}$$
 different samples.

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- There are $\binom{N}{n} = \frac{N!}{(N-n)!n!}$ different samples.
- Of those, $\binom{N-1}{n-1}$ contain each of the values y_1, y_2, \ldots, y_N .
- Clearly,

$$\sum (Y_1 + Y_2 + \ldots + Y_n) = \binom{N-1}{n-1} (y_1 + y_2 + \ldots + y_N)$$

where the sum in the left is taken over all $\binom{N}{n}$ different samples. Dividing by $\binom{N}{n}$ finishes the proof.

► Indeed,

$$\frac{\sum(Y_1 + Y_2 + \ldots + Y_n)}{\binom{N}{n}} = \frac{\binom{N-1}{n-1}(y_1 + y_2 + \ldots + y_N)}{\binom{N}{n}} = \frac{n}{N}(y_1 + y_2 + \ldots + y_N)$$

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► Therefore,

$$E[\overline{Y}] = \frac{\sum(Y_1 + \ldots + Y_n)/n}{\binom{N}{n}} = \frac{(y_1 + \ldots + y_N)}{N} = E[\overline{y}] = m$$

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The indicator variable method

We have

$$(Y_1 + Y_2 + \ldots + Y_n) = (y_1Z_1 + y_2Z_2 + \ldots y_NZ_N)$$

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where Z_i is a binary variable which takes value 1 if y_i belongs to a given sample.

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where Z_i is a binary variable which takes value 1 if y_i belongs to a given sample.

• The probability of that happening is n/N. Then,

$$E[(Y_1+Y_2+\ldots+Y_n)]=\frac{n}{N}(y_1+y_2+\ldots y_N),$$

which again gives the previous result $E[\overline{Y}] = \overline{y} = m$.

Population variance an quasi-variance

They are defined as:

$$\sigma^2 = \frac{\sum_{i=1}^{N} (y_i - \overline{y})^2}{N} \qquad \qquad \tilde{\sigma}^2 = \frac{\sum_{i=1}^{N} (y_i - \overline{y})^2}{N - 1}$$

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Similarly for sample analogues:

$$s^{2} = \frac{\sum_{i=1}^{n} (Y_{i} - \overline{Y})^{2}}{n} \qquad \qquad \tilde{s}^{2} = \frac{\sum_{i=1}^{n} (Y_{i} - \overline{Y})^{2}}{n-1}$$

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 Turns out some formulae are simpler in terms of quasi-variances.

Variance of \overline{Y} (I)

Theorem 2

In a finite population of size N, the estimator \overline{Y} of $m = \sum_{i=1}^{N} y_i / N$ based on a sample of size n < N without replacement Y_1, \ldots, Y_n has variance:

$$\operatorname{Var}[\overline{Y}] = \frac{\widetilde{\sigma}^2}{n} \left(1 - \frac{n}{N}\right)$$

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Factor

$$\left(1-\frac{n}{N}\right)$$

usually called "finite population correction factor" or "correction factor".

Finite Population Sampling

Variance of \overline{Y} (II)

Remarks:

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Variance of \overline{Y} (II)

Remarks:

▶ It is the same expression as in independent random sampling with i) σ^2 replaced by $\tilde{\sigma}^2$, and ii) corrected with the factor (1 - n/N).

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• If n = N, the variance $Var(\overline{Y})$ is 0 (why?).

Variance of \overline{Y} (II)

Remarks:

- It is the same expression as in independent random sampling with i) σ² replaced by σ̃², and ii) corrected with the factor (1 − n/N).
- If n = N, the variance $Var(\overline{Y})$ is 0 (why?).
- ► Formula covers middle ground between infinite populations (n/N = 0) and census sampling (n/N = 1).

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Finite Population Sampling

Variance of
$$\overline{Y}$$
 (III)

Proof

$$\begin{aligned} \operatorname{Var}(\overline{Y}) &= \operatorname{Var}\left(\frac{y_1 Z_i + \ldots + y_N Z_N}{n}\right) \\ &= \frac{1}{n^2} \left[\sum_{i=1}^N y_i^2 \operatorname{Var}(Z_i) + \sum_{i=1}^N \sum_{j \neq i} y_i y_j \operatorname{Cov}(Z_i, Z_j) \right] \end{aligned}$$

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• We only need expressions for $Var(Z_i)$ and $Cov(Z_i, Z_j)$.

Variance of \overline{Y} (IV)

• Since Z_i is binary with probability n/N,

$$\operatorname{Var}(Z_i) = (n/N)(1 - n/N).$$

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• But
$$E[Z_i Z_j] = P(Z_i = 1, Z_j = 1) = \frac{n(n-1)}{N(N-1)}$$
, so

$$\operatorname{Cov}(Z_i, Z_j) = \frac{n(n-1)}{N(N-1)} - \left(\frac{n}{N}\right)^2 = -\frac{n(1-n/N)}{N(N-1)}$$

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• Replacing in expression for $Var(\overline{Y})$ will lead to result.
Variance of \overline{Y} (V)

$$\operatorname{Var}(\overline{Y}) = \frac{1}{n^2} \left[\sum_{i=1}^{N} y_i^2 \underbrace{\operatorname{Var}(Z_i)}_{(n/N)(1-n/N)} + \sum_{i=1}^{N} \sum_{j \neq i} y_i y_j \underbrace{\operatorname{Cov}(Z_i, Z_j)}_{-\frac{n(1-n/N)}{N(N-1)}} \right]$$
$$= \frac{1}{n^2} \left(\frac{n}{N} \right) \left(1 - \frac{n}{N} \right) \left[\sum_{i=1}^{N} y_i^2 - \frac{1}{N-1} \sum_{i=1}^{N} \sum_{j \neq i} y_i y_j \right]$$

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Will rewrite expression in brackets.

Finite Population Sampling

Variance of \overline{Y} (VI)

Remark that,

$$\sum_{i=1}^{N} (y_i - m)^2 = \sum_{i=1}^{N} y_i^2 - \frac{\left(\sum_{i=1}^{N} y_i\right)^2}{N}$$
$$= \frac{N-1}{N} \left[\sum_{i=1}^{N} y_i^2 - \sum_{i=1}^{N} \sum_{j \neq i} \frac{y_i y_j}{N-1} \right]$$

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• The expression in square brackets in th r.h.s is therefore $\frac{N}{N-1}\sum_{i=1}^{N}(y_i - m)^2$.

Variance of \overline{Y} (VII)

We are now done!

$$\operatorname{Var}(\overline{Y}) = \frac{1}{n^2} \left(\frac{n}{N} \right) \left(1 - \frac{n}{N} \right) \left[\sum_{i=1}^N y_i^2 - \frac{1}{N-1} \sum_{i=1}^N \sum_{j \neq i} y_i y_j \right]$$
$$= \frac{1}{n} \left(1 - \frac{n}{N} \right) \frac{\sum_{i=1}^N (y_i - m)^2}{N-1}$$
$$= \left(1 - \frac{n}{N} \right) \frac{\tilde{\sigma}^2}{n}$$

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Sample size for given precision (I)

$$\delta = z_{\alpha/2} \sqrt{\frac{\tilde{\sigma}^2}{n} (1 - n/N)}$$

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Solving for n we obtain

$$n = \frac{N z_{\alpha/2}^2 \tilde{\sigma}^2}{N \delta^2 + \tilde{\sigma}^2 z_{\alpha/2}^2}$$

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In terms of the variance, it can be written as:

$$n = \frac{N z_{\alpha/2}^2 \sigma^2}{(N-1)\delta^2 + \sigma^2 z_{\alpha/2}^2}$$

Sample size for given precision (II)

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•
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 or σ^2 are required

Sample size for given precision (II)

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• We either replace an upper bound or conservative estimation for σ^2 .

Sample size for given precision (II)

- $\tilde{\sigma^2}$ or σ^2 are required.
- We either replace an upper bound or conservative estimation for σ^2 .

• Failing that, we estimate σ^2 or $\tilde{\sigma}^2$.

Finite Population Sampling

Why strata?

Sometimes we know something about the composition of the population, knowledge that can be put to use.

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Why strata?

- Sometimes we know something about the composition of the population, knowledge that can be put to use.
- **Example:** We might know that males and females have different spending in e.g. tobacco or cosmetics.
- To estimate average spending, it makes sense to sample males and females, and combine the estimations.
- Sometimes, the target quantity might be similar, but the variance quite different. Also makes sense to differentiate.

Finite Population Sampling

Example 1



Makes sense to estimate mean in each subpopulation

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Definitions and notation

► We assume the population is divided in *h* strata. Total size is N = N₁ + N₂ + ... + N_h.

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Definitions and notation

► We assume the population is divided in *h* strata. Total size is N = N₁ + N₂ + ... + N_h.

• The *i*-th stratum has a mean $m_i = \frac{1}{N_i} \sum_{j=1}^{N_i} y_{ij}$ and variance $\sigma_i^2 = \frac{1}{N_i} \sum_{j=1}^{N_i} (y_{ij} - m_i)^2$.

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Clearly,

$$m = \sum_{i=1}^{h} \left(\frac{N_i}{N}\right) m_i$$

$$\sigma^2 = \sum_{i=1}^{h} \frac{N_i}{N} \sigma_i^2 + \sum_{i=1}^{h} \frac{N_i}{N} (m_i - m)^2$$

Estimation of the mean

► The estimation of the mean sampling without replacement the whole population has variance $\frac{\tilde{\sigma}^2}{n}(1 - n/N)$.

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- ► Similarly, the estimation of the mean of each stratum has variance $\sigma_i^2 = \frac{\tilde{\sigma}_i^2}{n} (1 n_i / N_i)$.

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- Similarly, the estimation of the mean of each stratum has variance $\sigma_i^2 = \frac{\tilde{\sigma}_i^2}{n} (1 n_i / N_i)$.
- The variance of the global mean reconstituted from the estimated means of the strata is

$$\sigma_*^2 = \sum_{i=1}^h \left(\frac{N_i}{N}\right)^2 \frac{\tilde{\sigma}_i^2}{n_i} (1 - n_i/N_i)$$

Does the estimation of *m* improve?

Yes. If we sample each stratum in proportion to its size (i.e., n_i/N_i = n/N for all i), then:

$$\begin{split} \frac{\tilde{\sigma}^2}{n}(1-n/N) &- \sigma_*^2 = \\ & \left(1-\frac{n}{N}\right)\sum_{i=1}^h \left(\frac{N_i}{N}\right) \left[\frac{N_i-1}{N-1} - \frac{N_i}{N}\right]\frac{\tilde{\sigma}_i^2}{n_i} + \\ & \left(1-\frac{n}{N}\right)\frac{1}{n}\sum_{i=1}^h \frac{N_i}{N-1}(m_i-m)^2 \end{split}$$

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Yes. If we sample each stratum in proportion to its size (i.e., n_i/N_i = n/N for all i), then:

$$\frac{\tilde{\sigma}^2}{n}(1-n/N) - \sigma_*^2 = \left(1-\frac{n}{N}\right)\sum_{i=1}^h \left(\frac{N_i}{N}\right) \left[\frac{N_i-1}{N-1} - \frac{N_i}{N}\right]\frac{\tilde{\sigma}_i^2}{n_i} + \left(1-\frac{n}{N}\right)\frac{1}{n}\sum_{i=1}^h \frac{N_i}{N-1}(m_i-m)^2$$

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Marked Improvement when the m_i's very different.

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- Important contributions to the war effort as statistician (notably sequential analysis)
- Was consulted about aircraft armoring.

• Mark hits in B-29 bombers as they come back.



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- Pretty obvious! Will armor the most beaten areas.
- I didn't tell you to do that!
- Do you want us to protect the areas with no hits?
- That's exactly what I suggest!

Sample selection is ubiquitous!

 If you ask for volunteers in a field study, no chance you will get a truly random sample.

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- Never do!
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- A random sample is not a "grab set".
- Build a census, randomize properly, address the chosen units and no others.
- If you use systematic sampling (every *n*-th unit with random start), make sure no periodicities exist that will destroy randomness.