EKT-816 Lecture 2

Probability Review (2)

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- consistency: $\widehat{\theta} \longrightarrow_{p} \theta_{0}$
- if $E[\widehat{\theta}] = \theta_0$ we say $\widehat{\theta}$ is unbiased
- ullet efficiency or precision: say we have two estimators $\widehat{ heta}$ and $\widetilde{ heta}$
 - for now assume both are unbiased
 - if $V[heta] \leq V[heta]$, say that heta is more efficient than heta
- the mean square error of $\widehat{\theta}$ is $MSE(\widehat{\theta}) = E[(\widehat{\theta} \theta_0)^2]$
 - ullet easy to see that $\mathsf{MSE} = V[heta] + \mathsf{bias}^2$
 - often a trade-off between the two criteria
 - typically people seek unbiased estimators, but not always clear they are better in a MSE sense

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$$f(X_1, \ldots X_n, \theta) = g(T(X_1, \ldots X_n), \theta) \cdot h(X_1, \ldots X_n)$$

- e.g. $T = \sum_{i=1}^{n} X_i$ for normal data with known variance
- ullet then, a maximum likelihood estimator must be a function of T
- in fact, the Rao-Blackwell Theorem says (roughly) that any unbiased estimator which is not a function of the sufficient statistic has higher variance than "necessary"
 - more precisely, higher variance than the MLE, which hits the (Cramer-Rao) lower bound
 - intuition: if you base estimates on irrelevant information, you are sacrificing precision

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• we will not use an explicit likelihood framework much

- but, the idea of "sufficiency" is still useful
- in some situations, all of the relevant information can be reduced to some low-dimensional summaries
- see Chetty (2009) and Weyl (2019) for examples of how this idea connects theory and econometrics
- this idea also comes up in the guise of "selection (only) on observables"

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- say we have an iid sample of size n from a $U(0, \theta_0)$ distribution
 - the sample maximum is, in this case, sufficient for θ_0
 - in fact, can show the MLE is $\widehat{\theta} = \max\{X_1, \dots X_n\}$
- another estimator would be $\widetilde{\theta} = 2\overline{X}_n$
 - this is unbiased (show this!)
- which estimator has lower MSE? Which has lower variance?
- to derive the distribution of the sample maximum:
 - use the fact that $\max\{X_1,\ldots X_n\} \leq x$ if and only if each $X_i \leq x$
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some types of data you may encounter:

- cross-sectional
 - units from the population are surveyed once, and all at roughly the same time
 - a "snapshot" of the population

stratified or two-stage designs

- units are sampled randomly within certain pre-specified groups
- e.g region, race, sex

clustered designs

- often would be expensive to collect a simple random sample
- save on transport and labor costs by selecting clusters of units
- attempt to correct for the resulting correlations (why?)

panel or longitudinal designs: repeated observations on the same units

- rotating panels, where some units are "rotated" in and out of the survey
- retrospective histories
- synthetic panels: aggregate individuals to form a panel at cohort leve

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- suppose we have a population of N units, and we sample with unequal probabilities
 - let π_i be the probability that unit i is selected
 - in a simple random sample of size n << N, $\pi_i \approx n/N$, even if we sample without replacement
 - but we don't always want to sample each unit with equal probability
- let $w_i = (n\pi_i)^{-1}$ be the "design weight"; then the expected sum of the weights is

$$E\left[\sum_{i=1}^{n} w_{i}\right] = E\left[\sum_{i=1}^{N} t_{i} w_{i}\right]$$

$$\approx \sum_{i=1}^{N} (\pi_{i} n) w_{i} = N$$
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- intuition: weights are inversely proportional to the probabilities of inclusion
 - in a 1/100 sample, each included unit represents 100 others
- to estimate the population mean, we weight the observations by w_i , forming

$$\overline{x}_w = \sum_{i=1}^n w_i x_i \tag{2}$$

- exercise: show that $E[\overline{x}_w] = E[X]$
- we may want to deliberately oversample some groups to improve precision of conditional means
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- to understand why stratification is useful, think about trying to measure the national average of some X where there are two cities
- variance of a randomly sampled unit is

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- consider a stratified design where we take two independent samples from the two strata
 - size n_1 and n_2 respectively, with $n_1 + n_2 = r$
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$$V[\overline{x}^{STRAT}] = V\left[n^{-1}\sum_{i=1}^{n_1} x_i + n^{-1}\sum_{i=n_1+1}^{n} x_i\right]$$
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- say we choose $n_1/n = p$ and $n_2/n = 1 p$
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$$V[\overline{x}^{STRAT}] = n^{-1} \{ \rho \sigma_1^2 + (1 - \rho) \sigma_2^2 \} < n^{-1} V[X]$$

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- ullet suppose we have several clusters, indexed by c
- the distribution of the variable of interest, X, obeys the following:

$$x_{ic} = \mu + \alpha_c + \varepsilon_{ic} \tag{4}$$

- ullet here lpha and arepsilon are independent of each other, and both have mean zero
- let σ_{α}^{2} be the variance of the cluster-specific mean
- ullet $\sigma_arepsilon^2$ be the variance of the idiosyncratic error term $arepsilon_{lc}$
- ullet the mean of sample from k clusters, with m units per cluster, has variance

$$V[\overline{x}^{CLUST}] = \frac{\sigma^2}{km} \{ (m-1)\rho + 1 \}$$
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- here $\sigma^2 = \sigma_{\alpha}^2 + \sigma_{\varepsilon}^2$ is the overall variance of X
- $\rho = \sigma_{\alpha}^2/\sigma^2$ is the intercluster correlation coefficient
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- the distribution of the variable of interest, X, obeys the following:

$$x_{ic} = \mu + \alpha_c + \varepsilon_{ic} \tag{4}$$

- here α and ε are independent of each other, and both have mean zero
- let σ_{lpha}^2 be the variance of the cluster-specific mean
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- a hypothesis is a subset of the parameter space
 - if this is a single point, we say the hypothesis is simple
 - e.g. $H_0: \mu = 1$
 - otherwise, we say the hypothesis is complex or compound
 - e.g. $H_1: \mu \neq 1$
- we often designate one particular hypothesis as the "null hypothesis"
 - then, see if the data provides strong enough evidence against it
- the frequentist approach to hypothesis testing takes parameters as fixed and the data as random
 - thus $P(\mu=1|X)$ makes no sense, but $P(X|\mu=1)$ does
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- to perform a *test* of the null hypothesis we form a *test statistic*, say $\widehat{S}(X_1, \dots X_n)$
 - not always the estimator itself, e.g. the t-test
 - then, we need to compute (at least approximately) the distribution of \widehat{S} under the null
- suppose we know the distribution of \widehat{S} under H_0
 - now, we can set a *rejection region* R such that $P(S \in R|H_0)$ is "small"
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$$\alpha = \max_{\theta \in \mathcal{H}_0} P(\text{reject } H_0 | \theta)$$

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consider some extreme cases:

- could ignore the data and never reject: then you never make Type I errors
- similarly could always reject: never make Type II errors
- in general there is a tradeoff between size and power
- how well you can do, and how severe the tradeoff is, depends on the problem
 - in some "ill-posed" problems, you cannot beat the trivial test
 - i.e. ignore the data, generate a random number $U \sim U(0,1)$ and reject if $U < \alpha$
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References

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