EKT-816 Lecture 2

Probability Review (2)

Jesse Naidoo

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- consistency: $\widehat{\theta} \longrightarrow_{p} \theta_{0}$
- if $E[\widehat{\theta}] = \theta_0$ we say $\widehat{\theta}$ is unbiased
- \bullet $\it{efficiency}$ or $\it{precision}$: say we have two estimators $\widehat{\theta}$ and $\widetilde{\theta}$
 - for now assume both are unbiased
 - ullet 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 bettered 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
- 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
 - ullet the sample maximum is, in this case, sufficient for $heta_0$
 - in fact, can show the MLE is $\theta = \max\{X_1, \dots X_n\}$
- another estimator would be $\hat{\theta} = 2\overline{X}_n$
 - 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 $X_i \leq x$
 - by independence, the CDF of the sample maximum is the product of the individual CDFs

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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 (whv?)
- 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
 - a fraction p lives in city 1
 - the mean and variance of X (say, income) are μ_1 and σ_1^2 in city 1
 - ullet let $\mu= { extstyle p} \mu_1 + (1-{ extstyle p}) \mu_2$ be the national average
- the variance of a randomly sampled unit is

$$V[X_i] = p\sigma_1^2 + (1-p)\sigma_2^2 + p(\mu_1 - \mu)^2 + (1-p)(\mu_2 - \mu)^2$$

- notice that V[X] = E[V[X|S]] + V[E[X|S]]
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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 $n_1 + n_2 = n_1$
 - the variance of the sample mean is

$$V[\overline{x}^{STRAT}] = V\left[n^{-1}\sum_{i=1}^{n_1} x_i + n^{-1}\sum_{i=n_1+1}^{n} x_i\right]$$
$$= \left(\frac{n_1}{n}\right)^2 \frac{\sigma_1^2}{n_1} + \left(\frac{n_2}{n}\right)^2 \frac{\sigma_2^2}{n_2}$$
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- say we choose $n_1/n = p$ and $n_2/n = 1 p$
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$$V[\overline{x}^{STRAT}] = n^{-1} \{ p\sigma_1^2 + (1-p)\sigma_2^2 \} < n^{-1}V[X]$$

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- 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 zeroca
- ullet let σ_lpha^2 be the variance of the cluster-specific mean
- ullet $\sigma_arepsilon^{\epsilon}$ be the variance of the idiosyncratic error term $arepsilon_{ic}$
- 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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- ullet here $\sigma^2 = \sigma_lpha^2 + \sigma_arepsilon^2$ is the overall variance of X
- $m{\phi}=\sigma_{lpha}^{2}/\sigma^{2}$ is the intercluster correlation coefficient
- positive correlations within a cluster reduce the precision of your estimates
- effectively less information from each observation from the same cluster
- in limiting case $\rho = 1$, the effective sample size is k, not km

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

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

- here α and ε are independent of each other, and both have mean zero
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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-tes
 - then, we need to compute (at least approximately) the distribution of $\widehat{\mathcal{S}}$ 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"
 - given a rejection region R,

$$\alpha = \max_{\theta \in H_0} P(\text{reject } H_0 | \theta)$$

- is called the size of the test, or the significance level
- we take the maximum over H₀ (in case H₀ is a compound hypothesis) to be conservative
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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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Chetty, Raj. 2009. "Sufficient Statistics for Welfare Analysis: A Bridge Between Structural and Reduced-Form Methods." *Annual Review of Economics* 1 (1): 451–88. doi:10.1146/annurev.economics.050708.142910. Weyl, E. Glen. 2019. "Price Theory." *Journal of Economic Literature (Forthcoming)*, no. August.

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