

# EKT-816 Lecture 2

Probability Review (2)

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# Desirable Properties of Estimators

- *consistency*:  $\hat{\theta} \rightarrow_p \theta_0$
- if  $E[\hat{\theta}] = \theta_0$  we say  $\hat{\theta}$  is *unbiased*
- *efficiency* or *precision*: say we have two estimators  $\hat{\theta}$  and  $\tilde{\theta}$ 
  - for now assume both are unbiased
  - if  $V[\hat{\theta}] \leq V[\tilde{\theta}]$ , say that  $\hat{\theta}$  is *more efficient* than  $\tilde{\theta}$
- the *mean square error* of  $\hat{\theta}$  is  $\text{MSE}(\hat{\theta}) = E[(\hat{\theta} - \theta_0)^2]$ 
  - easy to see that  $\text{MSE} = V[\hat{\theta}] + \text{bias}^2$
  - 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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# Sufficient Statistics

- suppose there is a statistic  $T(X_1, \dots, X_n)$  such that the joint density factors as

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

- 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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## Example: Sufficiency and Comparison of Estimators

- 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  $\theta_0$
  - in fact, can show the MLE is  $\hat{\theta} = \max\{X_1, \dots, X_n\}$
- another estimator would be  $\tilde{\theta} = 2\bar{X}_n$ 
  - 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  $\max\{X_1, \dots, X_n\} \leq x$  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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# Sample Design

- 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 level



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  - cross-sectional:
    - ▶ units from the population are surveyed once, and all at roughly the same time.
    - ▶ a “snapshot” of the population
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  - differential attrition can be a serious problem
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# Weighting

- suppose we have a population of  $N$  units, and we sample with unequal probabilities
  - let  $\pi_i$  be the probability that unit  $i$  is selected
  - in a simple random sample of size  $n \ll N$ ,  $\pi_i \approx n/N$ , 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

$$\begin{aligned} E \left[ \sum_{i=1}^n w_i \right] &= E \left[ \sum_{i=1}^N t_i w_i \right] \\ &\approx \sum_{i=1}^N (\pi_i n) w_i = N \end{aligned} \tag{1}$$

- here  $t_i$  is the number of times unit  $i$  is included in the sample
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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

$$\bar{x}_w = \sum_{i=1}^n w_i x_i \quad (2)$$

- exercise: show that  $E[\bar{x}_w] = E[X]$
- we may want to deliberately oversample some groups to improve precision of conditional means
  - e.g. white or Indian South Africans
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# Stratification

- 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  $\mu_1$  and  $\sigma_1^2$  in city 1
  - let  $\mu = p\mu_1 + (1 - p)\mu_2$  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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# Stratification

- say we take subsamples of size  $n_1$  and  $n_2$  respectively, with  $n_1 + n_2 = n$ 
  - the variance of the sample mean is

$$\begin{aligned} V[\bar{x}^{STRAT}] &= V\left[n^{-1} \sum_{i=1}^{n_1} x_i + n^{-1} \sum_{i=n_1+1}^n x_i\right] \\ &= \left(\frac{n_1}{n}\right)^2 \frac{\sigma_1^2}{n_1} + \left(\frac{n_2}{n}\right)^2 \frac{\sigma_2^2}{n_2} \end{aligned} \quad (3)$$

- say we choose  $n_1/n = p$  and  $n_2/n = 1 - p$ 
  - then

$$V[\bar{x}^{STRAT}] = n^{-1} \{p\sigma_1^2 + (1-p)\sigma_2^2\} < n^{-1} V[X]$$

- as long as the means differ ( $\mu_1 \neq \mu_2$ ), stratification improves precision (why?)
- you can show that with several strata:
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- suppose we have several clusters, indexed by  $c$
- the distribution of the variable of interest,  $X$ , obeys the following:

$$x_{ic} = \mu + \alpha_c + \varepsilon_{ic} \quad (4)$$

- here  $\alpha$  and  $\varepsilon$  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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- suppose we know the distribution of  $\hat{S}$  under  $H_0$ 
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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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