AUA CS108, Statistics, Fall 2020 Lecture 31

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Contents

- ► MVUE
- ► Bias-Variance Decomposition of MSE

Recall again the B-V D:

$$\mathit{MSE}(\hat{\theta}, \theta) = \left(\mathit{Bias}(\hat{\theta}, \theta)\right)^2 + \mathit{Var}_{\theta}(\hat{\theta}).$$

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Well, in general, there will be a lot of Unbiased Estimators for the same Parameter.Say, if $\hat{\theta}_0$ and $\hat{\theta}_1$ are Unbiased Estimators of θ , then for any $\alpha \in [0,1]$, the Estimator

$$\hat{\theta}_{\alpha} = \alpha \cdot \hat{\theta}_1 + (1 - \alpha) \cdot \hat{\theta}_0$$

will be an Unbiased Estimator too.

So the idea is to restrict our attention to only Unbiased Estimators. In that case, since $Bias(\hat{\theta},\theta)=0$,

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- \triangleright $\hat{\theta}$ is Unbiased Estimator for θ ;
- $\hat{\theta}$ has the smallest Variance among all *Unbiased* Estimators of θ , i.e., for any Unbiased Estimator $\tilde{\theta}$,

$$Var_{\theta}(\hat{\theta}) \leq Var_{\theta}(\tilde{\theta}), \quad \forall \theta \in \Theta.$$

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- **>** strongly consistent, if $\hat{\theta}_n \stackrel{a.s.}{\longrightarrow} \theta$ for any $\theta \in \Theta$;
- weakly or Mean Square consistent, if $\hat{\theta}_n \xrightarrow{q.m.} \theta$ for any $\theta \in \Theta$, i.e., if

$$MSE(\hat{\theta}_n, \theta) = \mathbb{E}_{\theta}((\hat{\theta}_n - \theta)^2) \to 0 \qquad \forall \theta \in \Theta.$$

Example: Consider a Random Sample

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Then:

- \triangleright \hat{p} is a Biased Estimator for p;
- \triangleright \hat{p} is Consistent Estimator for p.

Proof: OTB

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Proof: OTB. Use the relation
$$\widehat{\sigma^2} = \frac{\sum_{k=1}^{n} (X_k)^2}{n} - \left(\frac{\sum_{k=1}^{n} X_k}{n}\right)^2$$
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And also, the universal measure for goodness is: an Estimator is good if it has a small MSE.

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Answer: No, in general. This is because, say,

- we can do a lot of resamplings even when our dataset is not big enough, but one large sample will not be available
- when taking a large sample, we will take each individual just once. But if we are doing resamplings, we can have the same individual in different samples.