# Statistics 67 Introduction to Probability and Statistics for Computer Science

Lecture notes for Statistics

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# From Probability ....

- To this point
  - probability as a measure of uncertainty
  - probabilities for events
    - \* axioms, probability rules, conditional probability, Bayes' rule
  - random variables as quantities of interest in an uncertain environment
  - probability distributions as descriptions of possible values for a random variable along with an assessment of how likely each value is to occur
    - \* discrete/continuous distributions
    - \* univariate/multivariate distributions
    - \* joint, marginal, conditional distributions
    - \* expected values (mean, variance, covariance)
  - sampling/simulation as ways of studying a population or distribution

## .... to Statistical Inference

- Goal for remainder of quarter is to use what we know about probability to help us analyze data in scientific studies
  - use a sample from the population to learn about characteristics of the population
  - a common approach is to assume that observed sample are independent observations from a population model (e.g., Poisson or normal)
  - estimate the parameter(s) of the assumed model(e.g., normal mean or binomial proportion)
  - check fit of the assumed probability model
  - draw conclusions based on the estimated parameters (if appropriate)

- Importance of how data are obtained
  - we don't discuss in detail here how our data are collected
  - for statistical methods to be valid we need the sample to be representative of the population we are studying
  - typically this involves the use of randomness or chance in selecting the sample to avoid biased selections
  - a simple random sample is the most basic approach and that is what we assume
  - more sophisticated methods (multistage sampling, cluster sampling) can be accommodated

- Estimand the quantity being estimated
- We can think of two types of estimands
  - Finite population summaries
    - \* mean of a finite population
    - \* variance of a finite population
  - Parameters in a mathematical model of a population (can think of as an infinite population)
    - \*  $\mu$  or  $\sigma^2$  in a Normal distribution
    - \*  $\lambda$  (mean = variance) of Poisson distribution
    - \* p in a binomial distribution
- For the most part we focus on parameters in a mathematical model of a population

# • Basic Approach

– suppose  $\theta$  is a parameter that we are interested in learning about from a random sample

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

- e.g.,  $\theta$  might be the mean of the population that we are interested in  $(\mu_X)$
- $\hat{\theta}$ , a point estimator, is some function of the data that we expect will approximate the true value of  $\theta$
- e.g., we might use  $\hat{\mu} = \bar{X}$  to estimate the mean of a population  $(\mu_X)$
- once we collect data and plug in we have a point estimate  $\bar{x}$
- point estimator is the random variable (or function)
   and point estimate is the specific instance

## • Two key questions are

- 1. How do we find point estimators?
- 2. What makes a good estimator?

## Point Estimation - basics

- Assume we have a sample of independent random variables  $X_1, X_2, ..., X_n$ , each assumed to have density f(x)
- We call this a random sample (or iid sample) from f(x)
- Assume the density is one of the families we have considered which depends on one or more parameters  $\theta$ ; we usually write the density as  $f(x|\theta)$
- Goal is to estimate  $\theta$ . Why?
  - $-f(x|\theta)$  is a description of the population
  - $-\theta$  is often an important scientific quantity (e.g., the mean or variance of the population)

## Method of moments

- Recall  $E(X^j)$  is the jth moment of the population (or of the distribution); it is a function of  $\theta$
- The jth moment of the sample is  $\frac{1}{n} \sum_{i} X_{i}^{j}$
- We can equate the sample moment and the population moment to identify an estimator
- Suppose that there are k parameters of interest (usually k is just one or two)
- Set first k sample moments equal to first k population moments to identify estimators
- This is known as the **method of moments** approach

# Method of moments

- Example: Poisson case
  - suppose  $X_1, X_2, \ldots, X_n$  are a random sample from the Poisson distribution with parameter  $\lambda$
  - recall that  $E(X_i) = \lambda$
  - the method of moments estimator is obtained by taking the first sample moment  $(\bar{X} = \frac{1}{n} \sum_i X_i)$  equal to the first population moment  $\lambda$  to yield  $\hat{\lambda} = \bar{X}$
  - $Var(X_i)$  is also equal to  $\lambda$  so it would also be possible to take the sample variance as an estimate of  $\lambda$  (thus method of moments estimates are not unique)

Method of moments

- Example: Normal case
  - suppose  $X_1, X_2, \ldots, X_n$  are a random sample from the normal distribution with parameters  $\mu$  and  $\sigma^2$
  - recall that  $E(X_i) = \mu$  and  $E(X_i^2) = \sigma^2 + \mu^2$
  - to find method of moments estimators we need to solve

$$\frac{1}{n} \sum_{i=1}^{n} X_i = \mu$$

$$\frac{1}{n} \sum_{i=1}^{n} X_i^2 = \sigma^2 + \mu^2$$

- results:

$$\hat{\mu}_{mom} = \frac{1}{n} \sum_{i=1}^{n} X_i = \bar{X}$$

$$\hat{\sigma}_{mom}^2 = \frac{1}{n} \sum_{i=1}^{n} (X_i - \bar{X})^2$$

- Method of moments summary
  - easy to use
  - generally not the best estimators
  - some ambiguity about which moments to use

## Maximum likelihood estimation

- The density of a single observation is  $f(x|\theta)$
- The joint density of our random sample is  $f(X_1, X_2, ..., X_n | \theta) = \prod_{i=1}^n f(X_i | \theta)$  (recall the  $X_i$ 's are independent)
- This joint density measures how likely a particular sample is (assuming we know  $\theta$ )
- Idea: look at the joint distribution as a function of  $\theta$  and choose the value of  $\theta$  that makes the observed sample as likely as possible
- Likelihood function =  $L(\theta|x_1,\ldots,x_n) = f(x_1,\ldots,x_n|\theta)$
- Maximum likelihood estimator  $\hat{\theta}_{mle}$  is the value of  $\theta$  that maximizes the likelihood function

## Maximum likelihood estimation

- To find the MLE:
  - solve  $dL/d\theta = 0$  to identify stationary point
  - check that we have a maximum (can use the 2nd derivative)
  - it is often easier to maximize the logarithm of the likelihood
    (which is equivalent to maximizing the likelihood)
  - in complex models it can be hard to find the maximum

Maximum likelihood estimation

- Example: Poisson case
  - suppose  $X_1, X_2, \ldots, X_n$  are a random sample from the Poisson distribution with parameter  $\lambda$
  - the joint distribution is

$$f(X_1, \dots, X_n | \lambda) = \prod_{i=1}^n \frac{e^{-\lambda} \lambda^{X_i}}{X_i!}$$

- the likelihood function is

$$L = f(X_1, \dots, X_n | \lambda) = e^{-n\lambda} \lambda^{\sum_i X_i} / (\prod_i X_i!)$$

- then

$$LogL = \sum_{i} X_{i} \ln \lambda - n\lambda - \ln(\prod_{i} X_{i}!)$$

$$dLogL/d\lambda = \sum_{i} X_{i}/\lambda - n = 0$$

which implies that  $\hat{\lambda} = \bar{X}$  is the maximum likelihood estimator

- second derivative of log likelihood confirms this estimate attains a maximum of the likelihood
- maximum likelihood and method of moments give the same estimator here

Maximum likelihood estimation

- Example: normal case
  - suppose  $X_1, X_2, \ldots, X_n$  are a random sample from the Normal distribution with mean  $\mu$ , variance  $\sigma^2$
  - $LogL = constant \frac{n}{2}\log\sigma^2 \frac{1}{2\sigma^2}\sum_{i=1}^n (X_i \mu)^2$
  - need to solve

$$\partial Log L/\partial \mu = \frac{1}{\sigma^2} \sum_{i=1}^n (X_i - \mu) = 0$$

$$\partial Log L/\partial \sigma^2 = -\frac{n}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_{i=1}^n (X_i - \mu)^2 = 0$$

- results (same estimators as method of moments)

$$\hat{\mu}_{mle} = \frac{1}{n} \sum_{i} X_{i} = \bar{X}$$

$$\hat{\sigma}_{mle}^{2} = \frac{1}{n} \sum_{i} (X_{i} - \bar{X})^{2}$$

- Maximum likelihood summary
  - more complex than method of moments
  - statistical theory (not covered) suggests that maximum likelihood estimates do well (especially with lots of data)

# Properties of point estimators

- Now have two methods for finding point estimators
- What makes for a good estimator?
  - suppose  $T(X_1, \ldots, X_n)$  is an estimator of  $\theta$
  - traditional approach to statistics asks how well T would do in repeated samples
  - key to studying estimators is to note that T is itself a random variable and we can study properties of its distribution
  - examples of good properties include
    - \* lack of bias
    - \* low variance

# Properties of point estimators

- Unbiasedness
  - estimator T is unbiased for  $\theta$  if  $E(T) = \theta$
  - unbiased means estimator is "right on average"
  - no guarantee that the estimate in one sample is good but unbiasedness tells us the estimator does well on average
  - example: in the normal case

$$E(\bar{X}) = \frac{1}{n} \sum_{i} E(X_i) = \mu$$

so  $\bar{X}$  is an unbiased estimate for  $\mu$ 

- Variance (Var  $T = E(T E(T))^2$ )
  - suppose we have two unbiased estimators
  - we should prefer the one with low variance
  - but low variance by itself is of limited use for example  $\hat{\theta} = T(X_1, \dots, X_n) = 6$  (estimator always estimates 6 regardless of the data) has low variance but will be a poor estimate if  $\theta$  is far from 6

# Properties of point estimators

- Mean squared error
  - natural to ask how well T does at estimating  $\theta$
  - a difficulty is that we need to know  $\theta$  in order to evaluate this
  - MSE =  $E(T \theta)^2$  is one way to judge how well an estimator performs
  - MSE depends on  $\theta$  but we may find that one estimator is better than another for every possible value of  $\theta$
  - it turns out that  $MSE = bias^2 + variance$ (where  $bias = E(T) - \theta$ )
  - this yields ... a bias-variance tradeoff
  - consider the example of estimating the normal mean
    - \*  $X_1$  is an unbiased estimator but has a lot of variance
    - \*  $\bar{X}$  is an unbiased estimator but has less variance (dominates  $X_1$ )
    - \* T=6 (a crazy estimator that always answers 6!!) has zero variance but lots of bias for some values of  $\theta$

# Properties of point estimators

- Large sample properties
  - natural to ask how well T does in large samples
  - consistency estimate tends to the correct value in large samples
  - efficiency estimate has smallest possible variance of any estimator in large samples
  - turns out that maximum likelihood estimators have these good large sample properties

# Bayesian estimation

- There is one alternative approach to point estimation that we introduce
- It differs from everything else we've done in that it allows us to use information from other sources
- Related to Bayes Theorem so known as Bayesian estimation
- Motivation
  - suppose we want to predict tomorrows temperature
  - a natural estimate is average of recent days temperatures (this is like using  $\bar{X}$ )
  - we have other knowledge (typical Southern
     California weather at this time of year)
  - natural to wonder if an estimator that combines information from history with current data will do better

## Bayesian estimation

- Three components to Bayesian point estimation
  - 1. Prior distribution  $g(\theta)$  describing uncertainty about  $\theta$  before any data is examined
  - 2. Likelihood / data distribution  $f(X_1, ..., X_n | \theta)$  summarizing the information in the data about  $\theta$  (assuming we have the right distribution)
  - 3. Posterior distribution  $p(\theta \mid X_1, \dots, X_n)$  is obtained by using Bayes Theorem to combine the prior distribution and the likelihood as  $p(\theta \mid X_1, \dots, X_n) = \frac{f(X_1, \dots, X_n \mid \theta)g(\theta)}{f(X_1, \dots, X_n)}$ . This posterior distribution describes the uncertainty about  $\theta$  after combining the information in the prior distribution and in the data
  - \* A final step is to define an estimator that summarizes the posterior distribution – most common to use the mean of the posterior distribution of  $\theta$  as the estimator

# Bayesian estimation

- Bernoulli trials example
  - assume  $X_1, \ldots, X_n$  are indep Bernoulli trials with probability of success  $\pi$
  - prior distribution for  $\pi$  is a uniform distribution between 0 and 1 (completely unsure about  $\pi$ ) so that  $g(\pi) = 1$  for  $0 < \pi < 1$
  - likelihood is  $L = \pi^{\sum_i X_i} (1 \pi)^{n \sum_i X_i}$
  - turns out that the posterior distribution is a known continuous distribution (the Beta distribution with parameters  $\sum_{i} X_{i} + 1$  and  $n \sum_{i} X_{i} + 1$ )
  - posterior mean (Bayesian point estimator) is  $\hat{\pi} = \frac{\sum_{i} X_{i} + 1}{n+2}$
  - note that this is different than  $\hat{\pi} = \bar{X}$  which would be the maximum likelihood estimator or the method of moments estimator
  - an interesting case: consider X=0 for which maximum likelihood estimate is  $\hat{\pi}=0$  and for which Bayes estimate is  $\hat{\pi}=1/(n+2)$

- Point estimation is an important first step in a statistical problem
- A key contribution of the field of statistics though is to supplement the point estimate with a measure of accuracy (e.g., the standard deviation of the estimator is such a measure)
- A common way to convey the estimate and the accuracy is through an interval estimate
- In other words we create an interval (based on the sample) which is likely to contain the true but unknown parameter value
- This interval is usually called a confidence interval (CI)
- There are a number of ways to create confidence intervals, we focus on a simple approach appropriate for large samples to illustrate the approach

#### Central Limit Theorem

- A key mathematical result that enables interval estimation (and other forms of statistical inference) is the **central limit theorem** (CLT)
- **Theorem:** Let  $X_1, \ldots, X_n$  be a random sample of size n from a distribution with mean  $\mu$  and variance  $\sigma^2$ . Then for large n,  $\bar{X} = \frac{1}{n} \sum_i X_i$  is approximately normal with mean  $\mu$  and variance  $\sigma^2/n$ .
- Note this means  $(\bar{X} \mu)/(\sigma/\sqrt{n})$  is approximately standard normal
- How big does n have to be? It depends on the population distribution
  - if the population distribution is itself normal, then the CLT holds for small samples (even n = 1)
  - if the population distribution is not too unusual,
     then the CLT holds for samples of 30 or more
  - if the population distribution is unusual (e.g., very long tails), then the CLT may require 100 or more observations

# Central Limit Theorem - example

- Example: The number of files stored in the home directory has mean  $\mu = 7$  and standard deviation  $\sigma = 5$ . (Note that this variable can not have a normal distribution because: (1) it is a discrete random variable; and (2) with that mean and s.d. the normal distribution would have substantial probability below zero.) What is the probability that a class of 50 students will store more than 400 files?
- First, we should note that the question about the total number of files is equivalent to asking for the probability that  $\bar{X}$  will be greater than 8.
- Then by CLT  $\bar{X}$  is approximately normal with mean 7 and s.d.  $5/\sqrt{50} = .707$
- Finally  $P(\bar{X} > 8) = P(\frac{\bar{X} \mu}{\sigma / \sqrt{n}} > \frac{8 7}{.707}) = P(Z > 1.41) = .0793$  (where Z is standard normal random variable)

# Central Limit Theorem - binomial proportion

- You may recall that we saw a result something like the CLT in talking about the normal approximation to the binomial distribution if np > 5 and n(1-p) > 5 then  $X \sim \text{Bin}(n,p)$  can be approximated by a normal random variable Y having mean np and variance np(1-p)
- This is equivalent to the CLT if we look at the proportion of successes X/n rather than the count of successes X
- To be specific, let  $W_1, \ldots, W_n$  be a random sample of Bernoulli trials (0/1 random variables) with probability of success p (hence mean is p and variance is p(1-p)) and let  $X = \sum_i W_i$  be the total number of successes in n trials. Then by the CLT  $\bar{W} = X/n$  is approximately normal with mean p and variance p(1-p)/n

# Central Limit Theorem - binomial proportion

- Example: Consider sampling light bulbs from a company which claims to produce only 2% defective light bulbs. What is the probability that a sample of 500 light bulbs would yield a defective proportion below 1%?
  - Let  $\overline{W}$  equal proportion of defectives in a sample of 500 light bulbs from a population with 2% defectives
  - By CLT  $\overline{W}$  is approximately normal (note that np = 10 and n(1-p) = 490) with mean .02 and variance (.02)(.98)/500 = .0000392
  - $-P(\bar{W} < .01) = P(\frac{\bar{W} p}{\sqrt{p(1-p)/n}} < \frac{.01 .02}{\sqrt{.0000392}})$ = P(Z < -1.60) = .0548

## Population mean

- Central Limit Theorem enables us to easily build a confidence interval for the mean of a population
- Assume  $X_1, \ldots, X_n$  are independent random variables with mean  $\mu$  and variance  $\sigma^2$
- Then  $\bar{X} = \frac{1}{n} \sum_{i} X_{i}$  (the sample mean) is the natural estimate of  $\mu$  (MLE, method of moments)
- We also know that  $\bar{X}$  is a random variable which has approximately a normal distribution,  $\bar{X} \sim N(\mu, \sigma^2/n)$
- It follows that  $\Pr(-1.96 < \frac{\bar{X} \mu}{\sigma/\sqrt{n}} < 1.96) \approx .95$
- Thus  $\bar{X} \pm 1.96\sigma/\sqrt{n}$  is an (approximate) 95% confidence interval for  $\mu$
- Note the above is an exact confidence interval if the population distribution of the X's is normal and an approximate confidence interval valid for large n if not

Population mean (cont'd)

- $\bar{X} \pm 1.96\sigma/\sqrt{n}$  is an (approximate) 95% confidence interval for  $\mu$  (based on CLT)
- Some variations/improvements
  - Different confidence level
    - \* We can get a different confidence level by using a suitable percentile of the standard normal distribution
    - \* e.g.,  $\bar{X} \pm 1.645\sigma/\sqrt{n}$  is an (approximate) 90% confidence interval for  $\mu$
  - Unknown population standard deviation
    - \* Results given so far require knowing the population standard deviation  $\sigma$
    - \* If  $\sigma$  is not known (it usually isn't) then we can use the sample standard deviation

$$s = \sqrt{\frac{1}{n-1} \sum_{i} (X_i - \bar{X})^2}$$
 as an estimate

- \* Then  $\bar{X} \pm 1.96s/\sqrt{n}$  is an approximate 95% confidence interval that should be good in large samples (now even larger than before .. say 100 observations or more)
- \* It turns out that it is possible to create a more exact 95% confidence interval in this case by replacing 1.96 with the relevant percentile of Student's t-distribution (not covered in this class)

# Binomial proportion

- Assume  $X_1, \ldots, X_n$  are independent Bernoulli trials with probability of success  $\pi$  (change from p now)
- Then  $\hat{\pi} = \frac{1}{n} \sum_{i} X_{i}$  = the sample proportion of successes is the natural estimate (MLE, method of moments)
- From central limit theorem we know that  $\hat{\pi}$  is approximately normal with mean  $\pi$  and s.d.  $\sqrt{\pi(1-\pi)/n}$
- It follows that  $Pr(-1.96 < \frac{\hat{\pi} \pi}{\sqrt{\pi(1-\pi)/n}} < 1.96) = .95$
- Thus any  $\pi$  for which the inequality above is satisfied is in a 95% confidence interval
- An alternative is replace the s.d. of  $\hat{\pi}$  by an estimate,  $\sqrt{\hat{\pi}(1-\hat{\pi})/n}$  and then note that  $\hat{\pi}\pm 1.96\sqrt{\hat{\pi}(1-\hat{\pi})/n}$  is an approximate 95% confidence interval for  $\pi$

# • General approach

- the previous two examples suggest a general approach
- suppose that we have a point estimator  $\hat{\theta}$  for a parameter  $\theta$
- $\hat{\theta}$  is a random variable with expected value typically approximately equal to  $\theta$  and with a standard deviation  $s.d.(\hat{\theta})$
- it follows that an approximate large-sample 95% confidence interval for  $\theta$  is given by  $\hat{\theta} \pm 1.96 \ s.d.(\hat{\theta})$  (sometimes we may need to estimate the s.d.)

# • Interpretation

- it is important to remember the interpretation of these confidence intervals
- the "confidence" belongs to the procedure; we have a procedure that creates intervals having the property that 95% of the confidence intervals contain the true values
- for any given instance the CI either contains the true value or not; our guarantee is only for average behavior in repeated samples

# Tests/Decisions

- Point estimates and interval estimates are important components of statistical inference
- Sometimes there is a desire however for a formal test or decision based on the value of a particular parameter
- For example:
  - We may want to assess whether  $\pi = 0.5$  in a binomial situation (or in other words we may want to ask if we have a fair coin)?
  - We may want to test whether  $\mu = 0$  (no change due to an intervention)?
  - We may want to compare average response in two groups to see if they are equal  $(\mu_1 = \mu_2)$ ?

## Statistical Tests - binomial case

- We illustrate the basic approach in the binomial setting
- Assume we sample n people at random from list of CS faculty in the U.S.
- Ask each whether their laptop runs Windows or Linux
- Observe 56% use Linux
- Can we conclude that a majority of CS faculty in the US prefer Linux for their laptop?
  - seems obvious that we can but ...
  - the difference between 56% and 50% may just be a fluke of the sample, the truth may be that the population is split 50/50

### Statistical Tests - binomial case

- The logic of statistical tests
  - let X denote the number of faculty preferring Linux
  - assume  $X \sim Bin(n, \pi)$ (note we use  $\pi$  instead of the usual p to avoid confusion later)
  - organize test in terms of null hypothesis (no effect, no difference) and alternative hypothesis (the difference we suspect may be present)
    - \* null  $H_o: \pi = 0.50$
    - \* alternative  $H_a: \pi > 0.50$
    - \* why use this formulation? easier to disprove things statistically than to prove them
  - we suspect  $H_o$  is false (and  $H_a$  is true) if  $X/n = \hat{\pi}$  is greater than 0.5. How much greater does it have to be?
  - approach: assume the null hypothesis is true and ask whether the observed data is as expected or is unusual

# Statistical Tests - general comments

- There are two slightly different (but related) approaches
  - significance tests assess the evidence against  $H_o$  with a p-value that measures how unusual the observed data are
  - **hypothesis tests** formally establish a decision rule for deciding between  $H_o$  and  $H_a$  to achieve desired goals (e.g., decide  $H_a$  is true if  $\hat{\pi} > c$  where cis chosen to control the probability of an error)
  - we focus on significance tests in this class

# Statistical Tests - general comments

- The key concept in significance tests is the *p*-value
- p-value = probability of observing data as or more extreme than the data we obtained if  $H_o$  is true
- Low *p*-values are evidence that either
  - (1)  $H_o$  is true and we saw an unusual event or
  - (2)  $H_o$  is not true
- The lower the p-value the more likely we are to conclude that  $H_o$  is not true
- Often use p < .05 as serious evidence against  $H_o$  but a strict cutoff is a BAD IDEA
- A couple of important points
  - the p-value DOES NOT measure the probability that  $H_o$  is true
  - even if p-value is small the observed failure of  $H_o$  may not be practically important

## Statistical Tests - binomial case

- Now return to binomial case and suppose that we have sampled 100 professors and find 56 use Linux, or in other words n = 100 and  $\hat{\pi} = .56$
- There are actually two ways to find the p-value: use the binomial distribution directly or, if n is large (as it is here) then we can use the CLT
- By the binomial distn ... let X be number of Linux supporters. Then under  $H_o$  we know  $X \sim \text{Bin}(100, .5)$  and  $P(X \geq 56) = .136$  (not in our table but can be computed)
- By the CLT ...

$$p\text{-value} = \Pr(\hat{\pi} \ge 0.56 \mid \pi = 0.5)$$

$$= \Pr(\frac{\hat{\pi} - 0.50}{\sqrt{.5(.5)/100}} \ge \frac{.56 - .50}{\sqrt{.5(.5)/100}})$$

$$\approx \Pr(Z \ge 1.2) = .115$$

(using the continuity correction we'd say  $p = P(\hat{\pi} \ge .555) = .136$ )

• Conclude: observed proportion .56 is higher than expected but could have happened by chance so can't conclude that there is a significant preference for Linux

#### Statistical Tests - binomial case

- Interpreting results
  - The p-value of .136 does not mean that  $H_o$  is true, it only means the current evidence is not strong enough to make us give it up
  - -p-value depends alot on sample size
    - ... with n = 200 and  $\hat{\pi} = .56$  we would have p = .045
    - ... with n=400 and  $\hat{\pi}=.56$  we would have p=.008

### Hypothesis Tests

- Significance tests focus on  $H_o$  and try to judge its appropriateness
- Hypothesis tests treat the two hypotheses more evenly and are thus used in more formal decision settings
  - hypothesis testing procedures trade off two types of errors
  - type I error = reject  $H_o$  if it is true
  - type II error = accept  $H_o$  if it is false
  - we can vary cutoff of test; if we increase cutoff to make it harder to reject  $H_o$  then we reduce type I errors but make more type II errors (and vice versa if we lower the cutoff)
- In practice hypothesis tests are very closely related to significance tests

### Relationship of tests to other procedures

- Tests and confidence intervals
  - confidence intervals provide a range of plausible values for a parameter
  - tests ask whether a specific parameter value seems plausible
  - these ideas are related ... suppose we have a 95% confidence interval for  $\pi$ 
    - \* if  $\pi=0.50$  is not in the confidence interval then our test will tend to reject the hypothesis that  $\pi=0.50$
- Tests and Bayesian inference
  - we have not emphasized the Bayesian approach to testing but there is one
  - to see how it might work, recall that the Bayesian approach yields a posterior distribution telling us, for example, the plausible values of  $\pi$  and how likely each is
  - the Bayesian posterior distribution can compute things like  $P(\pi > 0.5|\text{observed data})$  which seems to directly address what we want to know

# Decisions/Tests - general approach

- General setting: we have a hypothesis about a parameter  $\theta$ , say  $H_o: \theta = \theta_o$  (could be  $\pi$  in binomial or  $\mu$  in normal) and want to evaluate this null hypothesis against a suspected alternative  $H_a: \theta > \theta_o$
- A general approach:
  - obtain a suitable point estimate  $\hat{\theta}$  and use it to test the hypothesis (reject  $H_o$  if  $\hat{\theta}$  is far from  $\theta_o$ )
  - calculate *p*-value which is  $P(\hat{\theta} > \text{observed value})$ assuming  $H_o$  is true
  - this calculation requires distribution of  $\hat{\theta}$
  - distribution of  $\hat{\theta}$  will depend on specific example (e.g., binomial case above)
- Of course if alternative is  $\theta < \theta_o$  then p-value also uses "<"

## Decisions/Test - population mean

Example: Tests for  $\mu$  (the population mean)

- Natural estimate is  $\bar{X}$  (the sample mean)
- What do we know about the distribution of  $\bar{X}$  under  $H_o$ ?
  - If the population data are normal and  $\sigma$  is known, then  $\bar{X}$  is normal with mean  $\mu_o$  and s.d.  $\sigma/\sqrt{n}$
  - If the population data are normal and  $\sigma$  is not known, then  $\bar{X}$  is approximately normal with mean  $\mu_o$  and s.d.  $s/\sqrt{n}$  for large sample size
  - If sample size is large (no matter what the population data are), then  $\bar{X}$  is approximately normal with mean  $\mu_o$  and s.d.  $s/\sqrt{n}$
  - Only difference between the last two items is that might expect to need a "larger" sample size in the last case
- The above discussion leads to normal test of  $H_o: \mu = \mu_o$  with p-value  $= P(\bar{X} > \bar{x}) = P(Z > (\bar{x} \mu_o) / \frac{s}{\sqrt{n}})$  (with Z the usual standard normal distn)

## Decisions/Test - population mean

Example: Tests for  $\mu$  (the population mean) Some technical stuff (optional)

- When we don't know  $\sigma$  and plug in the estimate s, we should really adjust for this in our procedure
- It turns out that the proper adjustment (original discovered by a brewery worker!) is to use Student's *t*-distribution in place of the standard normal distribution
- Student's t-distribution is a distribution that looks something like the normal but has heavier tails (bigger values are possible). The t distribution is described by the number of degrees of freedom (how big a sample it is based on) with a large degrees of freedom corresponding more closely to a normal distribution
- Student's t-test of  $H_o: \mu = \mu_o$  would lead to p-value =  $P(\bar{X} > \bar{x}) = P(Z > (\bar{x} \mu_o) / \frac{s}{\sqrt{n}})$  where  $t_{n-1}$  is a random variable having Student's t-distribution with n-1 degrees of freedom
- For Stat 67 purposes ... just need to know that in large samples can use normal table and not worry about the Student's t-distribution

## Decisions/Test – population mean

### • Numerical example:

Suppose that the average database query response time is supposed to be 1 second or faster. We try 100 queries and observe an average response time of 1.05 seconds (with a standard deviation of .25 seconds). Can we conclude that the database does not meet its standard?

- frame question as a statistical test:

$$H_o: \mu = 1 \text{ vs } H_a: \mu > 1$$

- p-value

$$= P(Z \ge (1.05 - 1.00) / \frac{.25}{\sqrt{100}}) = P(Z \ge 2) = .023$$
 (if we use Student's t-test, then *p*-value = .024)

- reject  $H_o$  and conclude that the database is not performing as advertised
- note that the additional .05 seconds may not be practically important

## Decisions/Test - difference between two means

Note to self: If there is time, then do this slide and the next to show how testing handles harder problems

- A common situation is that we have two populations and we want to compare the means of the two populations
- Example (medical): suppose we have two treatments (drug A and drug B) and wish to compare average survival time of cancer patients given drug A  $(\mu_1)$  to average survival time of cancer patients given drug B  $(\mu_2)$
- Assuming we have data on the two populations
  - $-\bar{Y}_1 \bar{Y}_2$  is an estimator for  $\mu_1 \mu_2$
  - $-\bar{y}_1 \bar{y}_2$  is an estimate for  $\mu_1 \mu_2$
  - $\operatorname{Var}(\bar{Y}_1 \bar{Y}_2) = \sigma^2 \left( \frac{1}{n_1} + \frac{1}{n_2} \right)$
  - $S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2}{n_1+n_2-2}$  is a pooled estimator for common variance  $\sigma^2$
- Key result: under assumptions

$$t = \frac{\bar{Y}_1 - \bar{Y}_2 - (\mu_1 - \mu_2)}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}} \sim t_{n_1 + n_2 - 2}$$

• Again for Stat 67 don't worry about Student's t (for large samples can use normal distribution)

# Decisions/Test - difference between two means

- Confidence interval
  - 95% confidence interval for  $\mu_1 \mu_2$  assuming large samples is

$$\bar{Y}_1 - \bar{Y}_2 \pm 1.96 S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$

- Tests of hypotheses
  - null hypothesis  $H_o: \mu_1 = \mu_2$  (no difference)
  - alternative hypothesis  $H_a: \mu_1 \neq \mu_2$  (two-sided) or  $\mu_1 > \mu_2$  or  $\mu_1 < \mu_2$  (one-sided)
  - test statistic  $t = (\bar{Y}_1 \bar{Y}_2) / \left( S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}} \right)$
  - p-value = probability of obtaining a value of the test statistic as big or bigger than the observed value if  $H_o$  is true (need to use t-distribution or normal table if samples are large to find p-value)

### Probability and Statistical Modeling

#### • So far:

- Estimation
  - \* sample from population with assumed distribution
  - \* inference for mean or variance or other parameter
  - \* point or interval estimates
- Decisions / Tests
  - \* judge whether data are consistent with assumed population
  - \* judge whether two populations have equal means
- To apply statistical thinking in more complex settings (e.g., machine learning)
  - \* build a probability model relating observable data to underlying model parameters
  - \* use statistical methods to estimate parameters and judge fit of model

#### Introduction

- We use linear regression as a (relatively simple) example of statistical modeling
- Linear regression refers to a particular approach for studying the relationship of two or more quantitative variables variables
- Examples:
  - predict salary from education, years of experience, age
  - find effect of lead exposure on school performance
- Useful to distinguish between a functional or mathematical model

$$Y = g(X)$$
 (deterministic)  
and a structural or statistical model

$$Y = g(X) + \text{error}$$
 (stochastic)

Linear regression model

• The basic linear regression model

$$Y_i = \beta_o + \beta_1 x_i + \epsilon_i, \quad i = 1, \dots, n$$

- $-Y_i$  is the response or dependent variable
- $-x_i$  is the predictor, explanatory variable, independent variable
- $x_i$  is treated as a fixed quantity (i.e., is not a random variable)
- $-\epsilon_i$  is the error term or individual variation
- $-\epsilon_i$  are independent  $N(0,\sigma^2)$  random variables
- Key assumptions
  - linear relationship between Y and x
  - independent (uncorrelated) errors
  - constant variance errors
  - normally distributed errors

Interpreting the model

• Model can also be written as

$$Y_i \mid X_i = x_i \sim N(\beta_o + \beta_1 x_i, \sigma^2)$$

- mean of Y given X = x is  $\beta_o + \beta_1 x$  (known as the conditional mean)
- $-\beta_o$  is conditional mean when x=0
- $-\beta_1$  is the slope, measuring the change in the mean of Y for a 1 unit change in x
- $-\sigma^2$  measures variation of responses about the mean

Where does this model come from?

- This model may be plausible based on a physical or other argument
- The model may just be a convenient approximation
- One special case is worth mentioning:

  It turns out that if we believe that two random variables X and Y have a bivariate normal distribution (remember we saw this briefly), then the conditional distribution of Y given X is in fact a normal model with mean equal to a linear function of X and constant variance

### Estimation

- Maximum likelihood estimation
  - we can write down joint distn of all of the Y's, known as the likelihood function

$$L(\beta_o, \beta_1, \sigma^2 \mid Y_1, \dots, Y_n) = \prod_{i=1}^n N(Y_i \mid \beta_o + \beta_1 x_i, \sigma^2)$$

- we maximize this to get estimates  $\hat{\beta}_o, \hat{\beta}_1$
- turns out to be equivalent to ....
- Least squares estimation
  - choose  $\hat{\beta}_o$ ,  $\hat{\beta}_1$  to minimize  $g(\beta_o, \beta_1) = \sum_{i=1}^n (Y_i (\beta_o + \beta_1 x_i))^2$
  - least squares has a long history (even without assuming a normal distribution)
    - \* why squared errors? (convenient math)
    - \* why vertical distances? (Y is response)
  - result:

$$\hat{\beta}_{o} = \bar{Y} - \hat{\beta}_{1}\bar{x}$$

$$\hat{\beta}_{1} = \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})(Y_{i} - \bar{Y})}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}}$$

- predicted (or fitted) value for a case with  $X = x_i$  is  $\hat{Y}_i = \hat{\beta}_o + \hat{\beta}_1 x_i$
- residual (or error) is  $e_i = Y_i \hat{Y}_i$

Estimation - some details

• Least squares estimation: choose  $\hat{\beta}_o, \hat{\beta}_1$  to minimize

$$g(\beta_o, \beta_1) = \sum_{i=1}^{n} (Y_i - (\beta_o + \beta_1 x_i))^2$$

• Taking derivatives and setting them equal to zero yields normal equations

$$\beta_o n + \beta_1 \sum x_i = \sum Y_i$$

$$\beta_o \sum x_i + \beta_1 \sum x_i^2 = \sum x_i Y_i$$

• Solving these equations leads to answers on previous slide

Estimation of error variance

- Maximum likelihood estimate of  $\sigma^2$  is  $\frac{1}{n} \sum_i (Y_i \hat{Y}_i)^2 = \frac{1}{n} \sum_i e_i^2$
- It turns out that this estimate is generally too small
- A common estimate of  $\sigma^2$  is

$$s_e^2 = \frac{1}{n-2} \sum_{i=1}^n (Y_i - (\hat{\beta}_o + \hat{\beta}_1 x_i))^2 = \frac{1}{n-2} \sum_{i=1}^n e_i^2$$

which is used because the  $\frac{1}{n-2}$  makes this an unbiased estimate

Inference for  $\beta_1$ 

- There are many quantities of interest in a regression analysis
- We may be interested in learning about
  - the slope  $\beta_1$
  - the intercept  $\beta_o$
  - a particular fitted value  $\beta_o + \beta_1 x$
  - a prediction for an individual
- Time is limited so we discuss only drawing statistical conclusions about the slope

Inference for the slope,  $\beta_1$ 

• Begin by noting that our estimator of the slope is

$$\hat{\beta}_1 = \frac{\sum_{i=1}^n (x_i - \bar{x})(Y_i - \bar{Y})}{\sum_{i=1}^n (x_i - \bar{x})^2} = \frac{\sum_{i=1}^n (x_i - \bar{x})Y_i}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

•  $\hat{\beta}_1$  is a linear combination of normal random variables (the  $Y_i$ 's) so  $\hat{\beta}_1$  is normally distributed

$$E(\hat{\beta}_1) = \beta_1$$
  $Var(\hat{\beta}_1) = \frac{\sigma^2}{\sum_i (x_i - \bar{x})^2}$ 

- $\sigma^2$  is unknown; plug in estimate  $s_e^2$
- The estimated standard deviation of  $\hat{\beta}_1$  is then  $s_{\beta_1} = \sqrt{s_e^2/\sum_i (x_i \bar{x})^2}$
- Then for a large sample size we get an approximate 95% confidence interval for  $\beta_1$  is  $\hat{\beta}_1 \pm 1.96s_{\beta_1}$
- More exact confidence interval and test procedures (based on Student's t-distribution) are available but not discussed in this class

Model diagnostics - residuals

- We can check whether the linear regression model is a sensible model using the residuals
- Recall  $e_i = Y_i \hat{Y}_i = Y_i \hat{\beta}_o \hat{\beta}_1 x_i$
- $e_i$  is an approximation of the stochastic error  $(\epsilon_i)$  in our model
- Important properties
  - sum of residuals is zero hence a typical value is zero
  - variance of the residuals is approximately equal to one
  - if our model is correct the residuals should look something like a N(0,1) sample
  - we can look to see if there are patterns in the residuals that argue against our model

Diagnosing violations with residual plots

- Plot residuals versus predicted values and look for patterns
  - might detect nonconstant variance
  - might detect nonlinearity
  - might detect outliers
- Histogram or other display of residuals
  - might detect nonnormality
- Show sample pictures in class

Remedies for violated assumptions

- What if we find a problem?
- Sometimes the linear regression model will work with a "fix"
  - transform Y (use  $\log Y$  or  $\sqrt{Y}$  as the response)
  - add or modify predictors (perhaps add  $X^2$  to approximate a quadratic relationship)
- If no easy "fix", then we can consider more sophisticated models