## **Review of Week 5 Biostatistics**

For this handout, we consider a random sample of independent and identically distributed observations  $(X_1, X_2, ..., X_n)$ . The true mean of  $X_i$  in the population is  $\mu$  and true variance is  $\sigma^2$ . The sample mean is  $\bar{x} = (1/n) \sum_{i=1}^n X_i$ . The sample variance is  $s^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{x})^2$ ; and the sample standard deviation is  $s = \sqrt{s^2}$ .

- Sampling Distributions. In statistics, when we sample from a population, the individual observations in our sample are considered random variables. The mean of the observed observations  $\bar{x}$ , called the sample mean, is also a random variable. To understand this concept, think about the fact that we have only observed *one sample* out of the many different random samples that we could have observed.
- Central Limit Theorem. The CLT tells us about the behavior of the sample mean  $\bar{x}$  in large sample sizes. If we take a random sample of size n,  $(X_1, X_2, ..., X_n)$ , from a population with true mean  $\mu$  and standard deviation  $\sigma$ , the CLT says that, for large n,

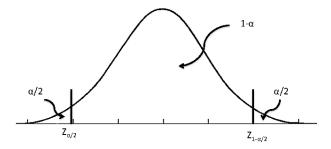
$$Z = \frac{\bar{x} - \mu}{\sigma / \sqrt{n}} \sim N(0, 1).$$

- Using the Central Limit Theorem, we can assume that, in large samples,  $\bar{x} \sim N(\mu, \sigma/\sqrt{n})$ .
- When the CLT applies, we can calculate the probability that  $\bar{x}$  lies between a and b very easily (when  $\sigma$  is known), now that we know the sampling distribution of  $\bar{x}$ .

Specifically,  $P(a < \bar{x} < b) = P(\bar{x} < b) - P(\bar{x} < a) = P(Z < \frac{\bar{x} - b}{\sigma/\sqrt{n}}) - P(Z < \frac{\bar{x} - a}{\sigma/\sqrt{n}})$ . To calculate these numbers, we can simply apply the normal function in Stata.

- **Predictive interval.** Predictive intervals are used to make predictions about random samples from a population, assuming the population parameters are known. Assume  $X_i \sim N(\mu, \sigma^2)$ .
  - A 95% predictive interval for  $X_i$  is  $\mu \pm Z_{0.975}\sigma$ .
  - A 90% predictive interval for  $X_i$  is  $\mu \pm Z_{0.95}\sigma$ .
  - A 95% predictive interval for  $\bar{x}$  is  $\mu \pm Z_{0.975}\sigma/\sqrt{n}$ .
  - A 95% predictive interval for  $\bar{x}$  is  $\mu \pm Z_{0.95}\sigma/\sqrt{n}$ .
  - For the 95% predictive interval, to calculate  $Z_{0.975}$ , use di invnormal (0.975).
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- More generally, for a  $1-\alpha$  confidence interval, to calculate  $Z_{1-\alpha/2}$ , use di invnormal  $(1-\alpha/2)$ . See the picture below, depicting a normal curve.



- Confidence interval. Confidence intervals are used to make inference about the population mean  $\mu$  using a random sample from the population. Assume  $X_i \sim N(\mu, \sigma^2)$ , where  $\mu$  is unknown.
  - When  $\sigma$  is known, a 95% confidence interval for  $\mu$  is  $\bar{x} \pm Z_{0.975} \sigma / \sqrt{n}$ .
  - When  $\sigma$  is known, a 99% confidence interval for  $\mu$  is  $\bar{x} \pm Z_{0.995} \sigma / \sqrt{n}$ .
  - When  $\sigma$  is unknown, a 95% confidence interval for  $\mu$  is  $\bar{x} \pm t_{0.975,(n-1)} s/\sqrt{n}$ .
  - When  $\sigma$  is unknown, 99% confidence interval for  $\mu$  is  $\bar{x} \pm t_{0.995,(n-1)} s / \sqrt{n}$ .
  - For the 95% confidence interval, to calculate  $t_{0.975,(n-1)}$ , use di invttail(n-1, 0.025) when  $\sigma$  is unknown.
  - For the 95% confidence interval, to calculate  $t_{0.95,(n-1)}$ , use di invttail(n-1, 0.05) when  $\sigma$  is unknown.
  - More generally, for a  $1-\alpha$  confidence interval, to calculate  $t_{1-\alpha/2,(n-1)}$ , use di invttail(n-1,  $\alpha/2$ ) when  $\sigma$  is unknown.

## • Confidence versus Predicive Intervals.

- Predictive intervals: Use the population distribution (known) to make predictions about a sample from the population (unknown). (This is the basis of probability.)
- Confidence intervals: Use a random sample from the population (known) to make inference about the population distribution (unknown). (This is the basis of statistics!)
- Hypothesis Testing. Using the random sample, we want to test  $H_0: \mu = \mu_0$  versus  $H_A: \mu \neq \mu_0$ .
  - One-sample Z-test. For large samples (CLT) OR when  $X_i \sim N(\mu, \sigma^2)$  and  $\sigma$  is known, the test-statistic is:

$$Z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}}$$

Under  $H_0$ ,  $Z \sim N(0,1)$ .

- \* For the one-sided test with alternative hypothesis  $H_a: \mu > \mu_0$ , we can calculate a p-value using the formula  $p = P(Z > Z^*)$ .
- \* For the one-sided test with alternative hypothesis  $H_a: \mu < \mu_0$ , we can calculate a p-value using the formula  $p = P(Z \le Z^*)$ .
- \* For the two-sided test with alternative hypothesis  $H_a: \mu \neq \mu_0$ , we can calculate a p-value using the formula  $p=2*P(Z<-|Z^*|)$ .
- One-sample t-test. When  $X_i \sim N(\mu, \sigma^2)$  and  $\sigma$  is unknown, the test-statistic is:

$$t = \frac{\bar{x} - \mu_0}{s / \sqrt{n}}$$

Under  $H_0$ ,  $t \sim t_{n-1}$  (the test statistic follows a t-distribution with n-1 degrees of freedom.)