

# Sec. 4.6

Day 10 (2/14)

## 1 Comments about statistical tests

### 1.1 Two-sided tests

Let  $X_1, X_2, \dots, X_n$  be a random sample from a normal distribution with mean  $\mu$  and standard deviation  $\sigma = 5000$ . Suppose that we want to test

$$H_0 : \mu = 30,000 \text{ versus } H_1 : \mu \neq 30,000$$

It makes sense that we should reject  $H_0$  if the sample mean is too small or too large, i.e.

$$\text{Reject } H_0 \text{ in favor of } H_1 \text{ if } \bar{X} \leq h \text{ or } \bar{X} \geq k$$

The benchmarks  $h$  and  $k$  are chosen so that the test achieves the desired size

$$\alpha = P_{H_0}(\bar{X} \leq h \text{ or } \bar{X} \geq k) = P_{H_0}(\bar{X} \leq h) + P_{H_0}(\bar{X} \geq k)$$

Suppose we choose to allocate  $\alpha$  equally, then

$$P_{H_0}(\bar{X} \leq h) = \alpha/2 \text{ and } P_{H_0}(\bar{X} \geq k) = \alpha/2$$

Under  $H_0$ , the statistic  $(\bar{X} - 30000)/(5000/\sqrt{n})$  is  $N(0, 1)$ . Since

$$\alpha/2 = P\left(\frac{\bar{X} - 30000}{5000/\sqrt{n}} \leq -z_{\alpha/2}\right) = P(\bar{X} \leq 30000 - z_{\alpha/2}5000/\sqrt{n}) \quad (1)$$

then  $h = 30000 - z_{\alpha/2}5000/\sqrt{n}$ . Similarly,

$$\alpha/2 = P\left(\frac{\bar{X} - 30000}{5000/\sqrt{n}} \geq z_{\alpha/2}\right) = P(\bar{X} \geq 30000 + z_{\alpha/2}5000/\sqrt{n}) \quad (2)$$

so that  $k = 30000 + z_{\alpha/2}5000/\sqrt{n}$ . For illustration, suppose  $\alpha = .05$  and  $n = 25$ , then  $h = 30000 - 1.965000/\sqrt{25} = 28040$  and  $k = 30000 + 1.965000/\sqrt{25} = 31960$ . We can write equations (1) and (2) together as

$$\alpha = P\left(\left|\frac{\bar{X} - 30000}{5000/\sqrt{n}}\right| \geq z_{\alpha/2}\right)$$

which gives a more compact expression for the decision rule

Reject  $H_0$  in favor of  $H_1$  if  $|\bar{X} - 30000| \geq z_{\alpha/2}5000/\sqrt{n}$

More generally, let  $X_1, X_2, \dots, X_n$  be a random sample from a distribution  $f$  with mean  $\mu$  and standard deviation  $\sigma$ . A large-sample size- $\alpha$  test for

$$H_0 : \mu = \mu_0 \text{ versus } H_1 : \mu \neq \mu_0$$

is given by the approximate decision rule

$$\text{Reject } H_0 \text{ in favor of } H_1 \text{ if } \left| \frac{\bar{X} - \mu_0}{S/\sqrt{n}} \right| \geq z_{\alpha/2}$$

This is equation (4.6.4) in your textbook.

## 1.2 Power function of the two-sided test

Now, suppose that the true value of  $\mu$  is in  $H_1$ . Recall that the power of the test is the probability that the test rejects at that value of  $\mu$ . From the critical regions in equations (1) and (2), the power function is

$$\begin{aligned} \gamma(\mu) &= P_\mu \left( \bar{X} \leq \mu_0 - z_{\alpha/2}\sigma/\sqrt{n} \right) + P_\mu \left( \bar{X} \geq \mu_0 + z_{\alpha/2}\sigma/\sqrt{n} \right) \\ &= P_\mu \left( \frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \leq \frac{\mu_0 - \mu - z_{\alpha/2}\sigma/\sqrt{n}}{\sigma/\sqrt{n}} \right) + P_\mu \left( \frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \geq \frac{\mu_0 - \mu + z_{\alpha/2}\sigma/\sqrt{n}}{\sigma/\sqrt{n}} \right) \\ &= \Phi \left( \frac{\sqrt{n}(\mu_0 - \mu)}{\sigma} - z_{\alpha/2} \right) + 1 - \Phi \left( \frac{\sqrt{n}(\mu_0 - \mu)}{\sigma} + z_{\alpha/2} \right) \end{aligned}$$

where  $\Phi(\cdot)$  is the standard normal cdf.

## 1.3 Relationship between two-sided test and confidence interval

Example:

$$\begin{aligned} \text{Reject } H_0 : \mu = 30000 \text{ if and only if } & \left| \frac{\bar{X} - 30000}{\sigma/\sqrt{n}} \right| > z_{\alpha/2} \\ & \text{if and only if } |\bar{X} - 30000| > z_{\alpha/2}\sigma/\sqrt{n} \\ & \text{if and only if } \bar{X} \text{ is farther than } z_{\alpha/2}\sigma/\sqrt{n} \text{ from } 30000 \\ & \text{if and only if } \bar{X} \pm z_{\alpha/2}\sigma/\sqrt{n} \text{ does not contain } 30000 \end{aligned}$$

In general, suppose we test

$$H_0 : \theta = \theta_0 \text{ versus } H_1 : \theta \neq \theta_0$$

then using the size- $\alpha$  test

$$\text{Reject } H_0 \text{ if } \left| \frac{\hat{\theta} - \theta_0}{\text{SE}} \right| > c_\alpha$$

is equivalent to

Reject  $H_0$  if the  $(1-\alpha)100\%$  confidence interval  $\hat{\theta} \pm c_\alpha \cdot \text{SE}$  does not contain 0.

Table 1: Summary of tests

Parameter	Estimate	SE	$c_\alpha$	Comment
$\mu$	$\bar{X}$	$s/\sqrt{n}$	$z_{\alpha/2}$	Large sample
	$\bar{X}$	$s/\sqrt{n}$	$t_{\alpha/2, n-1}$	Exact but needs normal population
$p$	$\hat{p}$	$\sqrt{\frac{\hat{p}(1-\hat{p})}{n}}$	$z_{\alpha/2}$	Large sample
$\mu_1 - \mu_2$	$\bar{X}_1 - \bar{X}_2$	$\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}$	$z_{\alpha/2}$	Large sample
		$\sqrt{S_p^2 \left( \frac{1}{n_1} + \frac{1}{n_2} \right)}$	$t_{\alpha/2, n_1+n_2-2}$	Needs normal, $\sigma_1^2 = \sigma_2^2$
$p_1 - p_2$	$\hat{p}_1 - \hat{p}_2$	$\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}$	$t_{\alpha/2, n^*}$	Needs normal
		$\sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{n_1} + \frac{\hat{p}_2(1-\hat{p}_2)}{n_2}}$	$z_{\alpha/2}$	Large sample
		$\sqrt{\hat{p}(1-\hat{p}) \left( \frac{1}{n_1} + \frac{1}{n_2} \right)}$		For $H_0 : p_1 = p_2$ where $\hat{p} = \frac{X_1+X_2}{n_1+n_2}$

R simulation of Welch versus pooled-t

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n1<-35           # Sample size for x
n2<-25           # Sample size for y
sigma1<-50       # Population sd for x
sigma2<-50       # Population sd for y
mu1<-40          # Population mean for x
mu2<-0           # Population mean for y
nsim<-10000     # Number of trials
pval1<-numeric(nsim) # Storage for p-value of pooled-t
pval2<-numeric(nsim) # Storage for p-value of Welch-t
for(i in 1:nsim){
  xsim<-rnorm(n1,mu1,sigma1) # Generate x-data
  ysim<-rnorm(n2,mu2,sigma2) # Generate y-data
  pval1[i]<-t.test(xsim,ysim,alternative="two.sided",var.equal=TRUE)$p.value
  pval2[i]<-t.test(xsim,ysim,alternative="two.sided",var.equal=FALSE)$p.value
}
cbind(mean(pval1<.05),mean(pval2<.05))

```

```

# Write a function
welch_sim<-function(nsim=10000, n1=30, n2=30, mu1=0, mu2=0, sigma1=1,sigma2=1){
pval1<-numeric(nsim)      # Storage for p-value of pooled-t
pval2<-numeric(nsim)      # Storage for p-value of Welch-t
for(i in 1:nsim){
xsim<-rnorm(n1,mu1,sigma1) # Generate x-data
ysim<-rnorm(n2,mu2,sigma2) # Generate y-data
pval1[i]<-t.test(xsim,ysim,alternative="two.sided",var.equal=TRUE)$p.value
pval2[i]<-t.test(xsim,ysim,alternative="two.sided",var.equal=FALSE)$p.value
}
# End of for() loop
return(c(mean(pval1<.05),mean(pval2<.05)))
}
# End of function()

# Call the function
welch_sim(10000,35,25,mu1=0,0,50,50)
welch_sim(10000,35,25,mu1=10,0,50,50)
welch_sim(10000,35,25,mu1=20,0,50,50)
welch_sim(10000,35,25,mu1=30,0,50,50)
welch_sim(10000,35,25,mu1=40,0,50,50)
# Use unequal variance
welch_sim(10000,35,25,mu1=0,0,20,50)
welch_sim(10000,35,25,mu1=0,0,50,20)

# Use for() loop to automate
muvec<-c(0,10,20,30,40)
outmat<-matrix(rep(0,2*length(muvec)), ncol=2)
for(j in 1:length(muvec)){
outmat[j,]<-welch_sim(10000,35,25,mu1=muvec[j],0,20,50)
}
outmat

```