

## 2.3 Power

Consider a hypothesis test about the parameter  $\theta$ :

$$egin{aligned} H_0: oldsymbol{ heta} &= oldsymbol{ heta}_0 \ H_a: oldsymbol{ heta} &> oldsymbol{ heta}_0 \end{aligned}$$

We let  $\beta = P(\text{fail to reject } H_0 | H_0 \text{ is false}) = P(\text{Type II error})$ , then Power =  $P( ext{reject } H_0 | H_0 ext{ is false}) = 1 - \beta.$ 

Power depends on the distance between the hypothesized value of the parameter  $\theta_0$  and the actual value  $\theta_1$ , so we can write  $1 - \beta(\theta_1)$ .

Why is power important?

1. If we have multiple statistical testing method for the same hypothesis, choose fest w/ wost pover.

For a few simple cases, you can derive a closed form expression of power.

All other cases, use Monte Carlo methods to <u>estimate</u> pover. **Example 2.4** Consider a one-sample z-test. Sample  $X_1, \ldots, X_n \stackrel{iid}{\sim} N(\mu, \sigma^2)$ . moun  $H_0: \mu = \mu_0$  using statisfic  $Z^* = \frac{\overline{X} - \mu_0}{\sigma / n}$  we reject  $H_0: \overline{f} \quad Z^* > Z_{1-\alpha}^{L}$  where  $H_a: \mu > \mu_0$ If  $\mu_0 = 5$  (hypothesized value) but the true near is  $\mu_1 = 6$ .

$$I - \beta = \rho(rged \#_0 | \#_0 \#_0 \#_0).$$

$$= \rho(Z^* > Z_{1-\alpha} - \frac{(\mu_1 - \mu_0)}{\Theta^6 / J_{n_3}})$$

$$Smallert Z where you will reject$$

So power is a function of

such it 
$$\rightarrow 1$$
. Significance level: as  $\alpha(T)$ , power  $T$  [trade off that type I and type I error]  
scare  $\rightarrow 2$ . Effect size =  $M_1 - M_0$ : as effect size  $T$ , power  $T$   
[3.] Sample size:  $\alpha s = nT$ , power  $T$   
can't  $\rightarrow 4$ . Variance:  $\alpha s$  variance  $V$ , power  $T$  (no control over this),  
this one. Notes:  $T$  as power =  $1 - \beta = T$ ,  $\beta(type I error) = \alpha T$ . For  
fixed  $n, 6 \notin M_1 - M_0$ , only way to increase power  $\xi$  that is  
 $T$  is only way to simultaneously increase power  $\xi$  that is  
 $T$  is

H,

## **2.4 MC Estimator of** $1 - \beta$

Assume  $X_1, \ldots, X_n \sim F(\theta_0)$  (i.e., assume  $H_0$  is true).

Then, we have the following hypothesis test –

$$egin{aligned} H_0: heta &= heta_0 \ H_a: heta &> heta_0 \end{aligned}$$

and the statistics  $T^*$ , which is a test statistic computed from data. Then we reject  $H_0$  if  $T^* >$  the critical value from the distribution of the test statistic.

$$T^* > \text{the critical value from the distribution of the test statistic.}$$
This leads to the following algorithm to estimate the power of the test  $(1 - \beta)$ 
(1) solect model, set up hypothesis test.  $T^{(m-s)}$  and  $T^{(m-s)}$ ,  $T^{(m-s)$ 

sotup

## Your Turn

Consider data generated from the following mixture distribution:

$$f(x)=(1-\epsilon)f_1(x)+\epsilon f_2(x),\quad x\in\mathbb{R}$$

where  $f_1$  is the pdf of a N(0, 1) distribution,  $f_2$  is the pdf of a N(0, 100) distribution, and  $\epsilon \in [0, 1]$ .

```
r_noisy_normal <- function(n, epsilon) {
z <- rbinom(n, 1, 1 - epsilon)
z*rnorm(n, 0, 1) + (1 - z)*rnorm(n, 0, 10)
}
n <- 100
data.frame(e = 0, sample = r_noisy_normal(n, 0)) %>%
rbind(data.frame(e = 0.1, sample = r_noisy_normal(n, 0.1))) %>%
rbind(data.frame(e = 0.6, sample = r_noisy_normal(n, 0.6))) %>%
rbind(data.frame(e = 0.9, sample = r_noisy_normal(n, 0.9))) %>%
ggplot() +
geom_histogram(aes(sample)) +
facet_wrap(.~e, scales = "free")
```



We will compare the power of various tests of normality. Let  $F_X$  be the distribution of a i.e. Ho says X is normally distributed and Ha says it isn't random variable X. We will consider the following hypothesis test,

$$H_0:F_x\in N \qquad ext{vs.} \qquad H_a:F_x
ot\in N,$$

where N denotes the family of univariate Normal distributions.

Recall Pearson's moment coefficient of skewness (See Example 2.2). and Corresponding sterness test for Normality.

We will compare Monte Carlo estimates of power for different levels of contamination (  $0 \le \epsilon \le 1$ ). We will use  $\alpha = 0.1$ , n = 100, and m = 100.

```
Jb_{i} = \frac{\frac{1}{n} \sum_{i=1}^{n} (x_{i} - \widehat{x})^{2}}{(\frac{1}{n} \sum_{i=1}^{n} (x_{i} - \widehat{x})^{2})^{3/2}} \sim N\left(0, \frac{6(n-2)}{(n+1)(n+3)}\right).
 # skewness statistic function
 skew <- function(x) {</pre>
   xbar <- mean(x)</pre>
   num <- mean((x - xbar)^3)
    denom <- mean((x - xbar)^2)
    num/denom^1.5
 }
                                                         Standy Lest for Normality
Ho: JB, = 0
 # setup for MC
 alpha <- .1
 n < -100
                                                           Ha: JB 70
 m < -100
 epsilon <- seq(0, 1, length.out = 200)</pre>
 var sqrt b1 <- 6*(n - 2)/((n + 1)*(n + 3)) \# adjusted variance for
   skewness test
 crit val <- qnorm(1 - alpha/2, 0, sqrt(var sqrt b1)) #crit value for
   the test
 empirical pwr <- rep(NA, length(epsilon)) #storage</pre>
 # estimate power for each value of epsilon
 for(j in 1:length(epsilon)) {
    # perform MC to estimate empirical power
   ## Your turn
 }
                                                         SE(1-\hat{p}) = \int \hat{p}(1-\hat{p})/n
 ## store empirical se
 empirical_se <- "Your Turn: fill this in"</pre>
 ## plot results --
 ## x axis = epsilon values
 ## y axis = empirical power
 ## use lines + add band of estimate +/- se
We can detect contamination levels between . 03 and . 2 at power ≥ 0.8
     when n=100.
E is like effect size.
```

Compare the power with n = 100 to the power with n = 10. Make a plot to compare the two for many values of  $\epsilon$ .