

Hypothesis testing

Jiachen Feng

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Explanation for the conclusion

```
alpha_0.6_m <- function(m){  
  1-pbinom(m-1,100,0.6)  
}  
  
##ggplot()+  
  ##stat_function(fun = alpha_0.6_m)+  
  ##xlim(1,100)  
  
for (i in 1:100) {  
  m <- 1  
  if(alpha_0.6_m(m)>0.05){  
    m <- i+1  
  }  
  if(alpha_0.6_m(m)<0.05){  
    print(m)  
    break  
  }  
}
```

```
## [1] 69
```

```
alpha_0.8_m <- function(m){  
  1-pbinom(m-1,100,0.8)  
}  
  
##ggplot()+  
  ##stat_function(fun = alpha_0.8_m)+  
  ##xlim(1,100)  
  
for (i in 1:100) {  
  m <- 1  
  if(alpha_0.8_m(m)>0.95){  
    m <- i+1  
  }  
  if(alpha_0.8_m(m)<0.95){  
    print(m-1)  
    break  
  }  
}
```

```
## [1] 73
```

$\alpha(p) = \sum_{m \leq k \leq n} b(n, p, k)$ gives the probability of a type 1 error. First, we need to find the smallest value for m that thwarts a type 1 error. In this case, p equals to 0.6, because the null hypothesis is true. Then, we need to find the largest value for m that thwarts a type 2 error. In this case, p equals to 0.8 here (this value is chosen arbitrarily). We need to find the largest value of m which makes $\beta(p) < 0.05$.

Replication and Explanation for Figure 3.7

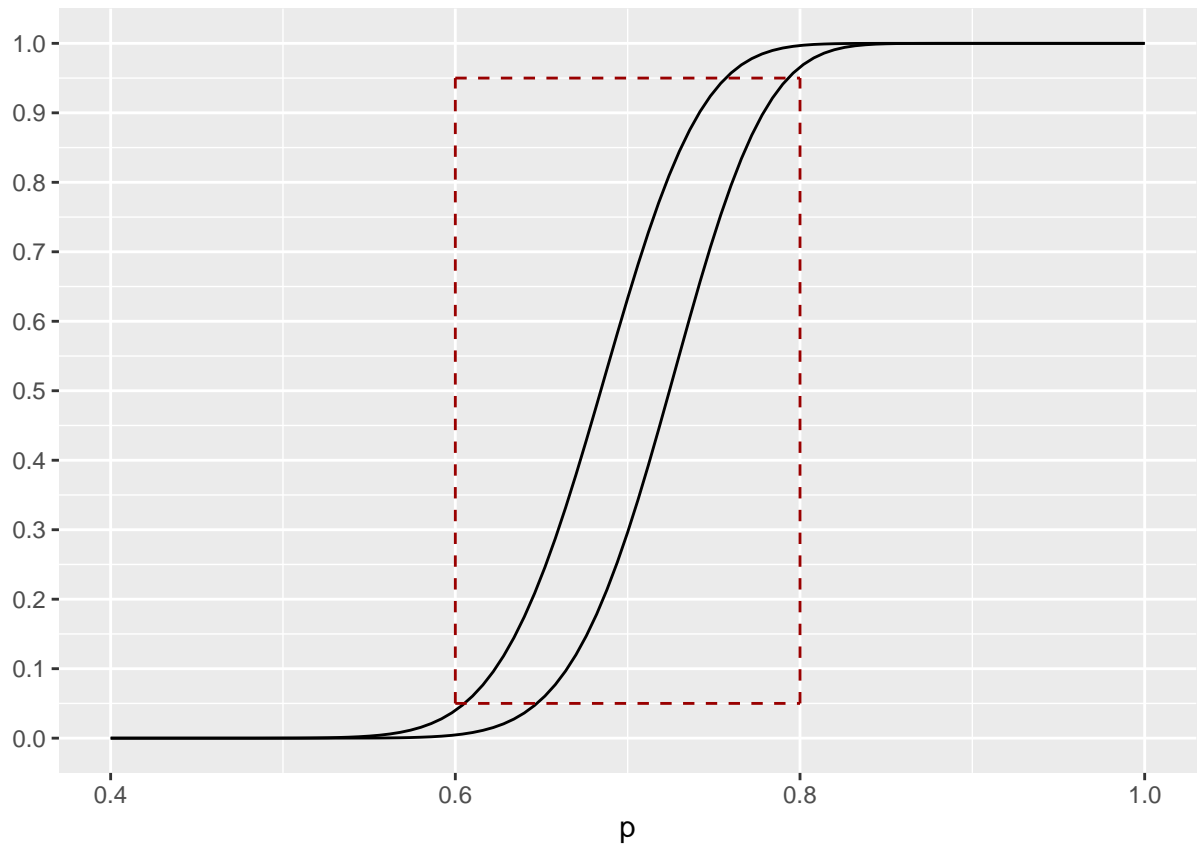
```
# Replicate

# define function for m=69
alpha_p_69 <- function(p){
  1-pbinom(68,100,p)
}

# define function for m=73
alpha_p_73 <- function(p){
  1-pbinom(72,100,p)
}

# plot
ggplot()+
  stat_function(fun = alpha_p_69)+
  xlim(.4,1)+
  stat_function(fun = alpha_p_73)+
  xlim(.4,1)+
  scale_y_continuous(breaks=seq(0, 1, 0.1))+
  geom_segment(aes(x=.6,xend=.8,y=.95,yend=.95),colour="#990000", linetype="dashed")+
  geom_segment(aes(x=.6,xend=.8,y=.05,yend=.05),colour="#990000", linetype="dashed")+
  geom_segment(aes(x=.6,xend=.6,y=.05,yend=.95),colour="#990000", linetype="dashed")+
  geom_segment(aes(x=.8,xend=.8,y=.05,yend=.95),colour="#990000", linetype="dashed")+
  xlab("p")+
  ylab(" ")

## Scale for 'x' is already present. Adding another scale for 'x', which will
## replace the existing scale.
```



As the figure declares, as m increases, the graph of α moves to the right. This means when m increases, it makes a type 1 error less likely.