

1. A group of researchers is running an experiment over the course of 30 months, with a single observation collected at the end of each month. Let X_1, \dots, X_{30} denote the observations for each month. From prior studies, the researchers know that

$$X_i \sim f_X(x),$$

but the mean μ_X is unknown, and they wish to conduct the following test

$$H_0 : \mu_X = 0$$

$$H_a : \mu_X > 0.$$

At month k , they have accumulated data X_1, \dots, X_k and they have the t -statistic

$$T_k = \frac{\bar{X} - 0}{S_k / \sqrt{n}}.$$

The initial plan was to test the hypotheses after all data was collected (at the end of month 30), at level $\alpha = 0.05$. However, conducting the experiment is expensive, so the researchers want to “peek” at the data at the end of month 20 to see if they can stop it early. That is, the researchers propose to check whether t_{20} provides statistically discernible support for the alternative. If it does, they will stop the experiment early and report support for the researcher’s alternative hypothesis. If it does not, they will continue to month 30 and test whether t_{30} provides statistically discernible support for the alternative.

- (a) What values of t_{20} provide statistically discernible support for the alternative hypothesis?

```
t.val = qt(.95, df = 19)
```

Values > 1.73

- (b) What values of t_{30} provide statistically discernible support for the alternative hypothesis?

```
t.val2 = qt(.95, df = 29)
```

Values > 1.70

- (c) Suppose $f_X(x)$ is a Laplace distribution with $a = 0$ and $b = 4.0$. Conduct a simulation study to assess the Type I error rate of this approach.

```
library(VGAM)

## Loading required package: stats4
## Loading required package: splines

#We know that the null distribution (Laplace) resembles the population distribution
type1.cnt = 0
for(i in 1:1000){
  dat = rlaplace(n = 30, scale = 4)
  p.20 = t.test(dat[1:20], alternative = "greater")
  p.30 = t.test(dat, alternative = "greater")
  if(p.20$statistic > t.val){ #If value is statistically discernible
    type1.cnt = type1.cnt + 1
  }
  else if(p.30$statistic > t.val2) #If value is statistically discernible
  {
    type1.cnt = type1.cnt + 1
  }
}
(proportion.type1 = type1.cnt/1000)

## [1] 0.077
```

Note: You can use the `rlaplace()` function from the `VGAM` package for R (Yee, 2010).

- (d) **Optional Challenge:** Can you find a value of $\alpha < 0.05$ that yields a Type I error rate of 0.05?
2. Perform a simulation study to assess the robustness of the T test. Specifically, generate samples of size $n = 15$ from the $\text{Beta}(10,2)$, $\text{Beta}(2,10)$, and $\text{Beta}(10,10)$ distributions and conduct the following hypothesis tests against the actual mean for each case (e.g., $\frac{10}{10+2}$, $\frac{2}{10+2}$, and $\frac{10}{10+10}$).

- (a) What proportion of the time do we make an error of Type I for a left-tailed test?

```
#Beta (10,2)
actual.mean1 = 10/(10+2)
actual.mean2 = 2/(2+10)
actual.mean3 = 10/(10+10)
total.b1 = 0
for(i in 1:1000){
  dat.b1 = rbeta(n = 15, shape1 = 10, shape2 = 2)
  b1 = t.test(dat.b1, alternative = "less", mu = actual.mean1)
  if(b1$p.value < .05){ #Because we know that the null should't be rejected
    total.b1 = total.b1 + 1
  }
}
#Beta (2,10)
(type1.error.b1 = total.b1/1000) #Proportion of type1 errors

## [1] 0.024

total.b2 = 0
for(i in 1:1000){
  dat.b2 = rbeta(n = 15, shape1 = 2, shape2 = 10)
  b2 = t.test(dat.b2, alternative = "less", mu = actual.mean2)
  if(b2$p.value < .05){
    total.b2 = total.b2 + 1
  }
}
(type1.error.b2 = total.b2/1000)

## [1] 0.085

#Beta (10,10)
total.b3 = 0
for(i in 1:1000){
  dat.b3 = rbeta(n = 15, shape1 = 10, shape2 = 10)
  b3 = t.test(dat.b3, alternative = "less", mu = actual.mean3)
  if(b3$p.value < .05){
    total.b3 = total.b3 + 1
  }
}
(type1.error.b3 = total.b3/1000)

## [1] 0.057
```

- (b) What proportion of the time do we make an error of Type I for a right-tailed test?

```
#Beta(10,2)
total.a1 = 0
for(i in 1:1000){
  dat.a1 = rbeta(n = 15, shape1 = 10, shape2 = 2)
  a1 = t.test(dat.a1, alternative = "greater", mu = actual.mean1)
  if(a1$p.value < .05){
    total.a1 = total.a1 + 1
  }
}
(type1.error.a1 = total.a1/1000)

## [1] 0.092

#Beta(2,10)
total.a2 = 0
for(i in 1:1000){
  dat.a2 = rbeta(n = 15, shape1 = 2, shape2 = 10)
  a2 = t.test(dat.a2, alternative = "greater", mu = actual.mean2)
  if(a2$p.value < .05){
    total.a2 = total.a2 + 1
  }
}
(type1.error.a2 = total.a2/1000)

## [1] 0.031

#Beta(10,10)
total.a3 = 0
for(i in 1:1000){
  dat.a3 = rbeta(n = 15, shape1 = 10, shape2 = 10)
  a3 = t.test(dat.a3, alternative = "greater", mu = actual.mean3)
  if(a3$p.value < .05){
    total.a3 = total.a3 + 1
  }
}
```

```

}
(type1.error.a3 = total.a3/1000)

## [1] 0.046

```

- (c) What proportion of the time do we make an error of Type I for a two-tailed test?

```

#####
#2.c

#Beta(10,2)
total.c1 = 0
for(i in 1:1000){
  datc1 = rbeta(n = 15, shape1 = 10, shape2 = 2)
  c1 = t.test(datc1, alternative = "two.sided", mu = actual.mean1)
  if(c1$p.value < .05){
    total.c1 = total.c1 + 1
  }
}
(type1.error.c1 = total.c1/1000)

## [1] 0.057

#Beta(2,10)
total.c2 = 0
for(i in 1:1000){
  dat.c2 = rbeta(n = 15, shape1 = 2, shape2 = 10)
  c2 = t.test(dat.c2, alternative = "two.sided", mu = actual.mean2)
  if(c2$p.value < .05){
    total.c2 = total.c2 + 1
  }
}
(type1.error.c2 = total.c2/1000)

## [1] 0.057

#Beta(10,10)
total.c3 = 0
for(i in 1:1000){
  dat.c3 = rbeta(n = 15, shape1 = 10, shape2 = 10)
  c3 = t.test(dat.c3, alternative = "two.sided", mu = actual.mean3)
  if(c3$p.value < .05){
    total.c3 = total.c3 + 1
  }
}
(type1.error.c3 = total.c3/1000)

## [1] 0.062

```

- (d) How does skewness of the underlying population distribution effect Type I error across the test types?

It impacts the Type 1 error for one sided tests. The sample size is not great enough for the central limit theorem to apply for the sample means from the simulations. This causes the skewed populations distributions to have sample mean distributions that are also skewed, which ultimately impacts our results for a one sided t-test over many simulations since the data is not normal. This is why the right skewed beta distribution has a large Type 1 error (greater than .05) when it does the less than one-sided t-test and why it has a smaller Type 1 error (less than .05) when it does the greater than one-sided t-test. For two-sided t tests, however, the large and small type1 errors on both sides compensate for one another which ultimately causes the Type 1 error to be similar to .05.

References

- Yee, T. W. (2010). The VGAM package for categorical data analysis. *Journal of Statistical Software*, 32(10):1-34.