

T test

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T test in R

For the Z-test, we can compute the type II error probability analytically. However for the t-test, the type II error probability cannot be given in a closed form because the test statistic contains the sample standard deviation S . What we can do, is to approximate the type II error probability by simulations.

General principle: if we do not know the probability p of a random event E , then we can generate a lot of independent experiments and count the relative frequency of E . This relative frequency will be our estimate of p . This strategy is called *Monte Carlo simulation* and is very common in computational probability.

```
#####
## tTestSim
## Monte Carlo simulation of type I and type II error probabilities of T test
## Input:
## N: number of iterations
## n: sample size
## mean: population mean
## mean0: mu_0 in the null hypothesis
## sigma: population standard deviation
## test: type of alternate hypothesis
## Output:
## y: Monte Carlo simulation of type I / type II error probability
#####
tTestSim<-function(N=1000,n=20,mean=0,
  mean0=1,sigma=1,alpha=.05,seedNum=1,test="lowerTail"){
  ## determine seed for reproducing a result for simulation
  set.seed(seedNum)
  ## result vector
  rejections <- numeric(N)

  for(i in 1:N){
    ## generate data from a normal distribution
    dataSet <- rnorm(n=n,mean=mean,sd=sigma)
    ## compute test statistic
    xbar <- mean(dataSet)
    s <- sd(dataSet)
    testStat <- (xbar-mean0)/(s/sqrt(n))

    if(test=="lowerTail"){
      ## H0: mu=mu0 vs Ha: mu < mu0
      ## RR testStat < -t_{alpha,n-1}
      if(testStat < -qt(p=alpha,df=n-1,lower.tail=FALSE)){
```

```

        rejections[i] <- 1
      }
    }
  }

  ## Reporting Results
  if(test=="lowerTail"){
    cat("Hypothesis Testing: H0: mu=",mean0," vs. Ha: mu < ",mean0,"\n")
    if(mean>=mean0){
      cat("Simulation: Type I Error Probability = ",mean(rejections),"\n")
    }else{
      cat("Simulation: Type II Error Probability",1-mean(rejections)," at mu = ",mean,"\n")
    }
  }
}

```

Let us now test this function:

```
tTestSim()
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu < 1
## Simulation: Type II Error Probability 0.004 at mu = 0
```

```
tTestSim(mean = 0.9)
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu < 1
## Simulation: Type II Error Probability 0.898 at mu = 0.9
```

As you can see, the function is incomplete as only the lower tail part is implemented. Please implement the other 2 cases as a homework.

```
#####
## tTestSim
## Monte Carlo simulation of type I and type II error probabilities of T test
## Input:
## N: number of iterations
## n: sample size
## mean: population mean
## mean0: mu_0 in the null hypothesis
## sigma: population standard deviation
## test: type of alternate hypothesis
## Output:
## y: Monte Carlo simulation of type I / type II error probability
#####
tTestSim<-function(N=1000,n=20,mean=0,
  mean0=1,sigma=1,alpha=.05,seedNum=1,test="lowerTail"){
  ## determine seed for reproducing a result for simulation
  set.seed(seedNum)
  ## result vector
  rejections <- numeric(N)

  for(i in 1:N){

```

```

## generate data from a normal distribution
dataSet <- rnorm(n=n,mean=mean,sd=sigma)
## compute test statistic
xbar <- mean(dataSet)
s <- sd(dataSet)
testStat <- (xbar-mean0)/(s/sqrt(n))

if(test=="lowerTail"){
  ## H0: mu=mu0 vs Ha: mu < mu0
  ## RR testStat < -t_{alpha,n-1}
  if(testStat < -qt(p=alpha,df=n-1,lower.tail=FALSE)){
    rejections[i] <- 1
  }
}else if(test=="upperTail"){
  ## H0: mu=mu0 vs Ha: mu > mu0
  ## RR testStat > t_{alpha,n-1}
  if(testStat > qt(p=alpha,df=n-1,lower.tail=FALSE)){
    rejections[i] <- 1
  }
}else if(test=="twoSided"){
  ## H0: mu=mu0 vs Ha: mu != mu0
  ## RR |testStat| >= t_{alpha/2,n-1}
  if(abs(testStat) >= qt(p=alpha/2,df=n-1,lower.tail=FALSE)){
    rejections[i] <- 1
  }
}
}

## Reporting Results
if(test=="lowerTail"){
  cat("Hypothesis Testing: H0: mu=",mean0," vs. Ha: mu < ",mean0,"\n")
  if(mean>=mean0){
    cat("Simulation: Type I Error Probability = ",mean(rejections),"\n")
  }else{
    cat("Simulation: Type II Error Probability",1-mean(rejections)," at mu = ",mean,"\n")
  }
}else if(test=="upperTail"){
  cat("Hypothesis Testing: H0: mu=",mean0," vs. Ha: mu > ",mean0,"\n")
  if(mean<=mean0){
    cat("Simulation: Type I Error Probability = ",mean(rejections),"\n")
  }else{
    cat("Simulation: Type II Error Probability",mean(rejections)," at mu = ",mean,"\n")
  }
}else if(test=="twoSided"){
  cat("Hypothesis Testing: H0: mu=",mean0," vs. Ha: mu != ",mean0,"\n")
  if(mean==mean0){
    cat("Simulation: Type I Error Probability = ",mean(rejections),"\n")
  }else{
    cat("Simulation: Type II Error Probability",mean(rejections)," at mu = ",mean,"\n")
  }
}
}

```

```
tTestSim(mean = 0.9, test="lowerTail")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu < 1  
## Simulation: Type II Error Probability 0.898 at mu = 0.9
```

```
tTestSim(mean = 0.9, test="upperTail")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu > 1  
## Simulation: Type I Error Probability = 0.014
```

```
tTestSim(mean = 0.9, test="twoSided")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu != 1  
## Simulation: Type II Error Probability 0.064 at mu = 0.9
```

```
tTestSim(mean = 1.1, test="lowerTail")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu < 1  
## Simulation: Type I Error Probability = 0.017
```

```
tTestSim(mean = 1.1, test="upperTail")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu > 1  
## Simulation: Type II Error Probability 0.095 at mu = 1.1
```

```
tTestSim(mean = 1.1, test="twoSided")
```

```
## Hypothesis Testing: H0: mu= 1 vs. Ha: mu != 1  
## Simulation: Type II Error Probability 0.061 at mu = 1.1
```