1. Exercise 1: One-Sample Z-Test

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o Task: Perform a one-sample Z-test to determine if the mean weight of a sample of
50 individuals is significantly different from 65 kg. Use a significance level of
0.05.
o Expected Output:
# Example data
sample_data <- rnorm(50, mean = 68, sd = 5) # Simulated sample data
# Population parameters (if known)
mu <- 65 # Population mean
sigma <- 5 # Population standard deviation
# Calculate Z-statistic
z stat <- (mean(sample data) - mu) / (sigma / sqrt(length(sample data)))
# Calculate p-value (two-tailed)
p_value <- 2 * (1 - pnorm(abs(z_stat)))</pre>
# Print results
cat("Z-statistic:", z stat, "\n")
cat("P-value:", p_value, "\n")
# Interpretation
if (p value < 0.05) {
cat("Reject null hypothesis: The sample mean is significantly different from",
mu, "\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the sample mean is different from", mu, "\n")
}
OUTPUT:
> sample<-rnorm(50, mean=68, sd=5)</pre>
> mu<-65
> sigma<-5
> z_stat<-(mean(sample)-mu)/(sigma/sqrt(length(sample_mean)))</pre>
> p<-2*(1-pnorm(abs(z_stat)))
> cat("Z-statistic:",z_stat,"\n")
Zat( Z-statistic: ",z_stat,
Z-statistic: 0.4580299
> cat("P value: ",p,"\n")
P value: 0.646931
> if(n<0.05);</pre>
> if(p<0.05){
+ cat("Reject null hypothesis: the sample mean is significantly different from",mu,"\n")
+ }else{
+ cat("Fail to reject null hypothesis: There is not enough evidence to conclude that the sample mean is different from",mu,"\n")
Fail to reject null hypothesis: There is not enough evidence to conclude t
hat the sample mean is different from 65
```

2. Exercise 2: Two-Sample Z-Test

```
o Task: Perform a two-sample Z-test to compare the mean scores of two groups:
Group A and Group B. Use the following data:
# Example data
groupA <- c(85, 89, 92, 78, 86, 88, 90, 82, 87, 84)
groupB <- c(80, 81, 85, 79, 83, 81, 84, 78, 82, 80)
# Calculate Z-statistic for two-sample test
z_stat <- (mean(groupA) - mean(groupB)) / sqrt(var(groupA)/length(groupA) +
var(groupB)/length(groupB))
# Calculate p-value (two-tailed)
p_value <- 2 * (1 - pnorm(abs(z_stat)))</pre>
# Print results
cat("Z-statistic:", z_stat, "\n")
cat("P-value:", p_value, "\n")
# Interpretation
if (p_value < 0.05) {
cat("Reject null hypothesis: The means of two groups are significantly different
\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the means of two groups are different \n")
}
OUTPUT:
> A<-c(85,89,92,78,86,88,90,82,87,84)
> B<-c(80,81,85,79,83,81,84,78,82,80)
> z_stat<-(mean(A)-mean(B))/sqrt(var(A)/length(A)+var(B)/length(B))</pre>
> p<-2*(1-pnorm(abs(z_stat)))
> cat("Z statistic:",z_stat,"\n")
Z statistic: 3.260958
> cat("P value:",p,"\n")
P value: 0.001110365
> if(p<0.05){
     cat("reject null hypothesis\n")
+ }else{
+ cat("fail to reject")
reject null hypothesis
```

```
Example 1: One-Sample t-Test
# Example data
sample_data <- c(12, 15, 18, 14, 16, 19, 17, 13, 15, 18, 16, 15, 17, 16, 14)
# Population parameters (if known)
mu <- 16 # Population mean (null hypothesis)
# Conduct one-sample t-test
t_test <- t.test(sample_data, mu = mu)
# Print test result
print(t_test)
# Interpretation
if (t_test$p.value < 0.05) {
cat("Reject null hypothesis: The sample mean is significantly different from",
mu, "\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the sample mean is different from", mu, "\n")
}
> sample < -c(12,15,18,14,16,19,17,13,15,18,16,15,17,16,14)
> mu < -16
> t_test<-t.test(sample,mu=mu)</pre>
> print(t_test)
         One Sample t-test
data:
        sample
t = -0.66144, df = 14, p-value = 0.5191
alternative hypothesis: true mean is not equal to 16
95 percent confidence interval:
 14.58580 16.74754
sample estimates:
mean of x
 15.66667
> if(t_test$p.value<0.05){</pre>
   cat("reject null hypothesis",mu,"\n")
+ }else{
     cat("fail to reject null hypothesis",mu,"\n")
fail to reject null hypothesis 16
```

```
Example 2: Two-Sample t-Test
# Example data
group1 <- c(72, 75, 78, 71, 74, 77, 76, 73, 75, 78)
group2 <- c(68, 71, 73, 69, 72, 70, 72, 67, 71, 74)
# Conduct two-sample t-test
t_test <- t.test(group1, group2)
# Print test result
print(t_test)
# Interpretation
if (t_test$p.value < 0.05) {
cat("Reject null hypothesis: The means of two groups are significantly different
\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the means of two groups are different \n")
}
 > grp1<-c(72,75,78,71,74,77,76,73,75,78)
> grp2<-c(68,71,73,69,72,70,72,67,71,74)
 > t_test<-t.test(grp1,grp2)</pre>
> print(t.test)
function (x, ...)
UseMethod("t.test")
 <bytecode: 0x000001c407b2a3d0>
 <environment: namespace:stats>
 > if(t_test$p.value<0.05){
+ cat("reject null hypothesis")</pre>
 + }else{
      cat("fail to reject null hypothesis")
 reject null hypothesis
```

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```
# Example data
sample_data <- rnorm(50, mean = 68, sd = 5) # Simulated sample data
# Population parameters (if known)
mu <- 65 # Population mean (null hypothesis)
# Conduct one-sample t-test
t_test <- t.test(sample_data, mu = mu)
# Print test result
print(t_test)
# Interpretation
if (t_test$p.value < 0.05) {
cat("Reject null hypothesis: The sample mean is significantly different from",
mu, "\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the sample mean is different from", mu, "\n")
}
 > # Example data
> sample_data <- rnorm(50, mean = 68, sd = 5) # Simulated sample data
> # Population parameters (if known)
 > mu <- 65 # Population mean (null hypothesis)
> # Conduct one-sample t-test
> t_test <- t.test(sample_data, mu = mu)
> # Print test result
> print(t_test)
               One Sample t-test
 data: sample_data
t = 4.5989, df = 49, p-value = 3.014e-05
alternative hypothesis: true mean is not equal to 65
95 percent confidence interval:
66.82854 69.66679
 sample estimates:
mean of x
68.24767
 > # Interpretation
> if (t_test$p.value < 0.05) {
+ cat("Reject null hypothesis: The sample mean is significantly different from",
+ _ _ mu, "\n")</pre>
 + } else "{
+   cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
+ the sample mean is different from", mu, "\n")
 Reject null hypothesis: The sample mean is significantly different from 65
```

```
Exercise 2: Two-Sample t-Test
o Task: Perform a two-sample t-test to compare the mean scores of two groups:
Group A and Group B. Use the following data:
# Example data
groupA <- c(85, 89, 92, 78, 86, 88, 90, 82, 87, 84)
groupB <- c(80, 81, 85, 79, 83, 81, 84, 78, 82, 80)
# Conduct two-sample t-test
t_test <- t.test(groupA, groupB)
# Print test result
print(t_test)
# Interpretation
if (t_test$p.value < 0.05) {
cat("Reject null hypothesis: The means of two groups are significantly different
\n")
} else {
cat("Fail to reject null hypothesis: There is not enough evidence to conclude that
the means of two groups are different \n")
}
> # Example data
> groupA <- c(85, 89, 92, 78, 86, 88, 90, 82, 87, 84)
> groupB <- c(80, 81, 85, 79, 83, 81, 84, 78, 82, 80)
> # Conduct two-sample t-test
> t_test <- t.test(groupA, groupB)
> # Print test result
> print(t_test)
           Welch Two Sample t-test
data: groupA and groupB
t = 3.261, df = 13.847, p-value = 0.005759
alternative hypothesis: true difference in means is not equal to 0
95 percent_confidence interval:
 1.639673 7.960327
sample estimates:
mean of x mean of
       86.1
> # Interpretation
> if (t_test$p.value < 0.05) {
+ cat("Reject null hypothesis: The means of two groups are significantly differen</pre>
  \n")
+ } else {
+ cat("Fail to reject null hypothesis: There is not enough evidence to conclude t
+ the means of two groups are different \n")
Reject null hypothesis: The means of two groups are significantly different
```