PS: One Sample Z test for TAs

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2023-08-09

df_water <- read.csv('C:/Users/Guti/OneDrive - University of Vermont/Curriculum/07_ Cursos/NR-2400/Prob

R practice.

Define p-values.

```
head(df water)
         region value
## 1 North West 1782.4
## 2 North East 1904.2
      Midlands 3070.1
## 3
## 4
        Anglian 1773.2
## 5
        Thames 1528.3
## 6
       Southern 1317.0
Define the parameters: mean and sd of the population, sample mean, sample size.
# Population mean (null hypothesis)
population_mean <- 1644
population_sd <- 497</pre>
# Calculate sample mean
sample_mean <- mean(df_water$value, na.rm=TRUE)</pre>
sample_mean
## [1] 1891.9
# Sample size
n <- length(df_water$value)</pre>
## [1] 10
Calculate the test statistic (z-score).
# Calculate the test statistic (z-score)
z_score <- (sample_mean - population_mean) / (population_sd / sqrt(n))</pre>
z_score
## [1] 1.577321
```

```
# Calculate p-value for two-tailed test
p_value_two_tail <- 2 * (1 - pnorm(abs(z_score)))</pre>
\# Calculate the p-value for the right-tailed test
p_value_right_tailed <- 1 - pnorm(z_score)</pre>
# Calculate the p-value for the left-tailed test
p_value_left_tailed <- pnorm(z_score)</pre>
```

Print estimated parameters.

```
# Print results
cat("Z-Score:", z_score)
## Z-Score: 1.577321
cat("Prob >|z|:", p_value_two_tail)
## Prob >|z|: 0.1147216
cat("Prob > z:", p_value_right_tailed)
## Prob > z: 0.05736082
cat("Prob < z:", p_value_left_tailed)</pre>
```

Prob < z: 0.9426392