Analysing Results of Questionnaire

aiswary-a

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0. Data Prep

0.1. Workspace Prep

0.2. Cleaning the Data

```
# insert CSV
df <-
    read_csv(
        "C:/Users/aisro/Desktop/UNI/BSc CogSci - 2nd Sem/Applied/VSCode + Git/Form Responses/CogSci_Applied
) # to be updated with final data, this is just a placeholder

# renaming the df columns
## NOTE: [1] pre-tool usage, [2] post-tool usage, [char] textual response, [num] numerical response</pre>
```

```
renaming_dfCols <-
  c(
    "timestamp",
    "age",
    "vote_eligibility",
    "preTool_biasPerception",
    "preTool_biasAwareness",
    "preTool_morality",
    "preTool_valueCommitment",
    "email",
    "postTool_biasPerception",
    "postTool_biasAwareness",
    "postTool_morality",
    "postTool_valueCommitment",
    "char_engagement",
    "num_decision_enjoyment",
    "num_likely_to_recommend",
    "num_ranking_agreement",
    "char_ranking_selfperception",
    "char_past_candidate_test",
    "char_future_candidate_test",
    "char_comment_tool_selfperception",
    "char_comment_tool_design",
    "char_comment_other",
    "gender",
    "empty"
colnames(df) <- renaming_dfCols</pre>
# deleting email column
df <- df %>%
  select(-email, -empty)
# removing first (test) row
df \leftarrow df[df$timestamp != "16/04/2024 14:07:03",]
# ID'ing non-numeric columnsss and making those all lowercase!
df <- df %>%
 mutate_if(~ !is.numeric(.), tolower)
# groups for stat tests!
# vars. of interest :))
groupSingles <- c(</pre>
  "preTool_biasPerception",
  "postTool_biasPerception",
  "preTool_biasAwareness",
  "postTool_biasAwareness",
  "preTool_morality",
  "postTool morality",
  "preTool_valueCommitment",
  "postTool_valueCommitment"
```

```
# defining the group pairings
groupPairs <- list(
   c("preTool_biasPerception", "postTool_biasPerception"),
   c("preTool_biasAwareness", "postTool_biasAwareness"),
   c("preTool_morality", "postTool_morality"),
   c("preTool_valueCommitment", "postTool_valueCommitment")
)</pre>
```

1. Demographics

```
print(paste("N =", nrow(df)))

## [1] "N = 16"

# age distribution
stats_dfAge <- df %>%
    summarise(mean = mean(age), sd = sd(age)) # insert age column

print(paste("The mean age:", round(stats_dfAge$mean, digits = 3)))

## [1] "The mean age: 28.438"

print(paste("The SD of the age:", round(stats_dfAge$sd, digits = 3)))

## [1] "The SD of the age: 14.278"

# gender distribution
stats_dfGender <- df %>%
    count(gender) # insert gender column

print(paste("The gender distribution of the sample:", stats_dfGender))

## [1] "The gender distribution of the sample: c(\"female\", \"male\")"
## [2] "The gender distribution of the sample: c(9, 7)"
```

2. Pre-Tool vs. Post-Tool Usage

2.1. Checking Assumptions

a. Normality

```
# Shapiro-Wilk test for Normality!
stat nResults <- list() # an empty list to store the results
# normality testing for-loop, going through each indexed column
for (i in seq_along(groupSingles)) {
  single <- groupSingles[i] # extract column name using index</pre>
  # taking the group and forcing it to numeric if necessary
  item <-
   as.numeric(df[[single]])
  # the Shapiro-Wilk normality test!
  shapiro_result <-</pre>
    shapiro.test(item)
  # storing the results
  test name <-
   paste("Shapiro-Wilk Test of Normality for", single)
  stat_nResults[[test_name]] <- shapiro_result</pre>
  # printing the result!
  cat(test_name, ":\n")
  print(shapiro_result)
  # performing the hypothesis test using results!
  if (!is.null(shapiro_result$p.value) &&
      shapiro_result$p.value > 0.05) {
    cat(
      "Fail to reject null hypothesis that data significantly differs from a normal distribution\n\n"
  } else if (!is.null(shapiro_result$p.value) &&
             shapiro_result$p.value <= 0.05) {</pre>
    cat(
      "Reject the null hypothesis that data significantly differs from a normal distribution\n\n"
   )
  } else {
    cat("Unable to compute p-value for the test.\n\n")
}
## Shapiro-Wilk Test of Normality for preTool_biasPerception :
##
##
  Shapiro-Wilk normality test
##
## data: item
## W = 0.8929, p-value = 0.06192
## Fail to reject null hypothesis that data significantly differs from a normal distribution
## Shapiro-Wilk Test of Normality for postTool_biasPerception :
##
## Shapiro-Wilk normality test
```

```
##
## data: item
## W = 0.92252, p-value = 0.2104
## Fail to reject null hypothesis that data significantly differs from a normal distribution
##
## Shapiro-Wilk Test of Normality for preTool_biasAwareness :
##
## Shapiro-Wilk normality test
##
## data: item
## W = 0.9227, p-value = 0.1864
## Fail to reject null hypothesis that data significantly differs from a normal distribution
##
## Shapiro-Wilk Test of Normality for postTool_biasAwareness :
##
## Shapiro-Wilk normality test
##
## data: item
## W = 0.95996, p-value = 0.6917
## Fail to reject null hypothesis that data significantly differs from a normal distribution
## Shapiro-Wilk Test of Normality for preTool_morality :
##
## Shapiro-Wilk normality test
##
## data: item
## W = 0.80769, p-value = 0.003452
## Reject the null hypothesis that data significantly differs from a normal distribution
## Shapiro-Wilk Test of Normality for postTool_morality :
## Shapiro-Wilk normality test
##
## data: item
## W = 0.79027, p-value = 0.002765
##
## Reject the null hypothesis that data significantly differs from a normal distribution
## Shapiro-Wilk Test of Normality for preTool_valueCommitment :
##
## Shapiro-Wilk normality test
##
## data: item
## W = 0.9269, p-value = 0.2176
## Fail to reject null hypothesis that data significantly differs from a normal distribution
## Shapiro-Wilk Test of Normality for postTool_valueCommitment :
##
## Shapiro-Wilk normality test
```

```
## \# data: item  
## \mathbb{W} = 0.88253, p-value = 0.05176  
## \mathbb{W} = 0.88253 that data significantly differs from a normal distribution
```

b. Homogeneity of Variance

```
# Levene's test for homoscedasticity (homogeneity of variance)
stat_lResults <- list() # empty list to store results</pre>
# Levene's test for-loop, going through each indexed column
for (pair in groupPairs) {
  # printing the pair to see the comparison
 print(pair)
  # extracting the pair!
  group1 <- df[[pair[1]]] # member no. 1 of pair!
  group2 <- df[[pair[2]]] # member no. 2 of pair!
  # performing Levene's test
  lResults <-
    leveneTest(group1, group2) # funky naming to deal with overwriting issues :/
  # storing results in the pre-established list
  test_name <- paste("Levene test between", pair[1], "and", pair[2])</pre>
  stat_lResults[[test_name]] <- lResults</pre>
  # see the result!
  cat(test_name, ":\n")
  print(lResults)
  # performing the hypothesis test using results!
  if (!is.null(lResults$`Pr(>F)`[1]) &&
     lResults$`Pr(>F)`[1] > 0.05) {
      "Fail to reject null hypothesis that there is no significant difference between the groups\n\n"
  } else if (!is.null(lResults\rightarrangle) \&&
             lResults$`Pr(>F)`[1] <= 0.05) {</pre>
    cat(
      "Reject the null hypothesis that there is no significant difference between the groups\n\n"
    )
  } else {
    cat("Unable to compute p-value for the test.\n\n")
```

```
## [1] "preTool_biasPerception" "postTool_biasPerception"
## Levene test between preTool_biasPerception and postTool_biasPerception :
```

```
## Levene's Test for Homogeneity of Variance (center = median)
        Df F value
                     Pr(>F)
##
## group 5
              8.85 0.002778 **
##
         9
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Reject the null hypothesis that there is no significant difference between the groups
## [1] "preTool_biasAwareness" "postTool_biasAwareness"
## Levene test between preTool_biasAwareness and postTool_biasAwareness :
## Levene's Test for Homogeneity of Variance (center = median)
        Df F value Pr(>F)
## group 5 0.5551 0.7322
## Fail to reject null hypothesis that there is no significant difference between the groups
## [1] "preTool_morality" "postTool_morality"
## Levene test between preTool_morality and postTool_morality :
## Levene's Test for Homogeneity of Variance (center = median)
        Df F value Pr(>F)
## group 2 1.5364 0.2546
        12
## Fail to reject null hypothesis that there is no significant difference between the groups
## [1] "preTool_valueCommitment" "postTool_valueCommitment"
## Levene test between preTool_valueCommitment and postTool_valueCommitment :
## Levene's Test for Homogeneity of Variance (center = median)
        Df F value Pr(>F)
## group 3 0.3847 0.7662
        11
## Fail to reject null hypothesis that there is no significant difference between the groups
```

2.2. Stat. Testing!

a. Paired t-Test

```
test_name <- paste("Paired t-test between", pair[1], "and", pair[2])</pre>
  stat_tResults[[test_name]] <- stat_tResults</pre>
  # seeing the result of our lovely test!
  cat(test_name, ":\n")
  print(stat_tResults)
  # performing the hypothesis test using results!
  if (!is.null(stat_tResults$p.value[1]) && stat_tResults$p.value[1] > 0.05) {
    cat("Fail to reject null hypothesis that there is no significant difference between the groups\n\n"
  } else if (!is.null(stat_tResults$p.value[1]) && stat_tResults$p.value[1] <= 0.05) {</pre>
    cat("Reject the null hypothesis that there is no significant difference between the groups \n'n")
    cat("Unable to compute p-value for the test.\n\n")
}
## Paired t-test between preTool_biasPerception and postTool_biasPerception :
## Paired t-test
##
## data: group1 and group2
## t = -1.8353, df = 14, p-value = 0.08779
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -1.15659390 0.08992723
## sample estimates:
## mean difference
        -0.5333333
##
##
## Fail to reject null hypothesis that there is no significant difference between the groups
## Paired t-test between preTool_biasAwareness and postTool_biasAwareness :
##
## Paired t-test
##
## data: group1 and group2
## t = -1.526, df = 14, p-value = 0.1493
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -2.084746 0.351413
## sample estimates:
## mean difference
        -0.8666667
##
##
## Fail to reject null hypothesis that there is no significant difference between the groups
## Paired t-test between preTool_morality and postTool_morality :
##
## Paired t-test
##
## data: group1 and group2
## t = 0.32323, df = 14, p-value = 0.7513
```

```
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -0.3756988 0.5090321
## sample estimates:
## mean difference
       0.06666667
##
## Fail to reject null hypothesis that there is no significant difference between the groups
## Paired t-test between preTool_valueCommitment and postTool_valueCommitment :
## Paired t-test
##
## data: group1 and group2
## t = 1.5718, df = 14, p-value = 0.1383
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## -0.145813 0.945813
## sample estimates:
## mean difference
##
              0 4
##
## Fail to reject null hypothesis that there is no significant difference between the groups
```

3. Unused Code I'm Too Paranoid to Delete

```
# making all text in df lowercase
df <- df %>%
  mutate(
   vote_eligibility = tolower(vote_eligibility),
    char_engagement = tolower(char_engagement),
   char_ranking_selfperception = tolower(char_engagement),
    char_past_candidate_test = tolower(char_past_candidate_test),
   char_future_candidate_test = tolower(char_future_candidate_test),
   char_comment_tool_selfperception = tolower(char_comment_tool_selfperception),
    char_comment_tool_design = tolower(char_comment_tool_design),
    char_comment_other = tolower(char_comment_other),
    gender = tolower(gender)
  )
# - - - -
# group pairings
gr_dfBiasPerception <- c(df$preTool_biasPerception,df$postTool_biasPerception)
gr_dfBiasAwareness <- c(df$preTool_biasAwareness, df$postTool_biasAwareness)</pre>
gr_dfMorality <- c(df$preTool_morality, df$postTool_morality)</pre>
gr_dfValCommitment <- c(df$preTool_valueCommitment, df$postTool_valueCommitment)
```