## KEN4258: Computational Statistics

Assignment 2

Aurélien Bertrand Bart van Gool Gaspar Kuper Ignacio Cadarso Quevedo Nikola Prianikov

March 2024

## Assignment 2

Link to our GitHub repository: https://github.com/nprianikov/compstats

1) Reproduce Figure 1 from (Candès et al. 2018).

```
library(tibble)
set.seed(101)
n <- 500
p <- 200
n_sim <- 10 # TODO: not showing anything with 10000
\# Generate X as a single AR(1) time-series of size p
generate_AR1 <- function(n, p) {</pre>
      X <- replicate(n, arima.sim(n=p, list(0.5)))</pre>
      return(t(X))
}
# Function to generate responses, fit a logistic regression model and return individual p-values
generate_responses <- function(n, p, prob = NULL) {</pre>
      X <- generate_AR1(n=n, p=p)</pre>
       if (is.null(prob)) {
             prob <- plogis(0.08 * rowSums(X[, 2:22]))</pre>
      Y <- rbinom(n, 1, prob)
      fit <- glm(Y ~ X, family = binomial(link="logit"))</pre>
      p_values \leftarrow summary(fit) coef[, "Pr(>|z|)"][2] # Take p-values for beta_1 only for the property of the proper
       return(p_values)
# Combine the lists into a single tibble
df_plot <- tibble(</pre>
      p_values_1 = c(replicate(n_sim, generate_responses(n=n, p=p, prob=0.5))),
      p_values_2 = c(replicate(n_sim, generate_responses(n=n, p=p)))
```

## 2) What is the problem that Figure 1 tries to illustrate?

Figure 1 illustrates the p-values for predictors in two different models. The models are given by the following functions:

- (1)  $Y|X_1,\ldots,X_p \sim \text{Bernoulli}(0.5)$
- (2)  $Y|X_1,\ldots,X_p \sim \text{Bernoulli}(\text{logit}(0.08(X_2+\ldots+X_{21})))$

Where  $(X_1, ..., X_p)$  are itself random variables generated by an AR(1) time series with AR coefficient 0.5, and p = 200. In other words, the results of the first simulation are completely independent from the 200 values of  $X_i$  while the second simulation is only dependent on  $(X_2, ..., X_{21})$ .

## 3) Propose a solution to address the problem.

```
# draw a new sample from the conditional distribution of \mathit{Xj} / \mathit{X-j} using a random number generator
# Not sure if this is the correct way to sample
simulate_Xj_given_Xj <- function(X, j) {</pre>
  Xj <- X[, j]
  X_minus_j <- X[, -j, drop = FALSE]</pre>
  # Fit a model to predict Xj from the rest of the data
  model <- glm(Xj ~ X_minus_j - 1, family = "gaussian")</pre>
  predicted_Xj <- predict(model, newdata = list(X_minus_j = X_minus_j))</pre>
  residuals sd <- sd(resid(model))</pre>
  # Generate new samples for Xj with added randomness
  new_Xj_samples <- predicted_Xj + rnorm(length(predicted_Xj), mean = 0, sd = residuals_sd)</pre>
  return(new_Xj_samples)
# Feature importance statistic function to test whether Xj and Y are conditionally independent.
compute_Tj <- function(X, y) {</pre>
  fit <- glm(y ~ X, family = binomial(link="logit"))</pre>
  coef <- coef(fit)</pre>
  return(abs(coef))
}
# Conditional Randomization Test for a specific j
conditional_randomization_test <- function(X, y, j, K = 200) {</pre>
  original_Tj <- compute_Tj(X, y)[j]</pre>
  greater count <- 0
  for (k in 1:K) {
    # Create a new data matrix by simulating the jth column and keeping the remaining columns the sa
    X_simulated <- X</pre>
    X_simulated[, j] <- simulate_Xj_given_Xj(X, j)</pre>
    # Compute Tj for the simulated data
    simulated_Tj <- compute_Tj(X_simulated, y)[j]</pre>
    # Increment count if simulated Tj is greater than or equal to the original Tj
    if (simulated_Tj >= original_Tj) {
      greater_count <- greater_count + 1</pre>
```

```
}
 # Calculate p-value
 p_value <- (1 + greater_count) / (K + 1)</pre>
 return(p_value)
{\it \# Conditional \ Randomization \ Test \ for \ entire \ matrix \ X}
conditional_randomization_test_all <- function(X, y, K = 1) {</pre>
 p_values <- numeric(ncol(X))</pre>
 for (j in 1:ncol(X)) {
   p_values[j] <- conditional_randomization_test(X, y, j, K)</pre>
 return(p_values)
n <- 500
p <- 200
X <- generate_AR1(n=n, p=p)</pre>
Y \leftarrow rbinom(n, 1, 0.5)
p_values <- conditional_randomization_test_all(X, Y)</pre>
print(p_values)
    [1] 1.0 0.5 0.5 1.0 0.5 0.5 1.0 0.5 1.0 0.5 0.5 1.0 1.0 1.0 1.0 1.0 1.0 0.5
    [19] 1.0 1.0 0.5 0.5 0.5 0.5 0.5 1.0 0.5 1.0 1.0 1.0 0.5 0.5 0.5 0.5 0.5 0.5
## [37] 0.5 0.5 0.5 1.0 1.0 0.5 1.0 0.5 0.5 0.5 0.5 0.5 1.0 1.0 1.0 0.5 1.0
## [55] 0.5 0.5 0.5 0.5 0.5 1.0 0.5 0.5 1.0 0.5 0.5 1.0 0.5 0.5 0.5 0.5 0.5 0.5 1.0
## [73] 1.0 0.5 1.0 1.0 1.0 1.0 1.0 0.5 1.0 1.0 0.5 0.5 1.0 0.5 1.0 0.5 1.0 1.0
## [91] 0.5 0.5 1.0 1.0 0.5 0.5 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 0.5 1.0 1.0 0.5
## [127] 0.5 0.5 0.5 0.5 1.0 0.5 0.5 0.5 1.0 0.5 1.0 1.0 1.0 1.0 1.0 1.0 0.5 1.0
## [145] 1.0 0.5 1.0 0.5 0.5 1.0 1.0 1.0 1.0 1.0 1.0 0.5 1.0 0.5 0.5 1.0 1.0 1.0
## [163] 0.5 0.5 0.5 1.0 0.5 1.0 1.0 1.0 1.0 0.5 0.5 1.0 1.0 1.0 0.5 1.0 0.5 1.0
## [181] 1.0 1.0 1.0 1.0 0.5 0.5 0.5 1.0 0.5 1.0 1.0 0.5 0.5 0.5 1.0 0.5 1.0 0.5
## [199] 1.0 0.5
```

- 4) Show that your solution fixes the problem.
- 5) Find a real dataset and apply your method.