mis_indep_of_Y

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2) Probability of Missing data doesn't depend on Response Y

- Similarly to above, suppose scientifc interest focuses on determining β_1 , β_2 , β_3 , β_4 in the linear model $y_i = \alpha + x_1\beta_1 + x_2\beta_2 + x_3\beta_3 + x_4\beta_4 + \epsilon_i$.
- Here, $\epsilon_i N(0, \sigma^2)$.
- The missingness model gives observations with $X_2 > \text{median}(X_2)$ a different probability of missingness in X_1, X_3, X_4 than for observations with $X_2 <= \text{median}(X_2)$

```
# Draw data from artificial model specified above
create.data <- function(alpha = 20, beta_1 = 1, beta_2 = 2, beta_3 = 3, beta_4 = 4,
                         sigma2 = 1, n = 200, run = 1) {
  set.seed(seed = run)
  x_1 <- rnorm(n)</pre>
  x 2 <- rnorm(n)
  x_3 <- rnorm(n)
  x 4 \leftarrow rnorm(n)
  y \leftarrow beta_1 * x_1 + beta_2 * x_2 + beta_3 * x_3 + beta_4 * x_4 + alpha + rnorm(n, sd = sqrt(sigma2))
  as.data.frame(cbind("Y" = y, "X_1" = x_1, "X_2" = x_2, "X_3" = x_3, "X_4" = x_4)
}
data <- create.data()</pre>
head(data)
                      X_1
                                 X_2
                                             X_3
## 1 20.96450 -0.6264538  0.4094018  1.0744410 -0.3410670
## 2 33.43197  0.1836433  1.6888733  1.8956548  1.5024245
## 3 23.63707 -0.8356286 1.5865884 -0.6029973 0.5283077
## 4 21.91777 1.5952808 -0.3309078 -0.3908678 0.5421914
## 5 13.36405 0.3295078 -2.2852355 -0.4162220 -0.1366734
## 6 18.32300 -0.8204684 2.4976616 -0.3756574 -1.1367339
MNAR.make.missing <- function(data, prob_missing_larger = 0.2,
                               prob_missing_smaller = 0.8){
  # Setting up the randomness categories for missingness in X_1, X_3, X_4
  higher <- data %>% filter(X_2 > median(X_2)) %>% select(X_1)
  rx1_larger <- rbinom(nrow(higher), 1, prob_missing_larger)</pre>
  rx1_smaller <- rbinom(nrow(data) - nrow(higher), 1, prob_missing_smaller)
  rx3_larger <- rbinom(nrow(higher), 1, prob_missing_larger)</pre>
  rx3_smaller <- rbinom(nrow(data) - nrow(higher), 1, prob_missing_smaller)
  rx4_larger <- rbinom(nrow(higher), 1, prob_missing_larger)</pre>
  rx4_smaller <- rbinom(nrow(data) - nrow(higher), 1, prob_missing_smaller)
  rx1 <- c(rx1_larger, rx1_smaller)</pre>
  rx3 <- c(rx3_larger, rx3_smaller)</pre>
  rx4 <- c(rx4_larger, rx4_smaller)</pre>
  data <- data %>%
```

Multiple Imputation

```
# Simulate multiple imputation, obtaining estimates and 95% confidence interval.
simulate MI2 <- function(runs = 100) {</pre>
  res \leftarrow array(NA, dim = c(5, runs, 3))
  times \leftarrow array(NA, dim = c(100, 1, 1))
  dimnames(res) <- list(c("Intercept", "X_1", "X_2", "X_3", "X_4"),</pre>
                         as.character(1:runs), c("estimate", "2.5%", "97.5%"))
  sim\ dataset \leftarrow as.data.frame(create.data(n = 200))
  for (run in 1:runs){
      # Note that time is only measured for the MI/imp steps
      # (i.e. filtering, predicting)
    missingness_sim_dataset <- MNAR.make.missing(sim_dataset, 0.2, 0.8)
    start_time <- Sys.time()</pre>
    imp_MI <- mice(missingness_sim_dataset, print = FALSE)</pre>
    fit \leftarrow with(imp_MI, lm(Y ~ X_1 + X_2 + X_3 + X_4))
    end_time <- Sys.time()</pre>
    tab <- summary(pool(fit), "all", conf.int = TRUE)</pre>
    res[1, run, ] <- as.numeric(tab[1, c("estimate", "2.5 %", "97.5 %")])
    res[2, run, ] <- as.numeric(tab[2, c("estimate", "2.5 %", "97.5 %")])
    res[3, run, ] <- as.numeric(tab[3, c("estimate", "2.5 %", "97.5 %")])
    res[4, run, ] <- as.numeric(tab[4, c("estimate", "2.5 %", "97.5 %")])
    res[5, run, ] <- as.numeric(tab[5, c("estimate", "2.5 %", "97.5 %")])
    times[run, 1, 1] <- as.numeric(end_time - start_time)</pre>
  }
 list(res, times)
}
# Run 100 iterations
res_MI2 <- simulate_MI2(100)</pre>
# Obtain confidence intervals & estimates for all coefficients, intercept.
apply(res_MI2[[1]], c(1, 3), mean, na.rm = TRUE)
##
              estimate
                                        97.5%
                               2.5%
## Intercept 20.132687 19.5481163 20.717258
## X_1
              1.093180 0.6973546 1.489006
```

	Raw_Bias	Percent_Bias	Coverage_Rate	Average_Width	RMSE
Intercept	0.133	0.663	0.95	1.169	0.233
X_1	0.093	9.318	0.98	0.792	0.161
X_2	-0.025	1.265	0.97	1.182	0.180
X_3	0.077	2.554	0.97	0.661	0.133
X_4	0.043	1.065	0.98	0.622	0.110

```
## X 2
              1.974707 1.3837934 2.565620
## X 3
              3.076629 2.7460809 3.407177
              4.042583 3.7316625 4.353504
## X_4
# Mean time for the multiple imputation instances
times <- res MI2[[2]]</pre>
Multiple_Imputation <- mean(times)</pre>
# Evaluating imputation method performance for estimating
# all parameters of interest.
res <- res_MI2[[1]]
true \leftarrow c(20, 1, 2, 3, 4)
Raw_Bias <- rowMeans(res[,, "estimate"]) - true</pre>
Percent_Bias <- 100 * abs((rowMeans(res[,, "estimate"]) - true)/ true)</pre>
Coverage_Rate <- rowMeans(res[,, "2.5%"] < true & true < res[,, "97.5%"])
Average_Width <- rowMeans(res[,, "97.5%"] - res[,, "2.5%"])
RMSE <- sqrt(rowMeans((res[,, "estimate"] - true)^2))</pre>
MI_measures <- data.frame(Raw_Bias, Percent_Bias, Coverage_Rate, Average_Width, RMSE)
knitr::kable(round(MI_measures, 3), align = "ccccc") %>% kable_styling()
```

Listwise Deletion

```
# Simulate listwise deletion, obtaining estimates and 95% confidence interval.
simulate_LD <- function(runs = 100){</pre>
  res \leftarrow array(NA, dim = c(5, runs, 3))
  dimnames(res) <- list(c("Intercept", "X_1", "X_2", "X_3", "X_4"),</pre>
                          as.character(1:runs), c("estimate", "2.5%", "97.5%"))
  times \leftarrow array(NA, dim = c(runs, 1, 1))
  sim\ dataset \leftarrow as.data.frame(create.data(n = 200))
  # Note that time is only measured for the LD/imp steps (i.e. filtering, predicting)
  for (run in 1:runs){
    missingness_sim_dataset <- MNAR.make.missing(sim_dataset, 0.2, 0.8)
    start_time <- Sys.time()</pre>
    filtered_sim_dataset <- missingness_sim_dataset %>%
      select(Y, X_1, X_2, X_3, X_4) %>%
      filter(!is.na(X_1), !is.na(X_3), !is.na(X_4))
    fit <- with(filtered_sim_dataset, lm(Y ~ X_1 + X_2 + X_3 + X_4))</pre>
    end_time <- Sys.time()</pre>
    times[run, 1, 1] <- as.numeric(end_time - start_time)</pre>
    # loop over each variable. Note we do the imputation just ONCE b/c LD is
    # deterministic.
    for (var in 1:5){
      edges <- as.numeric((confint(fit)[var,]))</pre>
      estimate <- as.numeric(fit$coefficients)[var]</pre>
```

	average-runtime
$Multiple_Imputation$	0.1891701
ListwiseDeletion	0.0083455

```
interval <- c(estimate, edges)</pre>
      res[var, run, ] <- interval</pre>
 }
  list(res, times)
}
result_LD <- simulate_LD()</pre>
# Obtain confidence intervals & estimates for all coefficients, intercept.
apply(result_LD[[1]], c(1, 3), mean, na.rm = TRUE)
##
              estimate
                              2.5%
                                        97.5%
## Intercept 20.114295 19.5798252 20.648765
              1.064777 0.7446178 1.384937
## X_1
## X_2
              1.966311 1.4257592 2.506864
## X_3
              3.014668 2.7245870 3.304749
              3.982521 3.7122312 4.252812
## X_4
# Evaluating imputation method performance for estimating
# all parameters of interest.
res <- result_LD[[1]]
true \leftarrow c(20, 1, 2, 3, 4)
Raw Bias <- rowMeans(res[,, "estimate"]) - true</pre>
Percent_Bias <- 100 * abs((rowMeans(res[,, "estimate"]) - true)/ true)</pre>
Coverage_Rate <- rowMeans(res[,, "2.5%"] < true & true < res[,, "97.5%"])
Average_Width <- rowMeans(res[,, "97.5%"] - res[,, "2.5%"])
RMSE <- sqrt(rowMeans((res[,, "estimate"] - true)^2))</pre>
LD_measures <- data.frame(Raw_Bias, Percent_Bias, Coverage_Rate, Average_Width, RMSE)
# Mean time for 100 instances of LD
times_LD <- result_LD[[2]]</pre>
mean(times_LD)
## [1] 0.008345499
ListwiseDeletion <- mean(times LD)</pre>
mean_times <- as.data.frame(rbind(Multiple_Imputation, ListwiseDeletion), col.names = "average_runtime"
colnames(mean_times) <- "average-runtime"</pre>
mean_times
##
                        average-runtime
## Multiple_Imputation
                            0.189170134
## ListwiseDeletion
                            0.008345499
knitr::kable(mean_times) %>% kable_styling(full_width = F)
knitr::kable(round(LD_measures, 3), align = "ccccc") %>% kable_styling()
```

	Raw_Bias	Percent_Bias	Coverage_Rate	Average_Width	RMSE
Intercept	0.114	0.571	0.98	1.069	0.216
X_1	0.065	6.478	1.00	0.640	0.134
X_2	-0.034	1.684	1.00	1.081	0.176
X_3	0.015	0.489	1.00	0.580	0.097
X_4	-0.017	0.437	1.00	0.541	0.086