# Testing different imputation methods on PUMS (MAR)

- Generative Adversarial Imputation Nets (GAIN)

```
# load dataset: df
load('../../Datasets/ordinalPUMS.Rdata')
# take 10,000 samples: df
set.seed(0)
n = 10000
sample <- sample(nrow(df), size = 10000)</pre>
df <- df[sample,]</pre>
# create MCAR scneario with 30% chance of missing: df_observed
missing_prob = 0.3
df observed <- df
missing_col = c(1,3,7,9,10,11)
# Make VEH and WKL MCAR
missing_col_MCAR = c(1,10)
for (col in missing col MCAR) {
 missing_ind <- rbernoulli(n,p = missing_prob)</pre>
  df_observed[missing_ind, col] <- NA</pre>
}
# Make the rest MAR
numeric_df = sapply(df, as.numeric)
normalized_df = t(t(numeric_df-1)/(apply(numeric_df, MARGIN = 2, FUN = max)-1))
missing_col_MAR = c(3,7,9,11)
fully_observed_col = c(2,4,5,6,8)
beta_NP = c(-0.05, -1.5, 0.6, -2, -0.05)
beta0_NP = -0.05
beta_SCHL = c(-3, 3, -0.75, 0.05, -0.2)
beta0 SCHL = 0.05
beta_AGEP = c(0.05, -0.2, 0.05, -1.25, 1)
beta0\_AGEP = -0.05
beta_PINCP = c(3, -0.05, -2.5, 0.05, -1)
beta0 PINCP = -0.05
# missing probability for NP
prob_NP = apply(t(t(normalized_df[, fully_observed_col])*beta_NP)+beta0_NP, MARGIN = 1, sum)
prob_NP = exp(prob_NP)/(exp(prob_NP)+1)
indicator = rbernoulli(n, p = prob_NP)
df_observed[indicator, missing_col_MAR[1]] <- NA</pre>
# missing probability for SCHL
prob_SCHL = apply(t(t(normalized_df[, fully_observed_col])*beta_SCHL)+beta0_SCHL, MARGIN = 1, sum)
prob_SCHL = exp(prob_SCHL)/(exp(prob_SCHL)+1)
indicator = rbernoulli(n, p = prob_SCHL)
df_observed[indicator, missing_col_MAR[2]] <- NA</pre>
```

```
# missing probability for AGEP
prob_AGEP = apply(t(t(normalized_df[, fully_observed_col])*beta_AGEP)+beta0_AGEP, MARGIN = 1, sum)
prob AGEP = \exp(\text{prob AGEP})/(\exp(\text{prob AGEP})+1)
indicator = rbernoulli(n, p = prob_AGEP)
df_observed[indicator, missing_col_MAR[3]] <- NA</pre>
# missing probability for PINCP
prob_PINCP = apply(t(t(normalized_df[, fully_observed_col])*beta_PINCP)+beta0_PINCP, MARGIN = 1, sum)
prob_PINCP = exp(prob_PINCP)/(exp(prob_PINCP)+1)
indicator = rbernoulli(n, p = prob_PINCP)
df_observed[indicator, missing_col_MAR[4]] <- NA</pre>
# 30.58% missing
apply(is.na(df_observed), MARGIN = 2, mean)
##
      VEH
              ΜV
                                   ENG MARHT
                                                 SCHL RACNUM
                                                                AGEP
                                                                        WKL PINCP
                      NP
                           RMSP
## 0.3030 0.0000 0.3121 0.0000 0.0000 0.0000 0.2814 0.0000 0.3355 0.3017 0.3011
```

#### Generative Adversarial Imputation Nets (GAIN)

reference: https://arxiv.org/abs/1806.02920

```
# Load imputed dataset
d1 = read.csv('../../GAIN/imputed_dataset/MAR_PUMS01_30percent_1.csv', header = FALSE, sep = ',')
d2 = read.csv('../../GAIN/imputed_dataset/MAR_PUMS01_30percent_2.csv', header = FALSE, sep = ',')
d3 = read.csv('../../GAIN/imputed_dataset/MAR_PUMS01_30percent_3.csv', header = FALSE, sep = ',')
d4 = read.csv('../../GAIN/imputed_dataset/MAR_PUMS01_30percent_4.csv', header = FALSE, sep = ',')
d5 = read.csv('../../GAIN/imputed_dataset/MAR_PUMS01_30percent_5.csv', header = FALSE, sep = ',')
colnames(d1) = colnames(df)
colnames(d2) = colnames(df)
colnames(d3) = colnames(df)
colnames(d4) = colnames(df)
imputed_df = rbind(d1, d2, d3, d4, d5)
```

Diagnostics

Assess bivariate joint distribution

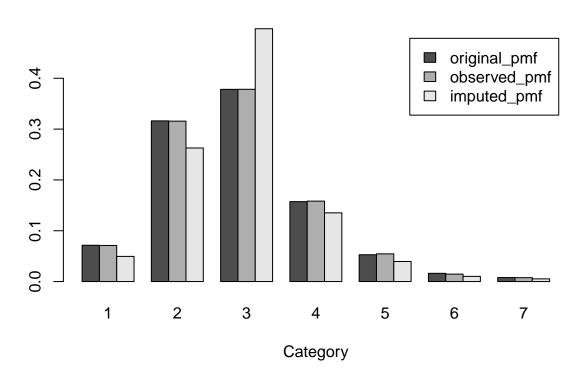
Assess bivariate joint distribution

```
# calculate rmse
numeric_df = sapply(df, as.numeric)
normalized_df = t(t(numeric_df-1)/(apply(numeric_df, MARGIN = 2, FUN = max)-1))
numeric_impute = sapply(d1, as.numeric)
normalized_impute = t(t(numeric_impute-1)/(apply(numeric_df, MARGIN = 2, FUN = max)-1))
missing_matrix = is.na(df_observed)

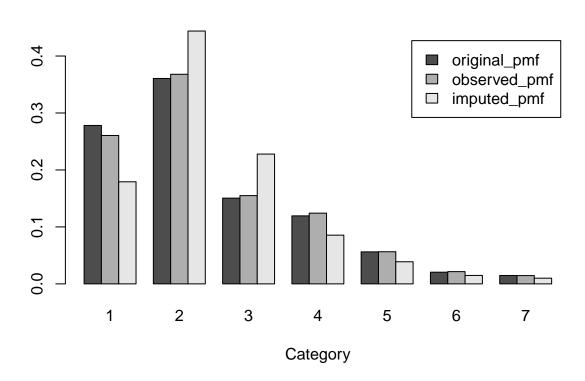
rmse = sqrt(sum((normalized_df[missing_matrix] - normalized_impute[missing_matrix])^2)/sum(missing_matrix)
```

## [1] 0.2568644

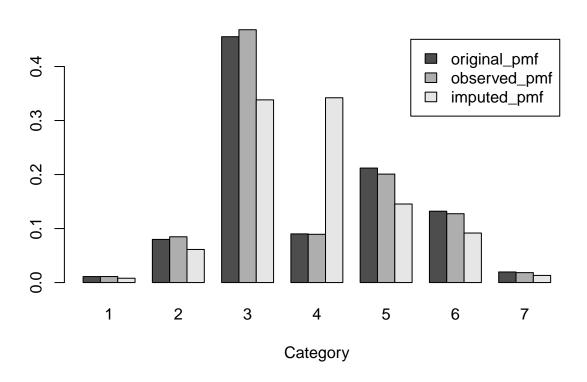
**MICE: VEH** 



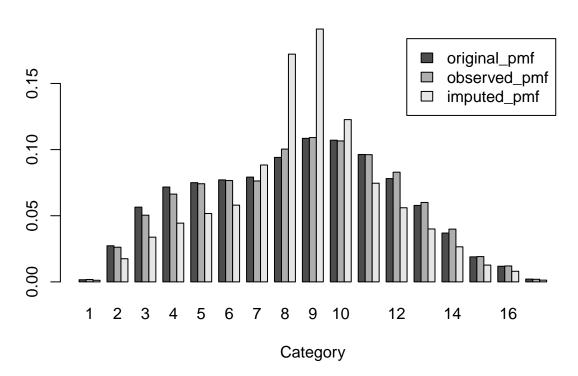
## MICE: NP



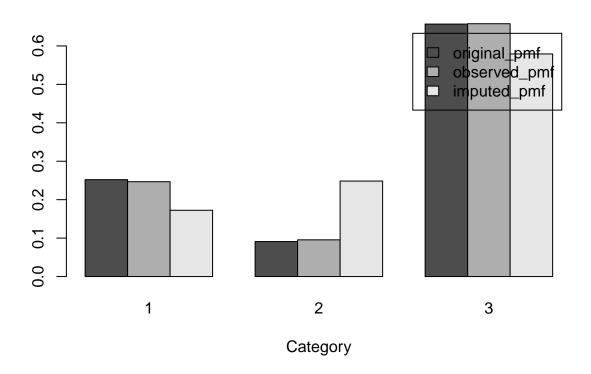
**MICE: SCHL** 



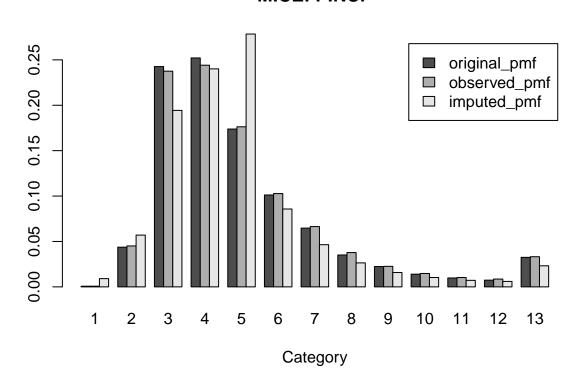
#### **MICE: AGEP**



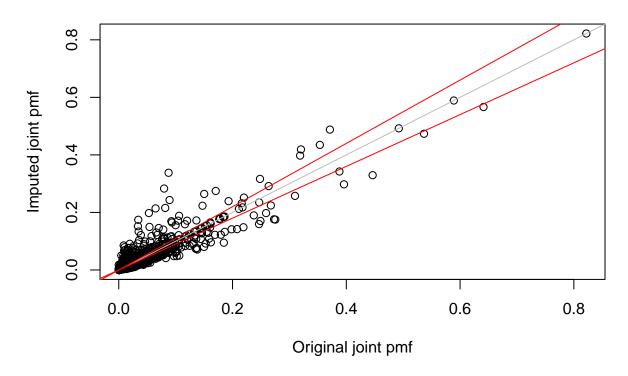
**MICE: WKL** 



#### **MICE: PINCP**



### **Bivariate pmf**



### **Trivariate pmf**

