# Testing different imputation methods on PUMS (MCAR) - RandomForest

```
# load dataset: df
load('../Datasets/ordinalPUMS.Rdata')

# take 10,000 samples: df
set.seed(0)
n = 10000
sample <- sample(nrow(df), size = 10000)
df <- df[sample,]

# create MCAR scneario with 30% chance of missing: df_observed
missing_prob = 0.3
df_observed <- df
missing_col = colnames(df)[c(1,3,5,7,9,11)]
for (col in missing_col) {
   missing_ind <- rbernoulli(n,p = missing_prob)
   df_observed[missing_ind, col] <- NA
}</pre>
```

#### missForest

```
df.imp <- missForest(df_observed, verbose = FALSE)
d1 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d2 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d3 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d4 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d5 <- df.imp$ximp</pre>
```

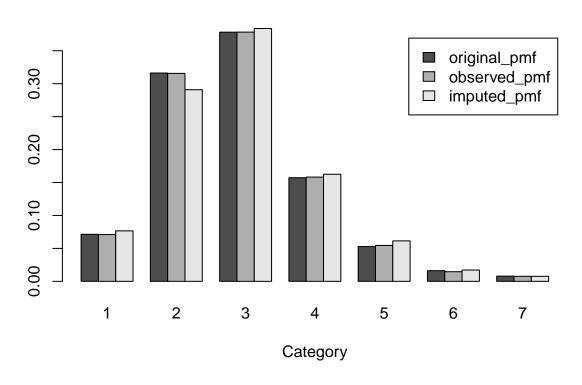
#### Diagnostics

```
for (var_index in c(1,3,5,7,9,11)) {
    y_original = df[,var_index]
    original_pmf = table(y_original)/length(y_original)

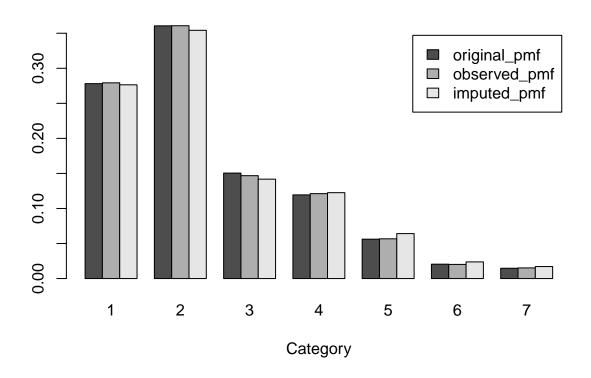
# Observed distribution
    missing_indicator = is.na(df_observed)[,var_index]
    y_observed = y_original[!missing_indicator]
    observed_pmf = table(y_observed)/length(y_observed)

# Extract variable from imputed data
    sample_estimate1 = table(d1[,var_index])/length(d1[,var_index])
    sample_estimate2 = table(d2[,var_index])/length(d2[,var_index])
    sample_estimate3 = table(d3[,var_index])/length(d3[,var_index])
    sample_estimate4 = table(d4[,var_index])/length(d4[,var_index])
    sample_estimate5 = table(d5[,var_index])/length(d5[,var_index])
```

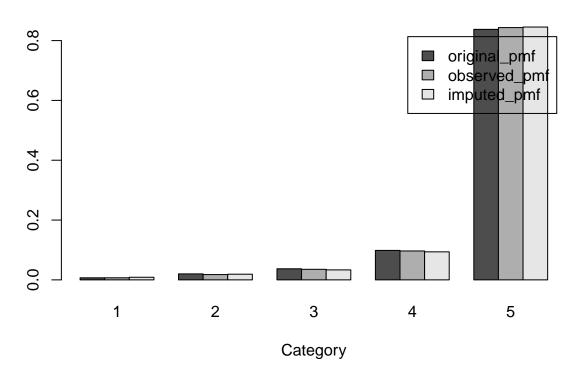
### **MICE: VEH**



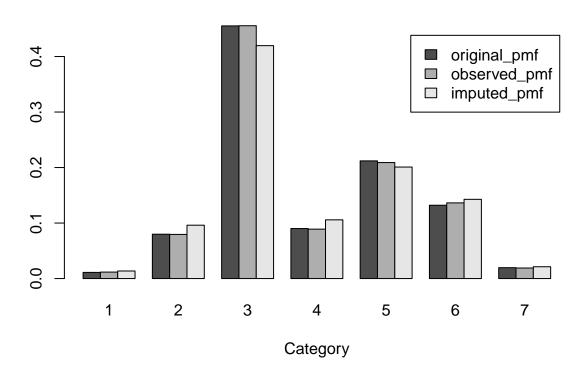
MICE: NP



**MICE: ENG** 



**MICE: SCHL** 



## **MICE: AGEP**

