# 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

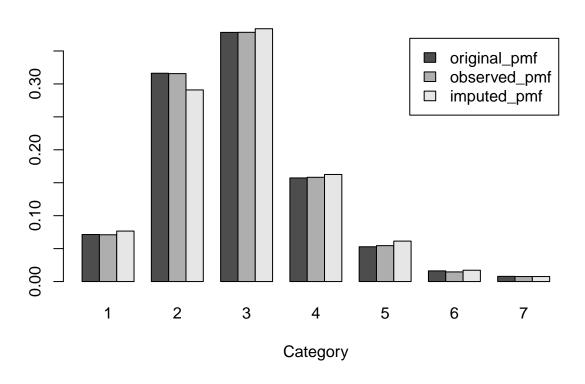
```
df.imp <- missForest(df_observed, verbose = FALSE)
d1 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d2 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d3 <- df.imp$ximp
df.imp <- missForest(df_observed, verbose = FALSE)
d4 <- df.imp$ximp
df.imp <- missForest(df_observed,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

#### Diagnostics

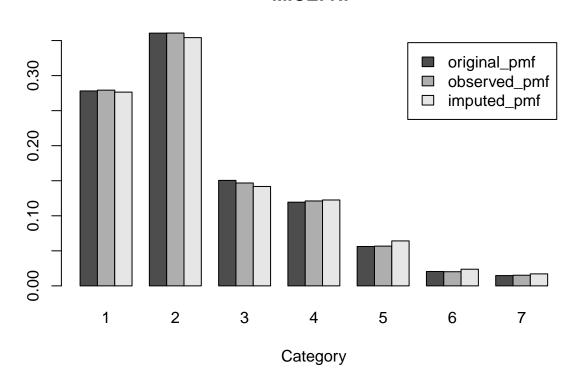
Assess bivariate joint distribution

Assess trivariate joint distribution

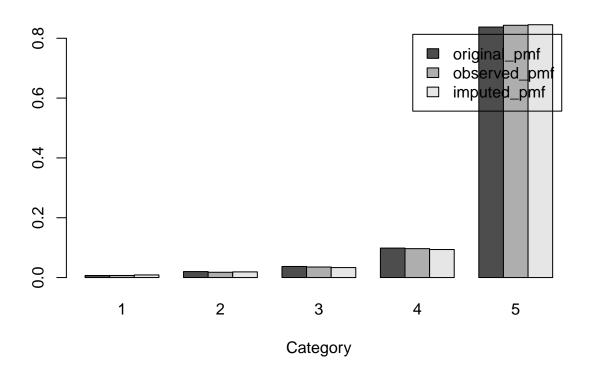
**MICE: VEH** 



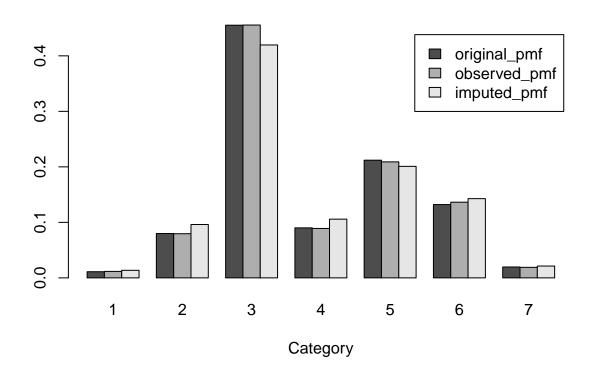
## MICE: NP



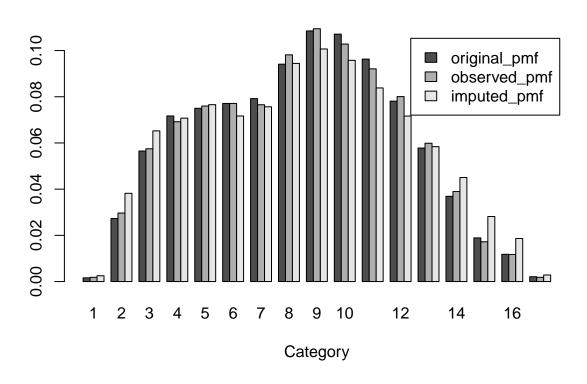
**MICE: ENG** 



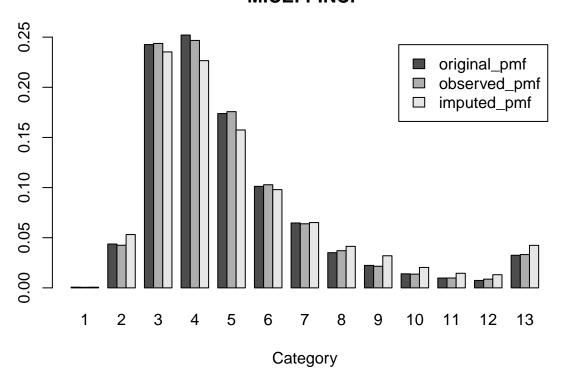
#### **MICE: SCHL**



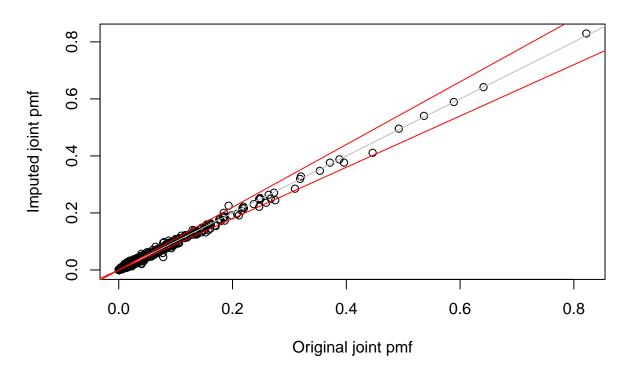
**MICE: AGEP** 







## Bivariate pmf, r square: 0.998



## Trivariate pmf , r square: 0.995

