# Testing different imputation methods on PUMS (MCAR)

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

# take 2000 samples: df
set.seed(0)
n = 3000
sample <- sample(nrow(df), size = n)
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>
```

#### Ordinal bayesian nonparametric model

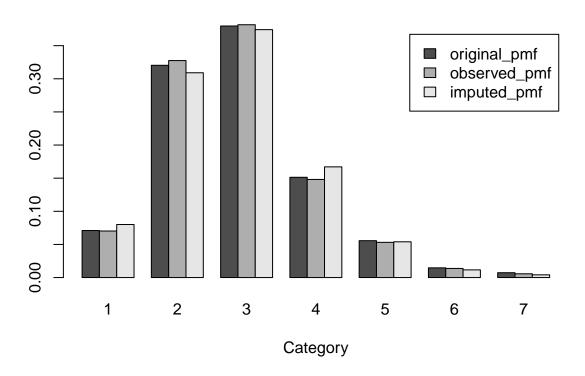
```
source("../probitBayes.R")
N = 40
Mon = 300
B = 300
thin.int = 1
# function(y, N = 40, Mon = 2000, B = 300, thin.int = 5, seed = 0)
output_list <- probitBayesImputation(df_observed, N, Mon, B, thin.int)
sampled_y <- output_list[['sampled_y']]
sampled_z <- output_list[['sampled_z']]</pre>
```

Diagnostics

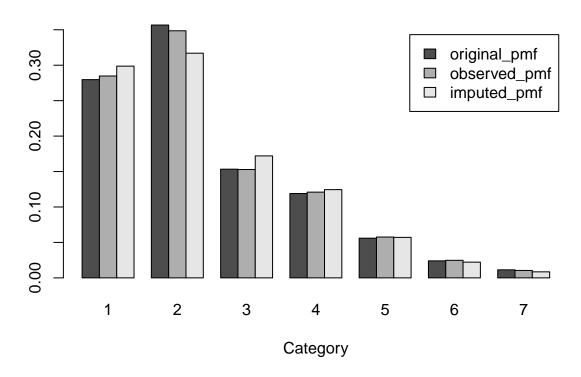
Assess bivariate joint distribution

Assess trivariate joint distribution

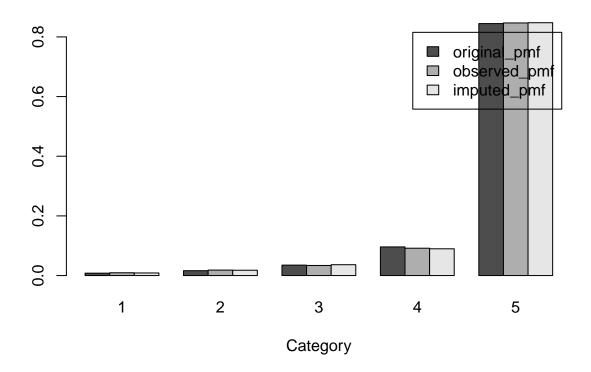
## **Blocked Gibbs Sampling Assessment: VEH**



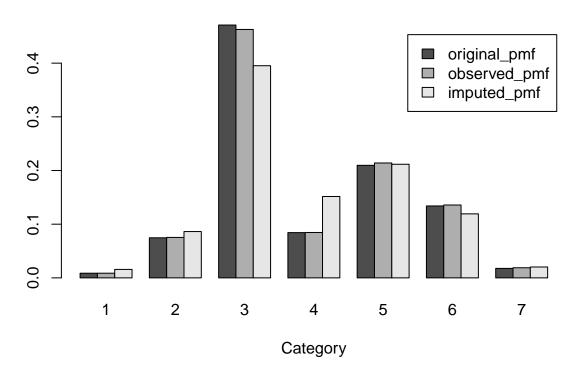
#### **Blocked Gibbs Sampling Assessment: NP**



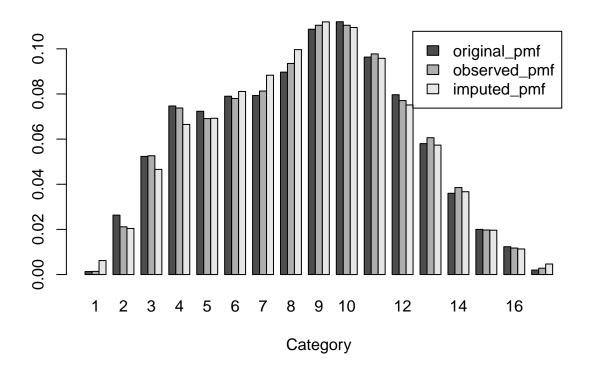
## **Blocked Gibbs Sampling Assessment: ENG**



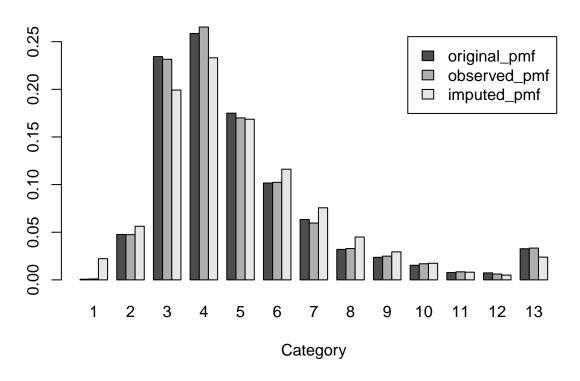
#### **Blocked Gibbs Sampling Assessment: SCHL**



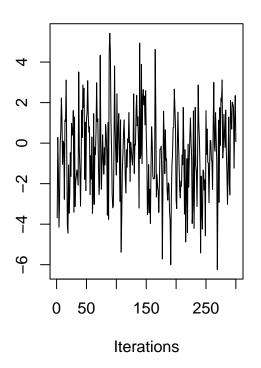
## **Blocked Gibbs Sampling Assessment: AGEP**



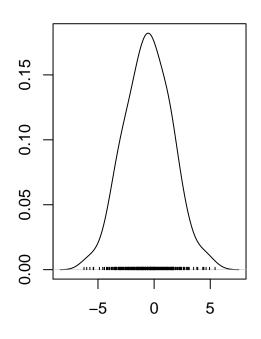
#### **Blocked Gibbs Sampling Assessment: PINCP**



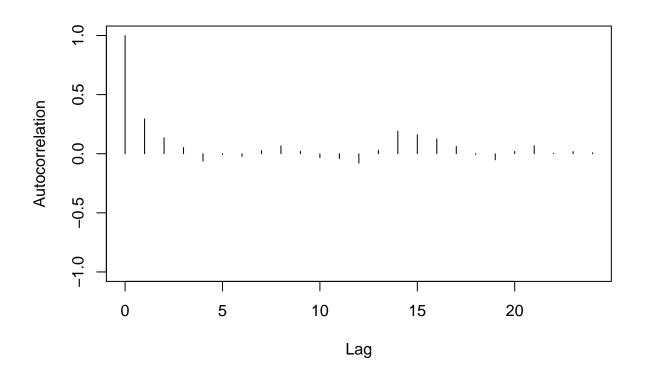
# Trace of var1



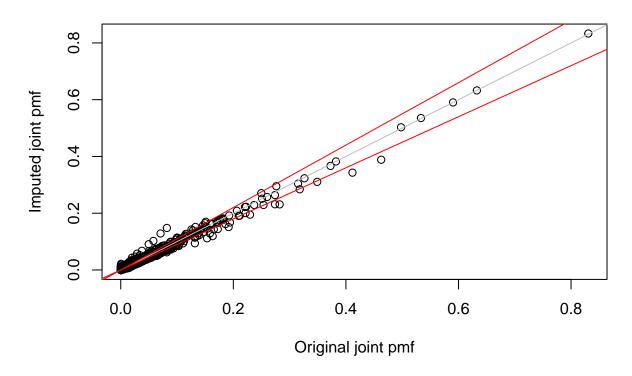
# Density of var1



N = 300 Bandwidth = 0.7072



## Bivariate pmf, r square: 0.994



## Trivariate pmf, r square: 0.99

