MCAR 45% missing - MICE

```r
# sample MCAR dataset from PUMS
source("../../utils/sampleMCAR.R")
n = 10000
missing_col = c(1,3,7,9,10,11)
missing_prob = 0.45
set.seed(1)

output_list <- sampleMCAR(n, missing_prob)
df <- output_list[['df']]
df_observed <- output_list[['df_observed']]
```

MICE

Create 5 imputed dataset

```r
library(mice)
```

```
##
## Attaching package: 'mice'
```

```
## The following objects are masked from 'package:base':
##
##     cbind, rbind
```

```r
imputed_df <-  mice(df_observed,m=5,print=F)
```

```
## Warning: Number of logged events: 150
```

Extract the 5 imputed dataset

```r
d1 <-  complete(imputed_df, 1)
d2 <-  complete(imputed_df, 2)
d3 <-  complete(imputed_df, 3)
d4 <-  complete(imputed_df, 4)
d5 <-  complete(imputed_df, 5)
imputed_sets = rbind(d1, d2, d3, d4, d5)
```

Diagnostics

Assess bivariate joint distribution

Assess trivariate joint distribution

```r
# 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_matri
rmse
```
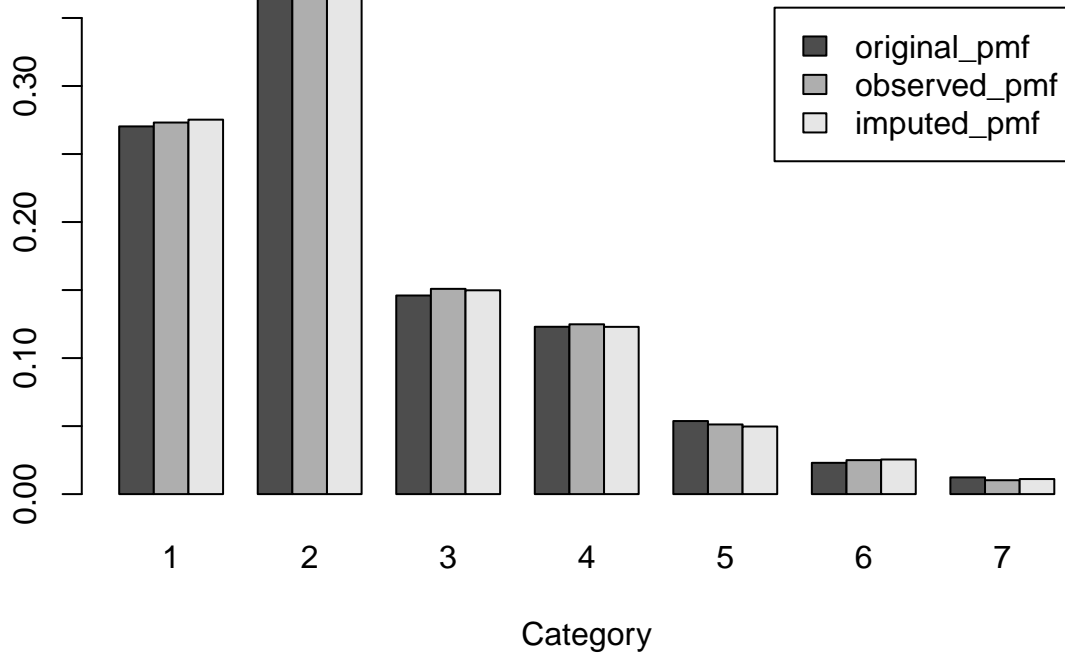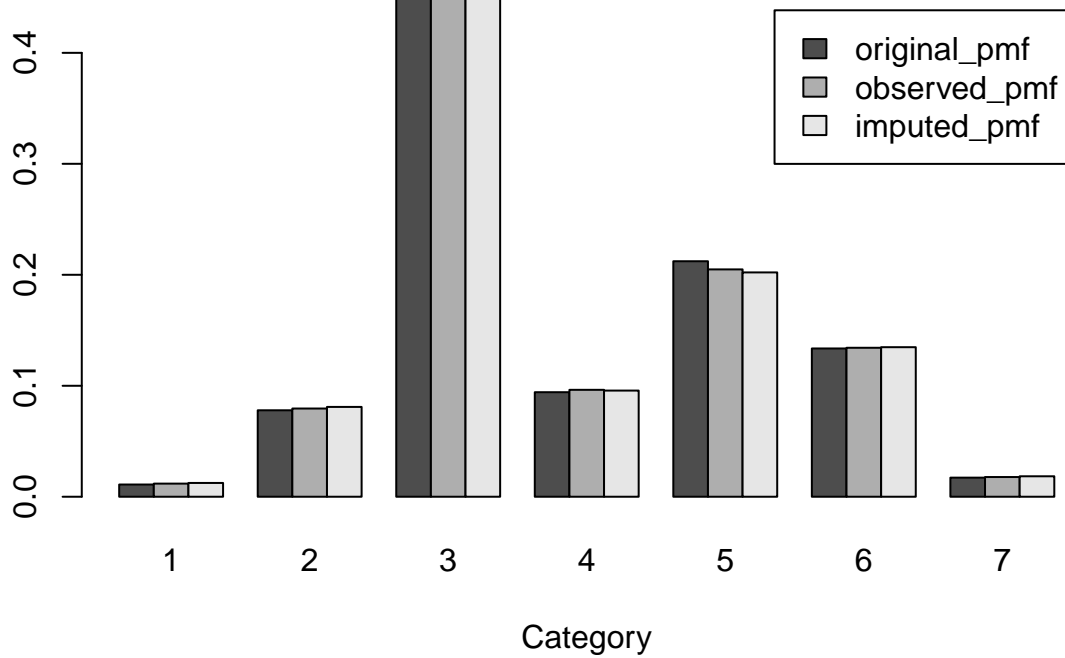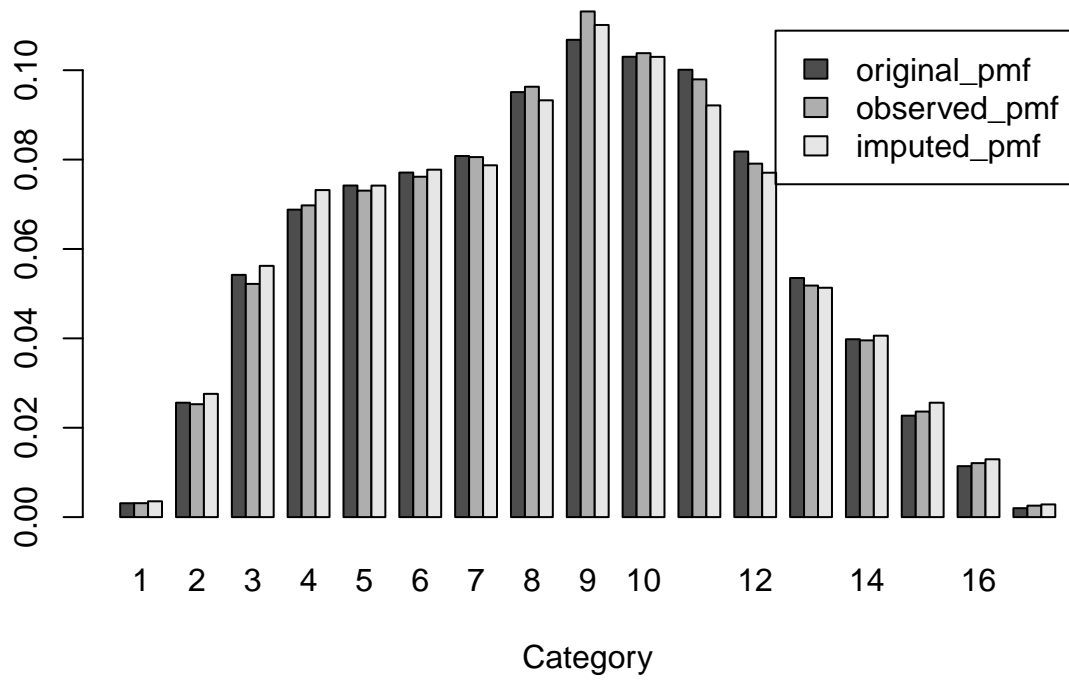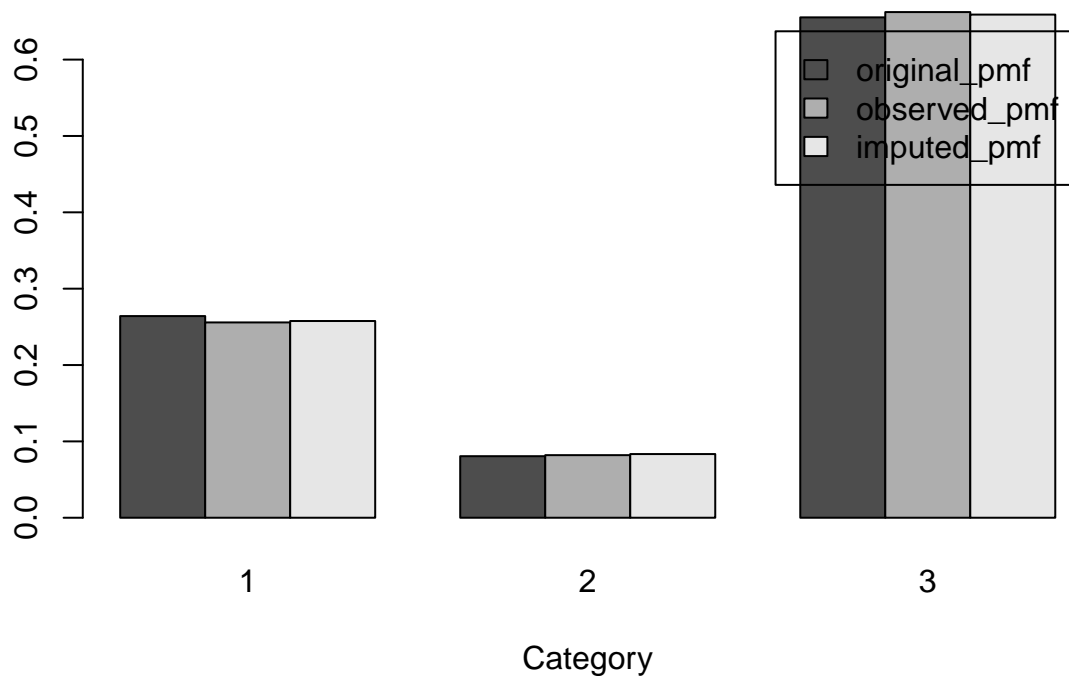
```
## [1] 0.3277459
```
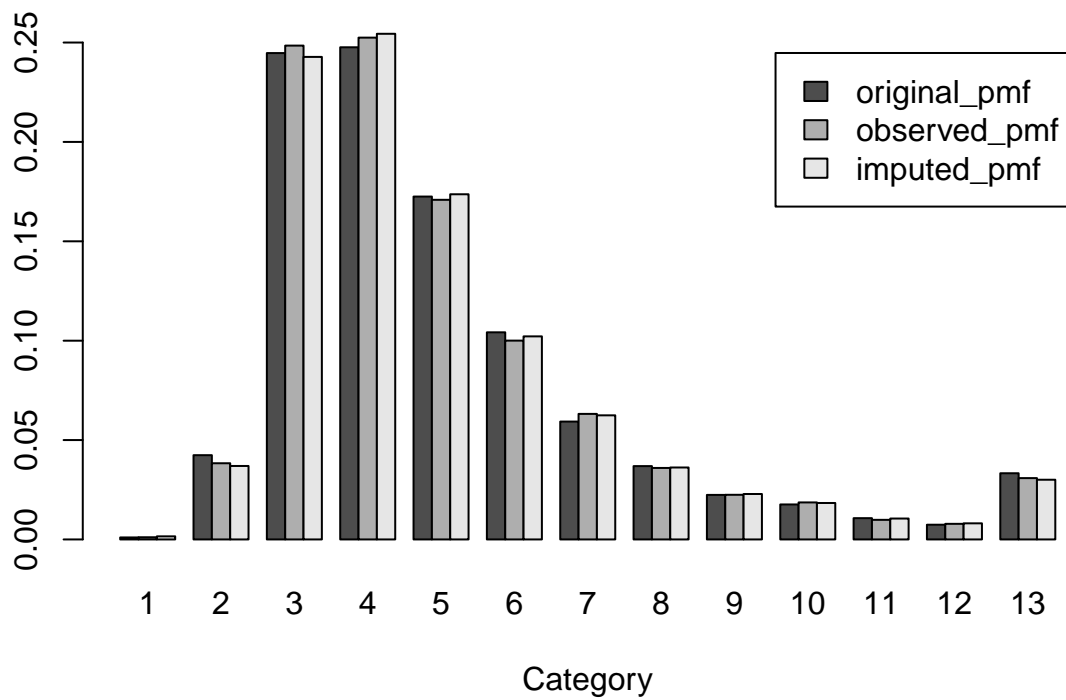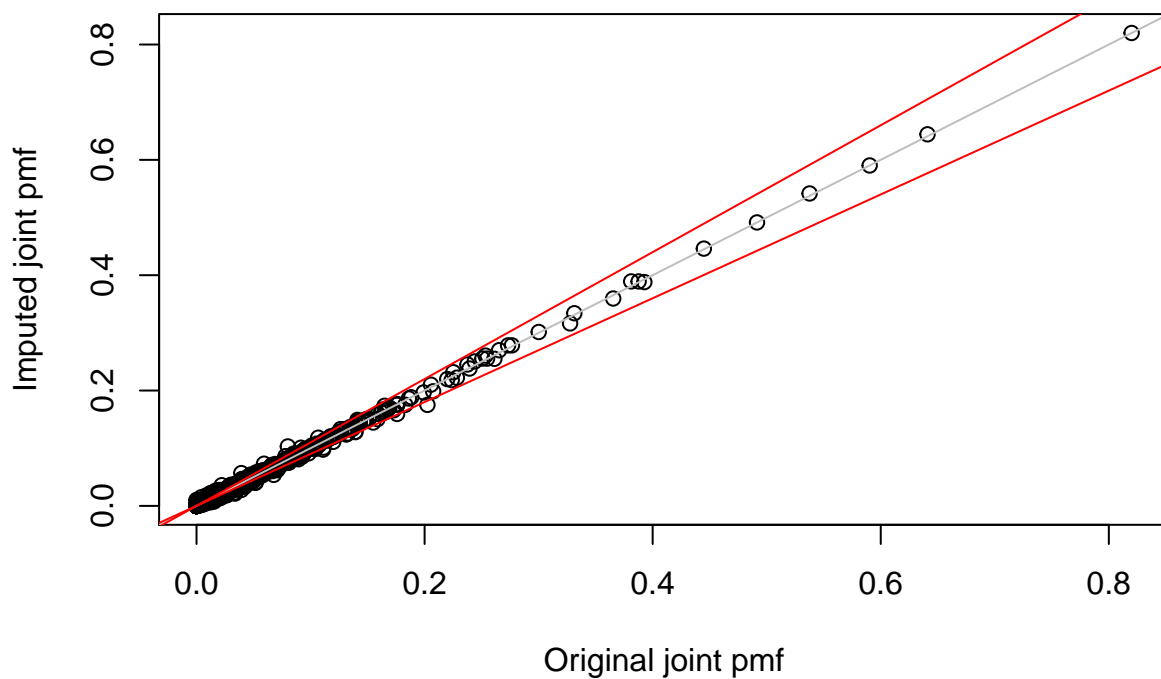
MICE: VEH
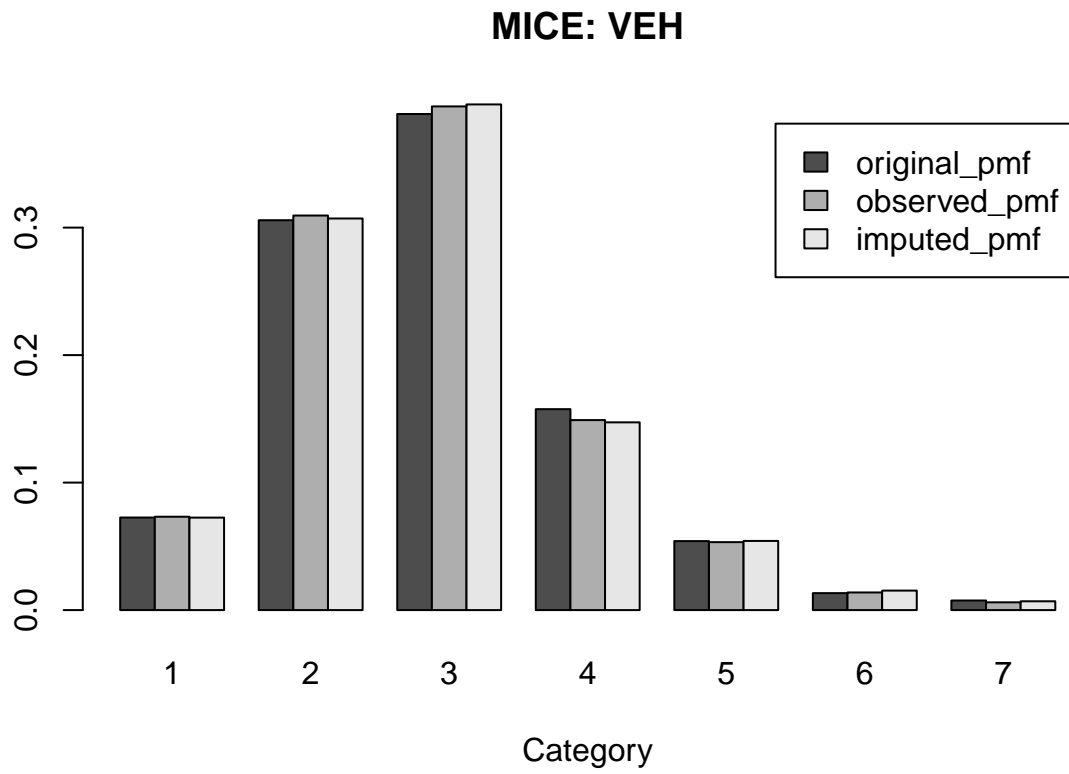
MICE: NP

MICE: SCHL

MICE: AGEP

MICE: WKL

MICE: PINCP

Bivariate pmf

Trivariate pmf