

# MAR 30% missing - Probit

---

```
# sample MCAR dataset from PUMS
source("../utils/sampleMAR30.R")
n = 3000
missing_col = c(1,3,7,9,10,11)
set.seed(2)

output_list <- sampleMAR30(n)
df <- output_list[['df']]
df_observed <- output_list[['df_observed']]

apply(is.na(df_observed), MARGIN = 2, mean)
```

```
##      VEH      MV      NP      RMSP      ENG      MARHT      SCHL      RACNUM
## 0.2926667 0.0000000 0.2816667 0.0000000 0.0000000 0.0000000 0.3213333 0.0000000
##      AGEP      WKL      PINCP
## 0.3330000 0.3100000 0.2890000
```

## 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']]
```

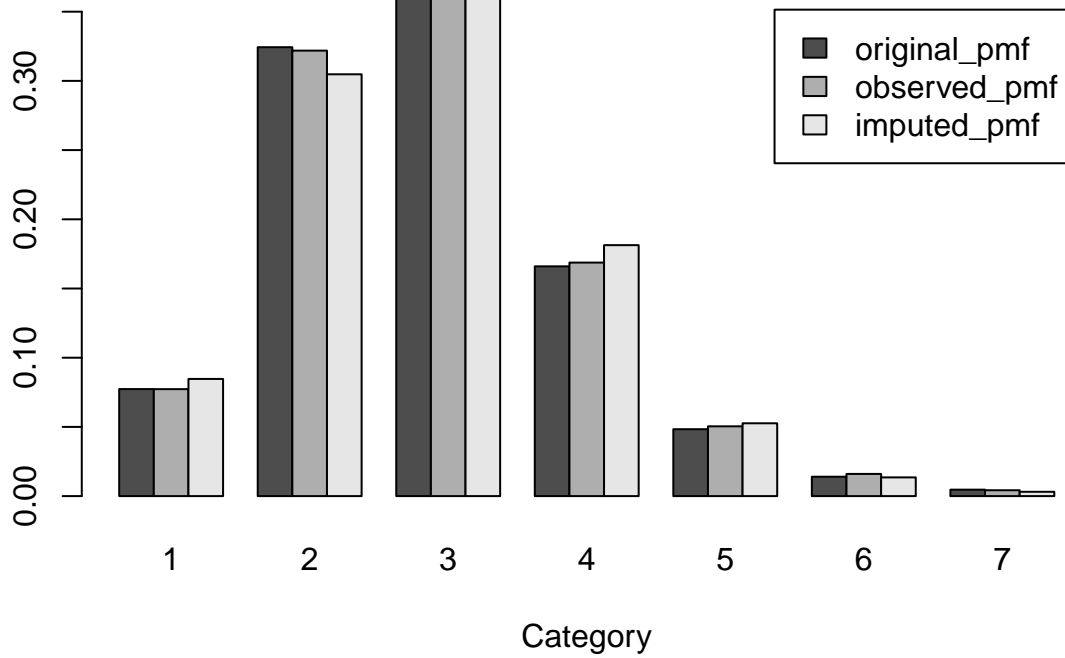
Diagnostics

Assess bivariate joint distribution

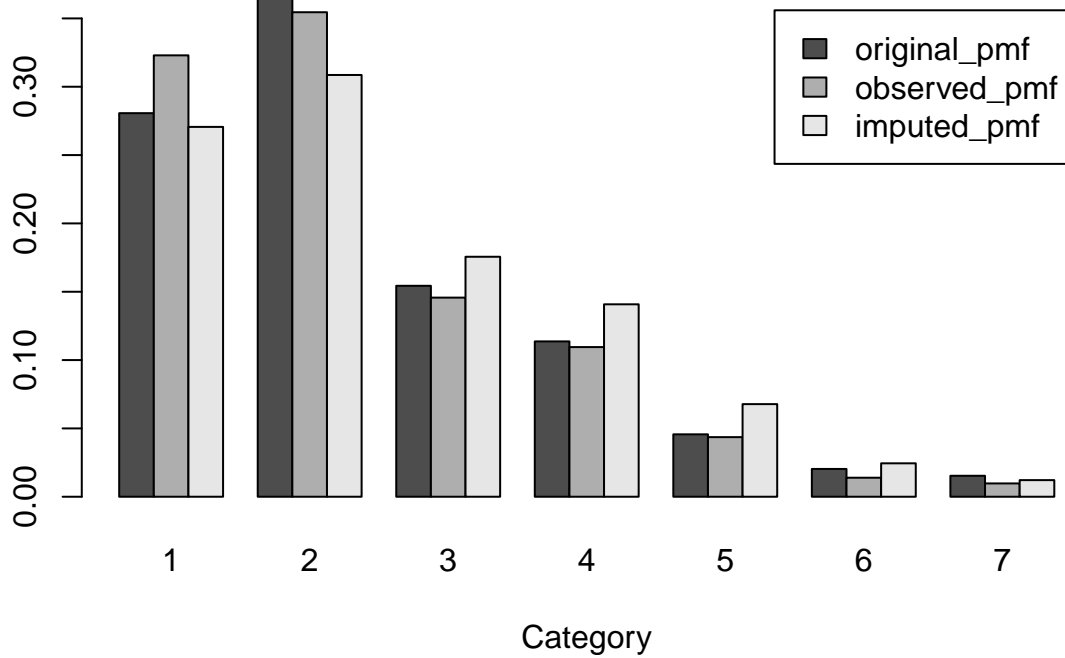
Assess trivariate joint distribution

---

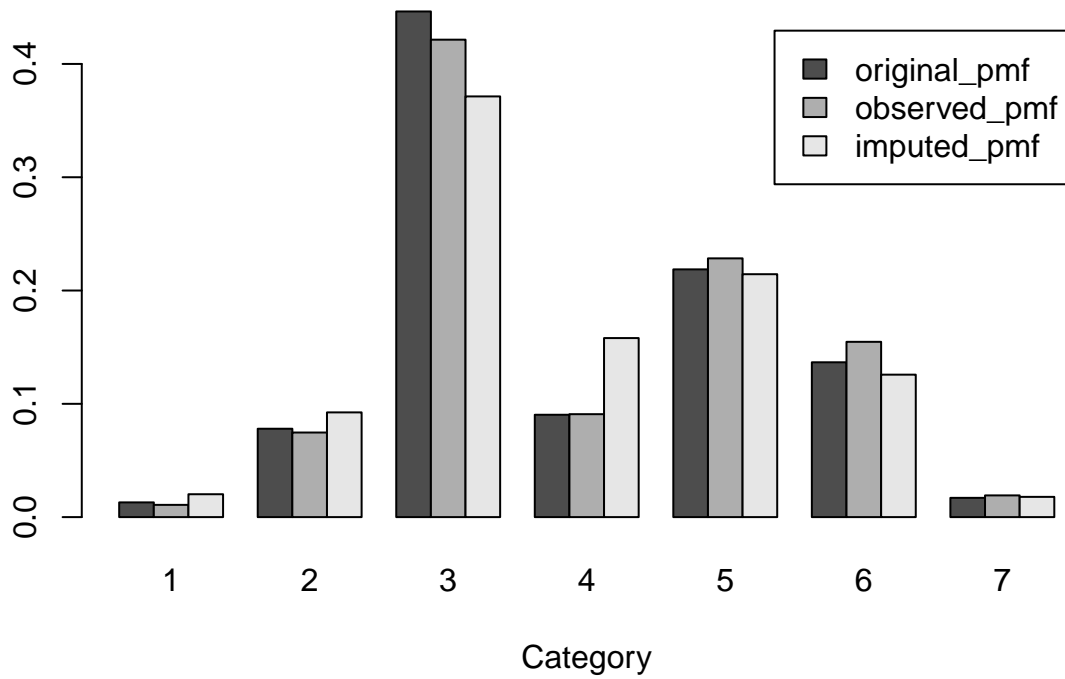
### Blocked Gibbs Sampling Assessment: VEH



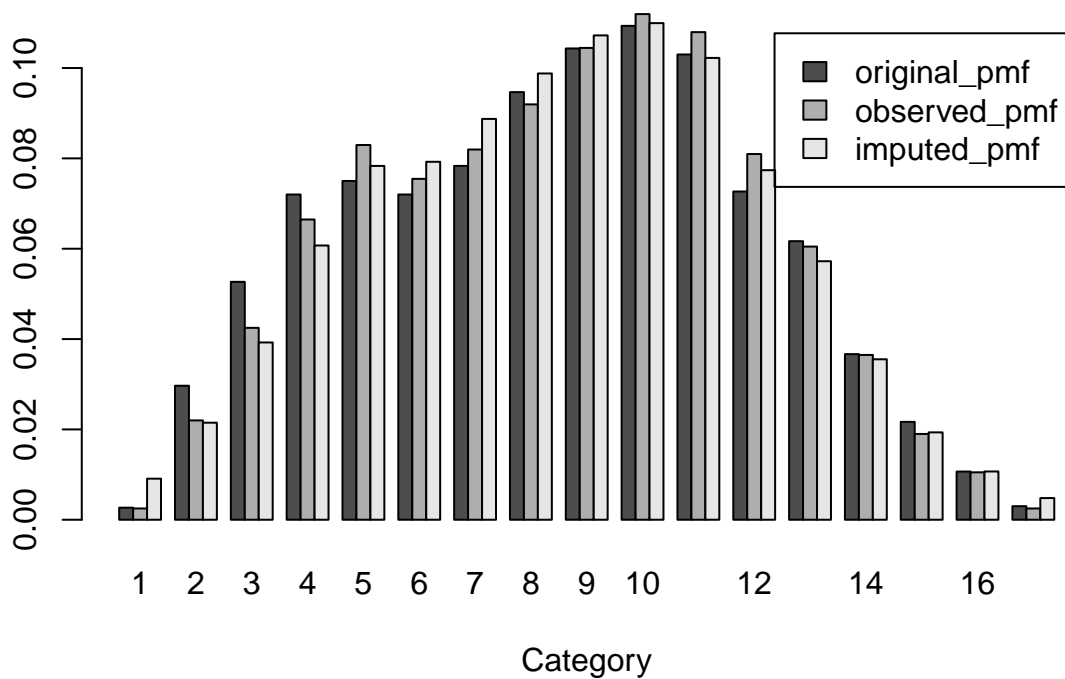
### Blocked Gibbs Sampling Assessment: NP



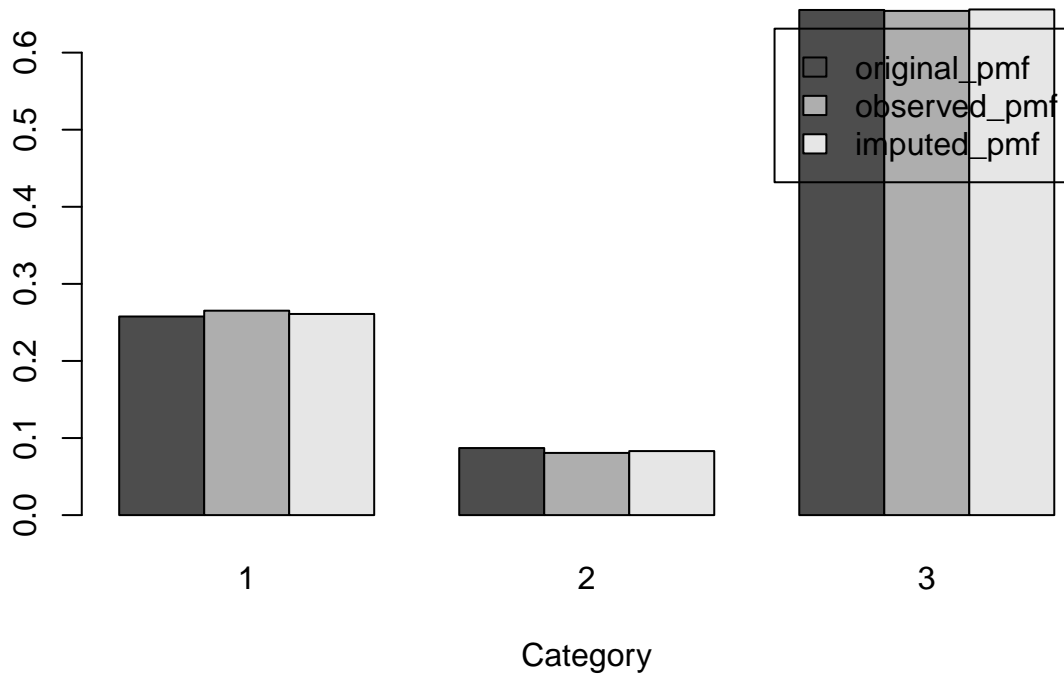
### Blocked Gibbs Sampling Assessment: SCHL



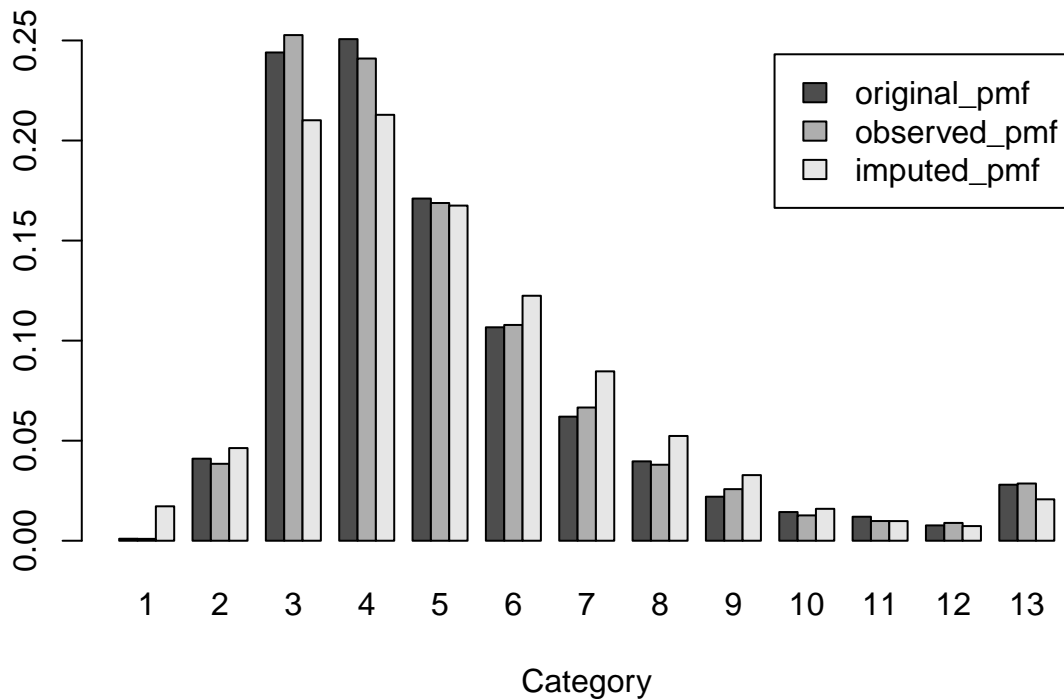
### Blocked Gibbs Sampling Assessment: AGEF



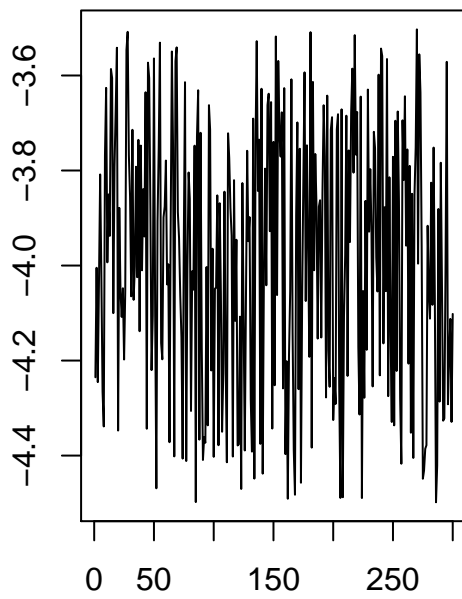
### Blocked Gibbs Sampling Assessment: WKL



### Blocked Gibbs Sampling Assessment: PINCP

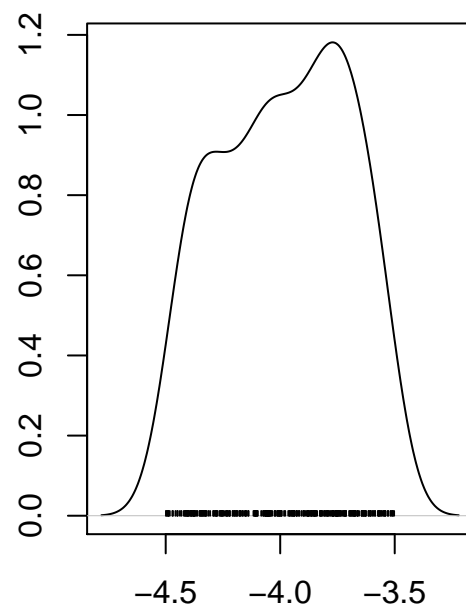


**Trace of var1**

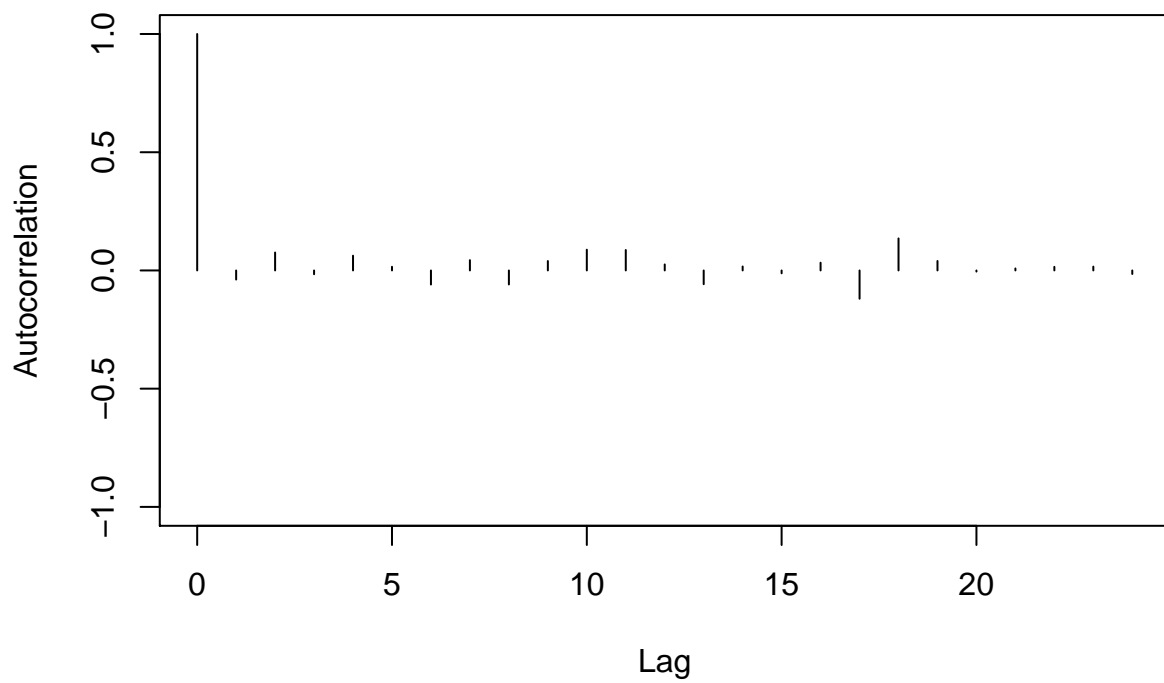


Iterations

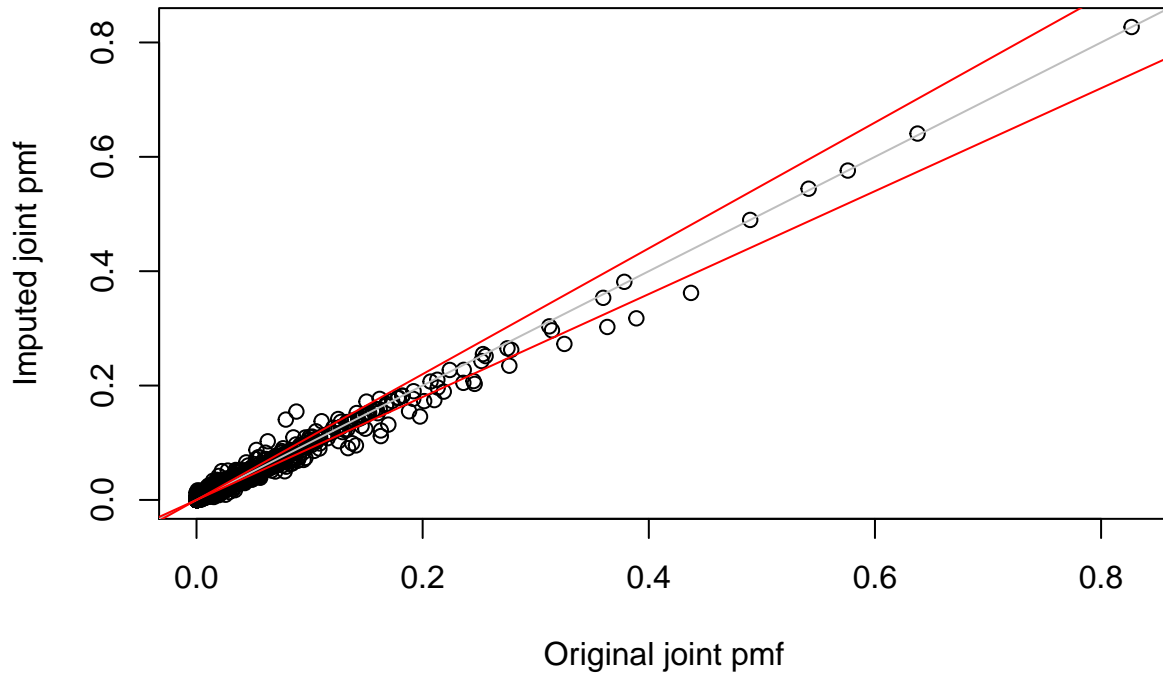
**Density of var1**



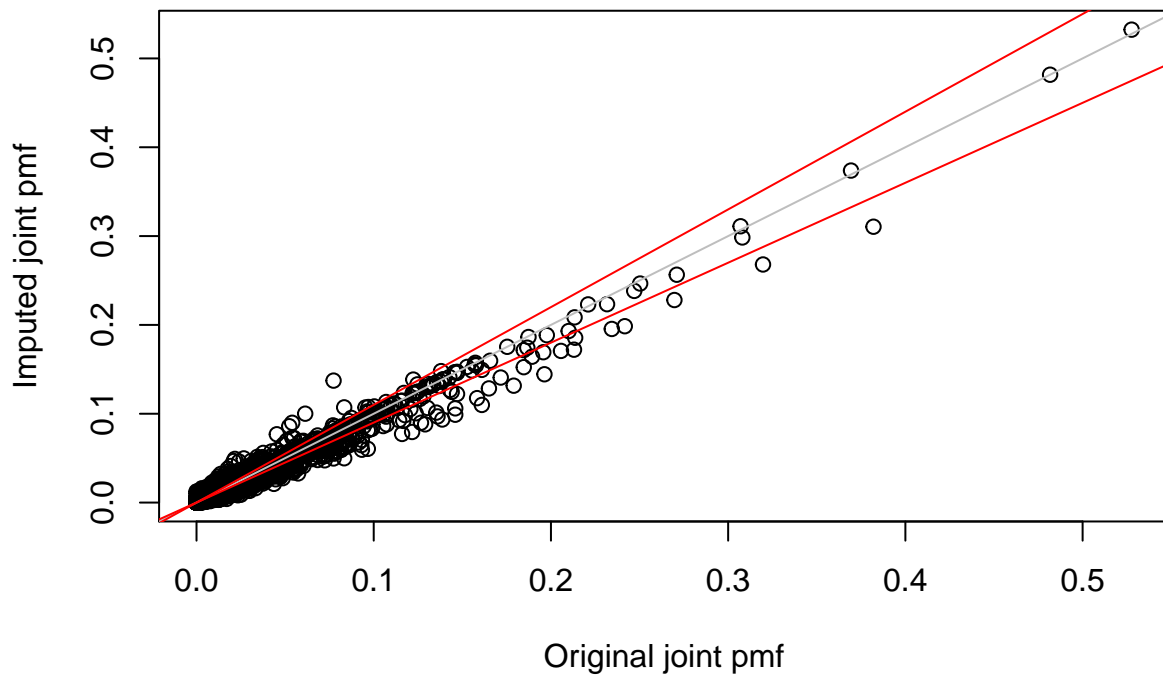
N = 300 Bandwidth = 0.09418



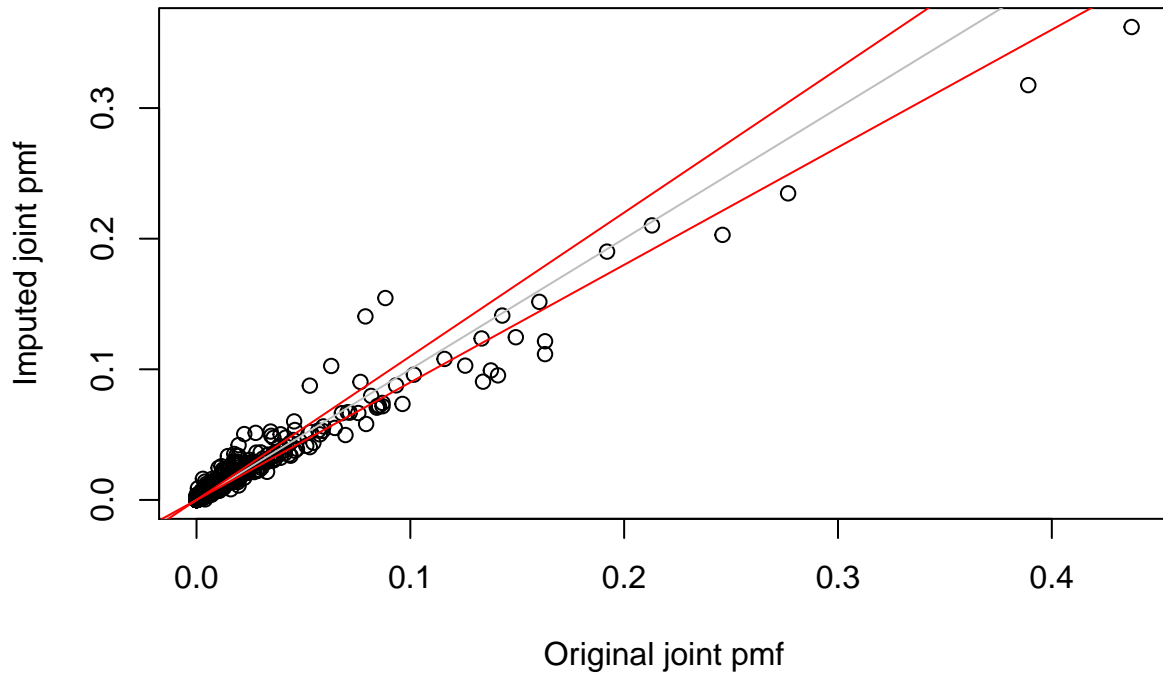
**Bivariate pmf**



**Trivariate pmf**



**Bivariate pmf SCHL**



**Bivariate pmf WKL**

