

# MCAR 30% missing - Probit

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```
# sample MCAR dataset from PUMS
source("../utils/sampleMCAR.R")
n = 3000
missing_col = c(1,3,7,9,10,11)
missing_prob = 0.3
set.seed(0)

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

## 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']]
```

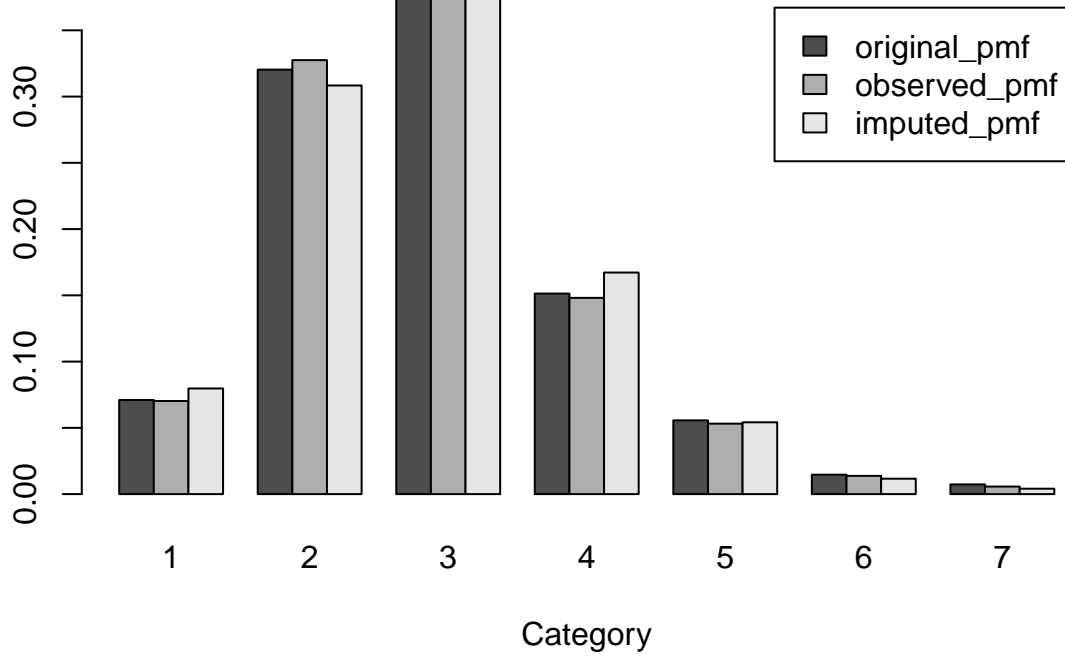
Diagnostics

Assess bivariate joint distribution

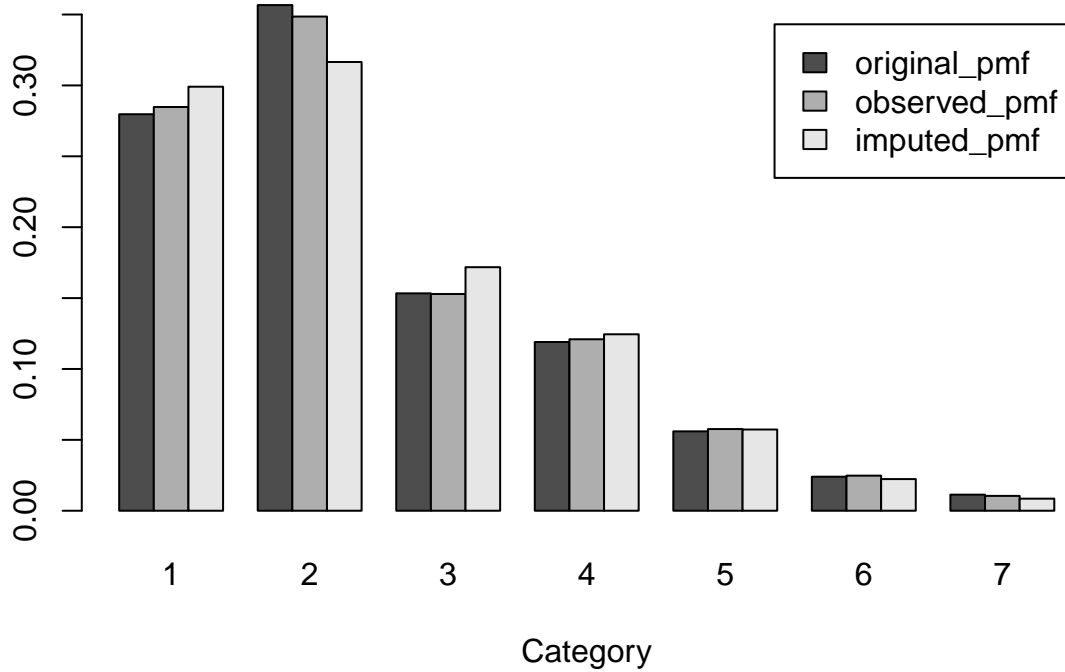
Assess trivariate joint distribution

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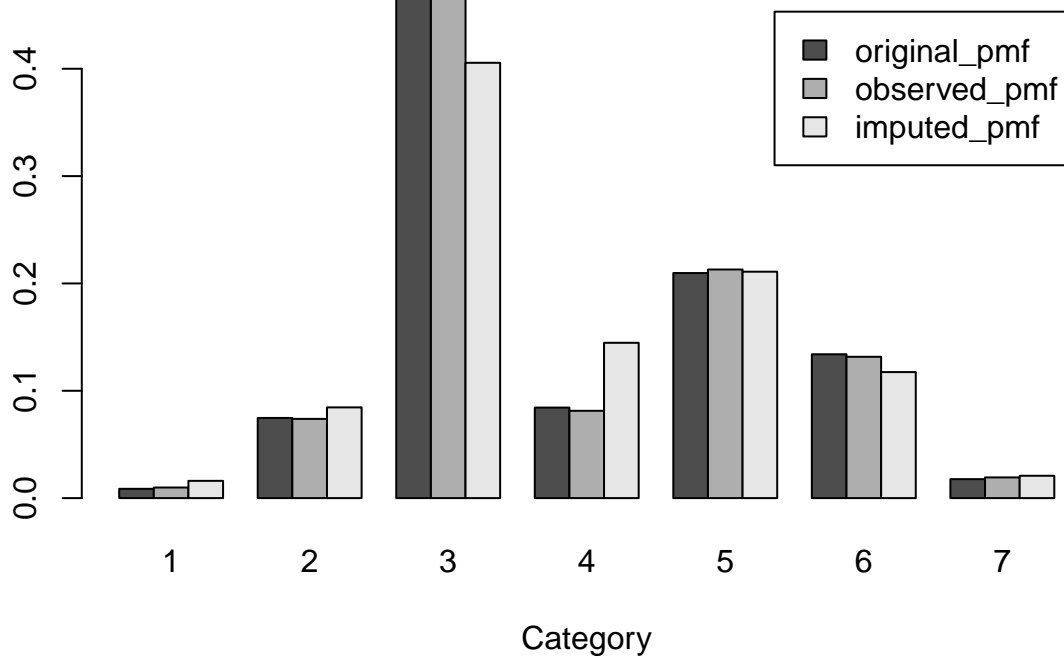
### Blocked Gibbs Sampling Assessment: VEH



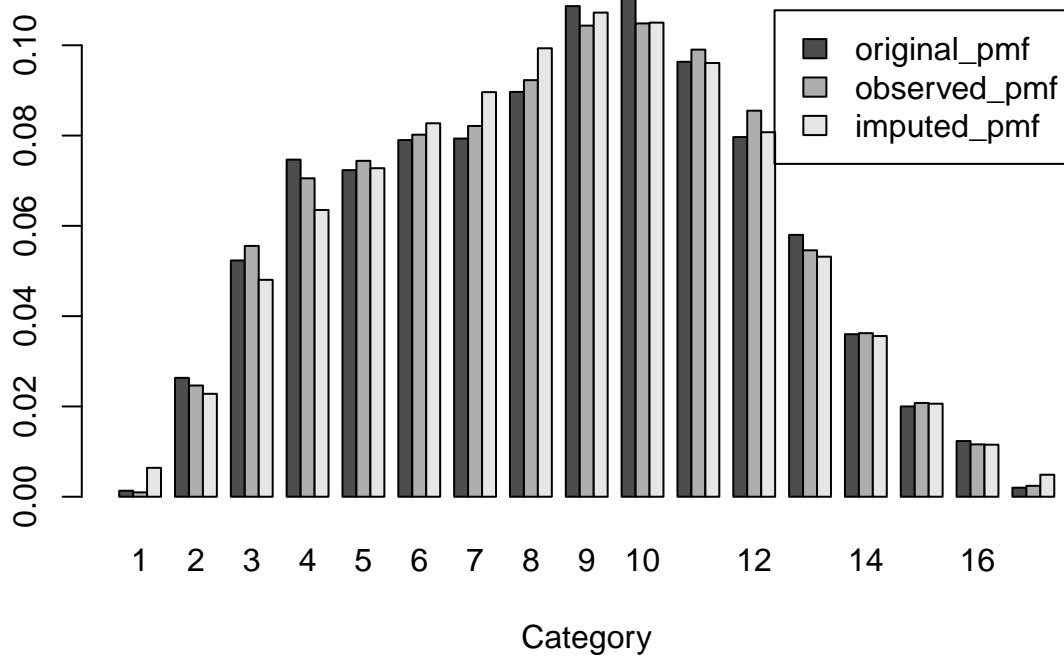
### Blocked Gibbs Sampling Assessment: NP



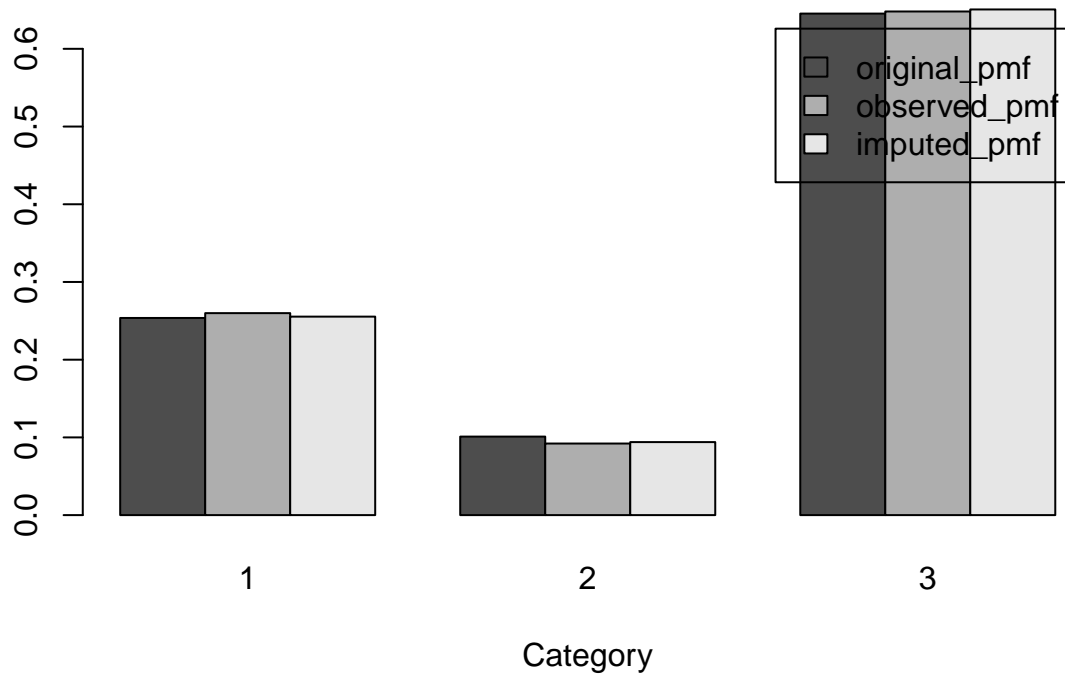
### Blocked Gibbs Sampling Assessment: SCHL



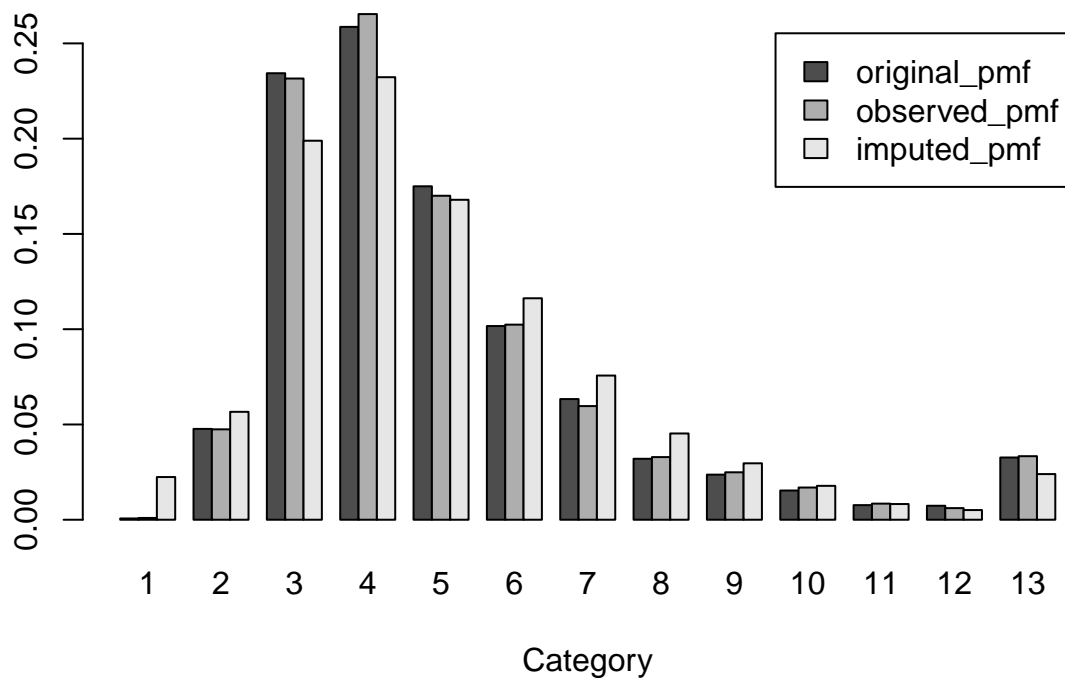
### Blocked Gibbs Sampling Assessment: AGEF



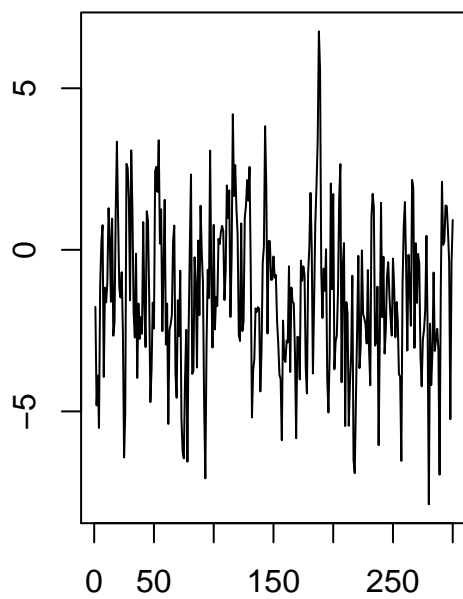
### Blocked Gibbs Sampling Assessment: WKL



### Blocked Gibbs Sampling Assessment: PINCP

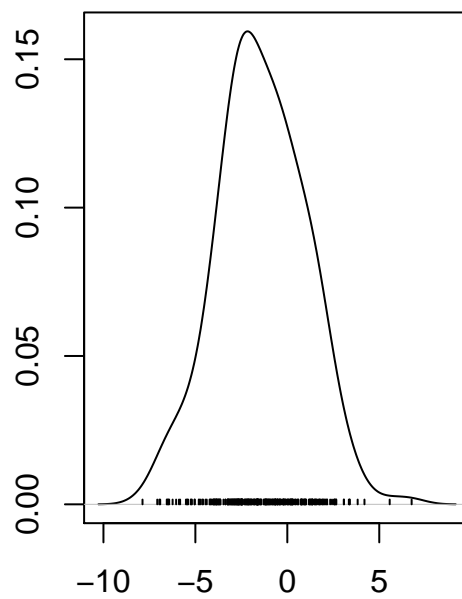


**Trace of var1**

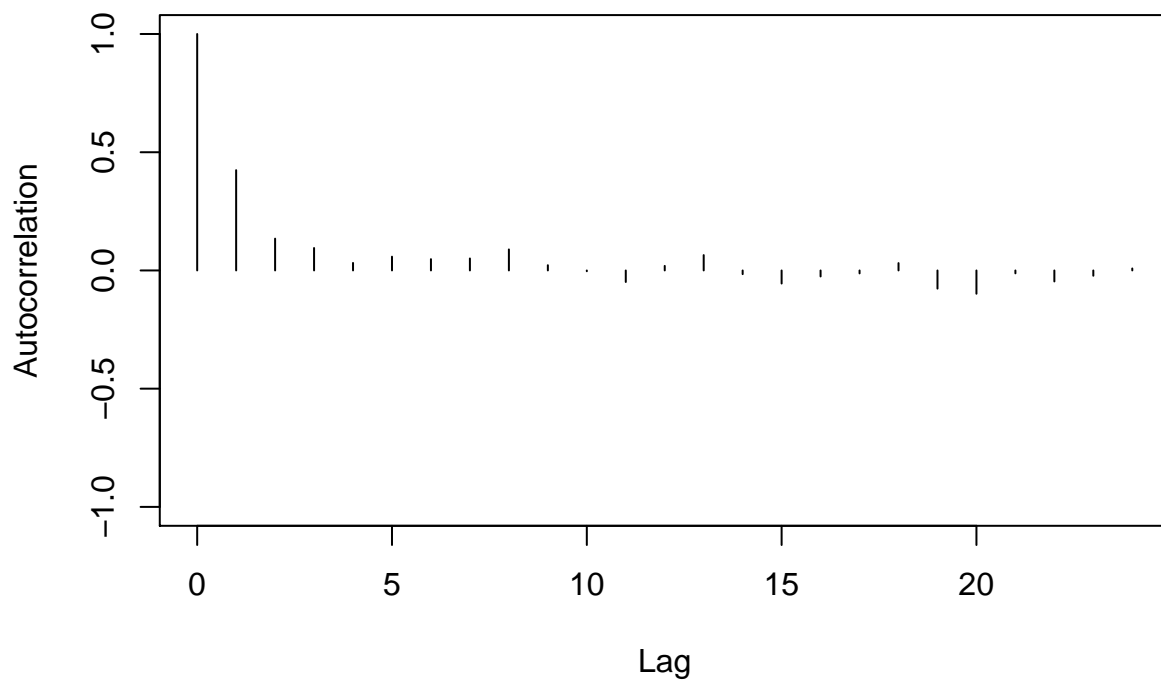


Iterations

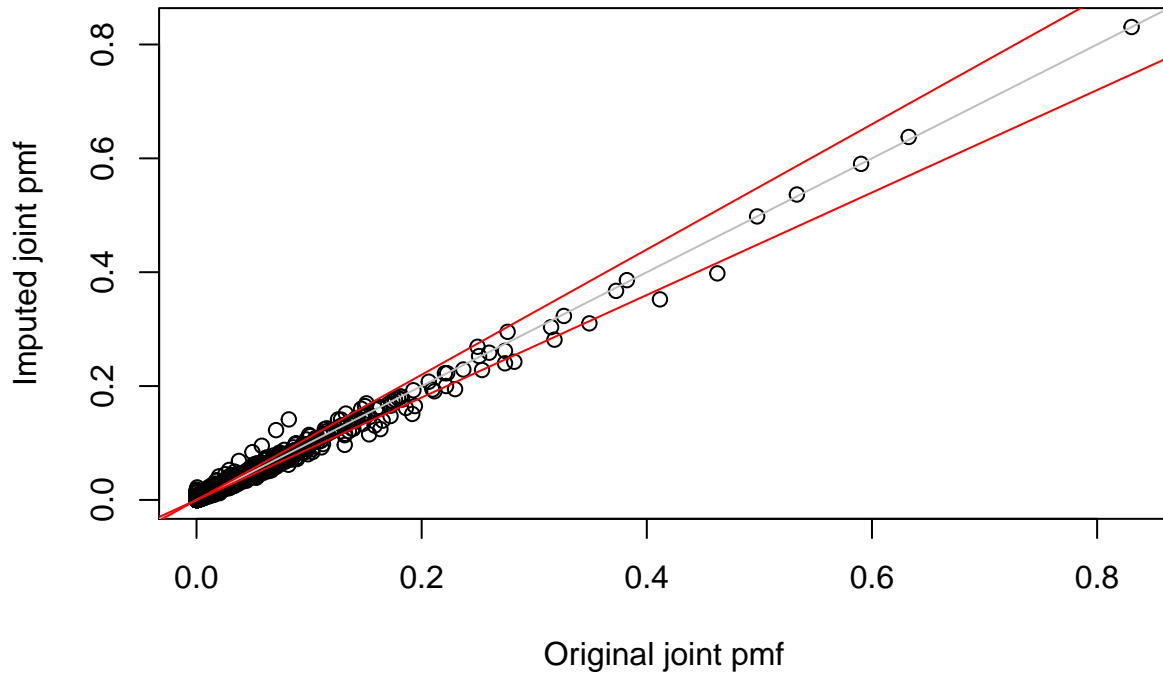
**Density of var1**



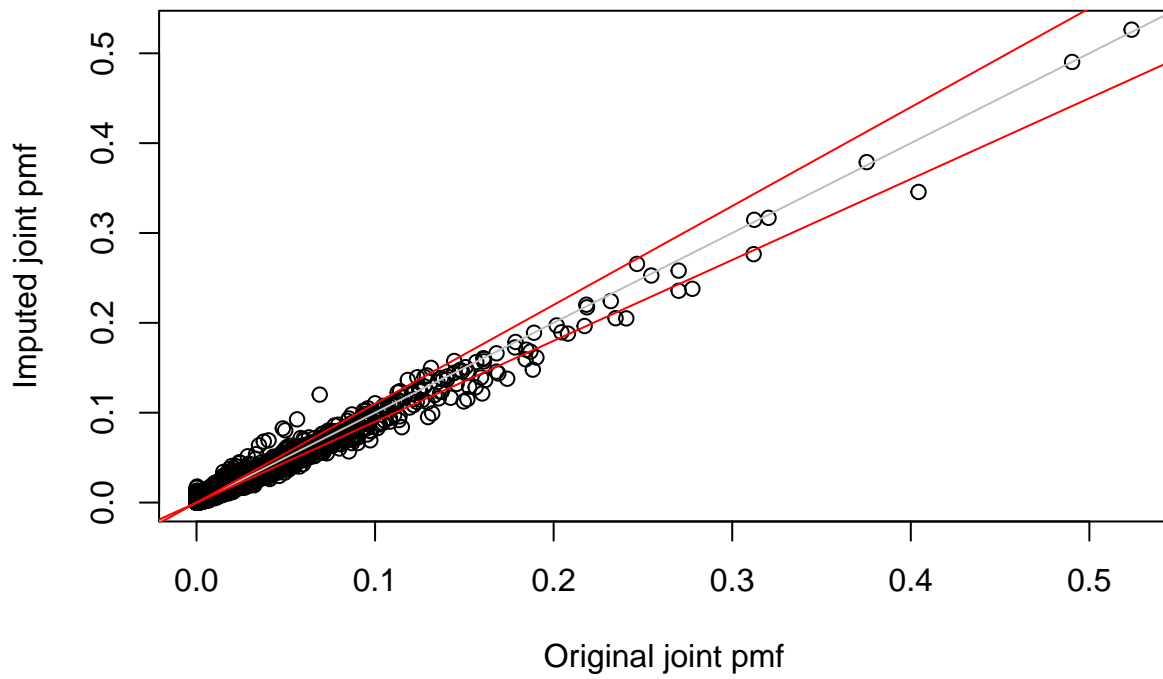
N = 300 Bandwidth = 0.7997



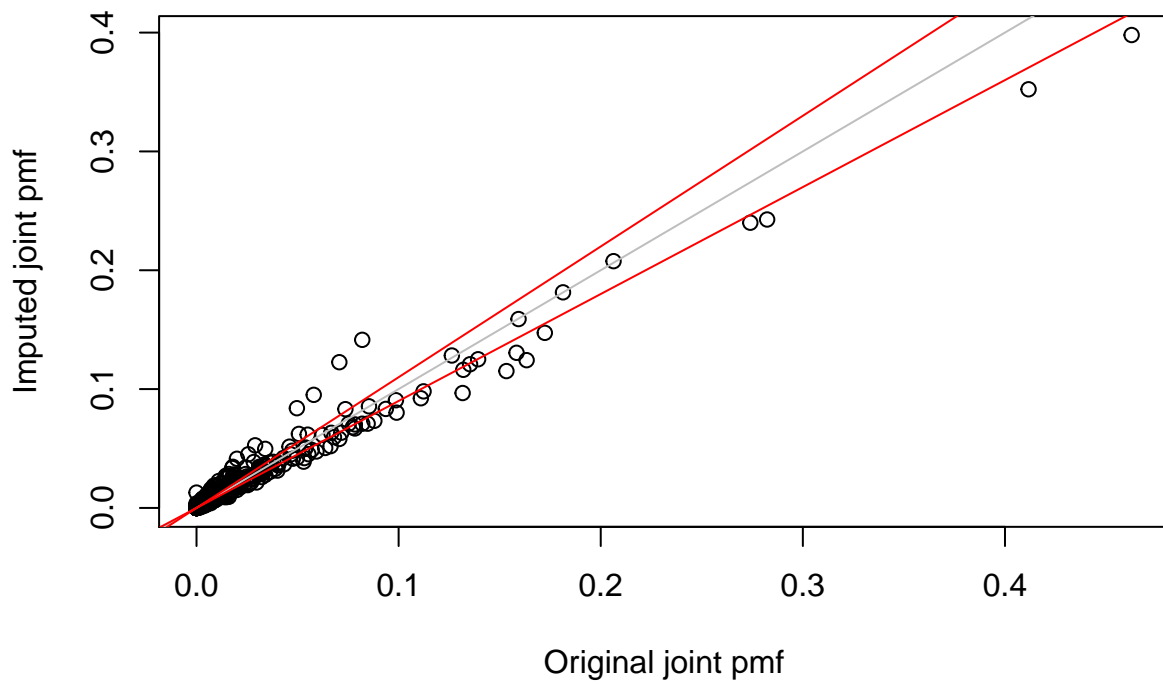
**Bivariate pmf**



**Trivariate pmf**



**Bivariate pmf SCHL**



**Bivariate pmf WKL**

