

Missing Values Alex

Alex

08/02/2022

```
load("cancer.rdata")
summary(cancer)
```

```
## Geography incidenceRate medIncome binnedInc
## Length:3047 Min. : 201.3 Min. : 22640 [22640, 34218.1] : 306
## Class :character 1st Qu.: 420.3 1st Qu.: 38883 (45201, 48021.6] : 306
## Mode :character Median : 453.5 Median : 45207 (54545.6, 61494.5] : 306
## Mean : 448.3 Mean : 47063 (42724.4, 45201] : 305
## 3rd Qu.: 480.9 3rd Qu.: 52492 (48021.6, 51046.4] : 305
## Max. :1206.9 Max. :125635 (51046.4, 54545.6] : 305
## (Other) :1214
## povertyPercent MedianAgeMale MedianAgeFemale AvgHouseholdSize
## Min. : 3.20 Min. :22.40 Min. :22.30 Min. :0.0221
## 1st Qu.:12.15 1st Qu.:36.35 1st Qu.:39.10 1st Qu.:2.3700
## Median :15.90 Median :39.60 Median :42.40 Median :2.5000
## Mean :16.88 Mean :39.57 Mean :42.15 Mean :2.4797
## 3rd Qu.:20.40 3rd Qu.:42.50 3rd Qu.:45.30 3rd Qu.:2.6300
## Max. :47.40 Max. :64.70 Max. :65.70 Max. :3.9700
##
## PercentMarried PctEmployed16_Over PctUnemployed16_Over PctPrivateCoverage
## Min. :23.10 Min. :17.60 Min. : 0.400 Min. :22.30
## 1st Qu.:47.75 1st Qu.:48.60 1st Qu.: 5.500 1st Qu.:57.20
## Median :52.40 Median :54.50 Median : 7.600 Median :65.10
## Mean :51.77 Mean :54.15 Mean : 7.852 Mean :64.35
## 3rd Qu.:56.40 3rd Qu.:60.30 3rd Qu.: 9.700 3rd Qu.:72.10
## Max. :72.50 Max. :80.10 Max. :29.400 Max. :92.30
## NA's :152
## PctEmpPrivCoverage PctPublicCoverage PctBlack PctMarriedHouseholds
## Min. :13.5 Min. :11.20 Min. : 0.0000 Min. :22.99
## 1st Qu.:34.5 1st Qu.:30.90 1st Qu.: 0.6207 1st Qu.:47.76
## Median :41.1 Median :36.30 Median : 2.2476 Median :51.67
## Mean :41.2 Mean :36.25 Mean : 9.1080 Mean :51.24
## 3rd Qu.:47.7 3rd Qu.:41.55 3rd Qu.:10.5097 3rd Qu.:55.40
## Max. :70.7 Max. :65.10 Max. :85.9478 Max. :78.08
##
## Edu18_24 deathRate
## Min. :1.487 Min. : 59.7
## 1st Qu.:2.206 1st Qu.:161.2
## Median :2.340 Median :178.1
## Mean :2.347 Mean :178.7
## 3rd Qu.:2.486 3rd Qu.:195.2
```

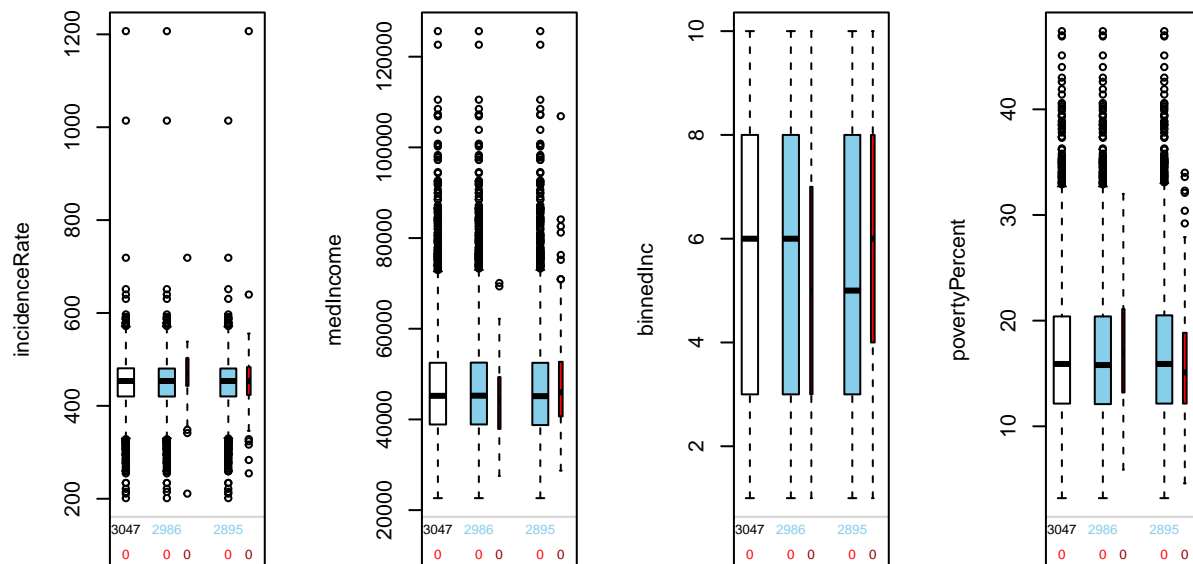
```
## Max.      :3.307   Max.      :362.8
##
```

There are some missing values in PctEmployed16_Over which need to be checked. Before that is checked though we should note that the outliers in AvgHouseholdSize should be treated as missing values as they are wrongly inputted (reference).

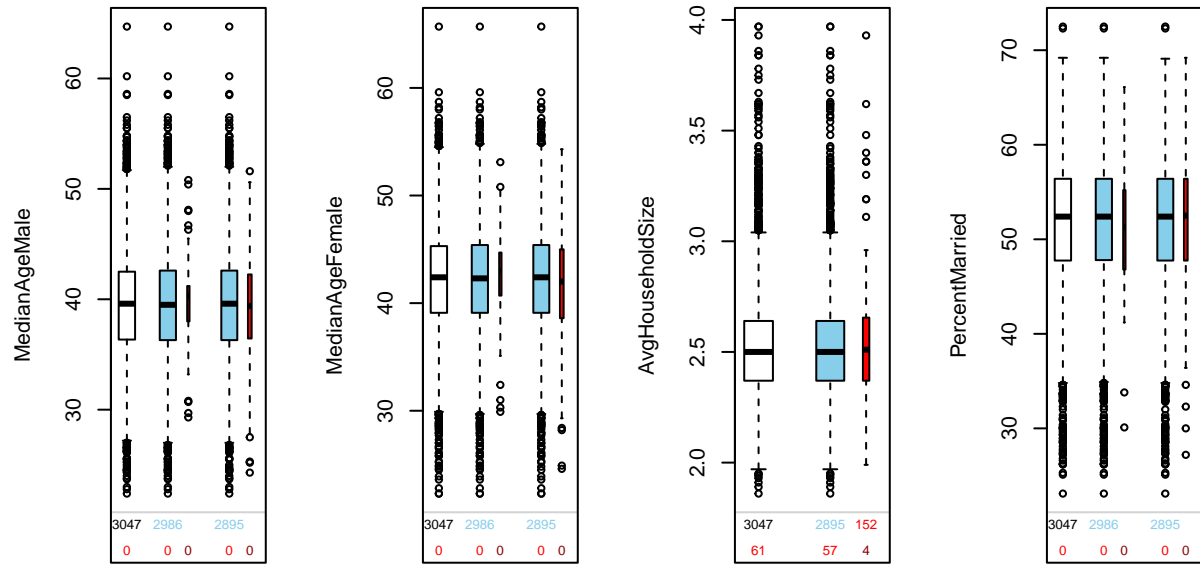
```
cancer1 <- cancer
cancer1$AvgHouseholdSize[cancer1$AvgHouseholdSize < 0.5] <- NA
```

Now I use the VIM package and the pbox() function to show that the missing data are all MCAR and can thus be easily dealt with.

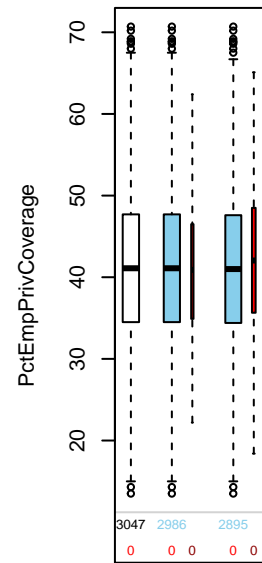
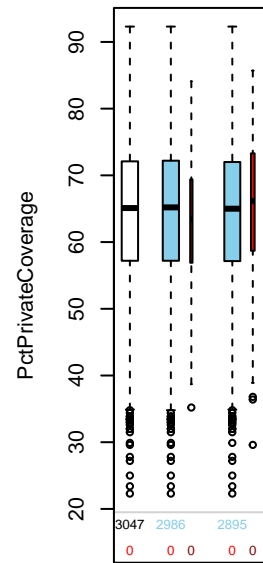
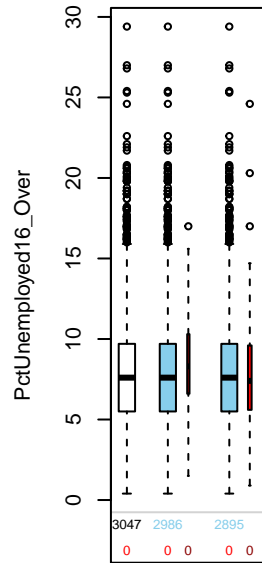
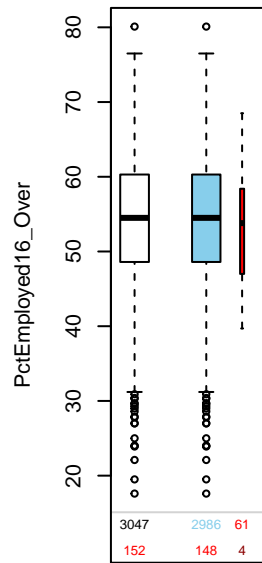
```
library(VIM)
par(mfrow = c(1,4))
for(i in 2:5){
  pbox(cancer1, pos = i)
}
```



```
par(mfrow = c(1,4))
for(i in 6:9){
  pbox(cancer1, pos = i)
}
```



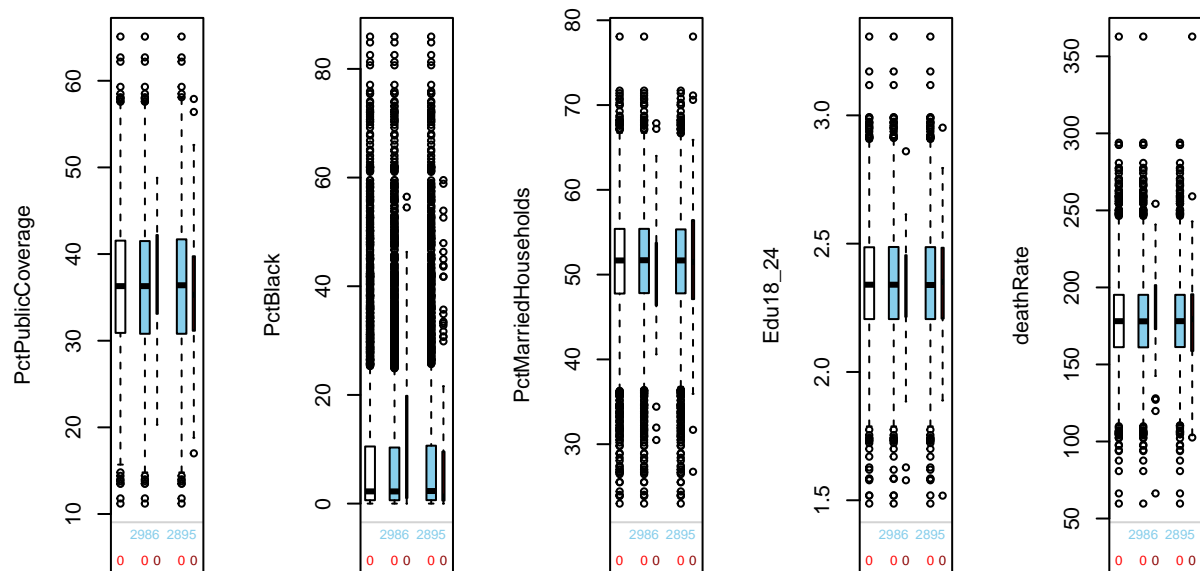
```
par(mfrow = c(1,4))
for(i in 10:13){
  pbox(cancer1, pos = i)
}
```



```

par(mfrow = c(1,5))
for(i in 14:18){
  pbox(cancer1, pos = i)
}

```



From the pbox plots we have evidence that the missing values interrogated are MCAR. Since the proportion of rows with missing values is small I recommend simply deleting the rows with missing values as it should not affect the validity of our analysis on the whole dataset as the smaller sample is still representative.

```
cancer2 <- na.omit(cancer1)
```

Our other option is to use the reference to fill in all our other missing values. Despite this being possible it is also un-necessary as the data is MCAR and won't significantly affect our analysis.