

Exploring PUMS for NJ

Use shapiro test to assess normality

```
shapiro = function(x) {  
  return(shapiro.test(x)$p.value)  
}  
  
ks = function(x) {  
  return(ks.test(x, "pnorm", mean = mean(x), sd = sd(x))$p.value)  
}
```

Read in NJ Pums data

```
pumsnj = read.csv("psam_h34.csv")  
  
# Some numbers for house values  
sum(!is.na(pumsnj$VALP)) #24423 non NA house values in NJ
```

```
## [1] 24423
```

```
mean(sapply(split(pumsnj, pumsnj$PUMA), function(x) {  
  sum(is.na(x$VALP))  
})) #average 211 people per PUMA in NJ
```

```
## [1] 211.4521
```

```
length((sapply(split(pumsnj, pumsnj$PUMA), function(x) {  
  sum(is.na(x$VALP))  
}))) #73 different PUMAS in NJ, but 21 counties
```

```
## [1] 73
```

```
# Some numbers for household income  
sum(!is.na(pumsnj$HINCP)) #33660 non NA house values in NJ
```

```
## [1] 33660
```

```
mean(sapply(split(pumsnj, pumsnj$PUMA), function(x) {  
  sum(is.na(x$HINCP))  
})) #average 85 people per PUMA in NJ
```

```
## [1] 84.91781
```

```
length((sapply(split(pumsnj, pumsnj$PUMA), function(x) {  
  sum(is.na(x$HINCP))  
}))) #73 different PUMAs in NJ, but 21 counties
```

```
## [1] 73
```

Normality within PUMAs? NO

```
# Examine normality of house values within PUMAs  
pvals = sapply(split(log(pumsnj$VALP), pumsnj$PUMA), shapiro)  
mean(pvals > 0.05, na.rm = T) #all rejected
```

```
## [1] 0
```

```
# Examine normality of household incomes within PUMAs  
pvals = sapply(split(log(pumsnj$HINCP + 1), pumsnj$PUMA), shapiro) # some are 0 so offset by 1 then log
```

```
## Warning in log(pumsnj$HINCP + 1): NaNs produced
```

```
mean(pvals > 0.05, na.rm = T) #all rejected
```

```
## [1] 0
```

Link PUMAs to FIPS in NJ (for now just assume PUMAs are smaller than FIPS)

Using equivalency at https://www2.census.gov/geo/pdfs/reference/puma/2010_PUMA_Equivalency_Summary_Levels.pdf

```
library(gdata)
```

```
## gdata: read.xls support for 'XLS' (Excel 97-2004) files ENABLED.
```

##

```
## gdata: read.xls support for 'XLSX' (Excel 2007+) files ENABLED.
```

##

```
## Attaching package: 'gdata'
```

```
## The following object is masked from 'package:stats':
```

##

```
##      nobs
```

```
## The following object is masked from 'package:utils':
```

##

```
##      object.size
```

```
## The following object is masked from 'package:base':
```

##

```
##      startsWith
```

```
equiv <- read.delim("PUMSEQ10_34.txt", stringsAsFactor = FALSE,
  header = F)
head(equiv)
```

##

## 1	795340177979500101	112887	62119Atlantic County (East)
## 2	79634017797950010100100882270	112887	62119Atlantic County (part)
## 3	79734017797950010100100882270000000000000000	0	0County subdivisions
## 4	798340177979500101001008822700000000000000009999999999999999	0	0County subdivisions
## 5	799340177979500101001008822700000000000000009999999999999999010101	0	0Census Tract 101.01
## 6	799340177979500101001008822700000000000000009999999999999999010102	0	0Census Tract 101.02

Messy. example: 796 34 017797950 0101 001 00882270 796

```
# state random puma
```

```
codes = unlist(lapply(strsplit(equiv[, 1][which(startsWith(equiv[,
1], "796") == T)], "\\s+"), function(x) {
  x[1]
}))
```

Extract PUMAs

```
pumas = as.numeric(substring(codes, 15, 18))
```

```

# Extract FIPS
fips = paste(substring(codes, 4, 5), substring(codes, 19, 21),
             sep = "")
# Dataframe matching PUMAs to FIPS
puma.fips.match = data.frame(pumas, fips)

# Add fips to previous dataframe
pumsnj$fips = puma.fips.match$fips[match(pumsnj$PUMA, puma.fips.match$pumas)]
length(unique(pumsnj$fips)) #2 counties of 21 excluded from pums sample?? Is this because multiple sma

## [1] 19

```

```

# Number of individuals sampled from each fips
sapply(split(log(pumsnj$VALP), pumsnj$fips), length)

```

```

## 34001 34003 34005 34007 34011 34013 34015 34017 34019 34021 34023 34025 34027
## 2585 3700 2324 2281 1210 3230 1360 2678 740 1538 3542 2917 2168
## 34029 34031 34035 34037 34039 34041
## 3098 1710 1356 811 2020 591

```

Normality within FIPS?? NO

```

# Examine normality of house values within fips
pvals = sapply(split(log(pumsnj$VALP), pumsnj$fips), shapiro)
mean(pvals > 0.05, na.rm = T)

```

```
## [1] 0
```

```

# Examine normality of household incomes within fips
pvals = sapply(split(log(pumsnj$HINCP + 1), pumsnj$fips), shapiro) # some are 0 so offset by 1 then lo.

```

```
## Warning in log(pumsnj$HINCP + 1): NaNs produced
```

```
mean(pvals > 0.05, na.rm = T)
```

```
## [1] 0
```

Visually assess normality

```

# Within PUMA
par(mfrow = c(2, 2))
sapply(split(pumsnj, pumsnj$PUMA), function(x) {
  hist(log(x$VALP))
})

```

Frequency

100

0

8 10 12 14 16

$\log(x\$VALP)$

Frequency

150

0

8 10 12 14 16

$\log(x\$VALP)$

Frequency

log(x\$VALP)

log(x\$VALP) Bin	Frequency
[7, 8)	0
[8, 9)	0
[9, 10)	0
[10, 11)	0
[11, 12)	10
[12, 13)	120
[13, 14)	50
[14, 15)	0
[15, 16)	5

Frequency

100

0

6 8 10 12 14

$\log(x\$SVALP)$

Frequency

log(x\$VALP)

Histogram of $\log(x\$VALP)$ for the 'x' variable. The x-axis is labeled $\log(x\$VALP)$ and ranges from 6 to 16. The y-axis is labeled 'Frequency' and ranges from 0 to 100. The distribution is highly right-skewed, with a peak frequency of approximately 150 for the bin $[12, 13)$.

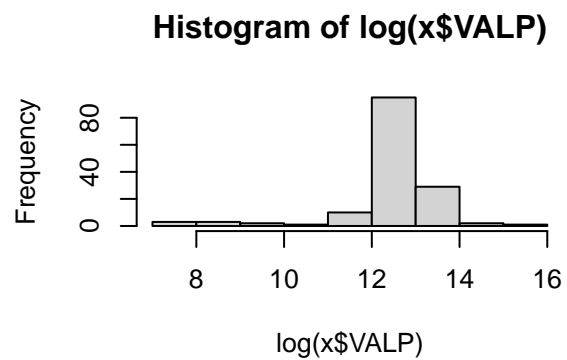
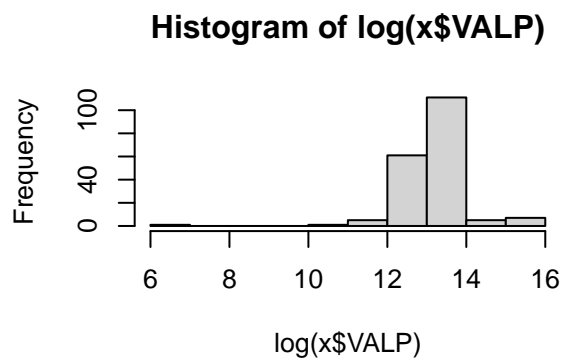
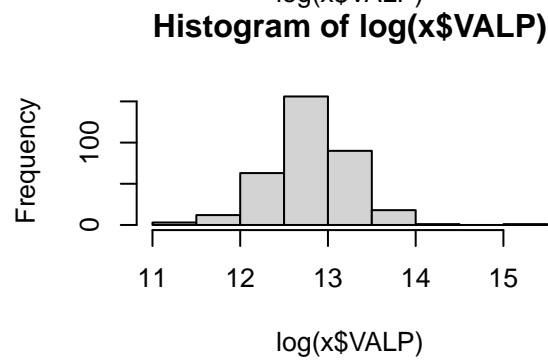
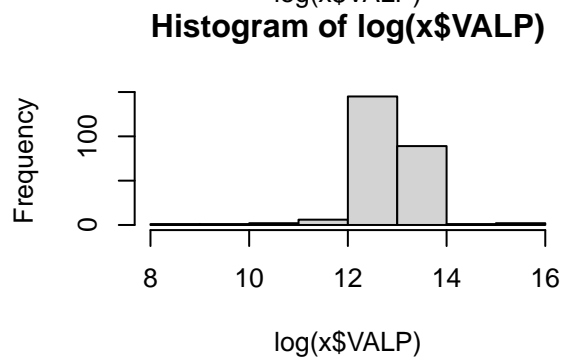
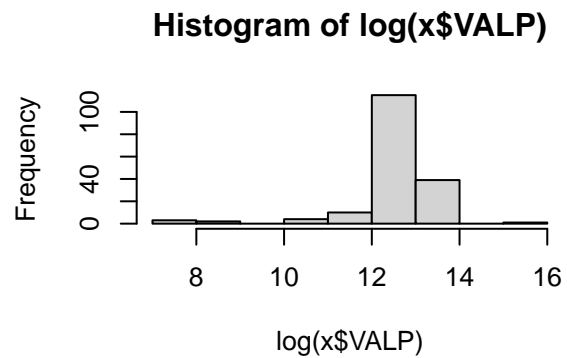
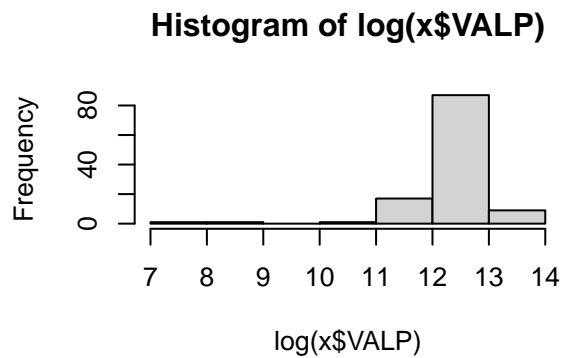
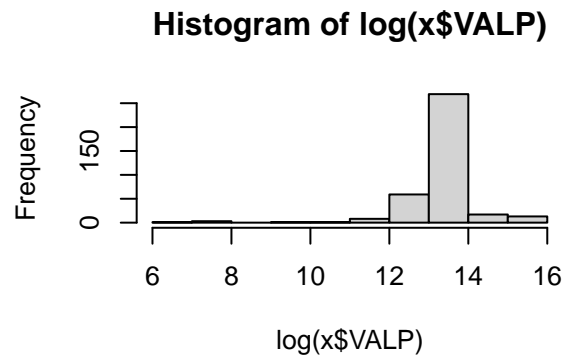
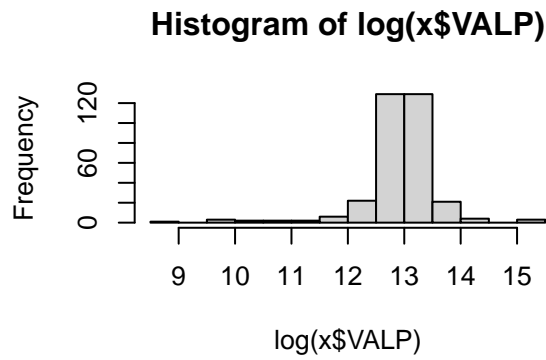
Frequency

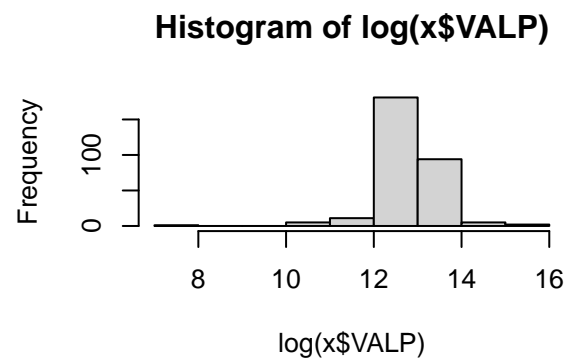
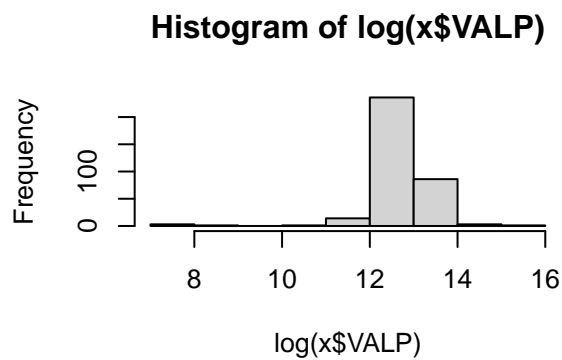
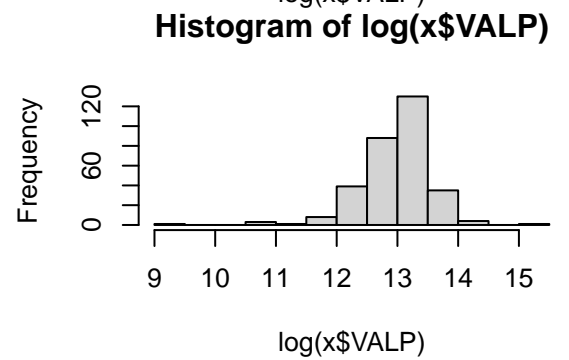
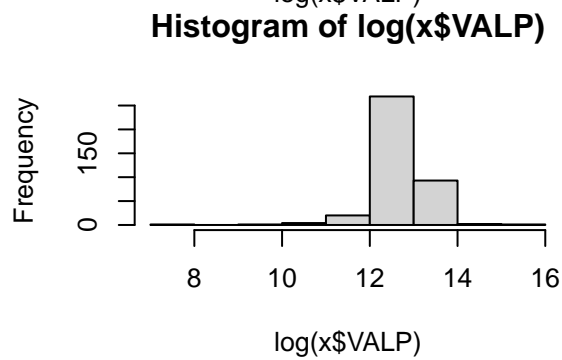
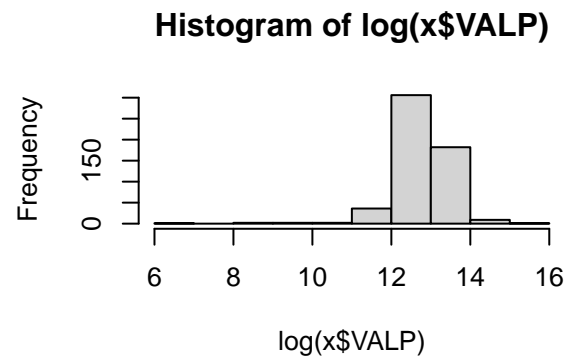
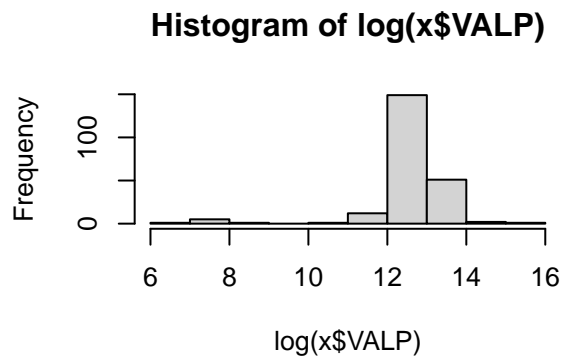
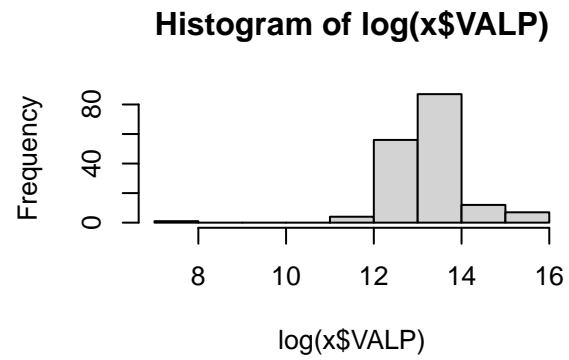
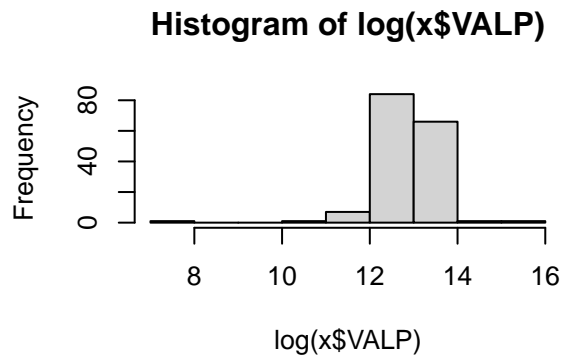
150

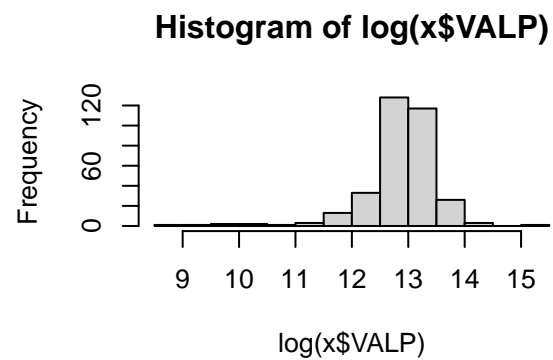
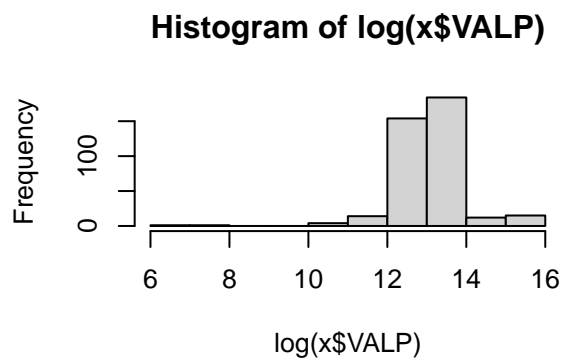
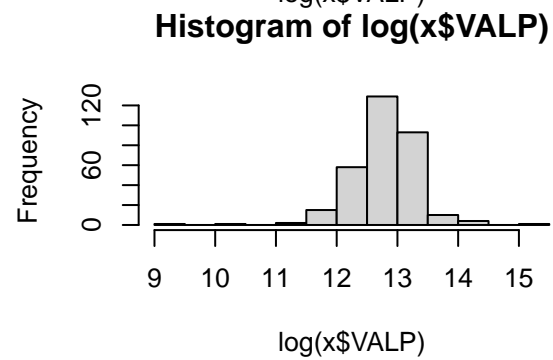
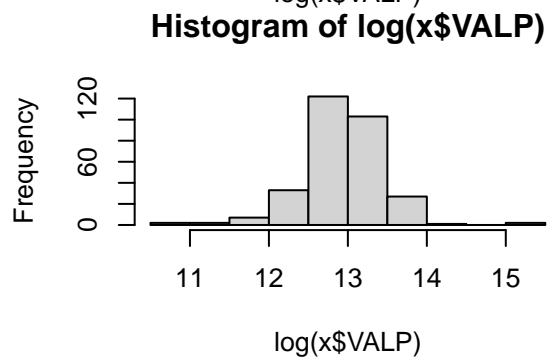
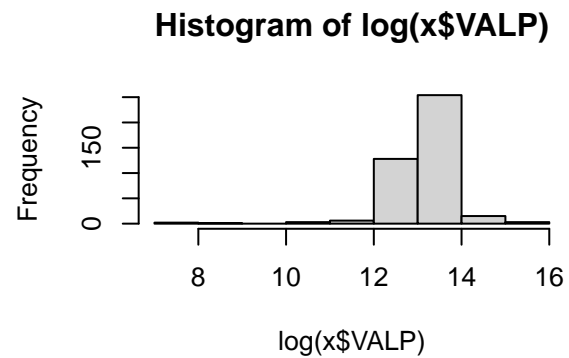
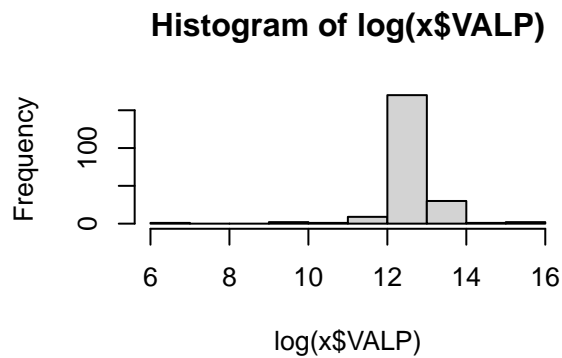
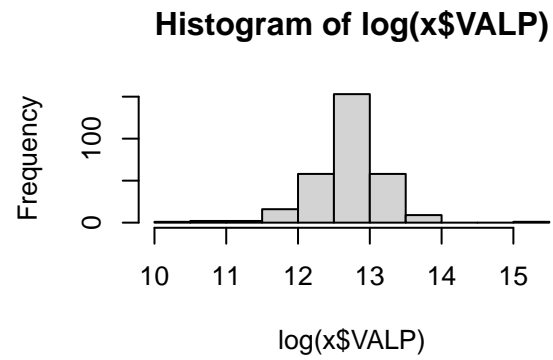
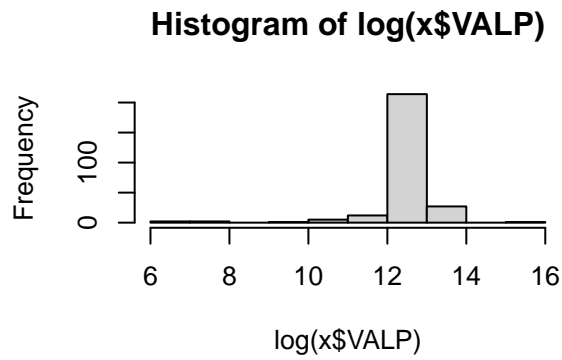
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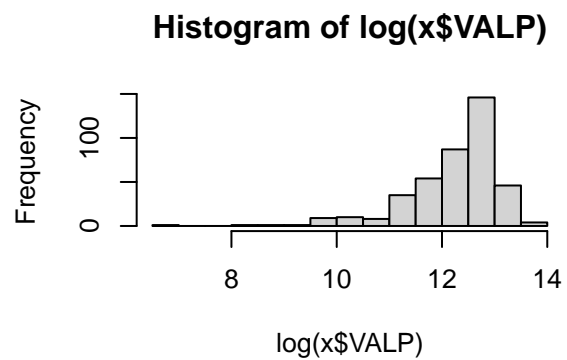
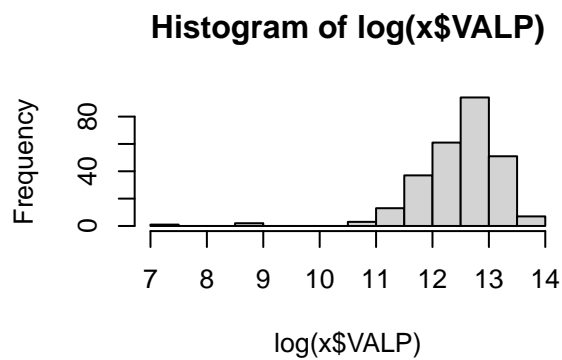
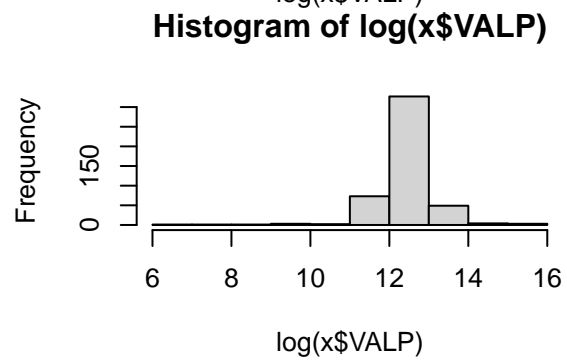
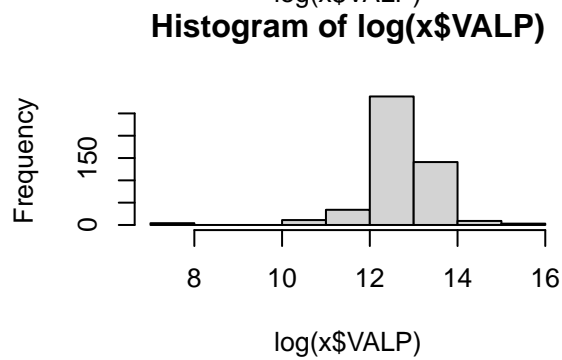
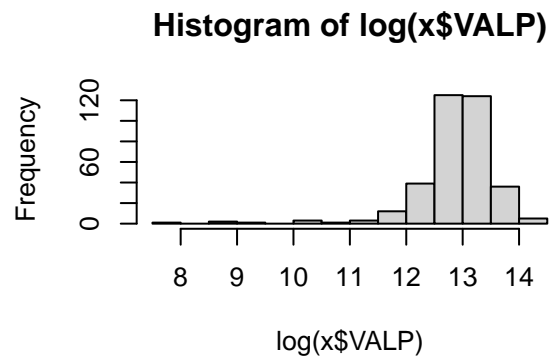
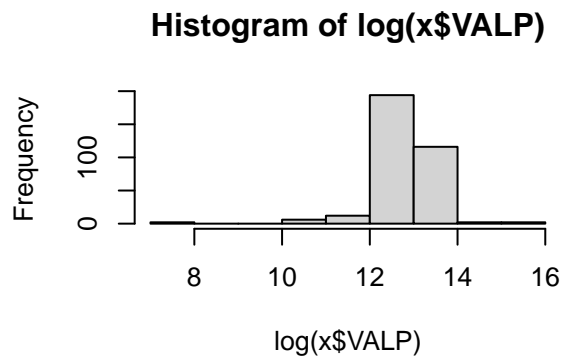
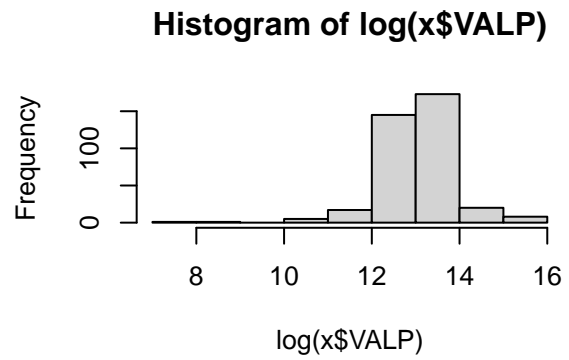
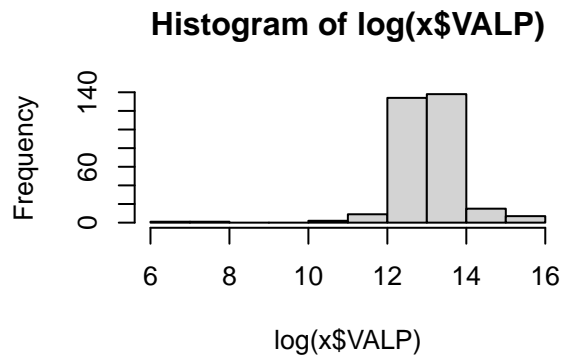
8 10 12 14 16

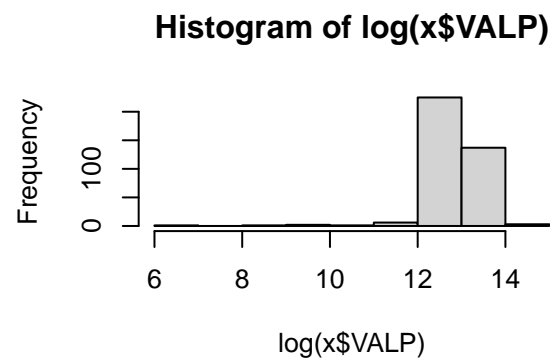
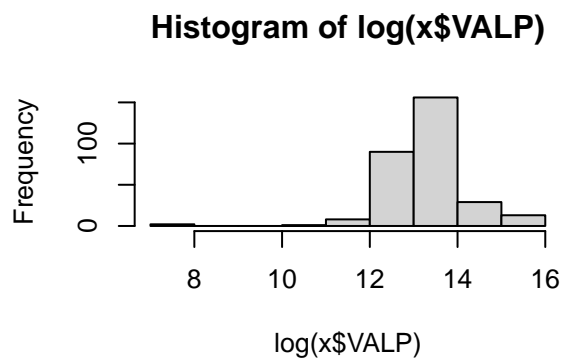
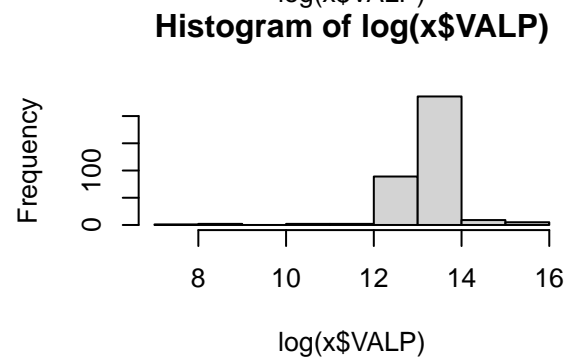
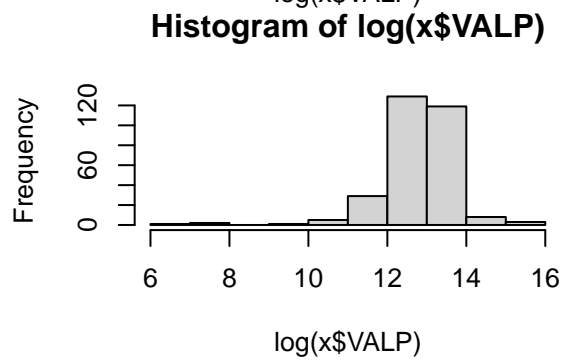
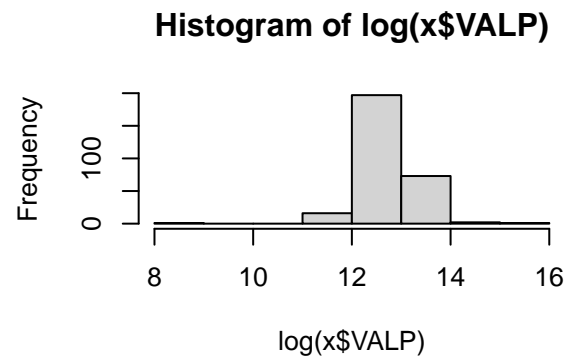
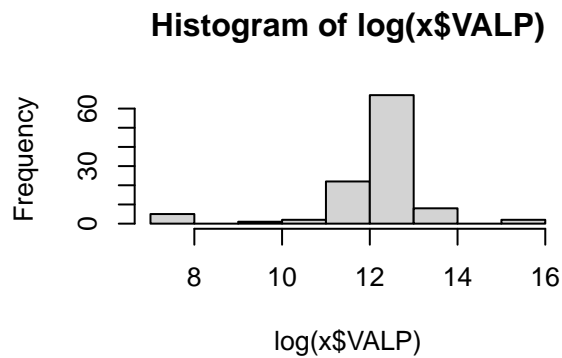
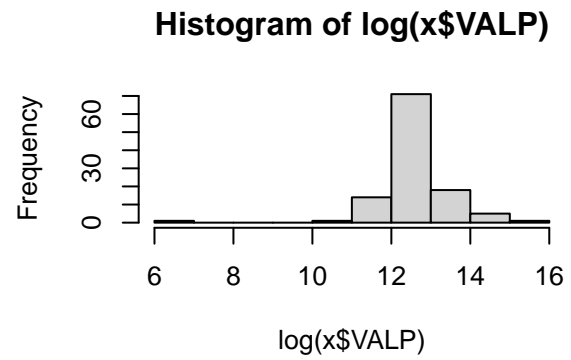
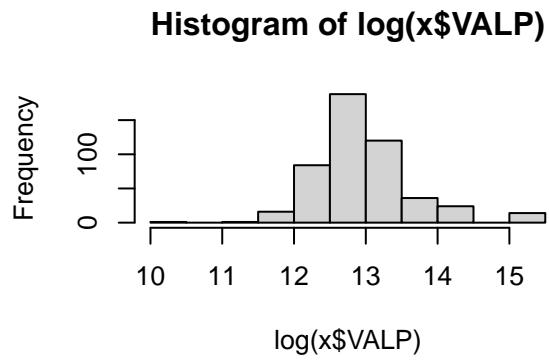
$\log(x\$SVALP)$

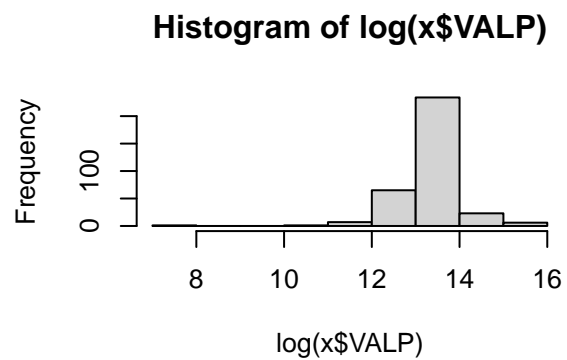
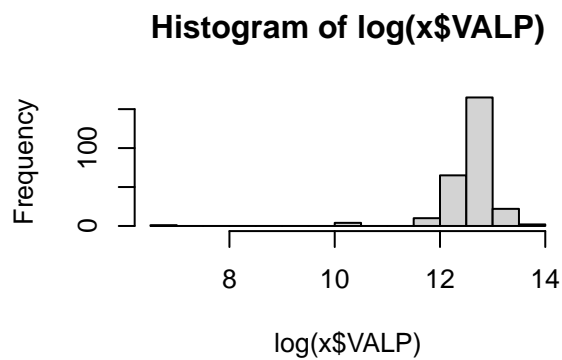
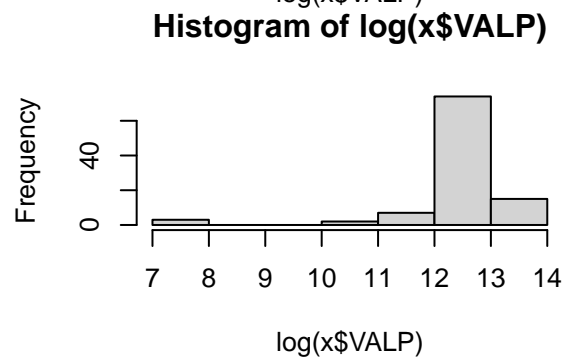
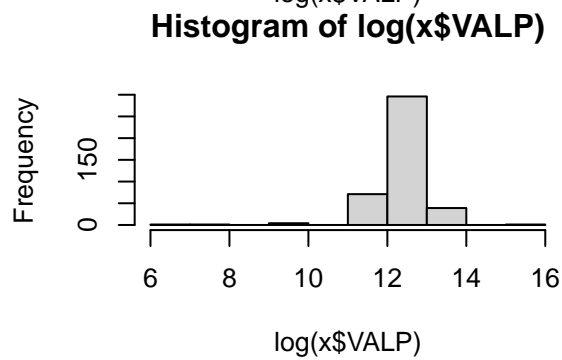
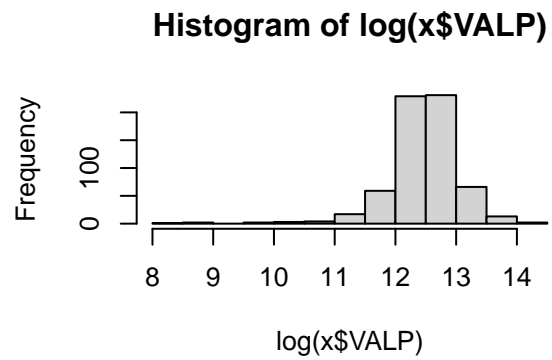
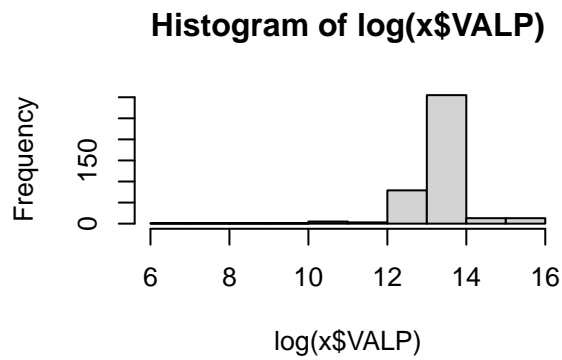
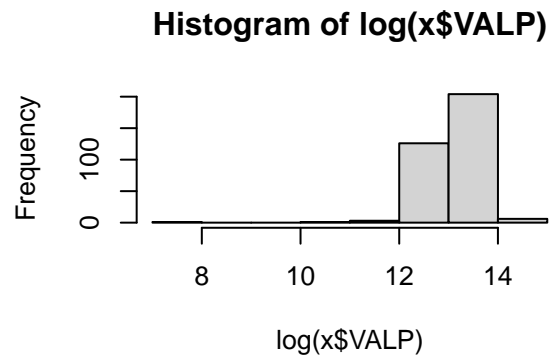
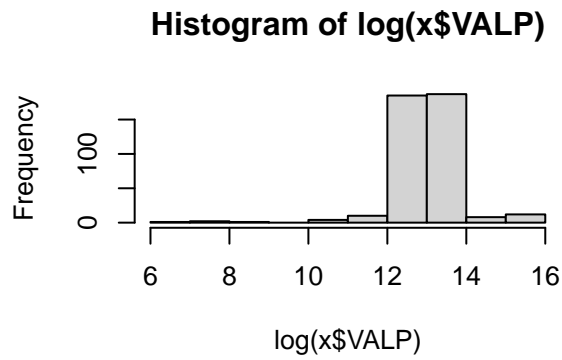


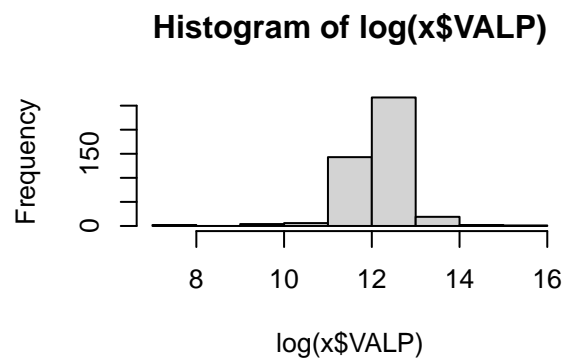
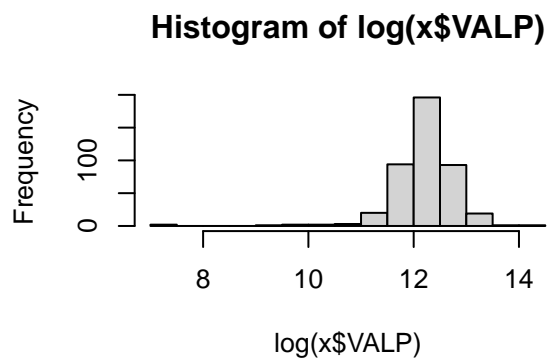
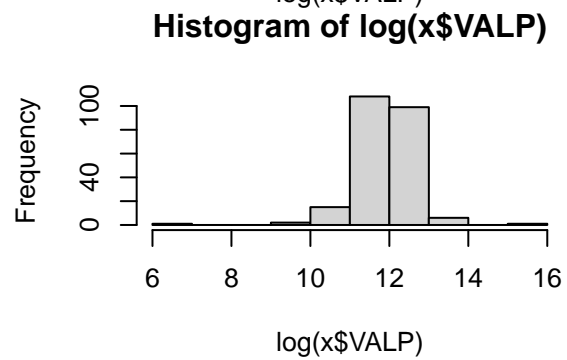
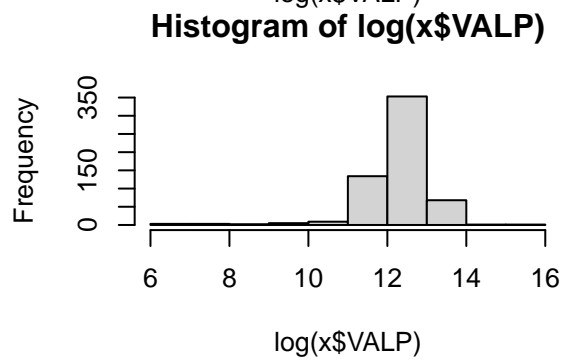
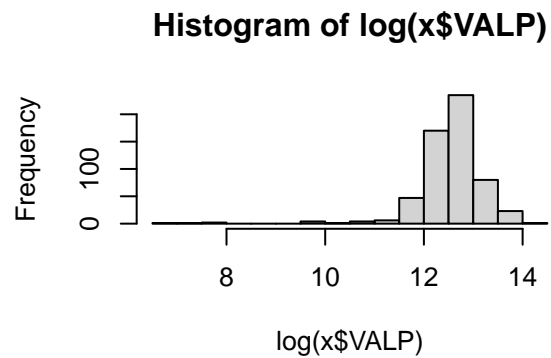
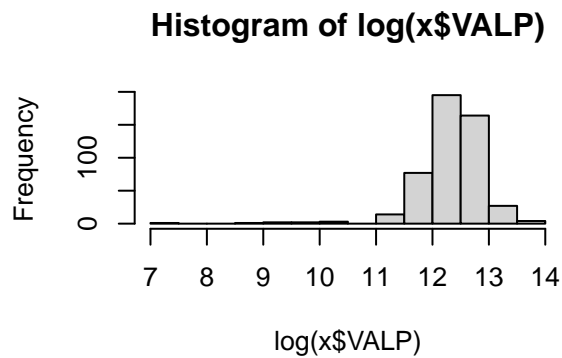
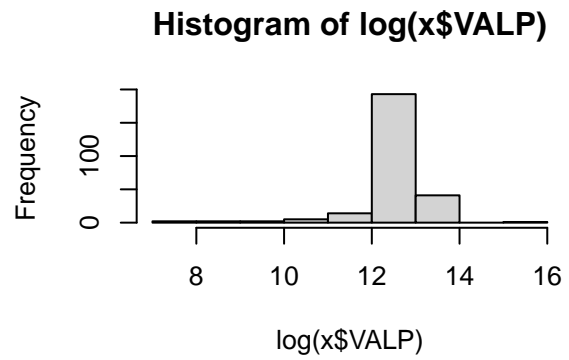
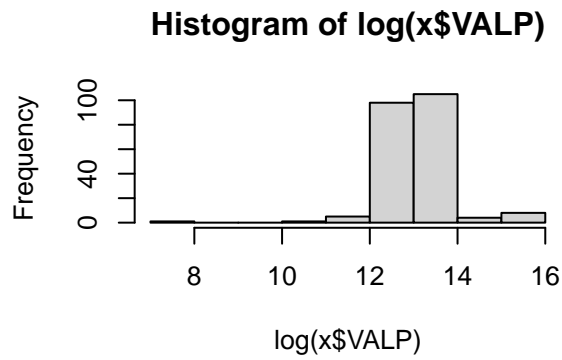


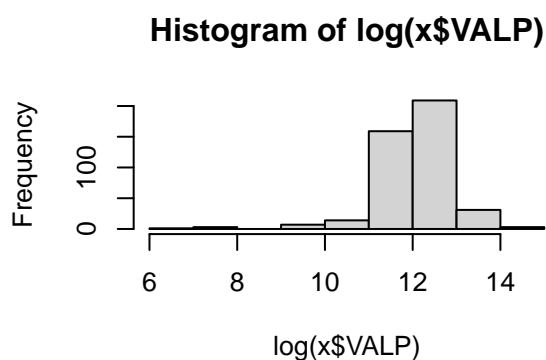
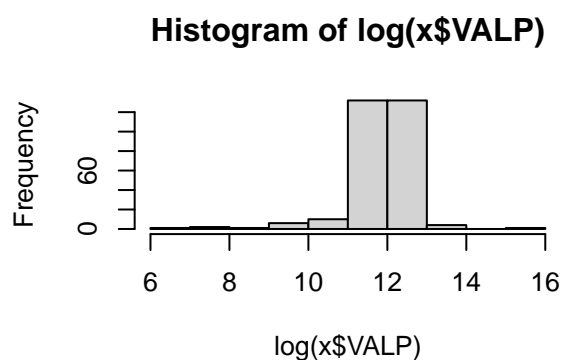
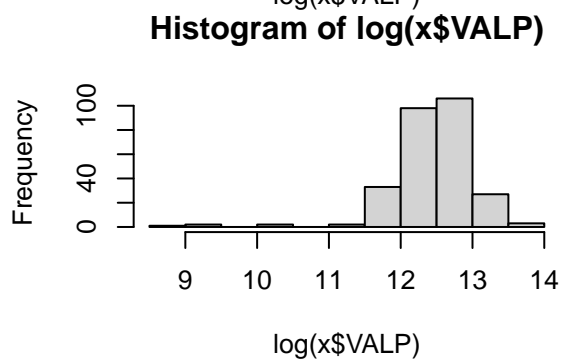
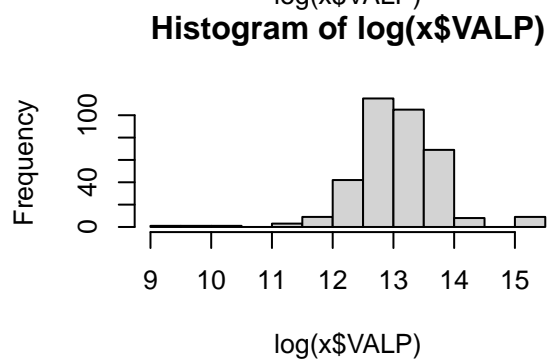
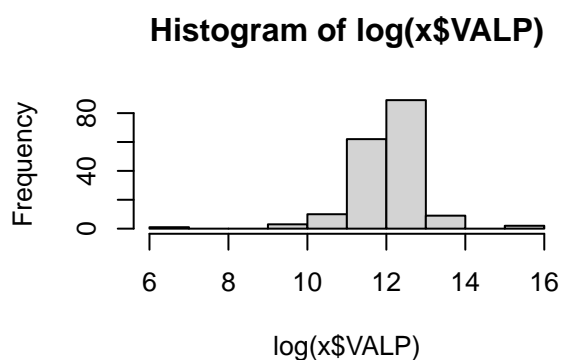
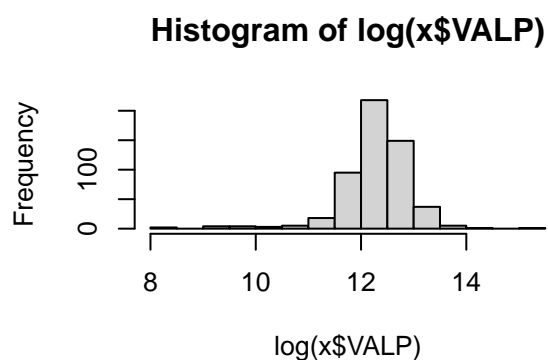
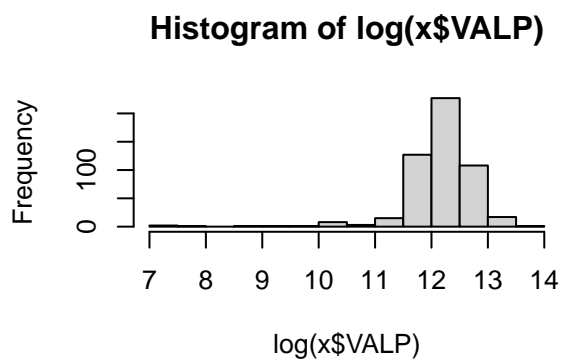
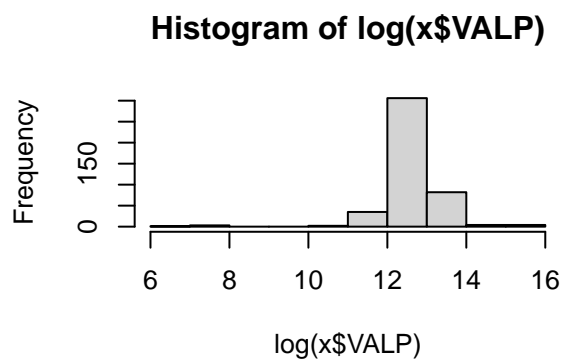












##	101	102	301	302	303
## breaks	integer,10	integer,10	integer,10	integer,10	integer,9
## counts	integer,9	integer,9	integer,9	integer,9	integer,8
## density	numeric,9	numeric,9	numeric,9	numeric,9	numeric,8

##	mids	numeric,9	numeric,9	numeric,9	numeric,9	numeric,8
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		304	305	306	307	308
##	breaks	integer,11	integer,10	integer,10	numeric,15	integer,11
##	counts	integer,10	integer,9	integer,9	integer,14	integer,10
##	density	numeric,10	numeric,9	numeric,9	numeric,14	numeric,10
##	mids	numeric,10	numeric,9	numeric,9	numeric,14	numeric,10
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		400	501	502	503	601
##	breaks	integer,8	integer,10	integer,9	numeric,10	integer,11
##	counts	integer,7	integer,9	integer,8	integer,9	integer,10
##	density	numeric,7	numeric,9	numeric,8	numeric,9	numeric,10
##	mids	numeric,7	numeric,9	numeric,8	numeric,9	numeric,10
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		602	701	702	703	800
##	breaks	integer,10	integer,10	integer,10	integer,11	integer,11
##	counts	integer,9	integer,9	integer,9	integer,10	integer,10
##	density	numeric,9	numeric,9	numeric,9	numeric,10	numeric,10
##	mids	numeric,9	numeric,9	numeric,9	numeric,10	numeric,10
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		901	902	903	904	905
##	breaks	integer,10	numeric,14	integer,10	integer,10	integer,11
##	counts	integer,9	integer,13	integer,9	integer,9	integer,10
##	density	numeric,9	numeric,13	numeric,9	numeric,9	numeric,10
##	mids	numeric,9	numeric,13	numeric,9	numeric,9	numeric,10
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		906	907	1001	1002	1003
##	breaks	numeric,12	integer,11	integer,10	numeric,11	numeric,14
##	counts	integer,11	integer,10	integer,9	integer,10	integer,13
##	density	numeric,11	numeric,10	numeric,9	numeric,10	numeric,13
##	mids	numeric,11	numeric,10	numeric,9	numeric,10	numeric,13
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		1101	1102	1103	1104	1105
##	breaks	integer,11	numeric,15	integer,11	integer,10	integer,10
##	counts	integer,10	integer,14	integer,10	integer,9	integer,9
##	density	numeric,10	numeric,14	numeric,10	numeric,9	numeric,9
##	mids	numeric,10	numeric,14	numeric,10	numeric,9	numeric,9
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		1106	1201	1202	1203	1204
##	breaks	numeric,15	integer,10	integer,11	numeric,15	numeric,16
##	counts	integer,14	integer,9	integer,10	integer,14	integer,15
##	density	numeric,14	numeric,9	numeric,10	numeric,14	numeric,15
##	mids	numeric,14	numeric,9	numeric,10	numeric,14	numeric,15
##	xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
##	equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##		1205	1301	1302	1401	1402
##	breaks	numeric,12	integer,11	integer,10	integer,9	integer,11

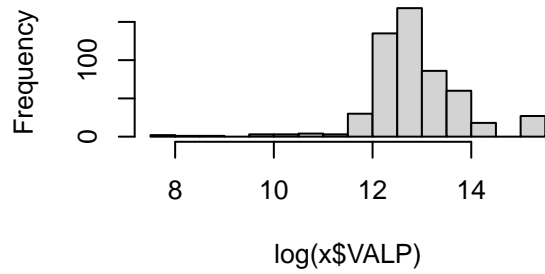
```

## counts    integer,11    integer,10    integer,9    integer,8    integer,10
## density   numeric,11    numeric,10    numeric,9    numeric,8    numeric,10
## mids      numeric,11    numeric,10    numeric,9    numeric,8    numeric,10
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           1403         1404         1501         1502         1503
## breaks    integer,10    integer,10    integer,10    integer,11    integer,9
## counts    integer,9     integer,9     integer,9     integer,10    integer,8
## density   numeric,9     numeric,9     numeric,9     numeric,10    numeric,8
## mids      numeric,9     numeric,9     numeric,9     numeric,10    numeric,8
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           1504         1600         1700         1800         1901
## breaks    integer,11    numeric,14    integer,11    integer,8     numeric,16
## counts    integer,10    integer,13    integer,10    integer,7     integer,15
## density   numeric,10    numeric,13    numeric,10    numeric,7     numeric,15
## mids      numeric,10    numeric,13    numeric,10    numeric,7     numeric,15
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           1902         1903         1904         2001         2002
## breaks    integer,10    integer,10    integer,10    numeric,15    numeric,17
## counts    integer,9     integer,9     integer,9     integer,14    integer,16
## density   numeric,9     numeric,9     numeric,9     numeric,14    numeric,16
## mids      numeric,9     numeric,9     numeric,9     numeric,14    numeric,16
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           2003         2101         2102         2103         2104
## breaks    integer,11    integer,11    numeric,16    integer,10    integer,11
## counts    integer,10    integer,10    integer,15    integer,9     integer,10
## density   numeric,10    numeric,10    numeric,15    numeric,9     numeric,10
## mids      numeric,10    numeric,10    numeric,15    numeric,9     numeric,10
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           2201         2202         2301         2302         2303
## breaks    numeric,15    numeric,16    integer,11    numeric,14    numeric,12
## counts    integer,14    integer,15    integer,10    integer,13    integer,11
## density   numeric,14    numeric,15    numeric,10    numeric,13    numeric,11
## mids      numeric,14    numeric,15    numeric,10    numeric,13    numeric,11
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE         TRUE         TRUE
##           2400         2500         2600
## breaks    integer,11    integer,10    numeric,17
## counts    integer,10    integer,9     integer,16
## density   numeric,10    numeric,9     numeric,16
## mids      numeric,10    numeric,9     numeric,16
## xname     "log(x$VALP)" "log(x$VALP)" "log(x$VALP)"
## equidist  TRUE         TRUE         TRUE

```

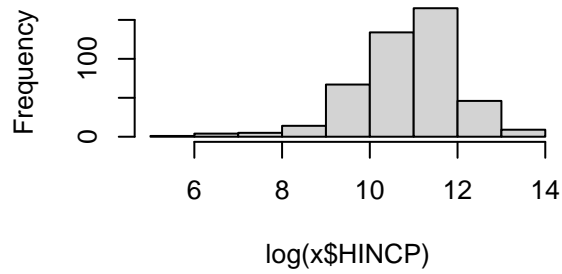
```
par(mfrow = c(2, 2))
```

Histogram of $\log(x\$VALP)$

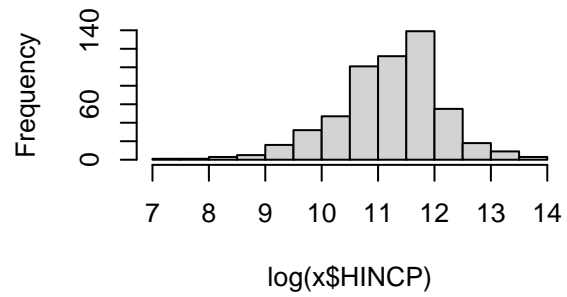


```
sapply(split(pumsnj, pumsnj$PUMA), function(x) {
  hist(log(x$HINCP))
})
```

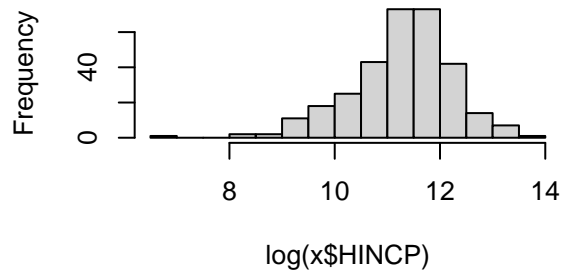
Histogram of $\log(x\$HINCP)$



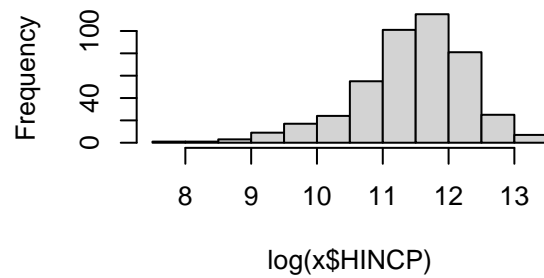
Histogram of $\log(x\$HINCP)$

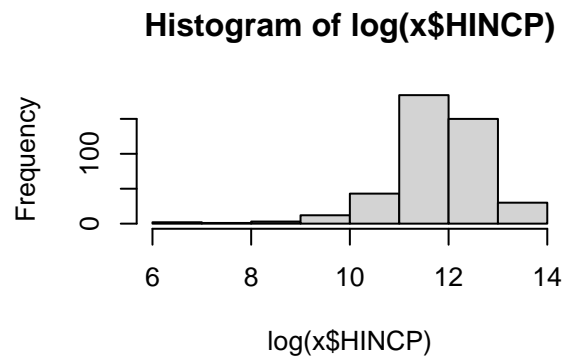
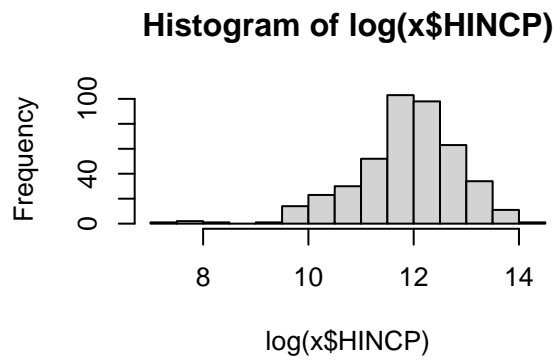
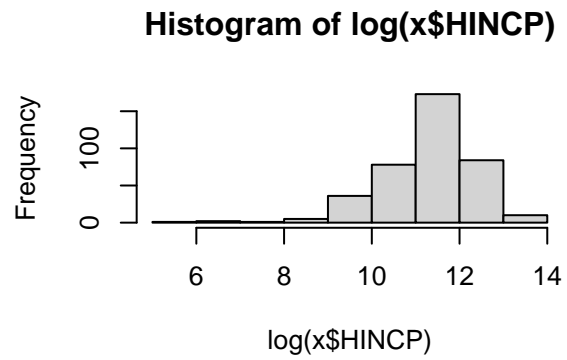
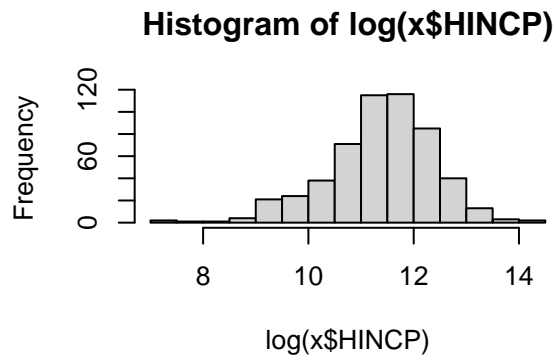


Histogram of $\log(x\$HINCP)$

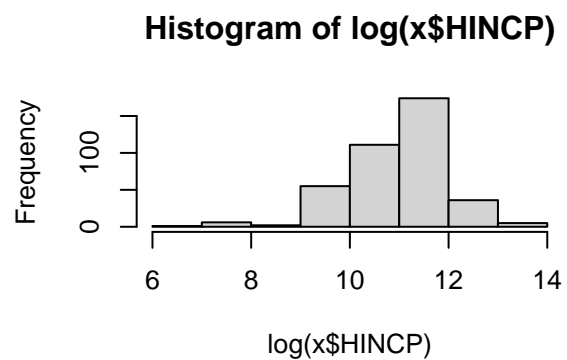
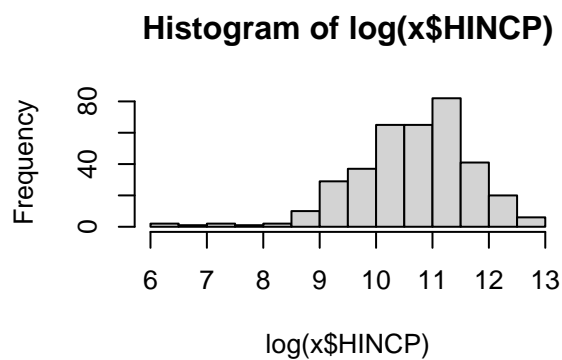
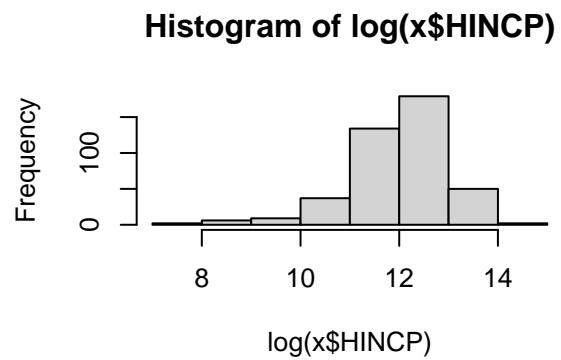
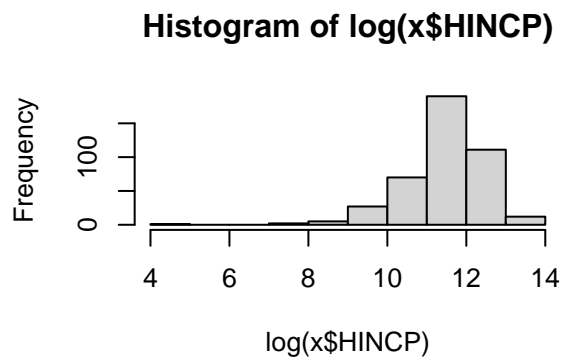


Histogram of $\log(x\$HINCP)$

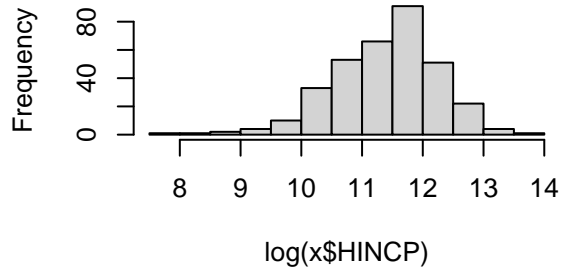




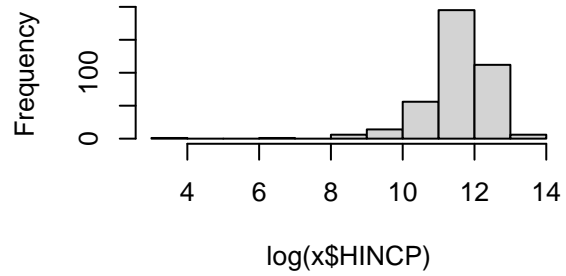
Warning in log(x\$HINCP): NaNs produced



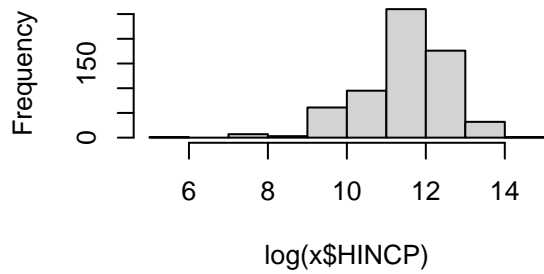
Histogram of log(x\$HINCP)



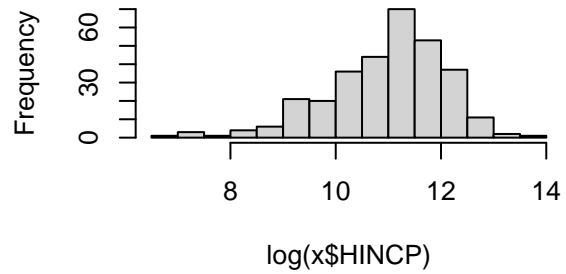
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

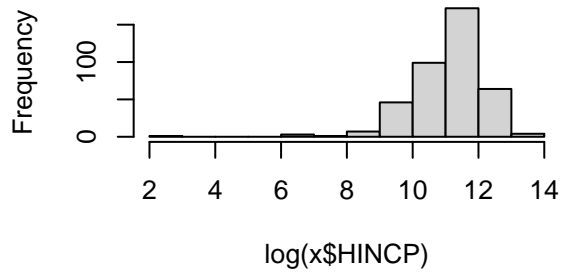


Histogram of log(x\$HINCP)

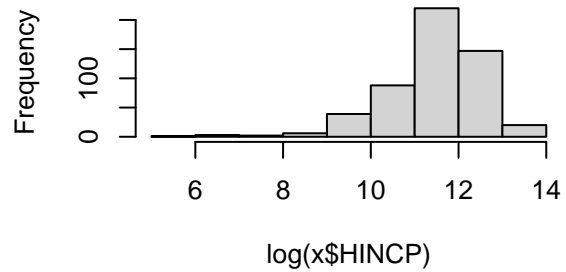


Warning in log(x\$HINCP): NaNs produced

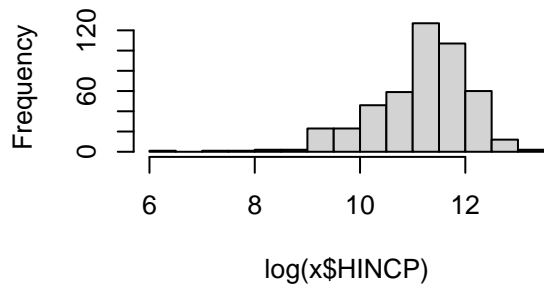
Histogram of log(x\$HINCP)



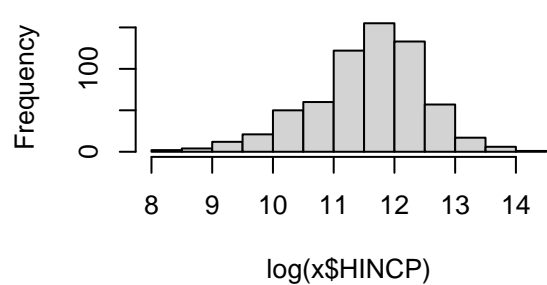
Histogram of log(x\$HINCP)



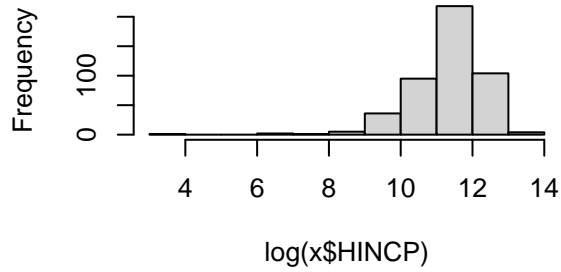
Histogram of log(x\$HINCP)



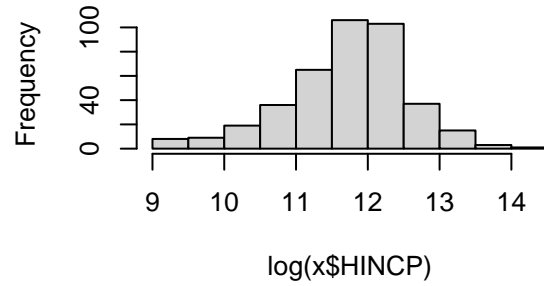
Histogram of log(x\$HINCP)



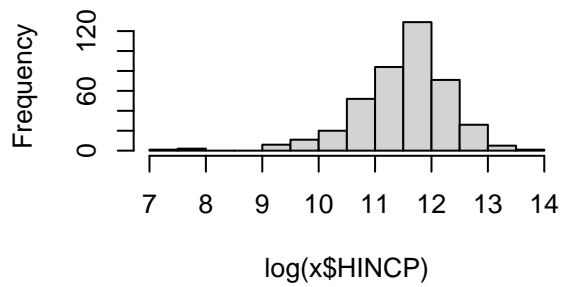
Histogram of log(x\$HINCP)



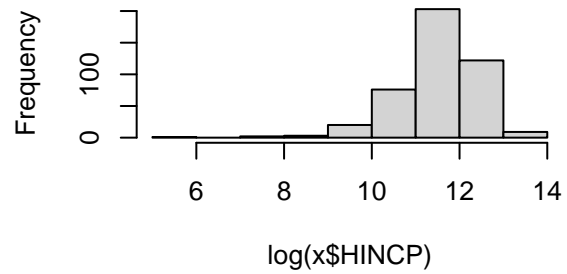
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

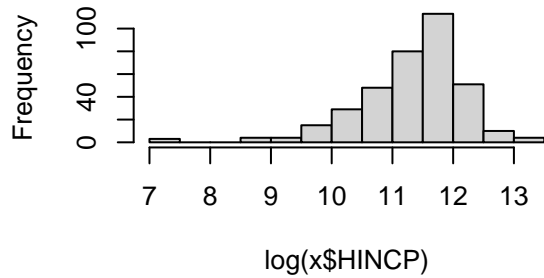


Histogram of log(x\$HINCP)

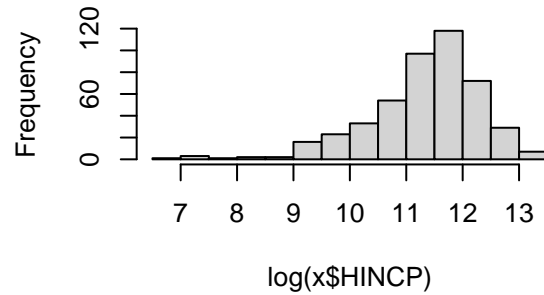


Warning in log(x\$HINCP): NaNs produced

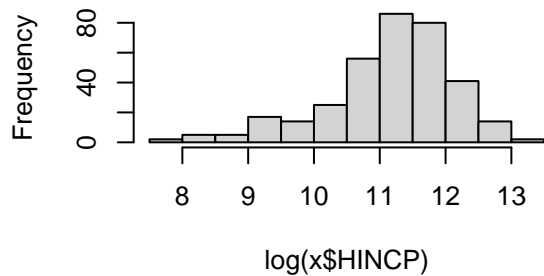
Histogram of log(x\$HINCP)



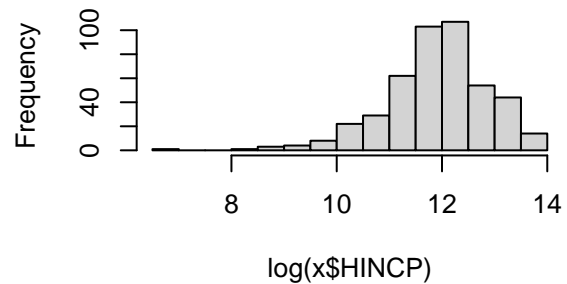
Histogram of log(x\$HINCP)



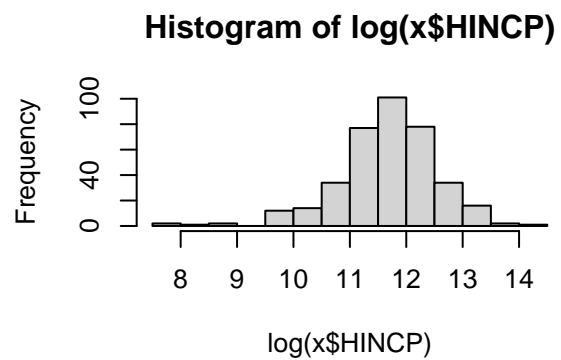
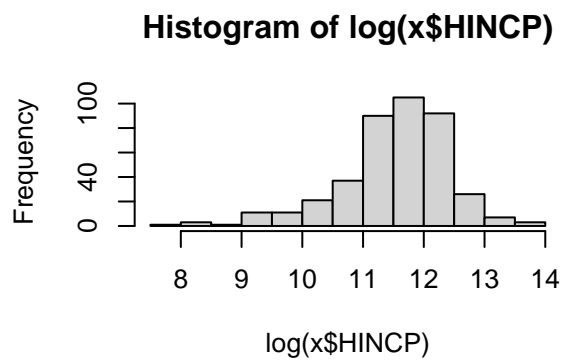
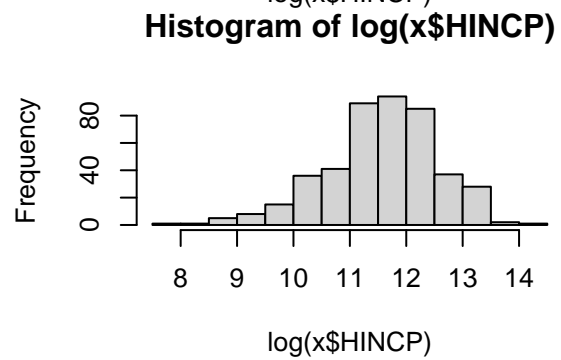
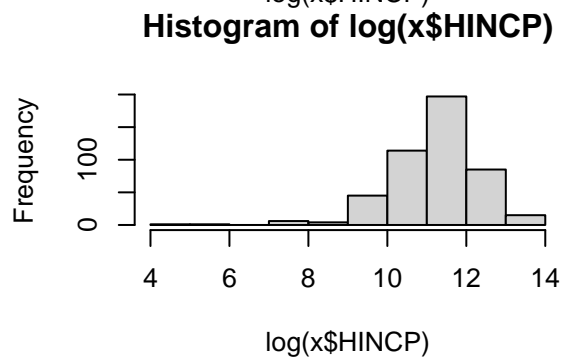
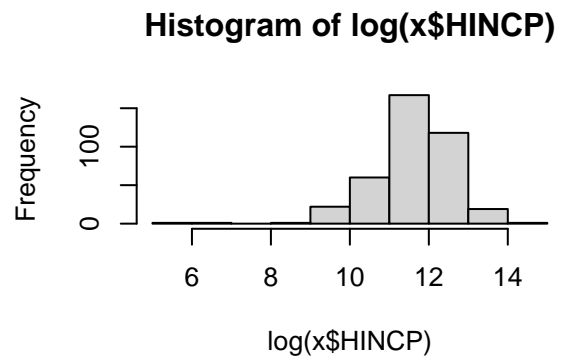
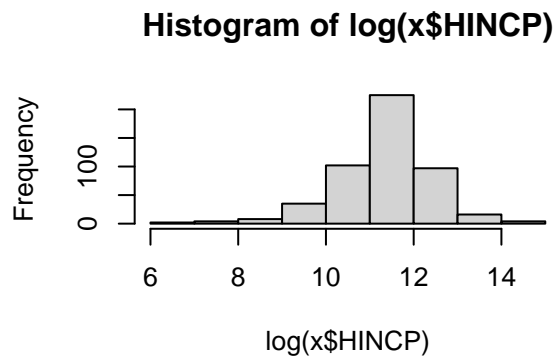
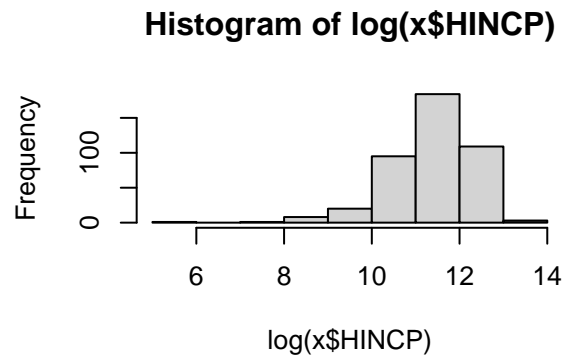
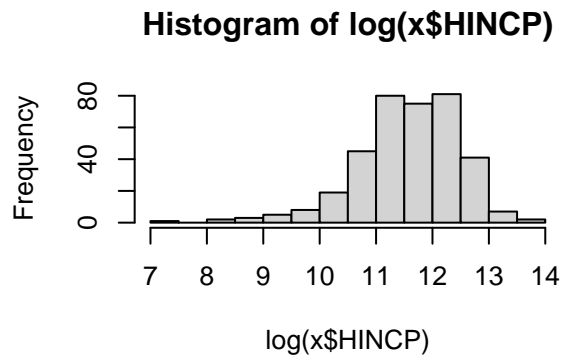
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

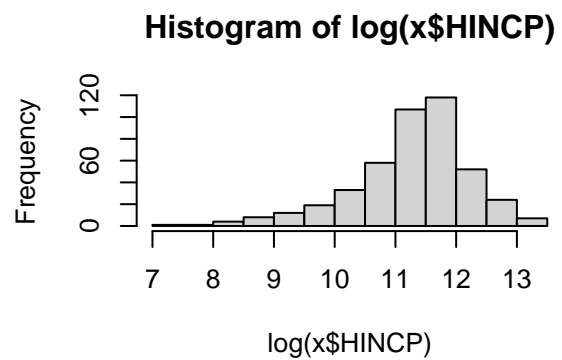
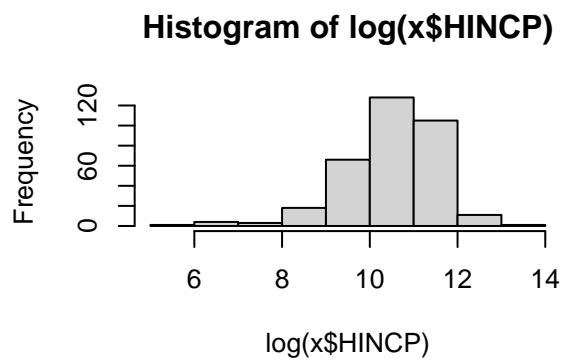
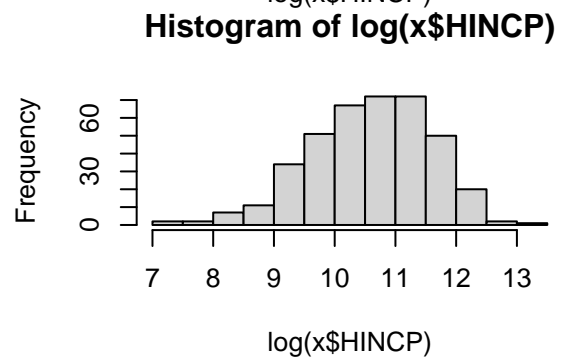
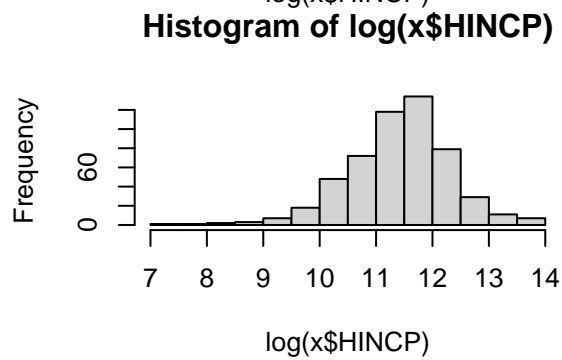
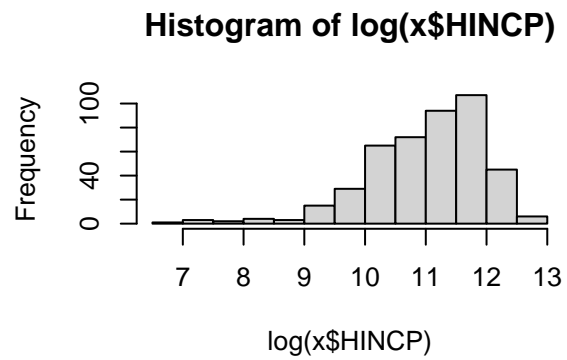
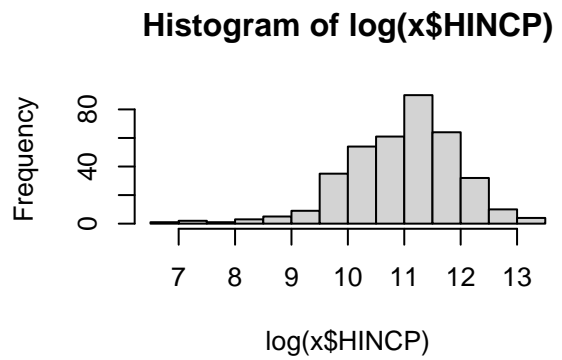
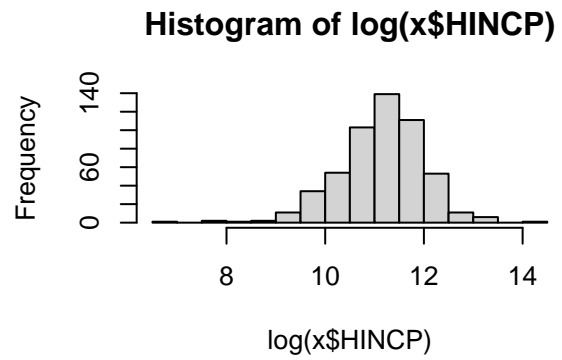
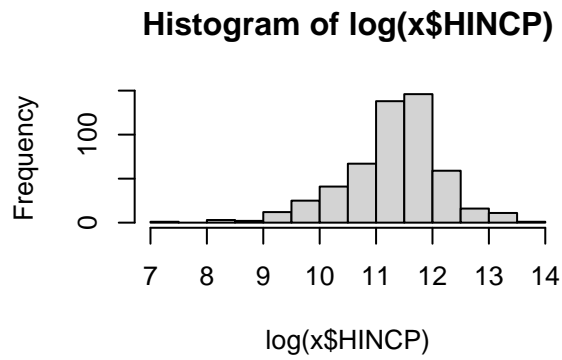


Warning in log(x\$HINCP): NaNs produced

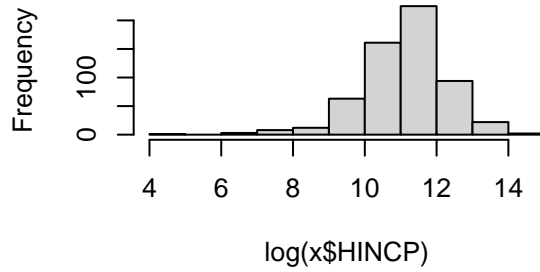


Warning in log(x\$HINCP): NaNs produced

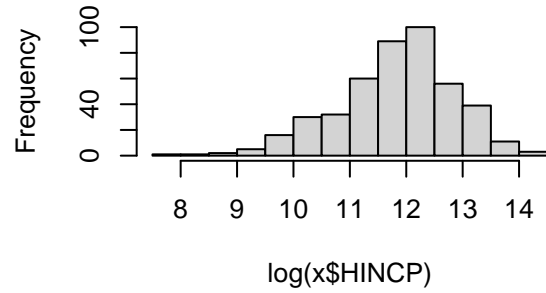
Warning in log(x\$HINCP): NaNs produced



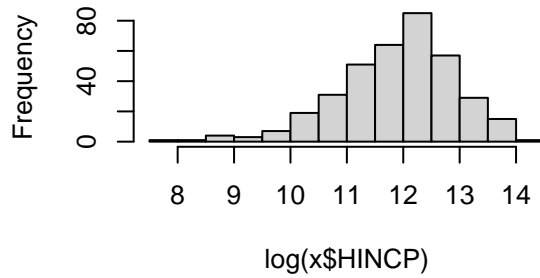
Histogram of log(x\$HINCP)



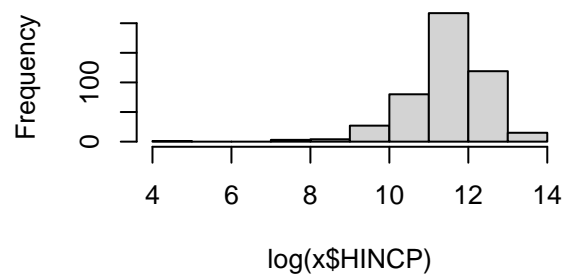
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

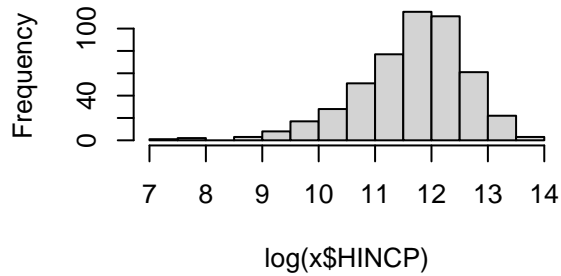


Histogram of log(x\$HINCP)

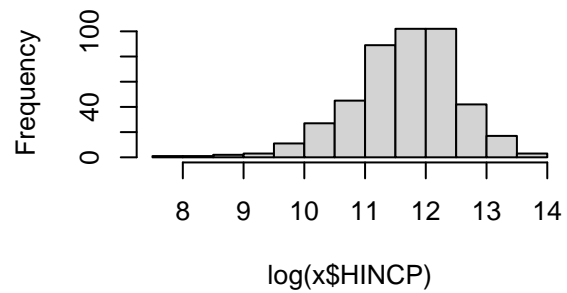


Warning in log(x\$HINCP): NaNs produced

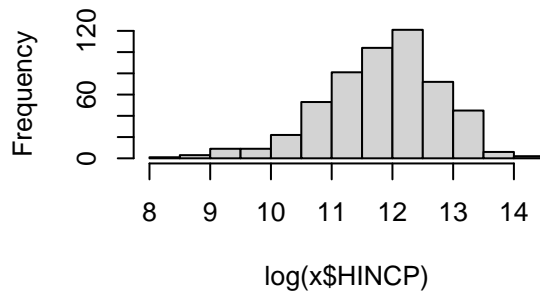
Histogram of log(x\$HINCP)



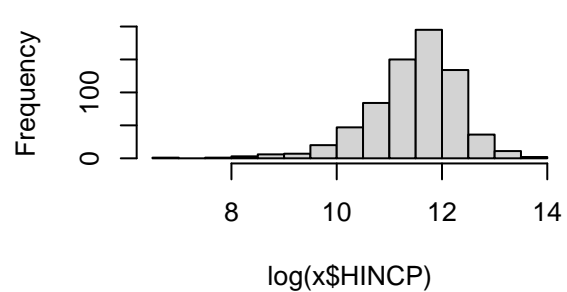
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

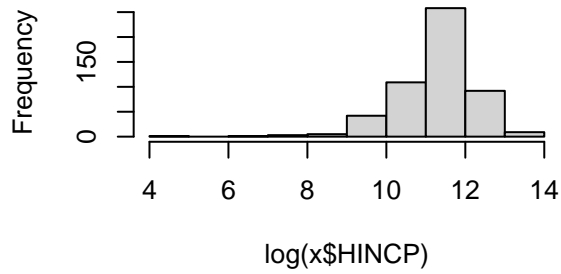


Histogram of log(x\$HINCP)

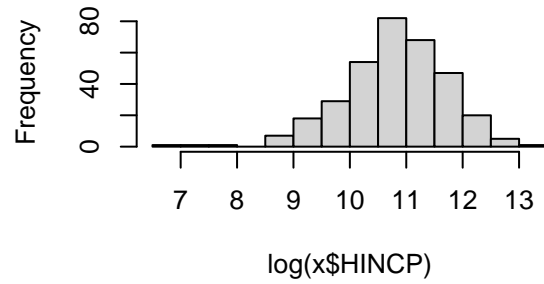


Warning in log(x\$HINCP): NaNs produced

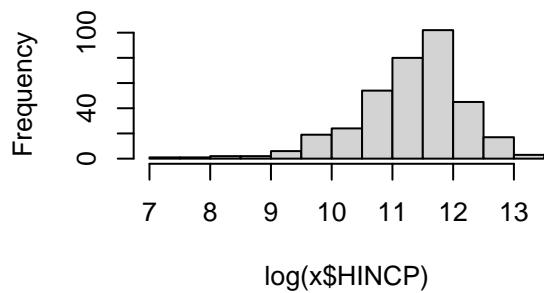
Histogram of log(x\$HINCP)



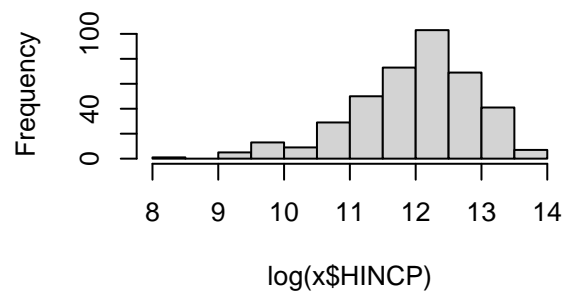
Histogram of log(x\$HINCP)



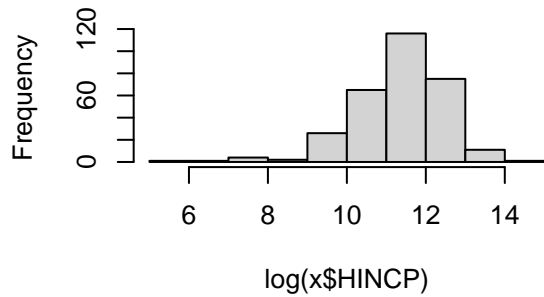
Histogram of log(x\$HINCP)



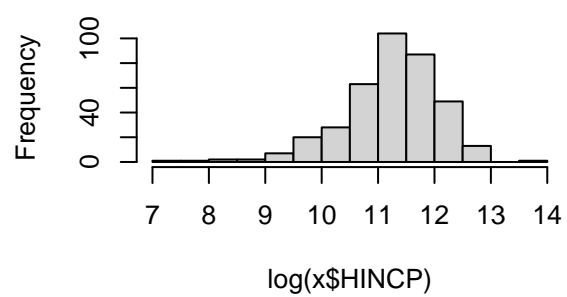
Histogram of log(x\$HINCP)



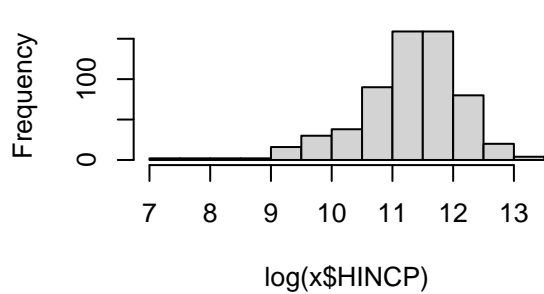
Histogram of log(x\$HINCP)



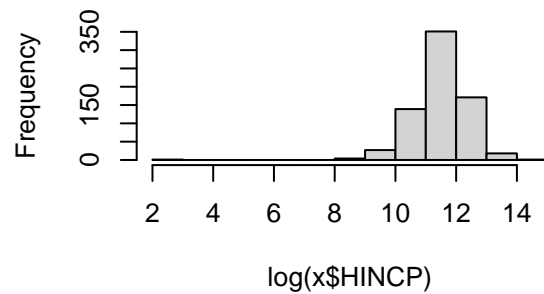
Histogram of log(x\$HINCP)

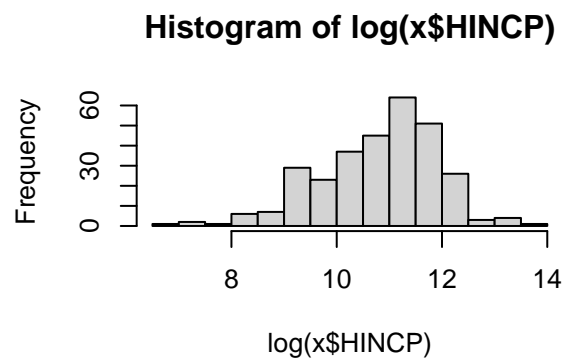
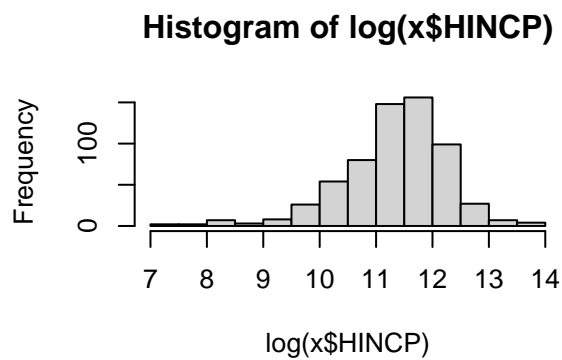
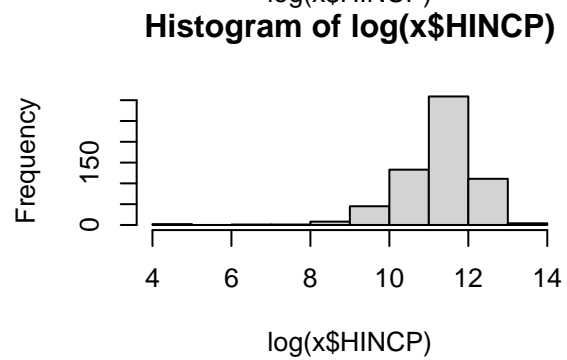
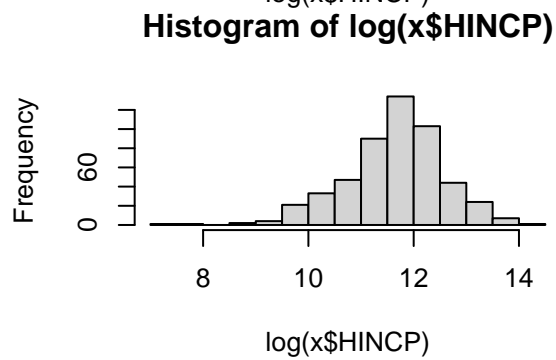
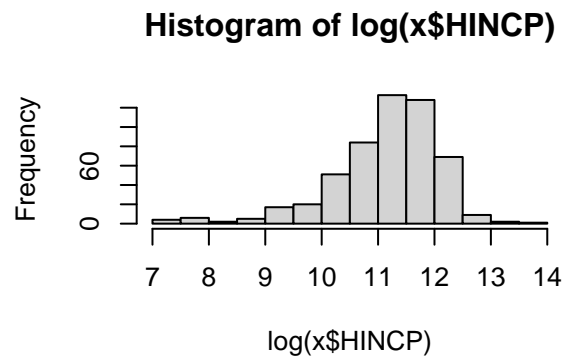
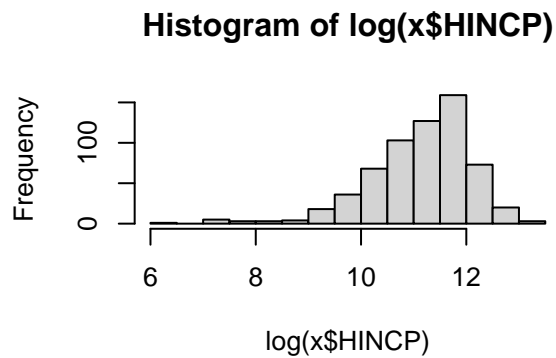
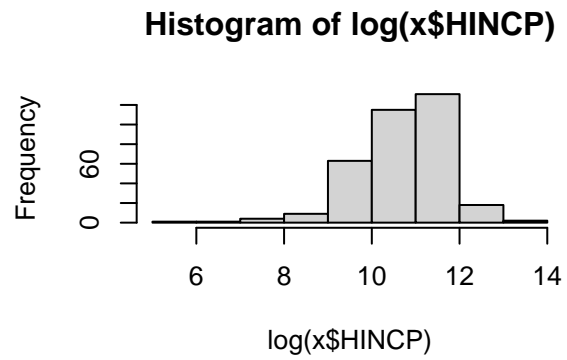
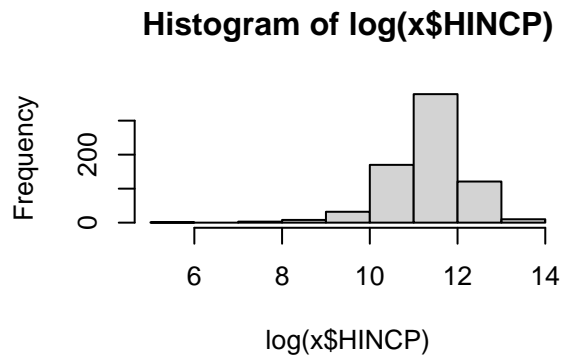


Histogram of log(x\$HINCP)



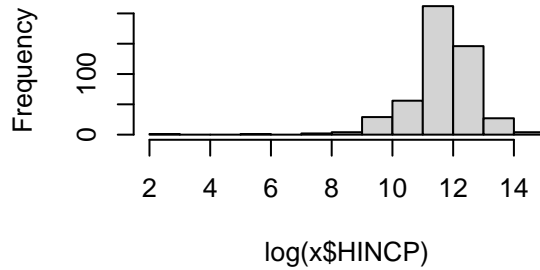
Histogram of log(x\$HINCP)



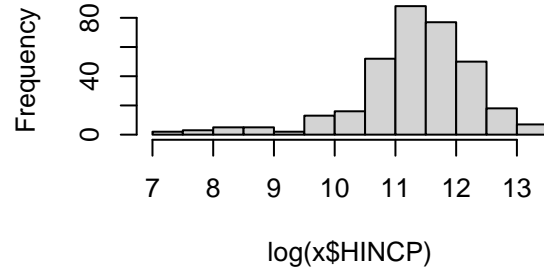


Warning in log(x\$HINCP): NaNs produced

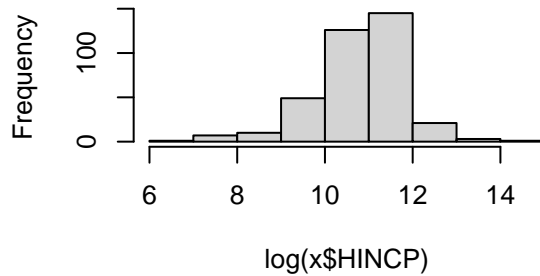
Histogram of log(x\$HINCP)



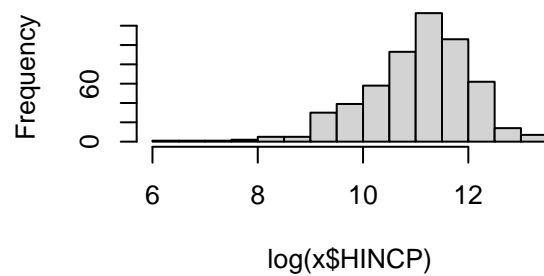
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)



##	101	102	301	302
## breaks	integer,10	numeric,15	numeric,16	numeric,13
## counts	integer,9	integer,14	integer,15	integer,12
## density	numeric,9	numeric,14	numeric,15	numeric,12
## mids	numeric,9	numeric,14	numeric,15	numeric,12
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	303	304	305	306
## breaks	numeric,16	integer,10	numeric,16	integer,9
## counts	integer,15	integer,9	integer,15	integer,8
## density	numeric,15	numeric,9	numeric,15	numeric,8
## mids	numeric,15	numeric,9	numeric,15	numeric,8
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	307	308	400	501
## breaks	integer,11	integer,9	numeric,15	integer,9
## counts	integer,10	integer,8	integer,14	integer,8
## density	numeric,10	numeric,8	numeric,14	numeric,8
## mids	numeric,10	numeric,8	numeric,14	numeric,8
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	502	503	601	602
## breaks	numeric,14	integer,12	integer,11	numeric,16
## counts	integer,13	integer,11	integer,10	integer,15
## density	numeric,13	numeric,11	numeric,10	numeric,15
## mids	numeric,13	numeric,11	numeric,10	numeric,15
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE

##	701	702	703	800
## breaks	integer,13	integer,10	numeric,16	numeric,14
## counts	integer,12	integer,9	integer,15	integer,13
## density	numeric,12	numeric,9	numeric,15	numeric,13
## mids	numeric,12	numeric,9	numeric,15	numeric,13
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	901	902	903	904
## breaks	integer,12	numeric,12	numeric,15	integer,10
## counts	integer,11	integer,11	integer,14	integer,9
## density	numeric,11	numeric,11	numeric,14	numeric,9
## mids	numeric,11	numeric,11	numeric,14	numeric,9
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	905	906	907	1001
## breaks	numeric,14	numeric,15	numeric,13	numeric,16
## counts	integer,13	integer,14	integer,12	integer,15
## density	numeric,13	numeric,14	numeric,12	numeric,15
## mids	numeric,13	numeric,14	numeric,12	numeric,15
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	1002	1003	1101	1102
## breaks	numeric,15	integer,10	integer,10	integer,11
## counts	integer,14	integer,9	integer,9	integer,10
## density	numeric,14	numeric,9	numeric,9	numeric,10
## mids	numeric,14	numeric,9	numeric,9	numeric,10
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	1103	1104	1105	1106
## breaks	integer,11	numeric,15	numeric,14	numeric,15
## counts	integer,10	integer,14	integer,13	integer,14
## density	numeric,10	numeric,14	numeric,13	numeric,14
## mids	numeric,10	numeric,14	numeric,13	numeric,14
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	1201	1202	1203	1204
## breaks	numeric,15	numeric,17	numeric,15	numeric,14
## counts	integer,14	integer,16	integer,14	integer,13
## density	numeric,14	numeric,16	numeric,14	numeric,13
## mids	numeric,14	numeric,16	numeric,14	numeric,13
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	1205	1301	1302	1401
## breaks	numeric,15	numeric,14	integer,10	numeric,14
## counts	integer,14	integer,13	integer,9	integer,13
## density	numeric,14	numeric,13	numeric,9	numeric,13
## mids	numeric,14	numeric,13	numeric,9	numeric,13
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	1402	1403	1404	1501
## breaks	integer,12	numeric,15	numeric,15	integer,11
## counts	integer,11	integer,14	integer,14	integer,10
## density	numeric,11	numeric,14	numeric,14	numeric,10
## mids	numeric,11	numeric,14	numeric,14	numeric,10

```

## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           1502            1503            1504            1600
## breaks     numeric,15      numeric,14        numeric,14        numeric,16
## counts     integer,14      integer,13        integer,13        integer,15
## density    numeric,14      numeric,13        numeric,13        numeric,15
## mids       numeric,14      numeric,13        numeric,13        numeric,15
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           1700            1800            1901            1902
## breaks     integer,11      numeric,15        numeric,14        numeric,13
## counts     integer,10      integer,14        integer,13        integer,12
## density    numeric,10      numeric,14        numeric,13        numeric,12
## mids       numeric,10      numeric,14        numeric,13        numeric,12
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           1903            1904            2001            2002
## breaks     integer,11      numeric,15        numeric,14        integer,14
## counts     integer,10      integer,14        integer,13        integer,13
## density    numeric,10      numeric,14        numeric,13        numeric,13
## mids       numeric,10      numeric,14        numeric,13        numeric,13
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           2003            2101            2102            2103
## breaks     integer,10      integer,10        numeric,16        numeric,15
## counts     integer,9       integer,9         integer,15        integer,14
## density    numeric,9       numeric,9         numeric,15        numeric,14
## mids       numeric,9       numeric,9         numeric,15        numeric,14
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           2104            2201            2202            2301
## breaks     numeric,16      integer,11        numeric,15        numeric,16
## counts     integer,15      integer,10        integer,14        integer,15
## density    numeric,15      numeric,10        numeric,14        numeric,15
## mids       numeric,15      numeric,10        numeric,14        numeric,15
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           2302            2303            2400            2500
## breaks     integer,14      numeric,14        integer,10        numeric,16
## counts     integer,13      integer,13        integer,9         integer,15
## density    numeric,13      numeric,13        numeric,9         numeric,15
## mids       numeric,13      numeric,13        numeric,9         numeric,15
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           2600
## breaks     integer,12
## counts     integer,11
## density    numeric,11
## mids       numeric,11
## xname      "log(x$HINCP)"
## equidist TRUE

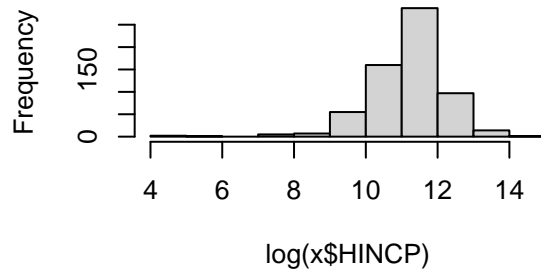
```

```

# Within FIPS
par(mfrow = c(2, 2))

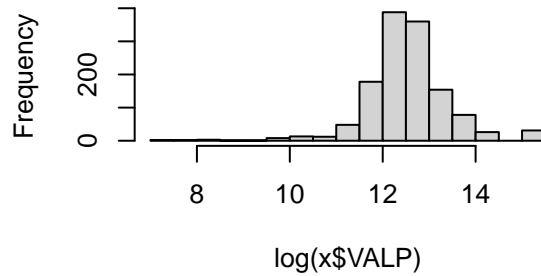
```

Histogram of log(x\$HINCP)

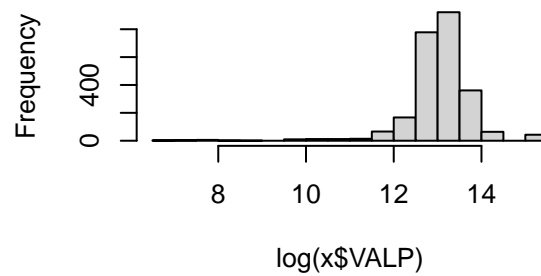


```
sapply(split(pumsnj, pumsnj$fips), function(x) {
  hist(log(x$VALP))
})
```

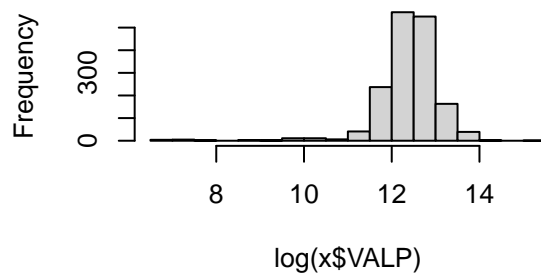
Histogram of log(x\$VALP)



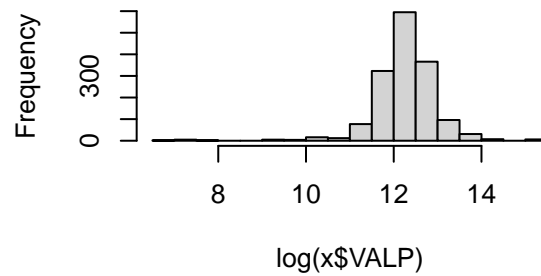
Histogram of log(x\$VALP)

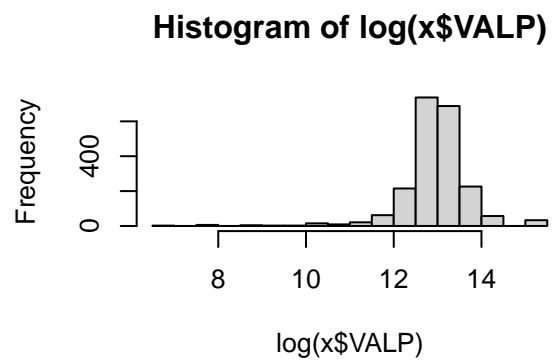
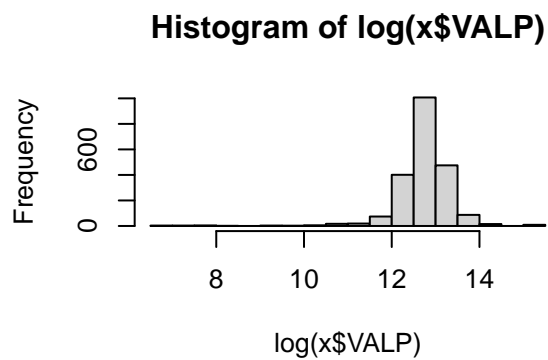
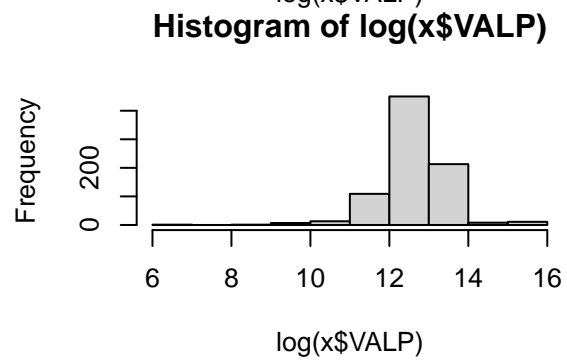
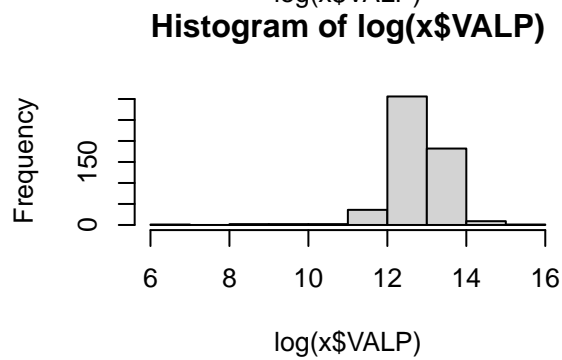
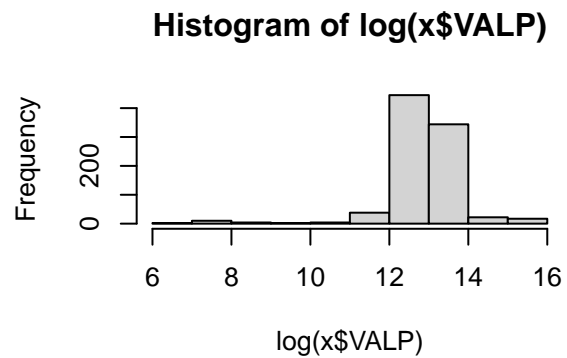
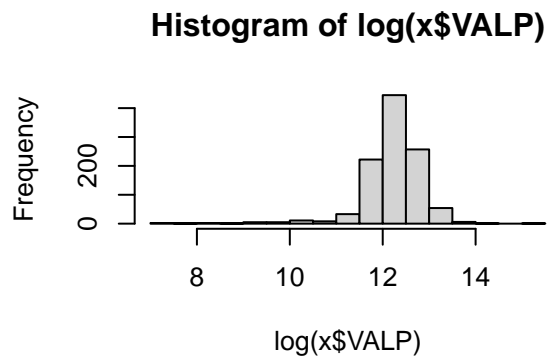
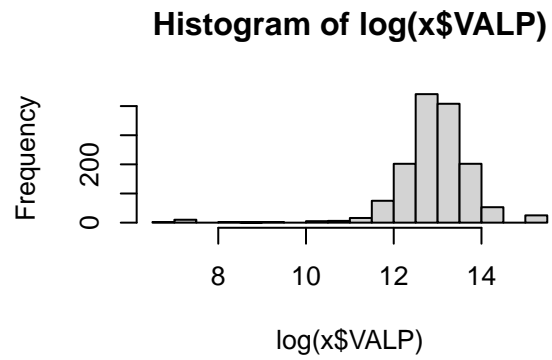
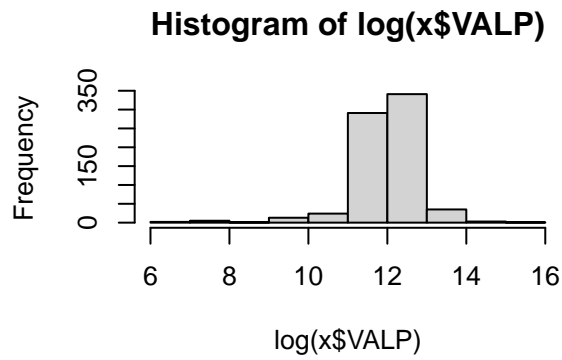


Histogram of log(x\$VALP)

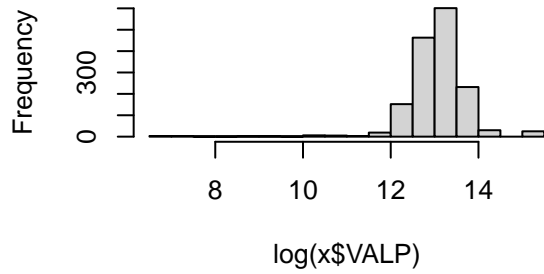


Histogram of log(x\$VALP)

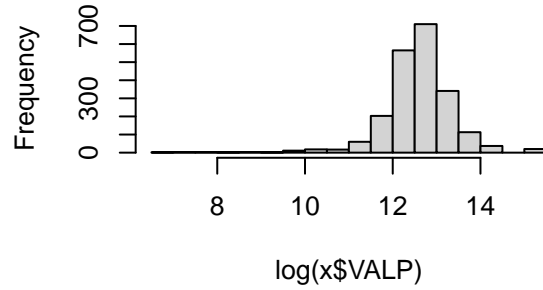




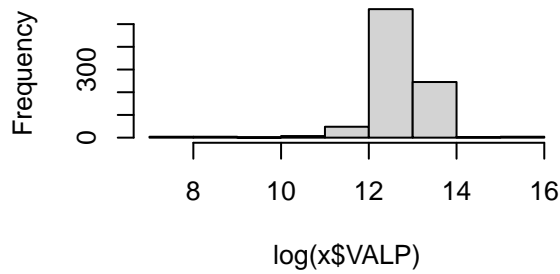
Histogram of log(x\$VALP)



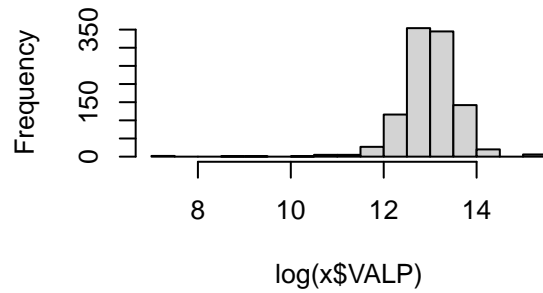
Histogram of log(x\$VALP)



Histogram of log(x\$VALP)

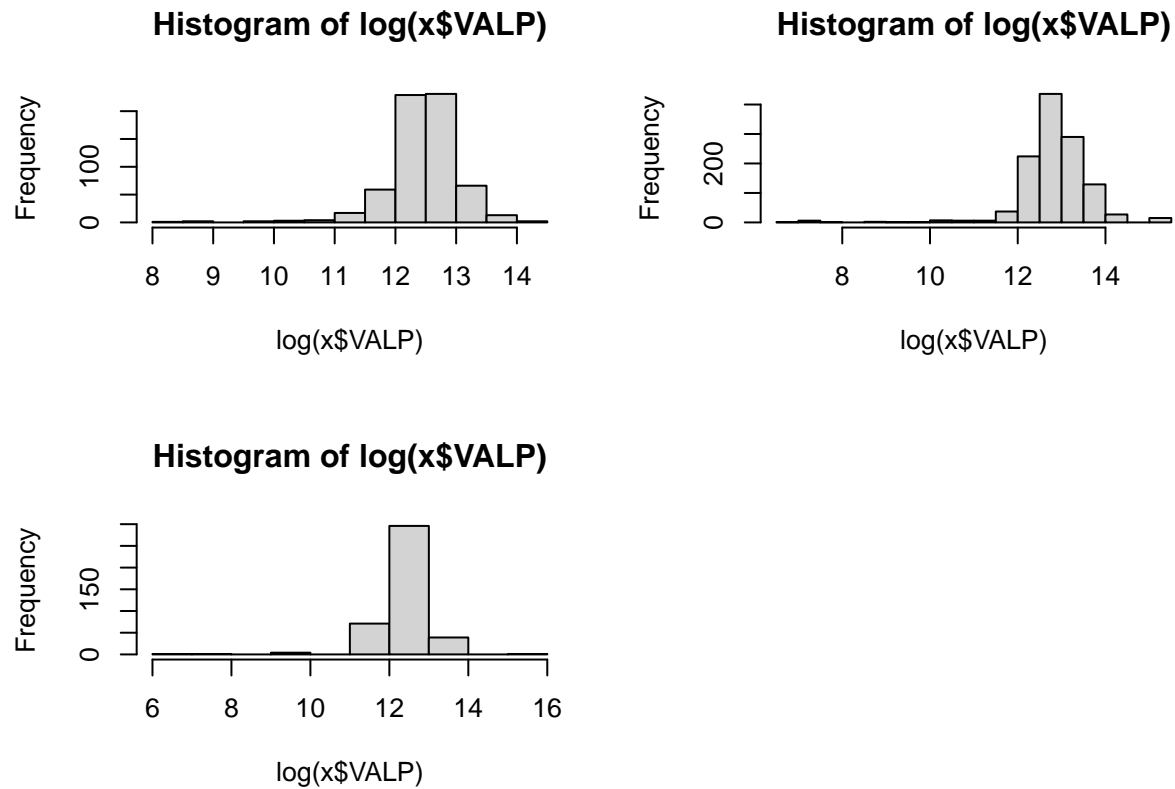


Histogram of log(x\$VALP)



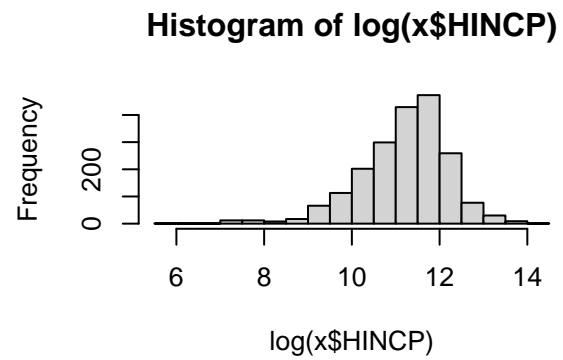
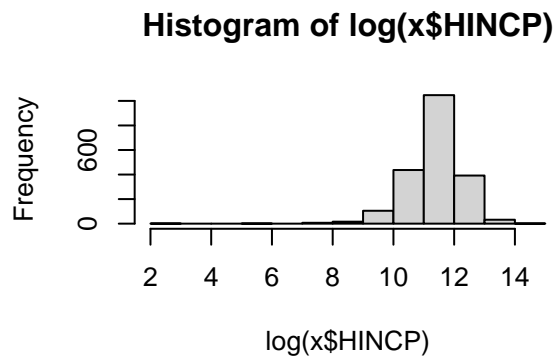
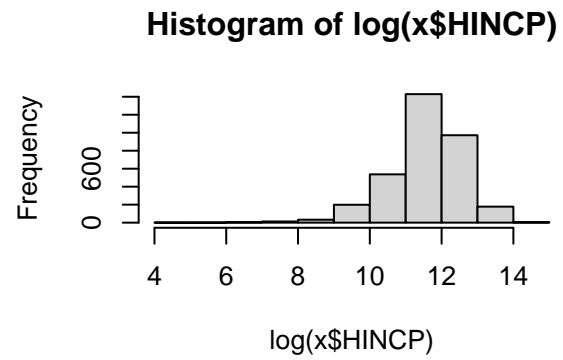
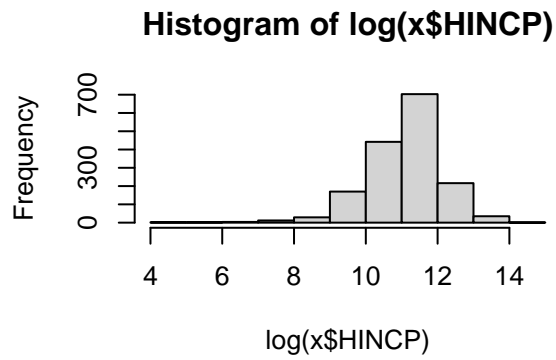
##	34001	34003	34005	34007	34011
## breaks	numeric,18	numeric,19	numeric,19	numeric,19	integer,11
## counts	integer,17	integer,18	integer,18	integer,18	integer,10
## density	numeric,17	numeric,18	numeric,18	numeric,18	numeric,10
## mids	numeric,17	numeric,18	numeric,18	numeric,18	numeric,10
## xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
## equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##	34013	34015	34017	34019	34021
## breaks	numeric,19	numeric,18	integer,11	integer,11	integer,11
## counts	integer,18	integer,17	integer,10	integer,10	integer,10
## density	numeric,18	numeric,17	numeric,10	numeric,10	numeric,10
## mids	numeric,18	numeric,17	numeric,10	numeric,10	numeric,10
## xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
## equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##	34023	34025	34027	34029	34031
## breaks	numeric,19	numeric,19	numeric,19	numeric,19	integer,10
## counts	integer,18	integer,18	integer,18	integer,18	integer,9
## density	numeric,18	numeric,18	numeric,18	numeric,18	numeric,9
## mids	numeric,18	numeric,18	numeric,18	numeric,18	numeric,9
## xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"
## equidist	TRUE	TRUE	TRUE	TRUE	TRUE
##	34035	34037	34039	34041	
## breaks	numeric,18	numeric,14	numeric,19	integer,11	
## counts	integer,17	integer,13	integer,18	integer,10	
## density	numeric,17	numeric,13	numeric,18	numeric,10	
## mids	numeric,17	numeric,13	numeric,18	numeric,10	
## xname	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	"log(x\$VALP)"	
## equidist	TRUE	TRUE	TRUE	TRUE	

```
par(mfrow = c(2, 2))
```

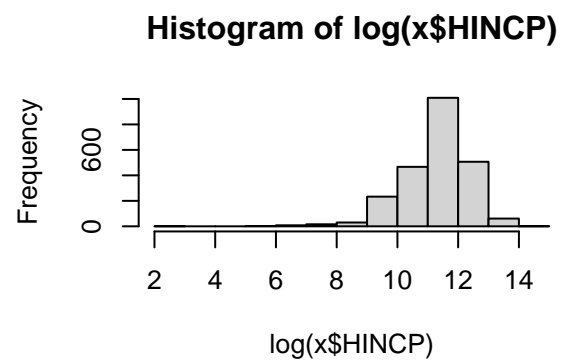
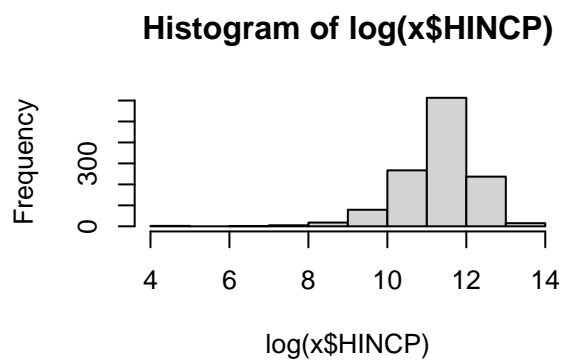
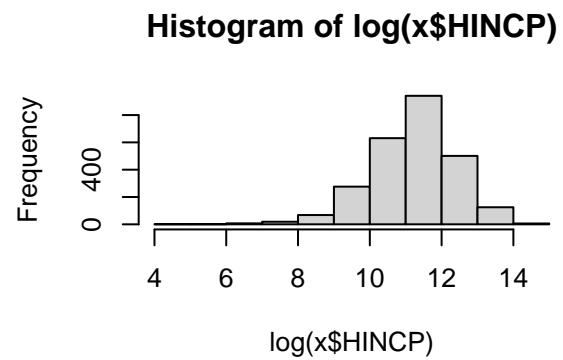
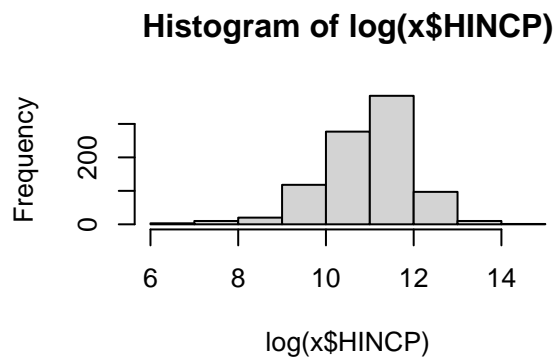


```
sapply(split(pumsnj, pumsnj$fips), function(x) {
  hist(log(x$HINCP))
})
```

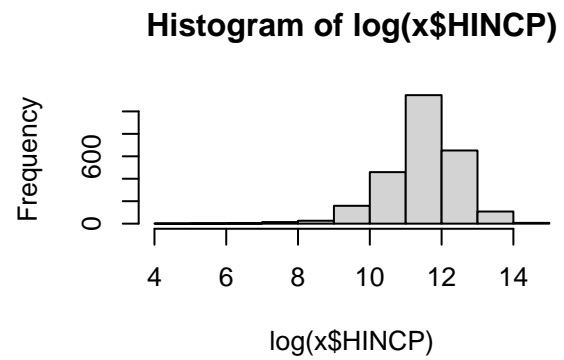
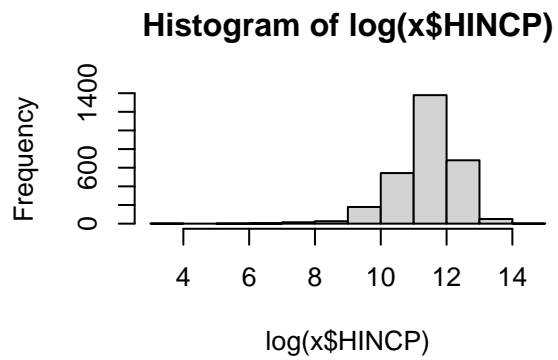
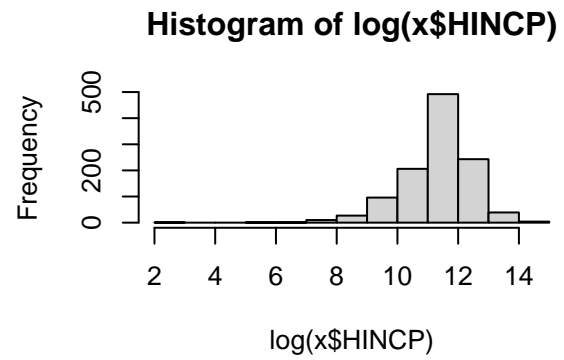
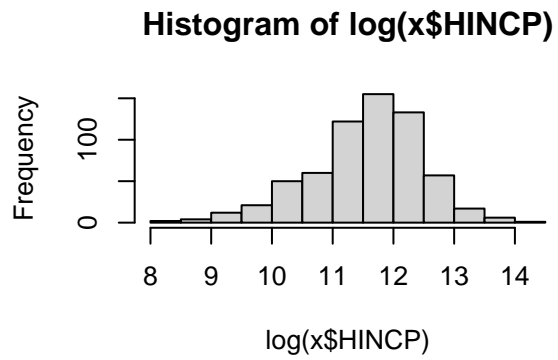
```
## Warning in log(x$HINCP): NaNs produced
```



Warning in log(x\$HINCP): NaNs produced

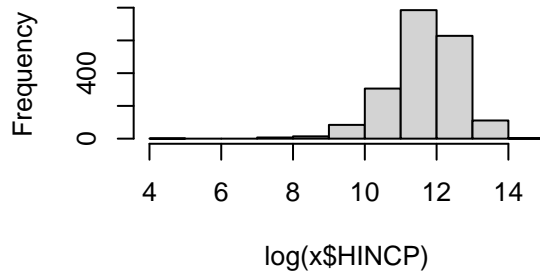


Warning in log(x\$HINCP): NaNs produced

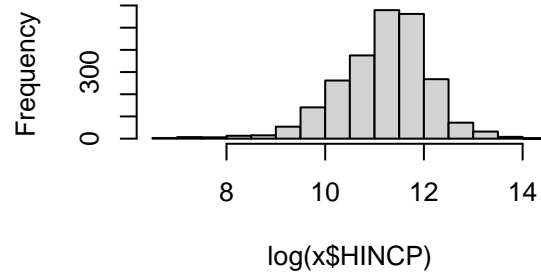


```
## Warning in log(x$HINCP): NaNs produced
## Warning in log(x$HINCP): NaNs produced
## Warning in log(x$HINCP): NaNs produced
## Warning in log(x$HINCP): NaNs produced
```

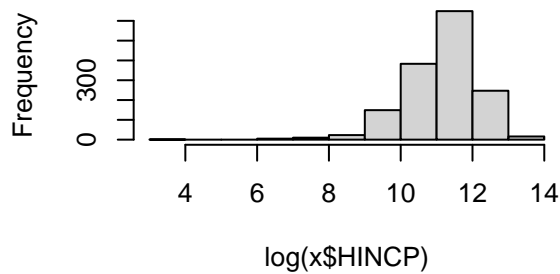
Histogram of log(x\$HINCP)



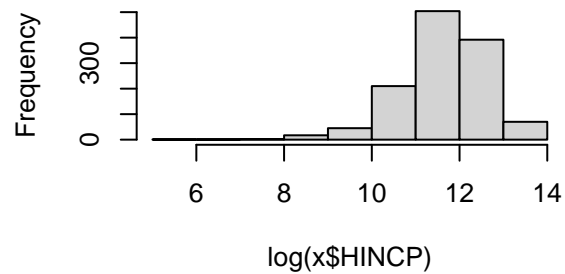
Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

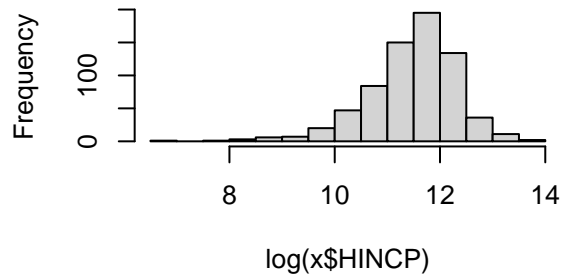


Warning in log(x\$HINCP): NaNs produced

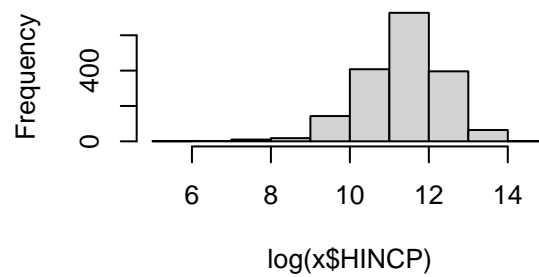
##	34001	34003	34005	34007
## breaks	integer,12	integer,12	integer,14	numeric,19
## counts	integer,11	integer,11	integer,13	integer,18
## density	numeric,11	numeric,11	numeric,13	numeric,18
## mids	numeric,11	numeric,11	numeric,13	numeric,18
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	34011	34013	34015	34017
## breaks	integer,10	integer,12	integer,11	integer,14
## counts	integer,9	integer,11	integer,10	integer,13
## density	numeric,9	numeric,11	numeric,10	numeric,13
## mids	numeric,9	numeric,11	numeric,10	numeric,13
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	34019	34021	34023	34025
## breaks	numeric,14	integer,14	integer,13	integer,12
## counts	integer,13	integer,13	integer,12	integer,11
## density	numeric,13	numeric,13	numeric,12	numeric,11
## mids	numeric,13	numeric,13	numeric,12	numeric,11
## xname	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"	"log(x\$HINCP)"
## equidist	TRUE	TRUE	TRUE	TRUE
##	34027	34029	34031	34035
## breaks	integer,12	numeric,17	integer,12	integer,10
## counts	integer,11	integer,16	integer,11	integer,9
## density	numeric,11	numeric,16	numeric,11	numeric,9
## mids	numeric,11	numeric,16	numeric,11	numeric,9

```
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE              TRUE
##           34037           34039           34041
## breaks     numeric,16      integer,11       integer,11
## counts      integer,15      integer,10       integer,10
## density     numeric,15      numeric,10       numeric,10
## mids         numeric,15      numeric,10       numeric,10
## xname      "log(x$HINCP)" "log(x$HINCP)" "log(x$HINCP)"
## equidist TRUE              TRUE              TRUE
```

Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)



Histogram of log(x\$HINCP)

