

Natsal-data-plots.R

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Thu May 05 16:31:07 2016

```
load("./mrp/data/cleaned-regn-input-mrpNatsal.RData")
```

```
Natsal0 <- Natsal
```

```
rsample <- sample(x=1:nrow(Natsal), prob=Natsal$total_wt, replace=TRUE)
```

```
Natsal <- Natsal[rsample,]
```

```
attach(Natsal)
```

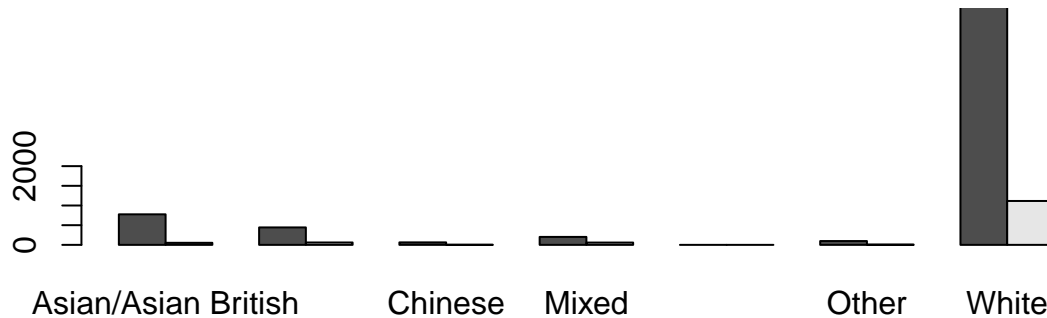
```
# http://stats.stackexchange.com/questions/94026/how-can-i-improve-the-predictive-power-of-this-logistic-regression-model
```

```
names(Natsal)
```

```
##      [1] "sin2"           "id"             "total_wt"
##      [4] "psu"            "strata"         "stratagrp"
##      [7] "stratagrp2"     "stratagrp3"     "rsex"
##     [10] "drink"          "drinkoft"       "manyalc2"
##     [13] "drinknum"       "smokenow"       "exsmoke"
##     [16] "bothmapa2"      "ynotboth2"      "notused"
##     [19] "fsterils"       "msterils"       "conpill"
##     [22] "condom"         "femidom"        "emcypill"
##     [25] "emcyiud"        "coil"           "mirena"
##     [28] "cap"            "injectns"       "gels"
##     [31] "rhythm"         "withdraw"       "implants"
##     [34] "otherfp"        "conpatch"       "ynotused"
##     [37] "yfsteril"       "ymsteril"       "yconpill"
##     [40] "ycondom"        "yfemidom"       "yemcypill"
##     [43] "yemcyiud"       "ycoil"          "ymirena"
##     [46] "ycap"           "yinjectn"       "ygels"
##     [49] "yrhythm"        "ywithdrw"       "yimplant"
##     [52] "yothrfp"        "yconptch"       "dblcon2"
##     [55] "usnotused"      "usfsteril"      "usmsteril"
##     [58] "usconpill"      "uscondom"       "usfemidom"
##     [61] "usemcypill"     "usemcyiud"      "uscoil"
##     [64] "usmirena"       "uscap"          "usinjectn"
##     [67] "usgels"         "usrhythm"       "uswithdrw"
##     [70] "usimplant"      "usother"        "usconptch"
##     [73] "sex4wks"        "cond4wk"        "yrcond"
##     [76] "nonocon"        "het5yrs"        "het1yr"
##     [79] "het3mnt"        "sam1yr"         "sam3mnt"
##     [82] "overlp5y"       "lbyear"         "lbyear2"
##     [85] "lbyear3"        "lbyear4"        "lbyear5"
##     [88] "lbyear6"        "lbyear7"        "lbyear8"
##     [91] "lbyear9"        "lbyear10"       "lbyear11"
##     [94] "lbyear12"       "lbyear13"       "lbyear14"
##     [97] "lbyear15"       "lbyear16"       "lbyear17"
```

```
## [100] "lbyear18"      "lbyear19"      "lbyear20"
## [103] "stdclin"       "chlamydia"     "gonorrh"
## [106] "warts"        "syphilis"      "trich"
## [109] "herpes"       "plichecrab"    "hepatitis"
## [112] "nsungu"       "epididy"       "dkstd"
## [115] "nostd"        "whnchlam"      "chtstwh1"
## [118] "chtstwy1"     "chtrtlgr"      "chlmtest"
## [121] "chltstwh"     "chltstwy"      "chlofref"
## [124] "drcan12m"     "income"        "ethnic"
## [127] "sexid"        "dage"          "rdob"
## [130] "afsex"        "afsexall"      "emergncy"
## [133] "yemergncy"    "sex4wks1"      "sex4wks2"
## [136] "hetnonew"     "totnewyr"      "nocondom"
## [139] "rnssecgp_6"   "rnssecgp_4"    "ethnicgrp"
## [142] "cttestly"     "sex.age"       "ethnic2"
## [145] "increasingdrinker" "student"      "age"
## [148] "sex"          "gor_1"         "gor"
## [151] "age.scaled"   "total_wt.int"
```

```
#x11()
par(mfrow=c(2,1))
tab <- table(cttestly, ethnicgrp)
barplot(tab, legend=levels((cttestly)), ylim=c(0,2000), beside=TRUE)
x <- barplot(prop.table(tab,2)*100, xaxt="n")#, legend=levels((cttestly)))
```

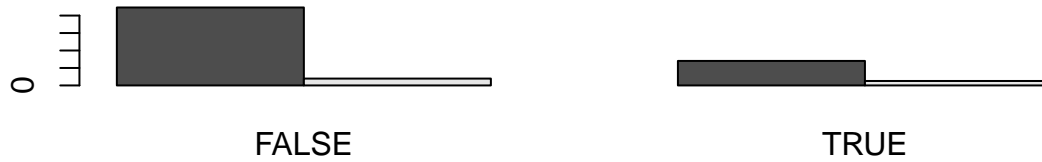


```

# labs <- paste(colnames(tab), "")
# text(cex=1, x=x-.5, y=-50.5, labs, xpd=TRUE, srt=45)

#x11()
tab <- table(cttestly, smokenow)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))

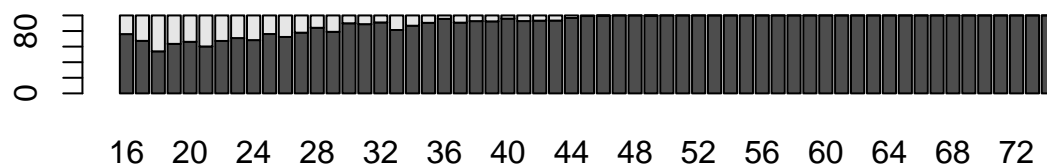
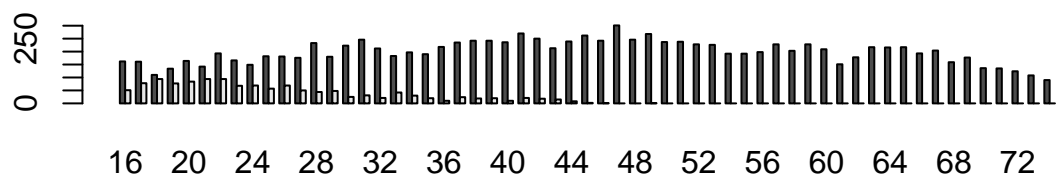
```



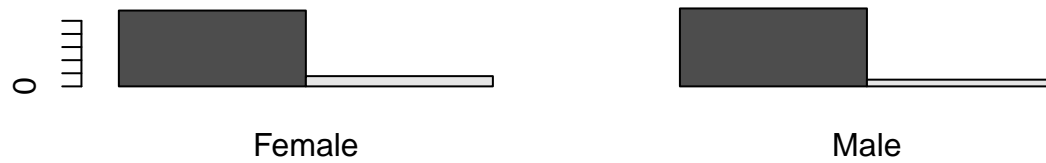
```

#x11()
tab <- table(cttestly, dage)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))

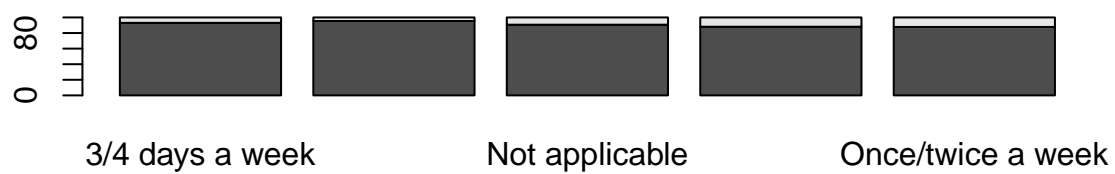
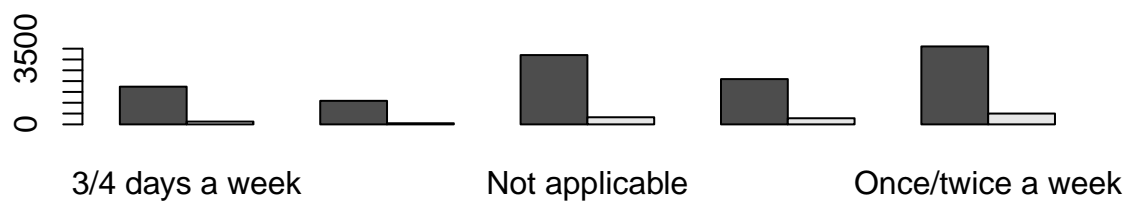
```



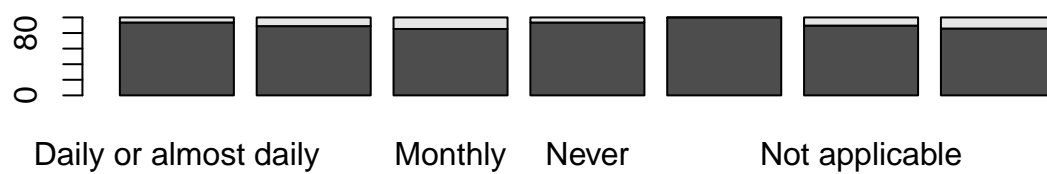
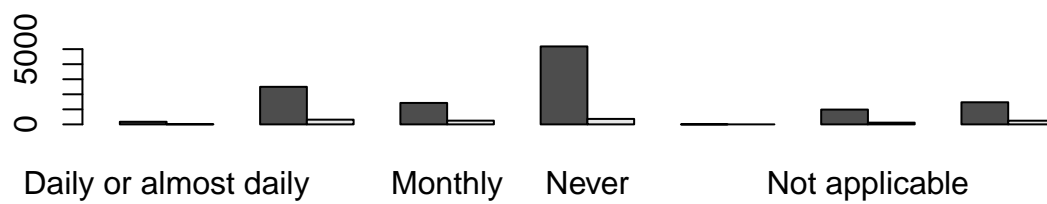
```
#x11()
tab <- table(cttestly, rsex)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
```



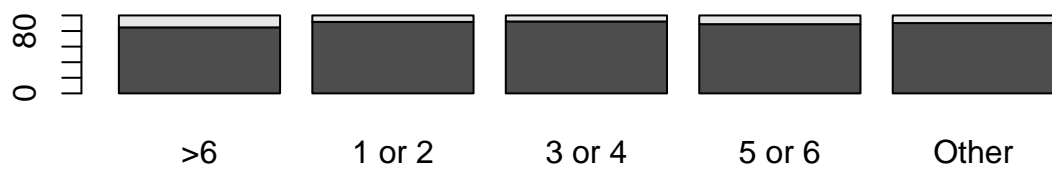
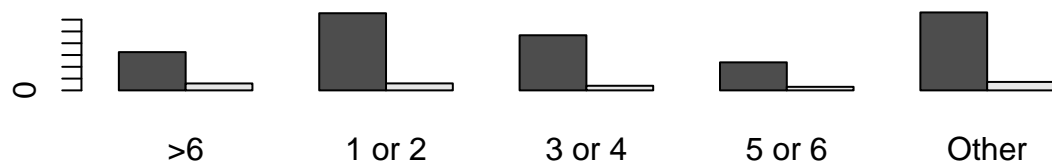
```
#x11()
tab <- table(cttestly, drinkoft)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
```



```
#x11()
tab <- table(cttestly, drinknum)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
```

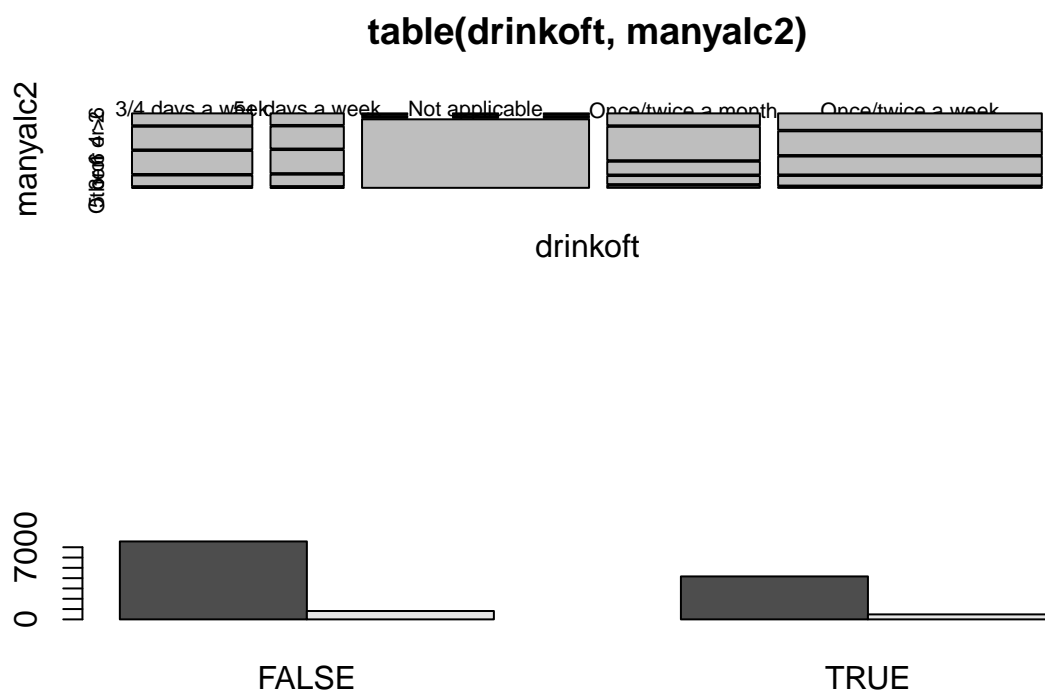


```
#x11()
tab <- table(cttestly, manyalc2)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
```

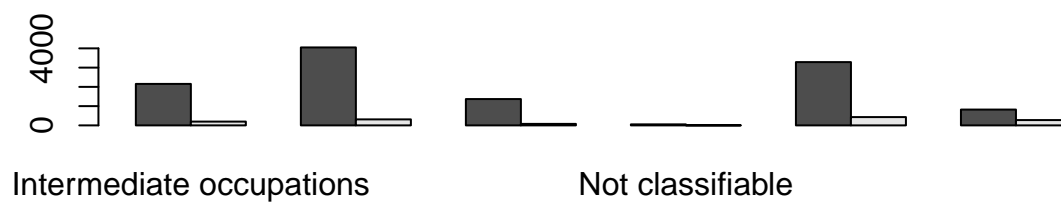


```
plot(table(drinkoft, manyalc2))

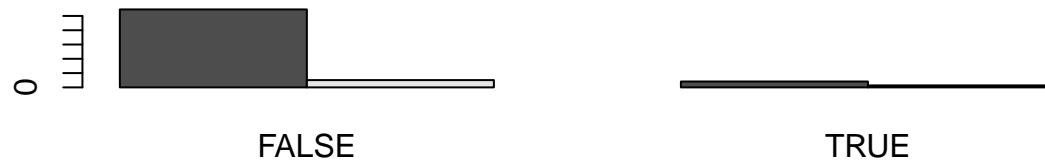
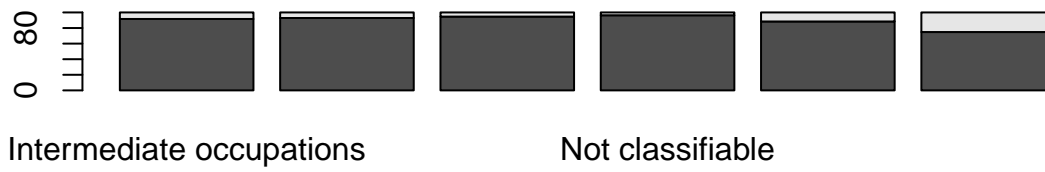
#x11()
tab <- table(cttestly, increasingdrinker)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
```

```
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
#x11()
tab <- table(cttestly, rnssecgp_4)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
```

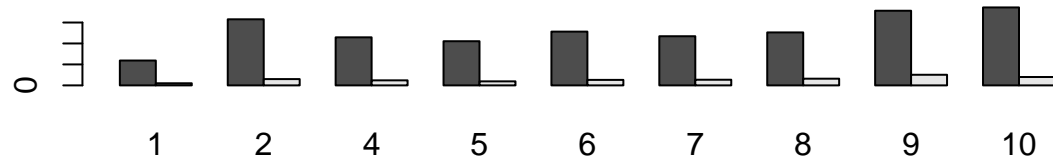


```
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))
#x11()
tab <- table(cttestly, student)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
```



```
barplot(prop.table(tab,2)*100)#, legend=levels((cttestly)))

#x11()
tab <- table(cttestly, gor)
barplot(tab, legend=levels((cttestly)), beside=TRUE)
```



```
barplot(prop.table(tab,2)*100)#, legend=levels((ctestly)))
```

