UKHLS exploration (weighed kin) 4.20

Reading and Cleaning data

Loading required package: splines Loading required package: survey Loading required package: grid Loading required package: Matrix

```
#houskeeping
  #Clear objects already in the environment - start with a clean slate
  rm(list=ls())
  #loading libraries
  library(tidyverse)
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr 1.1.2 v readr 2.1.4
v lubridate 1.9.2 v tidyr 1.3.0
v purrr 1.0.2
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
              masks stats::lag()
i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become
  library(svyVGAM)
Loading required package: VGAM
Loading required package: stats4
```

```
Attaching package: 'Matrix'
The following objects are masked from 'package:tidyr':
    expand, pack, unpack
Loading required package: survival
Attaching package: 'survey'
The following object is masked from 'package: VGAM':
    calibrate
The following object is masked from 'package:graphics':
    dotchart
  library(sjlabelled)
Warning: package 'sjlabelled' was built under R version 4.3.3
Attaching package: 'sjlabelled'
The following object is masked from 'package:forcats':
    as_factor
The following object is masked from 'package:dplyr':
    as_label
The following object is masked from 'package:ggplot2':
    as_label
  library(desctable)
Warning: package 'desctable' was built under R version 4.3.3
```

```
Loading required package: pander
Warning: package 'pander' was built under R version 4.3.3
Attaching package: 'desctable'
The following objects are masked from 'package:stats':
    chisq.test, fisher.test, IQR
  library(summarytools)
Warning: package 'summarytools' was built under R version 4.3.3
Attaching package: 'summarytools'
The following object is masked from 'package:sjlabelled':
    unlabel
The following object is masked from 'package:tibble':
    view
  library(naniar)
Warning: package 'naniar' was built under R version 4.3.3
  library(survey)
  library(svrep)
  library(Hmisc)
```

```
Attaching package: 'Hmisc'
The following objects are masked from 'package:summarytools':
    label, label<-
The following object is masked from 'package:survey':
    deff
The following objects are masked from 'package:dplyr':
    src, summarize
The following objects are masked from 'package:base':
    format.pval, units
  library(srvyr)
Attaching package: 'srvyr'
The following object is masked from 'package:Hmisc':
    summarize
The following object is masked from 'package:stats':
    filter
  library(marginaleffects)
  library(haven)
Attaching package: 'haven'
The following objects are masked from 'package:sjlabelled':
    as_factor, read_sas, read_spss, read_stata, write_sas, zap_labels
```

```
library(catregs)
  library(margins)
  library(modelsummary)
Attaching package: 'modelsummary'
The following object is masked from 'package:Hmisc':
    Mean
The following object is masked from 'package:VGAM':
    Max
  library(zoo)
Attaching package: 'zoo'
The following objects are masked from 'package:base':
    as.Date, as.Date.numeric
  library(mice)
Warning: package 'mice' was built under R version 4.3.3
Attaching package: 'mice'
The following object is masked from 'package:stats':
    filter
The following objects are masked from 'package:base':
    cbind, rbind
```

library(stargazer)

Please cite as: Hlavac, Marek (2022). stargazer: Well-Formatted Regression and Summary Statistics Tables. R package version 5.2.3. https://CRAN.R-project.org/package=stargazer library(texreg) Warning: package 'texreg' was built under R version 4.3.3 Version: 1.39.3 Date: 2023-11-09 Philip Leifeld (University of Essex) Author: Consider submitting praise using the praise or praise_interactive functions. Please cite the JSS article in your publications -- see citation("texreg"). Attaching package: 'texreg' The following object is masked from 'package:tidyr': extract library(VIM) Warning: package 'VIM' was built under R version 4.3.3 Loading required package: colorspace VIM is ready to use. Suggestions and bug-reports can be submitted at: https://github.com/statistikat/VIM/issues Attaching package: 'VIM'

The following object is masked from 'package: VGAM':

```
The following object is masked from 'package:datasets':
    sleep
  library(lattice)
  library(ggplot2)
  library(plyr)
You have loaded plyr after dplyr - this is likely to cause problems.
If you need functions from both plyr and dplyr, please load plyr first, then dplyr:
library(plyr); library(dplyr)
Attaching package: 'plyr'
The following objects are masked from 'package:srvyr':
    mutate, rename, summarise, summarize
The following objects are masked from 'package:Hmisc':
    is.discrete, summarize
The following objects are masked from 'package:dplyr':
    arrange, count, desc, failwith, id, mutate, rename, summarise,
    summarize
The following object is masked from 'package:purrr':
    compact
```

wine

library(dplyr)

Wave 1-5 egoalt (8-6) hhresp (8-6)

```
inpath<-"D:/r git projects/ox-R/final essay/UKHLS n BHPS stata/UKDA-6614-stata/stata/stata missval <- c(-9, -8, -7, -2, -1)
```

Wave 8~13 egoalt (9-13) hhresp (9-13)

```
#wave 8
h_indresp <- read_dta(file=paste0(inpath, "ukhls/h_indresp.dta"))</pre>
h_indresp<-h_indresp%>%dplyr::select(pidp, h_hidp, h_mastat_dv,
         h_age_dv, h_sex_dv, h_hiqual_dv,
        h_mastat_dv, h_racel_dv,h_fimnnet_dv,h_nchild_dv,h_urban_dv,h_gor_dv,h_jbstat,h_s
#wave 9
i_indresp <- read_dta(file=paste0(inpath, "ukhls/i_indresp.dta"))</pre>
i_indresp<-i_indresp%>%dplyr::select(pidp, i_hidp, i_mastat_dv,
         i_age_dv, i_sex_dv, i_hiqual_dv,
         i_mastat_dv, i_racel_dv,i_fimnnet_dv,i_nchild_dv,i_urban_dv,i_gor_dv,i_jbstat,i_s
#wave 10
j_indresp <- read_dta(file=paste0(inpath, "ukhls/j_indresp.dta"))</pre>
j_indresp<-j_indresp%>%dplyr::select(pidp, j_hidp, j_mastat_dv,
         j_age_dv, j_sex_dv, j_hiqual_dv,
         j_mastat_dv, j_racel_dv,j_fimnnet_dv,j_nchild_dv,j_urban_dv,j_gor_dv,j_jbstat,j_s
#wave 11
k_indresp <- read_dta(file=paste0(inpath, "ukhls/k_indresp.dta"))</pre>
k_indresp<-k_indresp%>%dplyr::select(pidp, k_hidp, k_mastat_dv,
         k_age_dv, k_sex_dv, k_hiqual_dv,
        k_mastat_dv, k_racel_dv,k_fimnnet_dv,k_nchild_dv,k_urban_dv,k_gor_dv,k_jbstat,k_s
#wave 12
l_indresp <- read_dta(file=paste0(inpath, "ukhls/l_indresp.dta"))</pre>
l_indresp<-l_indresp%>%dplyr::select(pidp, l_hidp, l_mastat_dv,
         l_age_dv, l_sex_dv, l_hiqual_dv,
         l_mastat_dv, l_racel_dv,l_fimnnet_dv,l_nchild_dv,l_urban_dv,l_gor_dv,l_jbstat,l_s
#wave 13
```

```
m_indresp <- read_dta(file=paste0(inpath, "ukhls/m_indresp.dta"))</pre>
m_indresp<-m_indresp%>%dplyr::select(pidp, m_hidp, m_mastat_dv,
       m_age_dv, m_sex_dv, m_hiqual_dv,
       m mastat dv, m racel dv,m fimnnet dv,m nchild dv,m urban dv,m gor dv,m jbstat,m s
h_egoalt<-read_dta(file=paste0(inpath, "ukhls/h_egoalt.dta"))</pre>
i_egoalt<-read_dta(file=paste0(inpath, "ukhls/i_egoalt.dta"))</pre>
j_egoalt<-read_dta(file=paste0(inpath, "ukhls/j_egoalt.dta"))</pre>
k_egoalt<-read_dta(file=paste0(inpath, "ukhls/k_egoalt.dta"))</pre>
l_egoalt<-read_dta(file=paste0(inpath, "ukhls/l_egoalt.dta"))</pre>
m_egoalt<-read_dta(file=paste0(inpath, "ukhls/m_egoalt.dta"))</pre>
h_hhresp<-read_dta(file=paste0(inpath, "ukhls/h_hhresp.dta")) %>%
dplyr::select(h_hidp,h_tenure_dv)
i_hhresp<-read_dta(file=paste0(inpath, "ukhls/i_hhresp.dta"))</pre>
 dplyr::select(i_hidp,i_tenure_dv)
j_hhresp<-read_dta(file=paste0(inpath, "ukhls/j_hhresp.dta"))</pre>
dplyr::select(j_hidp,j_tenure_dv)
k_hhresp<-read_dta(file=paste0(inpath, "ukhls/k_hhresp.dta"))</pre>
 dplyr::select(k_hidp,k_tenure_dv)
dplyr::select(l_hidp,l_tenure_dv)
dplyr::select(m_hidp,m_tenure_dv)
```

```
##remove all missing values in indresp
for (i in 1:5) {
 h_indresp<- h_indresp %>%
    mutate_all(., list(~na_if(., missval[i])))
for (i in 1:5) {
 i_indresp<- i_indresp %>%
    mutate_all(., list(~na_if(., missval[i])))
for (i in 1:5) {
  j_indresp<- j_indresp %>%
   mutate_all(., list(~na_if(., missval[i])))
for (i in 1:5) {
 k_indresp<- k_indresp %>%
   mutate_all(., list(~na_if(., missval[i])))
for (i in 1:5) {
  l_indresp<- l_indresp %>%
    mutate_all(., list(~na_if(., missval[i])))}
for (i in 1:5) {
 m_indresp<- m_indresp %>%
    mutate_all(., list(~na_if(., missval[i])))}
```

Getting the youths not living with parents in the next wave

Wave $1\sim5$

Wave 8~12

```
mothers <- h_egoalt %>%
  filter(h_relationship_dv %in% c(4) & h_asex==2) %>%
  select(pidp, apidp)
mothers$mother_pidp=mothers$apidp

h_egoalt <- h_egoalt %>%
```

```
full_join(mothers, by=c("pidp", "apidp"))
rm(mothers)
fathers <- h_egoalt %>%
  filter(h_relationship_dv %in% c(4) & h_asex==1) %>%
  select(pidp, apidp)
fathers$father_pidp=fathers$apidp
h_egoalt <- h_egoalt %>%
 full_join(fathers, by=c("pidp", "apidp"))
rm(fathers)
other_relatives <- h_egoalt %>%
 select(pidp, apidp)
other_relatives$other_relatives_pidp=other_relatives$apidp
h_egoalt <- h_egoalt %>%
  full_join(other_relatives, by=c("pidp", "apidp"))
rm(other_relatives)
#at least two relatives in the household structure who are not the focal individual's pare
test<-h_egoalt%>%aggregate(other_relatives_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","relativenum")</pre>
h_egoalt<-join_all(list(h_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent live with a child
child <- h_egoalt %>%
  filter(h_relationship_dv%in% c(9) ) %>%
  select(pidp, apidp)
child$child_pidp=child$apidp
h_{egoalt} \leftarrow h_{egoalt} \%
 full_join(child, by=c("pidp", "apidp"))
rm(child)
#count number of respondent children
test<-h_egoalt%>%aggregate(child_pidp~pidp,FUN=length)
```

```
colnames(test)<-c("pidp","childnum")</pre>
h_egoalt<-join_all(list(h_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(marriage)
mar <- h egoalt %>%
  filter(h relationship dv%in% c(1,2)) %>%
  select(pidp, apidp)
mar$mar_pidp=mar$apidp
h_egoalt <- h_egoalt %>%
  full_join(mar, by=c("pidp", "apidp"))
rm(mar)
#count number of respondent married partner
test<-h_egoalt%>%aggregate(mar_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_husbandwife")</pre>
h_egoalt<-join_all(list(h_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(cohabitation)
cohab <- h_egoalt %>%
  filter(h relationship dv%in% c(3)) %>%
  select(pidp, apidp)
cohab$cohab_pidp=cohab$apidp
h_egoalt <- h_egoalt %>%
  full_join(cohab, by=c("pidp", "apidp"))
rm(cohab)
#count number of cohabitee
test<-h_egoalt%>%aggregate(cohab_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_cohabitee")</pre>
h_egoalt<-join_all(list(h_egoalt,test),by="pidp",type="left")</pre>
rm(test)
# leave: not live with mothers
h_egoalt$separated_m <- labelled(</pre>
  ifelse((!is.na(h_egoalt$mother_pidp) & h_egoalt$h_enwstat==5), -8,
  ifelse((!is.na(h_egoalt$mother_pidp) & h_egoalt$h_anwstat==5), 3,
  ifelse((!is.na(h_egoalt$mother_pidp) &
          h_egoalt$h_anwstat==2 & h_egoalt$h_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(h_egoalt$mother_pidp) &
```

```
h_{\text{egoalt}}=2), c(3,4) & h_{\text{egoalt}}=2, 2,
  ifelse((!is.na(h_egoalt$mother_pidp) &
          h_egoalt$h_anwstat==2 & h_egoalt$h_enwstat==2), 1,
  ifelse((!is.na(h_egoalt$mother_pidp)& h_egoalt$h_anwstat==1), 0,
        ifelse((!is.na(h_egoalt$mother_pidp)), -9, NA))))))),
  label = "Whether living with biological mother at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated_m at next wave" = 1,
             "possibly separated_m at next wave" = 2,
             "mother not alive at next wave" = 3))
#leave: not live with fathers
h_egoalt$separated_f <- labelled(</pre>
  ifelse((!is.na(h_egoalt$father_pidp) & h_egoalt$h_enwstat==5), -8,
  ifelse((!is.na(h_egoalt$father_pidp) & h_egoalt$h_anwstat==5), 3,
  ifelse((!is.na(h_egoalt$father_pidp) &
          h_egoalt$h_anwstat==2 & h_egoalt$h_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(h_egoalt$father_pidp) &
          h_{\text{egoalt}}=2), c(3,4) & h_{\text{egoalt}}=2, 2,
  ifelse((!is.na(h egoalt$father pidp) &
          h_egoalt$h_anwstat==2 & h_egoalt$h_enwstat==2), 1,
  ifelse((!is.na(h_egoalt$father_pidp) & h_egoalt$h_anwstat==1), 0,
        ifelse((!is.na(h_egoalt$father_pidp)), -9, NA))))))),
  label = "Whether living with biological father at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated at next wave" = 1,
             "possibly separated at next wave" = 2,
             "father not alive at next wave" = 3))
#fill the separate indicator
h_egoalt<-h_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_m,.direction="updown")
h_egoalt<-h_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_f,.direction="updown")
#find out who separate from both parents
#not live with both parents
```

```
h_egoalt$separated<- labelled(</pre>
  ifelse((h_egoalt$separated_m%in%c(1)&h_egoalt$separated_f%in%c(1)),1,
  ifelse((h_egoalt$separated_m%in%c(1)& is.na(h_egoalt$separated_f)),1,#one
  ifelse((h egoalt$separated f%in%c(1)& is.na(h egoalt$separated m)),1,
  ifelse((h_egoalt$separated_f%in%c(1)& h_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((h_egoalt$separated_m%in%c(1)& h_egoalt$separated_f%in%c(3)),1,
  #possibly separated
  ifelse((h_egoalt$separated_m%in%c(2)&h_egoalt$separated_f%in%c(2)),2,
  ifelse((h_egoalt$separated_m%in%c(1)&h_egoalt$separated_f%in%c(2)),2,
  ifelse((h_egoalt\$separated_m\%in\%c(2)\&h_egoalt\$separated_f\%in\%c(1)), 2,\\
  ifelse((h_egoalt$separated_m%in%c(2)& is.na(h_egoalt$separated_f)),2,#one
  ifelse((h_egoalt$separated_f%in%c(2)& is.na(h_egoalt$separated_m)),2,
  ifelse((h_egoalt$separated_f%in%c(2)& h_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((h_egoalt$separated_m%in%c(2)& h_egoalt$separated_f%in%c(3)),1,
  #living together
  ifelse((h_egoalt$separated_m%in%c(0)|h_egoalt$separated_f%in%c(0)),0,
  #respondents dead
  ifelse((h_egoalt\$separated_m\%in\%c(-8)|h_egoalt\$separated_f\%in\%c(-8)),-8,
         ifelse((!is.na(h_egoalt$separated_m)|!is.na(h_egoalt$separated_f)),-9,NA))))))))
  label = "Whether living with at least one biological parent at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together with at least one parent at next wave" = 0,
             "separated from both parents at next wave" = 1,
             "possibly separated from both parents at next wave" = 2))
#living with both parents indicator
h_egoalt<-h_egoalt %>%group_by(pidp) %>%tidyr::fill(mother_pidp,.direction="updown")
h_egoalt<-h_egoalt %>%group_by(pidp) %>%tidyr::fill(father_pidp,.direction="updown")
#flags for living with mother or father this wave
h_egoalt<-h_egoalt%>%group_by(pidp)%>%mutate(livewithmother=ifelse(!is.na(h_egoalt$mother_
```

```
h_egoalt<-h_egoalt%>%group_by(pidp)%>%mutate(livewithfather=ifelse(!is.na(h_egoalt$father_
#living with both parents this wave
h_egoalt$liveboth<-c(0)</pre>
h egoalt$liveboth[h egoalt$livewithmother==1 & h egoalt$livewithfather==1]<-1
#a column for living with mother, father, or both this wave
h_egoalt$lvpa<-c(0)</pre>
h_egoalt$lvpa[h_egoalt$livewithmother==1]<-1</pre>
h_egoalt$lvpa[h_egoalt$livewithfather==1]<-2
h_egoalt$lvpa[h_egoalt$liveboth==1]<-3
h_egoalt$lvpa[h_egoalt$lvpa==0]<-NA
h_egoalt$lvpa<-factor(h_egoalt$lvpa, levels=c(1,2,3),labels=c("live with mother","live with
#select only for those who live with at least one parent this wave and at risk of leaving
h_egoalt<-h_egoalt[which(h_egoalt$lvpa%in%c("live with mother","live with father","live wi
#merge datafiles
h_indresp<-h_indresp%>%dplyr::select(pidp, h_hidp, h_mastat_dv,
         h_age_dv, h_sex_dv, h_hiqual_dv,
         h_mastat_dv, h_racel_dv,h_fimnnet_dv,h_nchild_dv,h_urban_dv,h_gor_dv,h_jbstat,h_s
h_egoaltt<-h_egoalt%>%dplyr::select(pidp,mother_pidp,father_pidp,separated_m,separated_f,s
#child levels
h_indrespc<-join_all(list(h_indresp,h_egoaltt), by='pidp', type='left')
h_indrespc<-h_indrespc %>% distinct_at(vars(pidp), .keep_all = TRUE)
h_indrespc<-h_indrespc[(which(!is.na(h_indrespc$mother_pidp)|!is.na(h_indrespc$father_pidp
#add parent levels
h_indresppt <-h_indresp
#join parental data by pidp
father<-as.data.frame(h_indrespc$father_pidp)</pre>
mother <- as.data.frame(h_indrespc$mother_pidp)
```

```
colnames(father)[1]<-"pidp"</pre>
colnames(mother)[1]<-"pidp"</pre>
#attached parental values
h indresppf<-join_all(list(father,h indresppt), by='pidp', type='left')
h_indresppm<-join_all(list(mother,h_indresppt), by='pidp', type='left')
#for housing tenure
h_indresppf<-join_all(list(h_indresppf,h_hhresp), by='h_hidp', type='left')
h_indresppm<-join_all(list(h_indresppm,h_hhresp), by='h_hidp', type='left')
#rename var to differentiate parents
h_indresppf <-
        rename_at(h_indresppf, vars(starts_with("h_")), ~str_replace(., "h_", "ip_"))
h_indresppm <-
        rename_at(h_indresppm, vars(starts_with("h_")), ~str_replace(., "h_", "im_"))
#join to the child-level data by parent's pidp
colnames(h_indresppf)[1]<-"father_pidp"</pre>
colnames(h_indresppm)[1]<-"mother_pidp"</pre>
h_indresppf<-join_all(list(h_indrespc,h_indresppf), by="father_pidp", type='left')</pre>
h_indresppm<-join_all(list(h_indrespc,h_indresppm), by="mother_pidp", type='left')
h_indrespp<-join_all(list(h_indresppm,h_indresppf), by="pidp", type='left')</pre>
h_indrespp<-h_indrespp %>% distinct_at(vars(pidp), .keep_all = TRUE)
rm(father,mother,h_egoaltt,h_indrespc,h_indresppf,h_indresppm,h_indresppt)
#drop duplicate columns
h_indrespp <- h_indrespp %>% select(-contains(".1"))
## add cross wave identifier
h_indrespp$from_wave_h<-1
```

```
mothers <- i_egoalt %>%
  filter(i_relationship_dv %in% c(4) & i_asex==2) %>%
  select(pidp, apidp)
mothers$mother_pidp=mothers$apidp
i_egoalt <- i_egoalt %>%
  full_join(mothers, by=c("pidp", "apidp"))
rm(mothers)
fathers <- i_egoalt %>%
  filter(i_relationship_dv %in% c(4) & i_asex==1) %>%
  select(pidp, apidp)
fathers$father_pidp=fathers$apidp
i_egoalt <- i_egoalt %>%
  full_join(fathers, by=c("pidp", "apidp"))
rm(fathers)
other_relatives <- i_egoalt %>%
  filter(!(i_relationship_dv%in% c(1,2,3,4,9))) %>%
  select(pidp, apidp)
other_relatives$other_relatives_pidp=other_relatives$apidp
i_egoalt <- i_egoalt %>%
  full_join(other_relatives, by=c("pidp", "apidp"))
rm(other_relatives)
#at least two relatives in the household structure who are not the focal individual's pare
test<-i_egoalt%>%aggregate(other_relatives_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","relativenum")</pre>
i_egoalt<-join_all(list(i_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent live with a child
child <- i_egoalt %>%
  filter(i_relationship_dv%in% c(9) ) %>%
  select(pidp, apidp)
```

```
child$child_pidp=child$apidp
i_egoalt <- i_egoalt %>%
  full_join(child, by=c("pidp", "apidp"))
rm(child)
#count number of respondent children
test<-i_egoalt%>%aggregate(child_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","childnum")</pre>
i_egoalt<-join_all(list(i_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(marriage)
mar <- i_egoalt %>%
  filter(i_relationship_dv%in% c(1,2)) %>%
  select(pidp, apidp)
mar$mar_pidp=mar$apidp
i_egoalt <- i_egoalt %>%
  full_join(mar, by=c("pidp", "apidp"))
rm(mar)
#count number of respondent married partner
test<-i_egoalt%>%aggregate(mar_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live with husbandwife")</pre>
i_egoalt<-join_all(list(i_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(cohabitation)
cohab <- i egoalt %>%
  filter(i_relationship_dv%in% c(3) ) %>%
  select(pidp, apidp)
cohab$cohab_pidp=cohab$apidp
i_egoalt <- i_egoalt %>%
  full_join(cohab, by=c("pidp", "apidp"))
rm(cohab)
#count number of cohabitee
test<-i_egoalt%>%aggregate(cohab_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_cohabitee")</pre>
i_egoalt<-join_all(list(i_egoalt,test),by="pidp",type="left")</pre>
rm(test)
# leave: not live with mothers
```

```
i_egoalt$separated_m <- labelled(</pre>
  ifelse((!is.na(i_egoalt$mother_pidp) & i_egoalt$i_enwstat==5), -8,
 ifelse((!is.na(i_egoalt$mother_pidp) & i_egoalt$i_anwstat==5), 3,
 ifelse((!is.na(i_egoalt$mother_pidp) &
          i_egoalt$i_anwstat==2 & i_egoalt$i_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(i egoalt$mother pidp) &
          i_egoalt$i_anwstat %in% c(3,4) & i_egoalt$i_enwstat==2), 2,
 ifelse((!is.na(i egoalt$mother pidp) &
          i_egoalt$i_anwstat==2 & i_egoalt$i_enwstat==2), 1,
  ifelse((!is.na(i_egoalt$mother_pidp)& i_egoalt$i_anwstat==1), 0,
        ifelse((!is.na(i_egoalt$mother_pidp)), -9, NA)))))),
 label = "Whether living with biological mother at the next wave",
 labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated m at next wave" = 1,
             "possibly separated_m at next wave" = 2,
             "mother not alive at next wave" = 3))
#leave: not live with fathers
i_egoalt$separated_f <- labelled(</pre>
  ifelse((!is.na(i_egoalt$father_pidp) & i_egoalt$i_enwstat==5), -8,
 ifelse((!is.na(i_egoalt$father_pidp) & i_egoalt$i_anwstat==5), 3,
 ifelse((!is.na(i_egoalt$father_pidp) &
          i_egoalt$i_anwstat==2 & i_egoalt$i_enwstat %in% c(3,4)), 2,
 ifelse((!is.na(i_egoalt$father_pidp) &
          i_egoalt$i_anwstat %in% c(3,4) & i_egoalt$i_enwstat==2), 2,
 ifelse((!is.na(i_egoalt$father_pidp) &
          i_egoalt$i_anwstat==2 & i_egoalt$i_enwstat==2), 1,
 ifelse((!is.na(i_egoalt$father_pidp) & i_egoalt$i_anwstat==1), 0,
        ifelse((!is.na(i_egoalt$father_pidp)), -9, NA))))))),
 label = "Whether living with biological father at the next wave",
 labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated at next wave" = 1,
             "possibly separated at next wave" = 2,
             "father not alive at next wave" = 3))
#fill the separate indicator
```

```
i_egoalt<-i_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_m,.direction="updown")
i_egoalt<-i_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_f,.direction="updown")
#find out who separate from both parents
#not live with both parents
i_egoalt$separated<- labelled(</pre>
  ifelse((i_egoalt$separated_m%in%c(1)&i_egoalt$separated_f%in%c(1)),1,
  ifelse((i_egoalt$separated_m%in%c(1)& is.na(i_egoalt$separated_f)),1,#one
  ifelse((i_egoalt$separated_f%in%c(1)& is.na(i_egoalt$separated_m)),1,
  ifelse((i_egoalt\$separated_f\%in\%c(\frac{1}{2})\& i_egoalt\$separated_m\%in\%c(\frac{3}{2})), \\ 1, \#deadth
  ifelse((i egoalt$separated m%in%c(1)& i egoalt$separated f%in%c(3)),1,
 #possibly separated
 ifelse((i_egoalt$separated_m%in%c(2)&i_egoalt$separated_f%in%c(2)),2,
  ifelse((i_egoalt$separated_m%in%c(1)&i_egoalt$separated_f%in%c(2)),2,
 ifelse((i_egoalt$separated_m%in%c(2)&i_egoalt$separated_f%in%c(1)),2,
  ifelse((i_egoalt$separated_m%in%c(2)& is.na(i_egoalt$separated_f)),2,#one
  ifelse((i_egoalt$separated_f%in%c(2)& is.na(i_egoalt$separated_m)),2,
  ifelse((i_egoalt$separated_f%in%c(2)& i_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((i_egoalt$separated_m%in%c(2)& i_egoalt$separated_f%in%c(3)),1,
 #living together
 ifelse((i_egoalt$separated_m%in%c(0)|i_egoalt$separated_f%in%c(0)),0,
 #respondents dead
  ifelse((i_egoalt$separated_m%in%c(-8)|i_egoalt$separated_f%in%c(-8)),-8,
         ifelse((!is.na(i_egoalt$separated_m)|!is.na(i_egoalt$separated_f)),-9,NA)))))))
 label = "Whether living with at least one biological parent at the next wave",
 labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together with at least one parent at next wave" = 0,
             "separated from both parents at next wave" = 1,
             "possibly separated from both parents at next wave" = 2))
```

#living with both parents indicator

```
i_egoalt<-i_egoalt %>%group_by(pidp) %>%tidyr::fill(mother_pidp,.direction="updown")
i_egoalt<-i_egoalt %>%group_by(pidp) %>%tidyr::fill(father_pidp,.direction="updown")
#flags for living with mother or father this wave
i_egoalt<-i_egoalt%>%group_by(pidp)%>%mutate(livewithmother=ifelse(!is.na(i_egoalt$mother_
i_egoalt<-i_egoalt%>%group_by(pidp)%>%mutate(livewithfather=ifelse(!is.na(i_egoalt$father_
#living with both parents this wave
i_egoalt$liveboth<-c(0)
i_egoalt$liveboth[i_egoalt$livewithmother==1 & i_egoalt$livewithfather==1]<-1
#a column for living with mother, father, or both this wave
i_egoalt$lvpa<-c(0)
i_egoalt$lvpa[i_egoalt$livewithmother==1]<-1</pre>
i_egoalt$lvpa[i_egoalt$livewithfather==1]<-2</pre>
i_egoalt$lvpa[i_egoalt$liveboth==1]<-3
i_egoalt$lvpa[i_egoalt$lvpa==0]<-NA
i_egoalt$lvpa<-factor(i_egoalt$lvpa, levels=c(1,2,3),labels=c("live with mother","live wit
#select only for those who live with at least one parent this wave and at risk of leaving
i_egoalt<-i_egoalt[which(i_egoalt$lvpa%in%c("live with mother","live with father","live wi
#merge datafiles
i_indresp<-i_indresp%>%dplyr::select(pidp, i_hidp, i_mastat_dv,
         i_age_dv, i_sex_dv, i_hiqual_dv,
         i_mastat_dv, i_racel_dv,i_fimnnet_dv,i_nchild_dv,i_urban_dv,i_gor_dv,i_jbstat,i_s
i_egoaltt<-i_egoalt%>%dplyr::select(pidp,mother_pidp,father_pidp,separated_m,separated_f,s
#child levels
i_indrespc<-join_all(list(i_indresp,i_egoaltt), by='pidp', type='left')</pre>
i_indrespc<-i_indrespc %>% distinct_at(vars(pidp), .keep_all = TRUE)
i_indrespc<-i_indrespc[(which(!is.na(i_indrespc$mother_pidp)|!is.na(i_indrespc$father_pidp
```

```
#add parent levels
i_indresppt <-i_indresp</pre>
#join parental data by pidp
father<-as.data.frame(i_indrespc$father_pidp)</pre>
mother<-as.data.frame(i_indrespc$mother_pidp)</pre>
colnames(father)[1]<-"pidp"</pre>
colnames(mother)[1]<-"pidp"</pre>
#attached parental values
i_indresppf<-join_all(list(father,i_indresppt), by='pidp', type='left')</pre>
i_indresppm<-join_all(list(mother,i_indresppt), by='pidp', type='left')</pre>
#for housing tenure
i_indresppf<-join_all(list(i_indresppf,i_hhresp), by='i_hidp', type='left')</pre>
i_indresppm<-join_all(list(i_indresppm,i_hhresp), by='i_hidp', type='left')</pre>
#rename var to differentiate parents
i_indresppf <-
        rename_at(i_indresppf, vars(starts_with("i_")), ~str_replace(., "i_", "ip_"))
i_indresppm <-
        rename_at(i_indresppm, vars(starts_with("i_")), ~str_replace(., "i_", "im_"))
#join to the child-level data by parent's pidp
colnames(i_indresppf)[1]<-"father_pidp"</pre>
colnames(i_indresppm)[1]<-"mother_pidp"</pre>
i_indresppf<-join_all(list(i_indrespc,i_indresppf), by="father_pidp", type='left')</pre>
i_indresppm<-join_all(list(i_indrespc,i_indresppm), by="mother_pidp", type='left')</pre>
i_indrespp<-join_all(list(i_indresppm,i_indresppf), by="pidp", type='left')</pre>
i_indrespp<-i_indrespp %>% distinct_at(vars(pidp), .keep_all = TRUE)
rm(father,mother,i_egoaltt,i_indrespc,i_indresppf,i_indresppm,i_indresppt)
#drop duplicate columns
i_indrespp <- i_indrespp %>% select(-contains(".1"))
```

```
## add cross wave identifier
i_indrespp$from_wave_i<-1</pre>
```

```
mothers <- j_egoalt %>%
  filter(j_relationship_dv %in% c(4) & j_asex==2) %>%
  select(pidp, apidp)
{\tt mothers\$mother\_pidp=mothers\$apidp}
j_egoalt <- j_egoalt %>%
  full_join(mothers, by=c("pidp", "apidp"))
rm(mothers)
fathers <- j_egoalt %>%
  filter(j_relationship_dv %in% c(4) & j_asex==1) %>%
  select(pidp, apidp)
fathers$father_pidp=fathers$apidp
j_egoalt <- j_egoalt %>%
  full_join(fathers, by=c("pidp", "apidp"))
rm(fathers)
other_relatives <- j_egoalt %>%
  filter(!(j_relationship_dv%in% c(1,2,3,4,9))) %>%
  select(pidp, apidp)
other_relatives$other_relatives_pidp=other_relatives$apidp
j_egoalt <- j_egoalt %>%
  full_join(other_relatives, by=c("pidp", "apidp"))
rm(other_relatives)
#at least two relatives in the household structure who are not the focal individual's pare
test<-j_egoalt%>%aggregate(other_relatives_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","relativenum")</pre>
j_egoalt<-join_all(list(j_egoalt,test),by="pidp",type="left")</pre>
rm(test)
```

```
#whether respondent live with a child
child <- j_egoalt %>%
  filter(j_relationship_dv%in% c(9) ) %>%
  select(pidp, apidp)
child$child_pidp=child$apidp
j_egoalt <- j_egoalt %>%
  full_join(child, by=c("pidp", "apidp"))
rm(child)
#count number of respondent children
test<-j_egoalt%>%aggregate(child_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","childnum")</pre>
j_egoalt<-join_all(list(j_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(marriage)
mar <- j_egoalt %>%
  filter(j_relationship_dv%in% c(1,2)) %>%
  select(pidp, apidp)
mar$mar_pidp=mar$apidp
j_egoalt <- j_egoalt %>%
  full_join(mar, by=c("pidp", "apidp"))
rm(mar)
#count number of respondent married partner
test<-j_egoalt%>%aggregate(mar_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_husbandwife")</pre>
j_egoalt<-join_all(list(j_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(cohabitation)
cohab <- j_egoalt %>%
  filter(j_relationship_dv%in% c(3) ) %>%
  select(pidp, apidp)
cohab$cohab_pidp=cohab$apidp
j_egoalt <- j_egoalt %>%
  full_join(cohab, by=c("pidp", "apidp"))
rm(cohab)
#count number of cohabitee
test<-j_egoalt%>%aggregate(cohab_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_cohabitee")</pre>
j_egoalt<-join_all(list(j_egoalt,test),by="pidp",type="left")</pre>
```

```
rm(test)
# leave: not live with mothers
j_egoalt$separated_m <- labelled(</pre>
  ifelse((!is.na(j_egoalt$mother_pidp) & j_egoalt$j_enwstat==5), -8,
  ifelse((!is.na(j egoalt$mother pidp) & j egoalt$j anwstat==5), 3,
  ifelse((!is.na(j_egoalt$mother_pidp) &
          j_egoalt$j_anwstat==2 & j_egoalt$j_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(j_egoalt$mother_pidp) &
          j_egoalt$j_anwstat %in% c(3,4) & j_egoalt$j_enwstat==2), 2,
  ifelse((!is.na(j_egoalt$mother_pidp) &
          j_egoalt$j_anwstat==2 & j_egoalt$j_enwstat==2), 1,
  ifelse((!is.na(j_egoalt$mother_pidp)& j_egoalt$j_anwstat==1), 0,
        ifelse((!is.na(j_egoalt$mother_pidp)), -9, NA)))))),
  label = "Whether living with biological mother at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated_m at next wave" = 1,
             "possibly separated_m at next wave" = 2,
             "mother not alive at next wave" = 3))
#leave: not live with fathers
j_egoalt$separated_f <- labelled(</pre>
  ifelse((!is.na(j_egoalt$father_pidp) & j_egoalt$j_enwstat==5), -8,
  ifelse((!is.na(j_egoalt$father_pidp) & j_egoalt$j_anwstat==5), 3,
  ifelse((!is.na(j_egoalt$father_pidp) &
          j_egoalt$j_anwstat==2 & j_egoalt$j_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(j_egoalt$father_pidp) &
          j_{\text{egoalt}}_{\text{j_enwstat}}  % c(3,4) & j_{\text{egoalt}}_{\text{j_enwstat}}  2,
  ifelse((!is.na(j_egoalt$father_pidp) &
          j_egoalt$j_anwstat==2 & j_egoalt$j_enwstat==2), 1,
  ifelse((!is.na(j_egoalt$father_pidp) & j_egoalt$j_anwstat==1), 0,
        ifelse((!is.na(j_egoalt$father_pidp)), -9, NA))))))),
  label = "Whether living with biological father at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated at next wave" = 1,
             "possibly separated at next wave" = 2,
```

```
"father not alive at next wave" = 3))
#fill the separate indicator
j_egoalt<-j_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_m,.direction="updown")
j_egoalt<-j_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_f,.direction="updown")
#find out who separate from both parents
#not live with both parents
j_egoalt$separated<- labelled(</pre>
  ifelse((j_egoalt$separated_m%in%c(1)&j_egoalt$separated_f%in%c(1)),1,
  ifelse((j_egoalt$separated_m%in%c(1)& is.na(j_egoalt$separated_f)),1,#one
  ifelse((j_egoalt$separated_f%in%c(1)& is.na(j_egoalt$separated_m)),1,
  ifelse((j_egoalt$separated_f%in%c(1)& j_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((j_egoalt$separated_m%in%c(1)& j_egoalt$separated_f%in%c(3)),1,
  #possibly separated
  ifelse((j_egoalt$separated_m%in%c(2)&j_egoalt$separated_f%in%c(2)),2,
  ifelse((j_egoalt$separated_m%in%c(1)&j_egoalt$separated_f%in%c(2)),2,
  ifelse((j\_egoalt\$separated\_m\%in\%c(2)\&j\_egoalt\$separated\_f\%in\%c(1)), 2,
  ifelse((j_egoalt$separated_m%in%c(2)& is.na(j_egoalt$separated_f)),2,#one
  ifelse((j_egoalt$separated_f%in%c(2)& is.na(j_egoalt$separated_m)),2,
  ifelse((j_egoalt$separated_f%in%c(2)& j_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((j_egoalt$separated_m%in%c(2)& j_egoalt$separated_f%in%c(3)),1,
  #living together
  ifelse((j_egoalt$separated_m%in%c(0)|j_egoalt$separated_f%in%c(0)),0,
  #respondents dead
  ifelse((j_egoalt$separated_m%in%c(-8)|j_egoalt$separated_f%in%c(-8)), -8,
         ifelse((!is.na(j_egoalt$separated_m)|!is.na(j_egoalt$separated_f)),-9,NA)))))))
  label = "Whether living with at least one biological parent at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together with at least one parent at next wave" = 0,
             "separated from both parents at next wave" = 1,
```

```
"possibly separated from both parents at next wave" = \frac{2}{1})
#living with both parents indicator
j_egoalt<-j_egoalt %>%group_by(pidp) %>%tidyr::fill(mother_pidp,.direction="updown")
j_egoalt<-j_egoalt %>%group_by(pidp) %>%tidyr::fill(father_pidp,.direction="updown")
#flags for living with mother or father this wave
j_egoalt<-j_egoalt%>%group_by(pidp)%>%mutate(livewithmother=ifelse(!is.na(j_egoalt$mother_
j_egoalt<-j_egoalt%>%group_by(pidp)%>%mutate(livewithfather=ifelse(!is.na(j_egoalt$father_
#living with both parents this wave
j_egoalt$liveboth<-c(0)</pre>
j_egoalt$liveboth[j_egoalt$livewithmother==1 & j_egoalt$livewithfather==1]<-1
#a column for living with mother, father, or both this wave
j_egoalt$lvpa<-c(0)</pre>
j_egoalt$lvpa[j_egoalt$livewithmother==1]<-1</pre>
j_egoalt$lvpa[j_egoalt$livewithfather==1]<-2</pre>
j_egoalt$lvpa[j_egoalt$liveboth==1]<-3</pre>
j_egoalt$lvpa[j_egoalt$lvpa==0]<-NA</pre>
j_egoalt$lvpa<-factor(j_egoalt$lvpa, levels=c(1,2,3),labels=c("live with mother","live wit
#select only for those who live with at least one parent this wave and at risk of leaving
j_egoalt<-j_egoalt[which(j_egoalt$lvpa%in%c("live with mother","live with father","live wi
#merge datafiles
j_indresp<-j_indresp%>%dplyr::select(pidp, j_hidp, j_mastat_dv,
         j_age_dv, j_sex_dv, j_hiqual_dv,
         j_mastat_dv, j_racel_dv,j_fimnnet_dv,j_nchild_dv,j_urban_dv,j_gor_dv,j_jbstat,j_s
j_egoaltt<-j_egoalt%>%dplyr::select(pidp,mother_pidp,father_pidp,separated_m,separated_f,s
#child levels
j_indrespc<-join_all(list(j_indresp,j_egoaltt), by='pidp', type='left')</pre>
j_indrespc<-j_indrespc %>% distinct_at(vars(pidp), .keep_all = TRUE)
```

```
#add parent levels
j_indresppt <-j_indresp</pre>
#join parental data by pidp
father<-as.data.frame(j_indrespc$father_pidp)</pre>
mother<-as.data.frame(j_indrespc$mother_pidp)</pre>
colnames(father)[1]<-"pidp"</pre>
colnames(mother)[1]<-"pidp"</pre>
#attached parental values
j_indresppf<-join_all(list(father,j_indresppt), by='pidp', type='left')</pre>
j_indresppm<-join_all(list(mother,j_indresppt), by='pidp', type='left')</pre>
#for housing tenure
j_indresppf<-join_all(list(j_indresppf,j_hhresp), by='j_hidp', type='left')
j_indresppm<-join_all(list(j_indresppm,j_hhresp), by='j_hidp', type='left')</pre>
#rename var to differentiate parents
j_indresppf <-</pre>
        rename_at(j_indresppf, vars(starts_with("j_")), ~str_replace(., "j_", "ip_"))
j_indresppm <-
        rename_at(j_indresppm, vars(starts_with("j_")), ~str_replace(., "j_", "im_"))
#join to the child-level data by parent's pidp
colnames(j_indresppf)[1]<-"father_pidp"</pre>
colnames(j_indresppm)[1]<-"mother_pidp"</pre>
j_indresppf<-join_all(list(j_indrespc,j_indresppf), by="father_pidp", type='left')</pre>
j_indresppm<-join_all(list(j_indrespc,j_indresppm), by="mother_pidp", type='left')
j_indrespp<-join_all(list(j_indresppm,j_indresppf), by="pidp", type='left')</pre>
j_indrespp<-j_indrespp %>% distinct_at(vars(pidp), .keep_all = TRUE)
rm(father,mother,j_egoaltt,j_indrespc,j_indresppf,j_indresppm,j_indresppt)
```

```
#drop duplicate columns
j_indrespp <- j_indrespp %>% select(-contains(".1"))

j_indrespp$from_wave_j<-1</pre>
```

```
mothers <- k_egoalt %>%
 filter(k_relationship_dv %in% c(4) & k_asex==2) %>%
  select(pidp, apidp)
mothers$mother_pidp=mothers$apidp
k_{egoalt} \leftarrow k_{egoalt} \%
 full_join(mothers, by=c("pidp", "apidp"))
rm(mothers)
fathers <- k_egoalt %>%
 filter(k_relationship_dv %in% c(4) & k_asex==1) %>%
  select(pidp, apidp)
fathers$father_pidp=fathers$apidp
k_egoalt <- k_egoalt %>%
  full_join(fathers, by=c("pidp", "apidp"))
rm(fathers)
other_relatives <- k_egoalt %>%
 select(pidp, apidp)
other_relatives$other_relatives_pidp=other_relatives$apidp
k_egoalt <- k_egoalt %>%
 full_join(other_relatives, by=c("pidp", "apidp"))
rm(other_relatives)
#at least two relatives in the household structure who are not the focal individual's pare
test<-k_egoalt%>%aggregate(other_relatives_pidp~pidp,FUN=length)
```

```
colnames(test)<-c("pidp","relativenum")</pre>
k_egoalt<-join_all(list(k_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent live with a child
child <- k egoalt %>%
  filter(k_relationship_dv%in% c(9) ) %>%
  select(pidp, apidp)
child$child_pidp=child$apidp
k_egoalt <- k_egoalt %>%
  full_join(child, by=c("pidp", "apidp"))
rm(child)
#count number of respondent children
test<-k_egoalt%>%aggregate(child_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","childnum")</pre>
k_egoalt<-join_all(list(k_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(marriage)
mar <- k egoalt %>%
  filter(k_relationship_dv%in% c(1,2) ) %>%
  select(pidp, apidp)
mar$mar_pidp=mar$apidp
k_egoalt <- k_egoalt %>%
  full_join(mar, by=c("pidp", "apidp"))
rm(mar)
#count number of respondent married partner
test<-k_egoalt%>%aggregate(mar_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_husbandwife")</pre>
k_egoalt<-join_all(list(k_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(cohabitation)
cohab <- k_egoalt %>%
  filter(k_relationship_dv%in% c(3) ) %>%
  select(pidp, apidp)
cohab$cohab_pidp=cohab$apidp
k_egoalt <- k_egoalt %>%
  full_join(cohab, by=c("pidp", "apidp"))
```

```
rm(cohab)
#count number of cohabitee
test<-k_egoalt%>%aggregate(cohab_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_cohabitee")</pre>
k_egoalt<-join_all(list(k_egoalt,test),by="pidp",type="left")</pre>
rm(test)
# leave: not live with mothers
k egoalt$separated m <- labelled(</pre>
  ifelse((!is.na(k_egoalt$mother_pidp) & k_egoalt$k_enwstat==5), -8,
  ifelse((!is.na(k_egoalt$mother_pidp) & k_egoalt$k_anwstat==5), 3,
  ifelse((!is.na(k_egoalt$mother_pidp) &
          k_{egoalt} anwstat==2 & k_{egoalt} enwstat %in% c(3,4)), 2,
  ifelse((!is.na(k_egoalt$mother_pidp) &
          k_{\text{egoalt}} = 0.00 k_egoaltk_{\text{enwstat}} = 0.00 k_egoaltk_{\text{enwstat}} = 0.00 k_egoaltk_{\text{enwstat}} = 0.00
  ifelse((!is.na(k_egoalt$mother_pidp) &
          k_egoalt$k_anwstat==2 & k_egoalt$k_enwstat==2), 1,
  ifelse((!is.na(k_egoalt$mother_pidp)& k_egoalt$k_anwstat==1), 0,
        ifelse((!is.na(k_egoalt$mother_pidp)), -9, NA))))))),
  label = "Whether living with biological mother at the next wave",
  labels = c("missing" = -9,
              "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated m at next wave" = 1,
              "possibly separated_m at next wave" = 2,
              "mother not alive at next wave" = 3))
#leave: not live with fathers
k_egoalt$separated_f <- labelled(</pre>
  ifelse((!is.na(k_egoalt$father_pidp) & k_egoalt$k_enwstat==5), -8,
  ifelse((!is.na(k_egoalt$father_pidp) & k_egoalt$k_anwstat==5), 3,
  ifelse((!is.na(k_egoalt$father_pidp) &
          k_egoalt$k_anwstat==2 & k_egoalt$k_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(k_egoalt$father_pidp) &
          k_{egoalt} anwstat in c(3,4) & k_{egoalt} enwstat==2), 2,
  ifelse((!is.na(k_egoalt$father_pidp) &
          k_egoalt$k_anwstat==2 & k_egoalt$k_enwstat==2), 1,
  ifelse((!is.na(k_egoalt$father_pidp) & k_egoalt$k_anwstat==1), 0,
        ifelse((!is.na(k_egoalt$father_pidp)), -9, NA))))))),
```

```
label = "Whether living with biological father at the next wave",
  labels = c("missing" = -9,
              "ego not alive at next wave" = -8,
              "living together at next wave" = 0, "separated at next wave" = 1,
              "possibly separated at next wave" = 2,
              "father not alive at next wave" = 3))
#fill the separate indicator
k_egoalt<-k_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_m,.direction="updown")
k_egoalt<-k_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_f,.direction="updown")
#find out who separate from both parents
#not live with both parents
k_egoalt$separated<- labelled(</pre>
  ifelse((k_egoalt$separated_m%in%c(1)&k_egoalt$separated_f%in%c(1)),1,
  ifelse((k_egoalt$separated_m%in%c(1)& is.na(k_egoalt$separated_f)),1,#one
  ifelse((k_egoalt$separated_f%in%c(1)& is.na(k_egoalt$separated_m)),1,
  ifelse((k_egoalt\$separated_f\%in\%c(\textcolor{red}{1})\& k_egoalt\$separated_m\%in\%c(\textcolor{red}{3})), \textcolor{red}{1}, \texttt{\#} deadth
  ifelse((k_egoalt$separated_m%in%c(1)& k_egoalt$separated_f%in%c(3)),1,
  #possibly separated
  ifelse((k_egoalt$separated_m%in%c(2)&k_egoalt$separated_f%in%c(2)),2,
  ifelse((k_egoalt$separated_m%in%c(1)&k_egoalt$separated_f%in%c(2)),2,
  ifelse((k_egoaltseparated_m\%in\%c(2)\&k_egoalt\\separated_f\%in\%c(1)),2,
  ifelse((k_egoalt$separated_m%in%c(2)& is.na(k_egoalt$separated_f)),2,#one
  ifelse((k_egoalt$separated_f%in%c(2)& is.na(k_egoalt$separated_m)),2,
  ifelse((k_egoalt\$separated_f\%in\%c(2)\& k_egoalt\$separated_m\%in\%c(3)), 1, \#deadth
  ifelse((k_egoalt$separated_m%in%c(2)& k_egoalt$separated_f%in%c(3)),1,
  #living together
  ifelse((k_egoaltseparated_m\%in\%c(0)|k_egoalt\\separated_f\%in\%c(0)),0,
  #respondents dead
  ifelse((k_egoalt\$separated_m\%in\%c(-8))k_egoalt\$separated_f\%in\%c(-8)),-8,
         ifelse((!is.na(k_egoalt$separated_m)|!is.na(k_egoalt$separated_f)),-9,NA))))))))
```

```
label = "Whether living with at least one biological parent at the next wave",
  labels = c("missing" = -9,
            "ego not alive at next wave" = -8,
            "living together with at least one parent at next wave" = 0,
            "separated from both parents at next wave" = 1,
            "possibly separated from both parents at next wave" = 2))
#living with both parents indicator
k_egoalt<-k_egoalt %>%group_by(pidp) %>%tidyr::fill(mother_pidp,.direction="updown")
k_egoalt<-k_egoalt %>%group_by(pidp) %>%tidyr::fill(father_pidp,.direction="updown")
#flags for living with mother or father this wave
k_egoalt<-k_egoalt%>%group_by(pidp)%>%mutate(livewithmother=ifelse(!is.na(k_egoalt$mother_
k_egoalt<-k_egoalt%%group_by(pidp)%>%mutate(livewithfather=ifelse(!is.na(k_egoalt$father_
#living with both parents this wave
k_egoalt$liveboth<-c(0)</pre>
#a column for living with mother, father, or both this wave
k_egoalt$lvpa<-c(0)
k_egoalt = [k_egoalt = 1] < -1
k_egoalt$lvpa[k_egoalt$livewithfather==1]<-2
k_egoalt$lvpa[k_egoalt$liveboth==1]<-3
k_egoalt$lvpa[k_egoalt$lvpa==0]<-NA
k_egoalt$lvpa<-factor(k_egoalt$lvpa, levels=c(1,2,3),labels=c("live with mother","live with
#select only for those who live with at least one parent this wave and at risk of leaving
k_egoalt<-k_egoalt[which(k_egoalt$lvpa%in%c("live with mother","live with father","live wi
#merge datafiles
k_indresp<-k_indresp%>%dplyr::select(pidp, k_hidp, k_mastat_dv,
        k_age_dv, k_sex_dv, k_hiqual_dv,
        k_mastat_dv, k_racel_dv,k_fimnnet_dv,k_nchild_dv,k_urban_dv,k_gor_dv,k_jbstat,k_s
k_egoaltt<-k_egoalt%>%dplyr::select(pidp,mother_pidp,father_pidp,separated_m,separated_f,s
```

```
#child levels
k_indrespc<-join_all(list(k_indresp,k_egoaltt), by='pidp', type='left')</pre>
k_indrespc<-k_indrespc %>% distinct_at(vars(pidp), .keep all = TRUE)
k_indrespc<-k_indrespc[(which(!is.na(k_indrespc$mother_pidp)|!is.na(k_indrespc$father_pidp
#add parent levels
k_indresppt <-k_indresp</pre>
#join parental data by pidp
father<-as.data.frame(k_indrespc$father_pidp)</pre>
mother<-as.data.frame(k_indrespc$mother_pidp)</pre>
colnames(father)[1]<-"pidp"</pre>
colnames(mother)[1]<-"pidp"</pre>
#attached parental values
k_indresppf<-join_all(list(father,k_indresppt), by='pidp', type='left')</pre>
k_indresppm<-join_all(list(mother,k_indresppt), by='pidp', type='left')</pre>
#for housing tenure
k_indresppf<-join_all(list(k_indresppf,k_hhresp), by='k_hidp', type='left')</pre>
k_indresppm<-join_all(list(k_indresppm,k_hhresp), by='k_hidp', type='left')</pre>
#rename var to differentiate parents
k_indresppf <-
        rename_at(k_indresppf, vars(starts_with("k_")), ~str_replace(., "k_", "ip_"))
k_indresppm <-
        rename_at(k_indresppm, vars(starts_with("k_")), ~str_replace(., "k_", "im_"))
#join to the child-level data by parent's pidp
colnames(k_indresppf)[1]<-"father_pidp"</pre>
colnames(k_indresppm)[1]<-"mother_pidp"</pre>
k_indresppf<-join_all(list(k_indrespc,k_indresppf), by="father_pidp", type='left')</pre>
k_indresppm<-join_all(list(k_indrespc,k_indresppm), by="mother_pidp", type='left')</pre>
k_indrespp<-join_all(list(k_indresppm,k_indresppf), by="pidp", type='left')</pre>
```

```
k_indrespp<-k_indrespp %>% distinct_at(vars(pidp), .keep_all = TRUE)

rm(father,mother,k_egoaltt,k_indrespc,k_indresppf,k_indresppm,k_indresppt)

#drop duplicate columns
k_indrespp <- k_indrespp %>% select(-contains(".1"))

k_indrespp$from_wave_k<-1</pre>
```

```
mothers <- l_egoalt %>%
 filter(l_relationship_dv %in% c(4) & l_asex==2) %>%
 select(pidp, apidp)
mothers$mother_pidp=mothers$apidp
l_egoalt <- l_egoalt %>%
 full_join(mothers, by=c("pidp", "apidp"))
rm(mothers)
fathers <- l egoalt %>%
 filter(l_relationship_dv %in% c(4) & l_asex==1) %>%
 select(pidp, apidp)
fathers$father_pidp=fathers$apidp
l_egoalt <- l_egoalt %>%
 full_join(fathers, by=c("pidp", "apidp"))
rm(fathers)
other_relatives <- l_egoalt %>%
 select(pidp, apidp)
other_relatives$other_relatives_pidp=other_relatives$apidp
l_egoalt <- l_egoalt %>%
 full_join(other_relatives, by=c("pidp", "apidp"))
```

```
rm(other_relatives)
#at least two relatives in the household structure who are not the focal individual's pare
test<-l_egoalt%>%aggregate(other_relatives_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","relativenum")</pre>
l_egoalt<-join_all(list(l_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent live with a child
child <- l_egoalt %>%
  filter(l_relationship_dv%in% c(9) ) %>%
  select(pidp, apidp)
child$child_pidp=child$apidp
l_egoalt <- l_egoalt %>%
  full_join(child, by=c("pidp", "apidp"))
rm(child)
#count number of respondent children
test<-l_egoalt%>%aggregate(child_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","childnum")</pre>
l_egoalt<-join_all(list(l_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(marriage)
mar <- l_egoalt %>%
  filter(l_relationship_dv%in% c(1,2)) %>%
  select(pidp, apidp)
mar$mar_pidp=mar$apidp
l_egoalt <- l_egoalt %>%
  full_join(mar, by=c("pidp", "apidp"))
rm(mar)
#count number of respondent married partner
test<-l_egoalt%>%aggregate(mar_pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_husbandwife")</pre>
l_egoalt<-join_all(list(l_egoalt,test),by="pidp",type="left")</pre>
rm(test)
#whether respondent cohabit with a partner(cohabitation)
cohab <- l_egoalt %>%
  filter(l_relationship_dv%in% c(3) ) %>%
```

```
select(pidp, apidp)
cohab$cohab_pidp=cohab$apidp
l_egoalt <- l_egoalt %>%
 full_join(cohab, by=c("pidp", "apidp"))
rm(cohab)
#count number of cohabitee
test<-l egoalt%>%aggregate(cohab pidp~pidp,FUN=length)
colnames(test)<-c("pidp","live_with_cohabitee")</pre>
l_egoalt<-join_all(list(l_egoalt,test),by="pidp",type="left")</pre>
rm(test)
# leave: not live with mothers
l_egoalt$separated_m <- labelled(</pre>
  ifelse((!is.na(l_egoalt$mother_pidp) & l_egoalt$l_enwstat==5), -8,
  ifelse((!is.na(l_egoalt$mother_pidp) & l_egoalt$l_anwstat==5), 3,
  ifelse((!is.na(l_egoalt$mother_pidp) &
          l_egoalt$l_anwstat==2 & l_egoalt$l_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(l_egoalt$mother_pidp) &
          l_egoalt$l_anwstat %in% c(3,4) & l_egoalt$l_enwstat==2), 2,
  ifelse((!is.na(l_egoalt$mother_pidp) &
          l egoalt$1 anwstat==2 & 1 egoalt$1 enwstat==2), 1,
  ifelse((!is.na(l_egoalt$mother_pidp)& l_egoalt$l_anwstat==1), 0,
        ifelse((!is.na(l_egoalt$mother_pidp)), -9, NA))))))),
  label = "Whether living with biological mother at the next wave",
  labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated_m at next wave" = 1,
             "possibly separated_m at next wave" = 2,
             "mother not alive at next wave" = 3))
#leave: not live with fathers
l_egoalt$separated_f <- labelled(</pre>
  ifelse((!is.na(l_egoalt$father_pidp) & l_egoalt$l_enwstat==5), -8,
  ifelse((!is.na(l_egoalt$father_pidp) & l_egoalt$l_anwstat==5), 3,
  ifelse((!is.na(l_egoalt$father_pidp) &
          l_egoalt$l_anwstat==2 & l_egoalt$l_enwstat %in% c(3,4)), 2,
  ifelse((!is.na(l_egoalt$father_pidp) &
          l_egoalt$l_anwstat %in% c(3,4) & l_egoalt$l_enwstat==2), 2,
  ifelse((!is.na(l_egoalt$father_pidp) &
```

```
l_egoalt$l_anwstat==2 & l_egoalt$l_enwstat==2), 1,
  ifelse((!is.na(l_egoalt$father_pidp) & l_egoalt$l_anwstat==1), 0,
        ifelse((!is.na(l_egoalt$father_pidp)), -9, NA)))))),
 label = "Whether living with biological father at the next wave",
 labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together at next wave" = 0, "separated at next wave" = 1,
             "possibly separated at next wave" = 2,
             "father not alive at next wave" = 3))
#fill the separate indicator
l_egoalt<-l_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_m,.direction="updown")
l_egoalt<-l_egoalt %>%group_by(pidp) %>%tidyr::fill(separated_f,.direction="updown")
#find out who separate from both parents
#not live with both parents
l_egoalt$separated<- labelled(</pre>
  ifelse((1 egoalt$separated m%in%c(1)&l egoalt$separated f%in%c(1)),1,
  ifelse((1_egoalt$separated_m%in%c(1)& is.na(1_egoalt$separated_f)),1,#one
  ifelse((l_egoalt$separated_f%in%c(1)& is.na(l_egoalt$separated_m)),1,
 ifelse((l_egoalt$separated_f%in%c(1)& l_egoalt$separated_m%in%c(3)),1,#deadth
  ifelse((1_egoalt$separated_m%in%c(1)& 1_egoalt$separated_f%in%c(3)),1,
 #possibly separated
 ifelse((1_egoalt$separated_m%in%c(2)&1_egoalt$separated_f%in%c(2)),2,
 ifelse((1_egoalt$separated_m%in%c(1)&l_egoalt$separated_f%in%c(2)),2,
 ifelse((l_egoalt$separated_m%in%c(2)&l_egoalt$separated_f%in%c(1)),2,
 ifelse((1 egoalt$separated m%in%c(2)& is.na(1 egoalt$separated f)),2,#one
  ifelse((1_egoalt$separated_f%in%c(2)& is.na(1_egoalt$separated_m)),2,
  ifelse((1_egoalt$separated_f%in%c(2)& 1_egoalt$separated_m%in%c(3)),1,#deadth
 ifelse((l_egoalt$separated_m%in%c(2)& l_egoalt$separated_f%in%c(3)),1,
 #living together
  ifelse((1_egoalt$separated_m%in%c(0)|1_egoalt$separated_f%in%c(0)),0,
```

```
#respondents dead
  ifelse((1_egoalt\$separated_m\%in\%c(-8))1_egoalt\$separated_f\%in\%c(-8)),-8,
         ifelse((!is.na(l_egoalt$separated_m)|!is.na(l_egoalt$separated_f)),-9,NA)))))))
 label = "Whether living with at least one biological parent at the next wave",
 labels = c("missing" = -9,
             "ego not alive at next wave" = -8,
             "living together with at least one parent at next wave" = 0,
             "separated from both parents at next wave" = 1,
             "possibly separated from both parents at next wave" = 2))
#living with both parents indicator
l_egoalt<-l_egoalt %>%group_by(pidp) %>%tidyr::fill(mother_pidp,.direction="updown")
l_egoalt<-l_egoalt %>%group_by(pidp) %>%tidyr::fill(father_pidp,.direction="updown")
#flags for living with mother or father this wave
l_egoalt<-l_egoalt%>%group_by(pidp)%>%mutate(livewithmother=ifelse(!is.na(l_egoalt$mother_
l_egoalt<-l_egoalt%>%group_by(pidp)%>%mutate(livewithfather=ifelse(!is.na(l_egoalt$father_
#living with both parents this wave
l_egoalt$liveboth<-c(0)</pre>
l_egoalt$liveboth[l_egoalt$livewithmother==1 & l_egoalt$livewithfather==1]<-1</pre>
#a column for living with mother, father, or both this wave
l_egoalt$lvpa<-c(0)</pre>
l_egoalt$lvpa[l_egoalt$livewithmother==1]<-1</pre>
l_egoalt$lvpa[l_egoalt$liveboth==1]<-3</pre>
l_egoalt$lvpa[l_egoalt$lvpa==0]<-NA</pre>
l_egoalt$lvpa<-factor(l_egoalt$lvpa, levels=c(1,2,3),labels=c("live with mother","live with
#select only for those who live with at least one parent this wave and at risk of leaving
l_egoalt<-l_egoalt[which(l_egoalt$lvpa%in%c("live with mother","live with father","live wi
#merge datafiles
l_indresp<-l_indresp%>%dplyr::select(pidp, l_hidp, l_mastat_dv,
        l_age_dv, l_sex_dv, l_hiqual_dv,
```

```
l_mastat_dv, l_racel_dv,l_fimnnet_dv,l_nchild_dv,l_urban_dv,l_gor_dv,l_jbstat,l_s
l_egoaltt<-l_egoalt%>%dplyr::select(pidp,mother_pidp,father_pidp,separated_m,separated_f,s
#child levels
l_indrespc<-join_all(list(l_indresp,l_egoaltt), by='pidp', type='left')</pre>
l_indrespc<-l_indrespc %>% distinct_at(vars(pidp), .keep_all = TRUE)
l_indrespc<-l_indrespc[(which(!is.na(l_indrespc$mother_pidp)|!is.na(l_indrespc$father_pidp
#add parent levels
l_indresppt <-l_indresp</pre>
#join parental data by pidp
father<-as.data.frame(l_indrespc$father_pidp)</pre>
mother<-as.data.frame(l_indrespc$mother_pidp)</pre>
colnames(father)[1]<-"pidp"</pre>
colnames(mother)[1]<-"pidp"</pre>
#attached parental values
l_indresppf<-join_all(list(father,l_indresppt), by='pidp', type='left')</pre>
l_indresppm<-join_all(list(mother,l_indresppt), by='pidp', type='left')</pre>
#for housing tenure
l_indresppf<-join_all(list(l_indresppf,l_hhresp), by='l_hidp', type='left')</pre>
l_indresppm<-join_all(list(l_indresppm,l_hhresp), by='l_hidp', type='left')</pre>
#rename var to differentiate parents
l_indresppf <-</pre>
        rename_at(l_indresppf, vars(starts_with("l_")), ~str_replace(., "l_", "ip_"))
        rename_at(l_indresppm, vars(starts_with("l_")), ~str_replace(., "l_", "im_"))
#join to the child-level data by parent's pidp
colnames(l_indresppf)[1]<-"father_pidp"</pre>
colnames(l_indresppm)[1]<-"mother_pidp"</pre>
```

```
l_indresppf<-join_all(list(l_indrespc,l_indresppf), by="father_pidp", type='left')
l_indresppm<-join_all(list(l_indrespc,l_indresppm), by="mother_pidp", type='left')
l_indrespp<-join_all(list(l_indresppm,l_indresppf), by="pidp", type='left')
l_indrespp<-l_indrespp %>% distinct_at(vars(pidp), .keep_all = TRUE)
rm(father,mother,l_egoaltt,l_indrespc,l_indresppf,l_indresppm,l_indresppt)
#drop duplicate columns
l_indrespp <- l_indrespp %>% select(-contains(".1"))
l_indrespp$from_wave_l<-1</pre>
```

Merging all

Coding variables

child level

```
#child level

#level of education completed
indrespb$edu<-c(0)</pre>
```

```
indrespb$edu[indrespb$hiqual_dv%in%c(9)]<-1 #low (everything below)
     indrespb\$edu[indrespb\$hiqual_dv\%in\%c(1,2)]<-3 #high (have a degree)
     indrespb$edu[indrespb$edu==0]<-NA
    print(attr(indrespb$hiqual_dv,"labels"))
                                                         inapplicable
                                                                                                              refusal
                                                                                                                                                don't know
                         missing
                           Degree Other higher degree
                                                                                                                                                    GCSE etc
                                                                                                      A-level etc
Other qualification
                                                No qualification
                                      5
     indrespb$edu<-factor(indrespb$edu,levels=c(1,2,3),labels=c('low',"middle","high"))</pre>
     #two-category education
     indrespb$edu2<-c(0)
     indrespb\$edu2[indrespb\$hiqual\_dv\%in\%c(1,2)] < -1 \ \#high \ (have a degree)
     indrespb$edu2[is.na(indrespb$hiqual_dv)]<-NA
     #family structure
     indrespb$family_status<-c(0)</pre>
     #coresident partner and child
     indrespb$family_status[indrespb$childnum>=1 & indrespb$live_with_husbandwife>=1]<-1
     indrespb$family_status[indrespb$childnum>=1 & indrespb$live_with_cohabitee>=1]<-1
     #coresident partner
     indrespb\$family\_status[indrespb\$live\_with\_cohabitee>= 1    \&indrespb\$childnum== 0] < -2  
     indrespb\$family\_status[indrespb\$live\_with\_husbandwife>=1 \& indrespb\$childnum==0]<-2 & indrespb\$childnum==0|<-2 & indrespb$childnum==0|<-2 & indrespb$child
     indrespb$family_status[indrespb$live_with_cohabitee>=1 &is.na(indrespb$childnum)]<-2
     indrespb$family_status[indrespb$live_with_husbandwife>=1 &is.na(indrespb$childnum)]<-2
     #coresident child
     indrespb$family_status[indrespb$childnum>=1 &indrespb$live_with_husbandwife==0 & indrespb$
     indrespb$family_status[indrespb$childnum>=1 &is.na(indrespb$live_with_husbandwife)& indres
     indrespb$family_status[indrespb$childnum>=1 &indrespb$live_with_husbandwife==0 & is.na(ind
     indrespb$family_status[indrespb$childnum>=1 &is.na(indrespb$live_with_husbandwife) & is.na
     #no coresident partner or child=0
     indrespb$family_status<-factor(indrespb$family_status,levels=c(0,1,2,3),labels=c("No cores
```

```
#age centered
indrespb$age_centered<-indrespb$age_dv-16
#age squared
indrespb$age_sq<-indrespb$age_centered**2
#logarithm of income
indrespb\$fimnnet\_dv[indrespb\$fimnnet\_dv<0]<-0
#add 1 to all income values to avoid -inf in log transformation
indrespb$incomeadd<-indrespb$fimnnet_dv+1</pre>
indrespb$lnincome<-log(indrespb$incomeadd)</pre>
indrespb<-indrespb%>%dplyr::select(-incomeadd)
#racel groups
indrespb$racel<-c(0)
indrespb\$racel[indrespb\$racel\_dv\%in\%c(1,2,3,4)] < -1 \ \#Whites
indrespb\$racel[indrespb\$racel\_dv\%in\%c(14,15)] <-2 \ \#African \ Caribbean
indrespb$racel[indrespb$racel_dv%in%c(9,10,11)]<-3 #Indiani,Pakistani, Bangladeshi</pre>
indrespb\$racel[indrespb\$racel\_dv\%in\%c(12,13)] < -4 \ \#0ther \ Asian
indrespb$racel[indrespb$racel==0]<-5 #Other
indrespb$racel<-factor(indrespb$racel,levels=c(1,2,3,4,5),labels=c("Whites","African Carib
indrespb$racel[is.na(indrespb$racel_dv)]<-NA</pre>
##two-categories
indrespb$racel2<-c(0)
indrespb\$racel2[indrespb\$racel\_dv\%in\%c(1,2,3,4)] <-1 \ \#Whites
indrespb$racel2[!(indrespb$racel_dv\%in\%c(1,2,3,4))]<-2 #non-Whites
indrespb$racel2[is.na(indrespb$racel_dv)]<-NA</pre>
indrespb$racel2<-factor(indrespb$racel2,levels=c(1,2),labels=c("Whites","non-Whites"))</pre>
#education status
indrespb$student<-c(0)
indrespb$student[indrespb$jbstat%in%c(7)]<-1 #full-time student
indrespb$student[is.na(indrespb$jbstat)]<-NA
#employment status
indrespb$unemployed<-c(0)
```

```
indrespb$unemployed[indrespb$jbstat%in%c(3)]<-1
  indrespb$unemployed[is.na(indrespb$jbstat)]<-NA</pre>
  #housing prices
  indrespb$houseprices<-c(0)</pre>
  indrespb$houseprices[indrespb$gor_dv%in%c(1,10,11,12)]<-1 #low
  indrespb$houseprices[indrespb$gor_dv%in%c(2,3,4,5)]<-2 #middle</pre>
  indrespb $houseprices[indrespb $gor_dv\%in\%c(6,7,8,9)] < -3 \# high
  indrespb$houseprices[is.na(indrespb$gor_dv)]<-NA
  indrespb$houseprices<-factor(indrespb$houseprices,levels=c(1,2,3),labels=c("low","middle",
  #urban,sex binary
  indrespb$sex<-c(0)
  indrespb$sex[indrespb$sex_dv==2]<-1 #females
  indrespb$sex[is.na(indrespb$sex_dv)]<-NA
  indrespburban<-c(0)
  indrespburban[indrespb$urban_dv==1]<-1 #urban areas
  indrespb$urban[is.na(indrespb$urban_dv)]<-NA
  indrespb<-indrespb%>%mutate(health=coalesce(sf1,scsf1))
Warning: `..1` and `..2` have conflicting value labels.
i Labels for these values will be taken from `..1`.
x Values: 5
  #living with both biological parents
  indrespb$liveboth<-c(0)</pre>
  indrespb$liveboth[indrespb$lvpa=="live with both parents"]<-1</pre>
  indrespb$liveboth[is.na(indrespb$lvpa)]<-NA</pre>
  #relative num for family structure
  indrespb$relative_num<-c(0)</pre>
  indrespb$relative_num[indrespb$relativenum>=2]<-1</pre>
  indrespb$relative_num<-factor(indrespb$relative_num,levels=c(0,1),labels=c("small hh","lar
parental level
  #level of education completed
  #Fathers:
```

```
indrespbpaedu<-c(0)
  indrespb paedu [indrespb ip\_hiqual\_dv\%in\%c (9)] < -1 \ \#low \ (everything \ below)
  indrespb$paedu[indrespb$ip_hiqual_dv%in%c(3,4,5)]<-2 #middle (completed A-level or seconda
  indrespb paedu [indrespb ip\_hiqual\_dv\%in\%c(1,2)] < -3 \ \#high \ (have \ a \ degree)
  indrespb$paedu[indrespb$paedu==0]<-NA
  print(attr(indrespb$ip_hiqual_dv,"labels"))
                                                                    don't know
           missing
                          inapplicable
                                                  refusal
                                                                     GCSE etc
            Degree Other higher degree
                                                A-level etc
                       No qualification
Other qualification
  indrespb$paedu<-factor(indrespb$paedu,levels=c(1,2,3),labels=c('low',"middle","high"))
  #Mothers:
  indrespb$maedu<-c(0)
  indrespb$maedu[indrespb$im_hiqual_dv%in%c(9)]<-1 #low (everything below)</pre>
  indrespb\macdu[indrespb\macdu[indrespb\macdu]-indrespb\macdu]--2 #middle (completed A-level or secondary)
  indrespb\$maedu[indrespb\$im\_hiqual\_dv\%in\%c(1,2)] < -3 \ \#high \ (have \ a \ degree)
  indrespb\maedu[indrespb\maedu==0]<-NA
  indrespb$maedu<-factor(indrespb$maedu,levels=c(1,2,3),labels=c('low',"middle","high"))
  #the highest education level between two parents
  indrespb<-transform(indrespb, parentedu = pmin(im_hiqual_dv, ip_hiqual_dv))</pre>
  indrespb$parentedu<-ifelse(is.na(indrespb$parentedu), indrespb$im_hiqual_dv, indrespb$pare</pre>
  indrespb$parentedu<-ifelse(is.na(indrespb$parentedu), indrespb$ip_hiqual_dv, indrespb$pare
  indrespb$parent_edu<-c(0)
  indrespb$parent_edu[indrespb$parentedu%in%c(9)]<-1 #low (everything below, the category "c
  indrespb parent_edu[indrespb parentedu \% in \% c(1,2)] < -3 \ \# high \ (have \ a \ degree)
  indrespb$parent_edu[indrespb$parentedu==0]<-NA
  indrespb$parent_edu<-factor(indrespb$parent_edu,levels=c(1,2,3),labels=c('low',"middle","h
  #two-category education
```

```
indrespb$parent_edu2<-c(0)
  indrespb parent_edu 2 [indrespb parentedu \% in \% c (1,2)] < -1 \ \# high \ (have a degree)
  indrespb$parent_edu2[is.na(indrespb$parentedu)]<-NA
  indrespb<-indrespb%>%dplyr::select(-parentedu)
  #general health of parents**
  indrespb<-indrespb%>%mutate(health_father=coalesce(ip_sf1, ip_scsf1))
Warning: `..1` and `..2` have conflicting value labels.
i Labels for these values will be taken from `..1`.
x Values: 5
  indrespb<-indrespb%>%mutate(health_mother=coalesce(im_sf1, im_scsf1))
Warning: `..1` and `..2` have conflicting value labels.
i Labels for these values will be taken from `..1`.
x Values: 5
  #least healthy parent
  indrespb<-transform(indrespb, parenthealth = pmax(health_father, health_mother))</pre>
  indrespb$parenthealth<-ifelse(is.na(indrespb$parenthealth), indrespb$health_father, indrespb$parenthealth
  indrespb$parenthealth<-ifelse(is.na(indrespb$parenthealth), indrespb$health_mother, indres
  #parent's age (youngest)
  indrespb <- transform(indrespb, parentage = pmin(ip_age_dv</pre>
  , im_age_dv))
  indrespb$parentage<-ifelse(is.na(indrespb$parentage), indrespb$ip_age_dv, indrespb$parentage</pre>
  indrespb$parentage<-ifelse(is.na(indrespb$parentage), indrespb$im_age_dv, indrespb$parentage</pre>
  #parent's income (sum and logged)
  indrespb$palnincome<-indrespb$ip_fimnnet_dv+indrespb$im_fimnnet_dv
  \verb|indrespb$palnincome| [indrespb$palnincome<0]<-0|
  #add 1 to all income values to avoid -inf in log transformation
  indrespb$paincomeadd<-indrespb$palnincome+1
  indrespb$palnincome<-log(indrespb$paincomeadd)</pre>
  indrespb<-indrespb%>%dplyr::select(-paincomeadd)
  #parent's income (quartiles)
```

```
\verb|indrespb$paincome<-indrespb$| ip_fimnnet_dv+indrespb$| im_fimnnet_dv+indrespb$| im_fimnet_dv+indrespb$| im_fimntet_dv+indrespb$| im_fimntet_dv
       print(quantile(indrespb$paincome, prob=c(.25,.5,.75), type=1,na.rm=T))
                                     50%
                                                              75%
            25%
1928.18 2946.67 4192.75
       indrespb$paincomeq<-c(0)
       indrespb$paincomeq[indrespb$paincome>=0 & indrespb$paincome<1928.18]<-1</pre>
       indrespb$paincomeq[indrespb$paincome>=1928.18 & indrespb$paincome<2946.67]<-2
       indrespb$paincomeq[indrespb$paincome>=2946.67 & indrespb$paincome<4192.75 ]<-3
       indrespb$paincomeq[indrespb$paincome>=4192.75]<-4
       #parental tenure
       indrespb$mateure<-c(0)</pre>
       indrespb$mateure[indrespb$im_tenure_dv%in%c(1,2)]<-1 #ownership</pre>
       indrespb\mateure[indrespb\mateure_dv\%in\%c(3,4,5)] < -2 \mbox{ \#social rent}
       indrespb\mateure[indrespb\mateure_dv\%in\%c(6,7,8)] < -3 \mbox{ \#private rent n other}
       indrespb$mateure[indrespb$mateure==0]<-NA
       indrespb$mateure<-factor(indrespb$mateure,levels=c(1,2,3),labels=c("ownership","social ren
       indrespb$pateure<-c(0)
       indrespb$pateure[indrespb$ip_tenure_dv%in%c(1,2)]<-1 #ownership</pre>
       indrespb$pateure[indrespb$ip_tenure_dv%in%c(3,4,5)]<-2 #social rent
       indrespbpateure[indrespb$ip_tenure_dv%in%c(6,7,8)]<-3 #private rent n other
       indrespb$pateure[indrespb$pateure==0]<-NA
       indrespb$pateure<-factor(indrespb$pateure,levels=c(1,2,3),labels=c("ownership","social ren
       #the parents who have the most adv housing
       indrespb <- transform(indrespb, housingtenure = pmin(ip_tenure_dv, im_tenure_dv))</pre>
       indrespb$housingtenure<-ifelse(is.na(indrespb$housingtenure), indrespb$ip_tenure_dv, indre</pre>
       indrespb$housingtenure<-ifelse(is.na(indrespb$housingtenure), indrespb$im_tenure_dv, indre</pre>
       indrespb$housing_tenure<-c(0)
       \verb|indrespb$housing_tenure[indrespb$housingtenure%in%c(1,2)]<-1 #ownership|
       indrespb\$housing\_tenure[indrespb\$housingtenure\%in\%c(3,4,5)] < -2 \ \#social \ rent
       indrespb$housing_tenure[indrespb$housingtenure%in%c(6,7,8)]<-3 #private rent n other
       indrespb$housing_tenure[indrespb$housing_tenure==0]<-NA
       indrespb \$ housing\_tenure < -factor (indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c(1,2,3), labels = c("ownership indrespb \$ housing\_tenure, levels = c("o
```

```
indrespb<-indrespb%>%dplyr::select(-housingtenure)

#cross_wave n interaction variables
#age * parental lnincome
indrespb$paincomeXage<-indrespb$palnincome*indrespb$age_centered
#age * parental lnincome quartiles
indrespb$paincomeqXage<-indrespb$paincomeq*indrespb$age_centered

#cross wave identifiers
indrespb$wave<-c(0)
indrespb$wave[indrespb$from_wave_h==1]<-1
indrespb$wave[indrespb$from_wave_i==1]<-2
indrespb$wave[indrespb$from_wave_j==1]<-3
indrespb$wave[indrespb$from_wave_k==1]<-4
indrespb$wave[indrespb$from_wave_l==1]<-5

indrespb$wave<-factor(indrespb$wave,levels=c(1,2,3,4,5),labels=c("wave 8","wave 9","wave 1
#indrespb<-indrespb%>%dplyr::select(-from_wave_h,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_i,-from_wave_j,-from_wave_k,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-from_wave_i,-f
```

Dependent variables

```
indrespb$leave<-c(0)
indrespb$leave[indrespb$separated%in%c(1,2)]<-1

#wave h leavers
hleave<-indrespb[which(indrespb$leave==1&indrespb$from_wave_h==1),]
#join with i indresp for information
hleave<-join_all(list(hleave,i_indresp), by='pidp', type='left')

hleave$out<-c(0)
#leaving for marriage
hleave$out[hleave$i_mastat_dv%in%c(2,3)& hleave$i_jbstat!=7]<-1

#leaving for cohabitation
hleave$out[hleave$i_mastat_dv%in%c(10) & hleave$i_jbstat!=7]<-2

#leaving as a full-time student
hleave$out[hleave$i_jbstat%in%c(7)]<-3</pre>
```

```
#leaving for other reasons=0
#wave i leavers
ileave<-indrespb[which(indrespb$leave==1&indrespb$from_wave_i==1),]
#join with j indresp for information
ileave<-join_all(list(ileave, j_indresp), by='pidp', type='left')</pre>
ileave$out<-c(0)</pre>
#leaving for marriage
ileave$out[ileave$j_mastat_dv%in%c(2,3) & ileave$j_jbstat!=7]<-1</pre>
#leaving for coiabitation
ileave$out[ileave$j_mastat_dv%in%c(10) & ileave$j_jbstat!=7]<-2</pre>
#leaving as a full-time student
ileave\out[ileave j_jbstat\%in\%c(7)] < -3
#leaving for other reasons=0
#wave j leavers
jleave<-indrespb[which(indrespb$leave==1&indrespb$from_wave_j==1),]
#join with k indresp for information
jleave<-join_all(list(jleave, k_indresp), by='pidp', type='left')</pre>
jleave$out<-c(0)</pre>
#leaving for marriage
jleave$out[jleave$k_mastat_dv%in%c(2,3) & jleave$k_jbstat!=7]<-1</pre>
#leaving for coiabitation
jleave$out[jleave$k_mastat_dv%in%c(10) & jleave$k_jbstat!=7]<-2</pre>
#leaving as a full-time student
{\tt jleave\$out[jleave\$k\_jbstat\%in\%c(7)]<-3}
#leaving for other reasons=0
#wave k leavers
kleave<-indrespb[which(indrespb$leave==1&indrespb$from_wave_k==1),]</pre>
#join with 1 indresp for information
```

```
kleave<-join_all(list(kleave, l_indresp), by='pidp', type='left')</pre>
kleave$out<-c(0)</pre>
#leaving for marriage
kleave$out[kleave$1_mastat_dv%in%c(2,3) & kleave$1_jbstat!=7]<-1</pre>
#leaving for coiabitation
\label{leavesum} $$ kleave = 1_mastat_dv\%in\%c(10) \& kleave = 7] < -2 $$
#leaving as a full-time student
kleave$out[kleave$l_jbstat%in%c(7)]<-3</pre>
#leaving for other reasons=0
#wave 1 leavers
lleave<-indrespb[which(indrespb$leave==1&indrespb$from_wave_l==1),]</pre>
#join with m indresp for information
lleave<-join_all(list(lleave, m_indresp), by='pidp', type='left')</pre>
leave out < -c(0)
#leaving for marriage
lleave$out[lleave$m_mastat_dv%in%c(2,3) & lleave$m_jbstat!=7]<-1</pre>
#leaving for coiabitation
lleave\out[lleave\m_mastat_dv\%in\%c(10) \& lleave\m_jbstat!=7]<-2
#leaving as a full-time student
lleave$out[lleave$m_jbstat%in%c(7)]<-3</pre>
#leaving for other reasons=0
out<-rbind.fill(hleave,ileave,jleave,kleave,lleave)</pre>
out<-out%>%dplyr::select(pidp,wave,out)
indrespb<-join_all(list(indrespb,out), by=c("pidp", "wave"), type='left')</pre>
#mar
indrespb\mar<-c(0)
indrespb$mar[indrespb$out==1]<-1</pre>
#cohab
indrespb$cohab<-c(0)
indrespb$cohab[indrespb$out==2]<-1
```

```
#student
indrespb$student<-c(0)
indrespb$student[indrespb$out==3]<-1
#other
indrespb$other<-c(0)
indrespb$other[indrespb$out==0]<-1</pre>
```

Model building

Wave 1-5

Wave 8-12

```
indrespby<-indrespb[which(indrespb$age_centered%in%c(0:14)),]
indrespbfy<-indrespb[which(indrespb$age_centered%in%c(0:14)&indrespb$sex==1),]
indrespbmy<-indrespb[which(indrespb$age_centered%in%c(0:14)&indrespb$sex==0),]

mar_bf<-glm(mar~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)

cohab_bf<-glm(cohab~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)

student_bf<-glm(student~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)

Warning: glm.fit: algorithm did not converge

all_bf<-glm(leave-factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)

#paincomeXage effects more strongly for men mar_bm<-glm(mar~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)

Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred

cohab_bm<-glm(cohab~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)</pre>
```

```
student_bm<-glm(student~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing
Warning: glm.fit: algorithm did not converge
  all_bm<-glm(leave~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenur
  mar_b<-glm(mar~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure)+</pre>
  cohab_b<-glm(cohab~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenu
  student_b<-glm(student~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_
Warning: glm.fit: algorithm did not converge
  other_b<-glm(other~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenu
  all_b<-glm(leave~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure
  #how many boomerang
  counts <- indrespb%>%dplyr::count(pidp,separated)
  counts2<-data.frame(unclass(table(counts$pidp,counts$separated)))</pre>
  counts2<-counts2%>%mutate(leave=coalesce(X1, X2))
  counts2<-counts2%>%mutate(missing_dead=coalesce(X.9 , X.8))
  test<-counts2$X0+counts2$leave
  prop.table(table(test))
test
0.13213945 0.79771114 0.07014941
  0.2086468
[1] 0.2086468
```

Plotting results

```
convertModel <- function(model) {</pre>
    tr <- createTexreg(</pre>
      coef.names = names(model$coef),
      coef = as.numeric(summary(model)$coefficients[,1]),
      se = as.numeric(summary(model)$coefficients[,2]),
      pvalues = as.numeric(summary(model)$coefficients[,4]),
      gof.names = c("N","AIC"),
      gof = c(nobs(all_b),AIC(model)),
      gof.decimal = c(F,F)
    )
  }
  #compare males and females in wave 8-12
  p1<-huxtablereg(lapply(list(all_bf,all_bm,mar_bf,mar_bm,cohab_bf,cohab_bm,student_bf,student_bf)
  #full and nested models for all_b
  nest1<-glm(leave~age_centered,data=indrespby,family=binomial(link="logit"))</pre>
  nest2<-glm(leave~age_centered+factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(h
  p2<-huxtablereg(lapply(list(nest1,nest2,all_b), convertModel))</pre>
  nest1<-glm(leave~age_centered,data=indrespby,family=binomial(link="logit"))</pre>
  nest2<-glm(leave~age_centered+factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(h
  all_b<-glm(leave~factor(paincomeq)+paincomeqXage+factor(parent_edu2)+factor(housing_tenure
Summarize all leavers (all, men and women)
  print(summary(all_b))
Call:
glm(formula = leave ~ factor(paincomeq) + paincomeqXage + factor(parent_edu2) +
```

```
factor(housing_tenure) + parenthealth + parentage + factor(liveboth) +
age_centered + factor(sex) + lnincome + factor(edu2) + factor(family_status) +
factor(racel2) + factor(student) + factor(unemployed) + factor(urban) +
factor(houseprices) + factor(wave), family = binomial(link = "logit"),
data = indrespby)
```

Coefficients:

COEIICIEILS.				
	Estimate	Std. Error	z value	
(Intercept)	-2.356393	0.292362	-8.060	
factor(paincomeq)1	-0.402442	0.156161	-2.577	
<pre>factor(paincomeq)2</pre>	-0.401644	0.156386	-2.568	
<pre>factor(paincomeq)3</pre>	-0.410963	0.165258	-2.487	
factor(paincomeq)4	-0.274153	0.178886	-1.533	
paincomeqXage	0.022434	0.004958	4.525	
factor(parent_edu2)1	0.151985	0.064109	2.371	
factor(housing_tenure)social rent	-0.078708	0.086685	-0.908	
<pre>factor(housing_tenure)private rent n other</pre>	0.216275	0.117673	1.838	
parenthealth	-0.102420	0.030417	-3.367	
parentage	-0.012507	0.005530	-2.262	
factor(liveboth)1	-0.234701	0.119145	-1.970	
age_centered	0.111682	0.012476	8.951	
factor(sex)1	0.210578	0.059393	3.545	
lnincome	0.067006	0.014339	4.673	
factor(edu2)1	0.437977	0.067080	6.529	
<pre>factor(family_status)coresident partner and child</pre>	0.735003	0.290754	2.528	
<pre>factor(family_status)coresident partner</pre>	0.517030	0.179192	2.885	
<pre>factor(family_status)coresident child</pre>	0.535401	0.190600	2.809	
factor(racel2)non-Whites	-1.026362	0.088813	-11.556	
factor(student)1	18.417807	152.783828	0.121	
factor(unemployed)1	0.011589	0.109376	0.106	
factor(urban)1	-0.183862	0.069373	-2.650	
factor(houseprices)middle	0.236792	0.077460	3.057	
factor(houseprices)high	0.010229	0.077154	0.133	
factor(wave)wave 9	-0.175781	0.087754	-2.003	
factor(wave)wave 10	-0.174380	0.087982	-1.982	
factor(wave)wave 11	-0.236152	0.091840	-2.571	
factor(wave)wave 12	-0.039302	0.089298	-0.440	
	Pr(> z)			
(Intercept)	7.64e-16 *	* *		
factor(paincomeq)1	0.009963 *>	k		
factor(paincomeq)2	0.010220 *			
factor(paincomeq)3	0.012890 *			
factor(paincomeq)4	0.125384			

```
paincomeqXage
                                                   6.03e-06 ***
factor(parent_edu2)1
                                                   0.017753 *
factor(housing_tenure)social rent
                                                   0.363887
factor(housing_tenure)private rent n other
                                                   0.066072 .
                                                   0.000759 ***
parenthealth
                                                   0.023721 *
parentage
factor(liveboth)1
                                                   0.048852 *
age_centered
                                                    < 2e-16 ***
factor(sex)1
                                                   0.000392 ***
Inincome
                                                   2.97e-06 ***
factor(edu2)1
                                                   6.61e-11 ***
factor(family_status)coresident partner and child 0.011474 *
factor(family_status)coresident partner
                                                   0.003910 **
factor(family_status)coresident child
                                                   0.004969 **
factor(racel2)non-Whites
                                                    < 2e-16 ***
factor(student)1
                                                   0.904049
factor(unemployed)1
                                                   0.915618
factor(urban)1
                                                   0.008041 **
factor(houseprices)middle
                                                   0.002236 **
factor(houseprices)high
                                                   0.894526
factor(wave)wave 9
                                                   0.045166 *
factor(wave)wave 10
                                                   0.047479 *
factor(wave)wave 11
                                                   0.010131 *
factor(wave)wave 12
                                                   0.659849
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 10123 on 17966 degrees of freedom
Residual deviance: 8771 on 17938 degrees of freedom
  (3505 observations deleted due to missingness)
AIC: 8829
Number of Fisher Scoring iterations: 14
  print(summary(all_bf))
Call:
glm(formula = leave ~ factor(paincomeq) + paincomeqXage + factor(parent_edu2) +
```

```
factor(housing_tenure) + parenthealth + parentage + factor(liveboth) +
age_centered + lnincome + factor(edu2) + factor(family_status) +
factor(racel2) + factor(student) + factor(unemployed) + factor(urban) +
factor(houseprices) + factor(relative_num) + factor(wave),
family = binomial(link = "logit"), data = indrespbfy)
```

Coefficients:

COEIIICIEIICS.			
	Estimate	Std. Error	
(Intercept)	-1.914644	0.410905	-4.660
<pre>factor(paincomeq)1</pre>	-0.532449	0.204472	-2.604
<pre>factor(paincomeq)2</pre>	-0.400107	0.202321	-1.978
<pre>factor(paincomeq)3</pre>	-0.524362	0.215743	-2.430
factor(paincomeq)4	-0.357552	0.233980	-1.528
paincomeqXage	0.019623	0.006757	2.904
<pre>factor(parent_edu2)1</pre>	0.189051	0.087185	2.168
factor(housing_tenure)social rent	-0.135790	0.117232	-1.158
<pre>factor(housing_tenure)private rent n other</pre>	0.196210	0.158309	1.239
parenthealth	-0.086504	0.041103	-2.105
parentage	-0.020240	0.007692	-2.631
factor(liveboth)1	-0.010996	0.149699	-0.073
age_centered	0.139613	0.017085	8.172
lnincome	0.056882	0.019623	2.899
factor(edu2)1	0.437052	0.091059	4.800
<pre>factor(family_status)coresident partner and child</pre>	1.153586	0.401672	2.872
<pre>factor(family_status)coresident partner</pre>	0.588224	0.235833	2.494
<pre>factor(family_status)coresident child</pre>	0.594785	0.197580	3.010
factor(racel2)non-Whites	-0.939593	0.121385	-7.741
factor(student)1	18.368027	177.697805	0.103
factor(unemployed)1	0.194617	0.154065	1.263
factor(urban)1	-0.230512	0.093359	-2.469
factor(houseprices)middle	0.110194	0.103210	1.068
factor(houseprices)high	-0.098233	0.103754	-0.947
factor(relative_num)large hh	0.018360	0.089281	0.206
factor(wave)wave 9	-0.043029	0.116494	-0.369
factor(wave)wave 10	-0.308088	0.123243	-2.500
factor(wave)wave 11	-0.285484	0.126067	-2.265
factor(wave)wave 12	0.022144	0.119082	0.186
	Pr(> z)		
(Intercept)	3.17e-06 *>	* *	
<pre>factor(paincomeq)1</pre>	0.00921 *>	k	
factor(paincomeq)2	0.04798 *		
factor(paincomeq)3	0.01508 *		
factor(paincomeq)4	0.12648		

```
paincomeqXage
                                                    0.00368 **
factor(parent_edu2)1
                                                    0.03013 *
factor(housing_tenure)social rent
                                                    0.24674
factor(housing_tenure)private rent n other
                                                    0.21519
parenthealth
                                                    0.03533 *
                                                    0.00850 **
parentage
factor(liveboth)1
                                                    0.94144
age_centered
                                                   3.04e-16 ***
lnincome
                                                    0.00375 **
factor(edu2)1
                                                   1.59e-06 ***
factor(family_status)coresident partner and child 0.00408 **
factor(family_status)coresident partner
                                                    0.01262 *
factor(family_status)coresident child
                                                    0.00261 **
factor(racel2)non-Whites
                                                   9.89e-15 ***
factor(student)1
                                                    0.91767
factor(unemployed)1
                                                    0.20651
factor(urban)1
                                                    0.01355 *
factor(houseprices)middle
                                                    0.28567
factor(houseprices)high
                                                    0.34375
factor(relative_num)large hh
                                                    0.83707
factor(wave)wave 9
                                                    0.71186
factor(wave)wave 10
                                                    0.01243 *
factor(wave)wave 11
                                                    0.02354 *
factor(wave)wave 12
                                                    0.85248
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 5590.2 on 9201 degrees of freedom
Residual deviance: 4727.3 on 9173 degrees of freedom
  (1831 observations deleted due to missingness)
AIC: 4785.3
Number of Fisher Scoring iterations: 14
  print(summary(all_bm))
Call:
glm(formula = leave ~ factor(paincomeq) + paincomeqXage + factor(parent_edu2) +
```

```
factor(housing_tenure) + parenthealth + parentage + factor(liveboth) +
age_centered + lnincome + factor(edu2) + factor(family_status) +
factor(racel2) + factor(student) + factor(unemployed) + factor(urban) +
factor(houseprices) + factor(relative_num) + factor(wave),
family = binomial(link = "logit"), data = indrespbmy)
```

Coefficients:

COEIIICIEIICS.			
	Estimate	Std. Error	
(Intercept)	-2.486324	0.460937	-5.394
<pre>factor(paincomeq)1</pre>	-0.176593	0.250522	-0.705
<pre>factor(paincomeq)2</pre>	-0.314968	0.253015	-1.245
<pre>factor(paincomeq)3</pre>	-0.234535	0.265526	-0.883
factor(paincomeq)4	-0.119077	0.285112	-0.418
paincomeqXage	0.025759	0.007370	3.495
<pre>factor(parent_edu2)1</pre>	0.147720	0.095943	1.540
factor(housing_tenure)social rent	-0.018444	0.130174	-0.142
<pre>factor(housing_tenure)private rent n other</pre>	0.285033	0.177757	1.603
parenthealth	-0.123966	0.045540	-2.722
parentage	-0.006569	0.008545	-0.769
factor(liveboth)1	-0.590391	0.202333	-2.918
age_centered	0.080510	0.018533	4.344
lnincome	0.081096	0.021108	3.842
factor(edu2)1	0.418824	0.100647	4.161
<pre>factor(family_status)coresident partner and child</pre>	0.441990	0.453697	0.974
<pre>factor(family_status)coresident partner</pre>	0.445541	0.283465	1.572
<pre>factor(family_status)coresident child</pre>	-13.279130	364.914609	-0.036
factor(racel2)non-Whites	-1.090580	0.136672	-7.980
factor(student)1	18.645759	300.727126	0.062
factor(unemployed)1	-0.158447	0.156248	-1.014
factor(urban)1	-0.087908	0.105153	-0.836
factor(houseprices)middle	0.407995	0.119357	3.418
factor(houseprices)high	0.153327	0.117696	1.303
factor(relative_num)large hh	-0.183542	0.098733	-1.859
factor(wave)wave 9	-0.336046	0.135445	-2.481
factor(wave)wave 10	-0.007020	0.126459	-0.056
factor(wave)wave 11	-0.158245	0.134755	-1.174
factor(wave)wave 12	-0.096663	0.136490	-0.708
	Pr(> z)		
(Intercept)	6.89e-08 *	* *	
factor(paincomeq)1	0.480874		
factor(paincomeq)2	0.213184		
factor(paincomeq)3	0.377081		
factor(paincomeq)4	0.676203		

```
paincomeqXage
                                                   0.000474 ***
factor(parent_edu2)1
                                                   0.123642
factor(housing_tenure)social rent
                                                   0.887325
factor(housing_tenure)private rent n other
                                                   0.108824
                                                   0.006486 **
parenthealth
parentage
                                                   0.442071
factor(liveboth)1
                                                   0.003524 **
age_centered
                                                   1.40e-05 ***
lnincome
                                                   0.000122 ***
factor(edu2)1
                                                   3.16e-05 ***
factor(family_status)coresident partner and child 0.329959
factor(family_status)coresident partner
                                                   0.116004
factor(family_status)coresident child
                                                   0.970972
factor(racel2)non-Whites
                                                   1.47e-15 ***
factor(student)1
                                                   0.950561
factor(unemployed)1
                                                   0.310546
factor(urban)1
                                                   0.403152
factor(houseprices)middle
                                                   0.000630 ***
factor(houseprices)high
                                                   0.192663
factor(relative_num)large hh
                                                   0.063031 .
factor(wave)wave 9
                                                   0.013099 *
factor(wave)wave 10
                                                   0.955729
factor(wave)wave 11
                                                   0.240269
factor(wave)wave 12
                                                   0.478821
___
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 4510.4 on 8764 degrees of freedom Residual deviance: 3992.5 on 8736 degrees of freedom

(1674 observations deleted due to missingness)

AIC: 4050.5

Number of Fisher Scoring iterations: 14

Check assumptions

```
view(dfSummary(indrespb))
```

Switching method to 'browser'

 ${\tt Output\ file\ written:\ C:\ Vsers\ Think\ AppData\ Local\ Temp\ RtmpoRPwqW\ file\ 4dec\ 18e9659d.html}$