

DEM 7223 - Event History Analysis - Cox Proportional Hazards Model Part 2

Corey S. Sparks, PhD

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Review of Cox Regression Assumptions

- Although the Cox PH model offers an attractive alternative to parametric models, especially when ties are present, the assumptions of the model need to be assessed
- The primary assumption we are concerned with are:
- The time-constant covariate effect, i.e. the effect does not vary with time
- Grambsch and Therneau (1994) derived a method for checking the proportionality assumption for the Cox model using the residuals from the Cox model fit
- First we need to look at the various kinds of residuals from the Cox model and their properties

Cox Model Residuals

The basic principle for the construction of residuals is:

- Observed value – Predicted value
- This lets you get an idea of how well you are modeling your data with your covariates
- In hazard models, this principle is a bit more difficult because of censoring in the data, but despite this, we can define several types of model residuals and use them to diagnose problems with the model.

Schoenfeld Residuals

- These are used to test the proportionality assumption of the model
- If there are p covariates and n observations, with observed duration, censoring indicators and covariates, then the Schoenfeld residual is defined as the observed - expected value of a covariate at a particular *failure time*
- Plotting these residuals versus time should show any time dependency in the covariate, which violates the proportionality assumption of the model.

Martingale residuals

- These residuals can be thought of as the difference between an expected event occurring and an actual event occurring
- These use the censoring indicator for each observation and the estimate of the cumulative hazard function
- These residuals are given by:

$$M_i(t) = \delta_i(t) - H_i(t)$$

- Where $H_i(t)$ is the cumulative hazard function
- This residual is derived from counting process theory and represents the difference between the observed count of failures at any time, minus those that are predicted by the cumulative hazard function.
- Martingale residuals are also useful for assessing the functional form of a covariate
- Meaning, is the effect linear or quadratic
- A plot of the martingale residuals against the values of the covariate will indicate if the covariate is being modeled correctly
- If a line fit to these residuals is a straight line, then the covariate has been modeled effectively, if it is curvilinear, you may need to enter the covariate as a quadratic
- This is usually not a problem for binary covariates

Testing non-proportional effects

- As mentioned before, Grambsch and Therneau (1994) derived a method for checking the proportionality assumption for the Cox model using the residuals from the Cox model fit
- Now that we know what these residuals are, we can see the test
- Their test is equivalent to regressing the Schoenfeld residual on time for each covariate
- If there is a significant trend (correlation) between the residual and time, then non-proportionality is likely.

Model stratification

- One common method for dealing with non proportionality of hazards is via model stratification.
- If one of the covariates exhibits non-proportionality we can re-specify the model so that each group will have its own baseline hazard rate
- The effect of the other covariates in the model is assumed to behave the same in both groups!
- This creates the model:

$$h_{is}(t) = h_{0s} \exp(x' \beta)$$

- which allows for different baseline hazard rates for each of the s strata, which should control for their unequal hazards of experiencing the event.
- This procedure is slightly different than fitting separate models for each level of the stratification variable
- This method will allow the effects of the covariates to vary between strata
- Unfortunately, when we split the data and run separate models for each level, we lose the ability to discuss “between level” effects, since each analysis is run on a different sample

Data examples

This example will illustrate how to examine the fit of the Cox Proportional hazards model to a discrete-time (longitudinal) data set and examine various model diagnostics to evaluate the overall model fit. The data example uses data from the [ECLS-K](#). Specifically, we will examine the transition into poverty between kindergarten and third grade.

```

#Load required libraries
library(foreign)
library(survival)
library(car)
library(survey)
library(eha)
library(tidyverse)
options(survey.lonely.psu = "adjust")

eclskk5<-readRDS("C:/Users/ozd504/OneDrive - University of Texas at San Antonio/classes/de
names(eclskk5)<-tolower(names(eclskk5))
#get out only the variables I'm going to use for this example
myvars<-c( "childid", "x_chsex_r", "x_raceth_r", "x1kage_r", "x4age",
          "x5age", "x6age", "x7age", "x2povty", "x4povty_i", "x6povty_i",
          "x8povty_i", "x12parled_i", "s2_id", "w6c6p_6psu",
          "w6c6p_6str", "w6c6p_20")
eclskk5<-eclskk5[,myvars]

# time varying variables
eclskk5$age_1<-ifelse(eclskk5$x1kage_r== -9, NA, eclskk5$x1kage_r/12)
eclskk5$age_2<-ifelse(eclskk5$x4age== -9, NA, eclskk5$x4age/12)
#for the later waves, the NCES group the ages into ranges of months,
#so 1= <105 months, 2=105 to 108 months.
#So, I fix the age at the midpoint of the interval they give,
#and make it into years by dividing by 12

eclskk5$age_3<-ifelse(eclskk5$x5age== -9, NA, eclskk5$x5age/12)

eclskk5$pov_1<-ifelse(eclskk5$x2povty==1,1,0)
eclskk5$pov_2<-ifelse(eclskk5$x4povty_i==1,1,0)
eclskk5$pov_3<-ifelse(eclskk5$x6povty_i==1,1,0)

#Time constant variables
#Recode race with white, non Hispanic as reference using dummy vars
eclskk5$race_rec<-Recode (eclskk5$x_raceth_r, recodes="1 = 'nhwhite';2='nhblack';3:4='hisp
eclskk5$male<-Recode(eclskk5$x_chsex_r, recodes="1=1; 2=0; -9=NA")
eclskk5$mlths<-Recode(eclskk5$x12parled_i, recodes = "1:2=1; 3:9=0; else = NA")
eclskk5$mgths<-Recode(eclskk5$x12parled_i, recodes = "1:3=0; 4:9=1; else =NA")

```

Now, I need to form the transition variable, this is my event variable, and in this case it will

be 1 if a child enters poverty between the first wave of the data and the third grade wave, and 0 otherwise.

NOTE I need to remove any children who are already in poverty age wave 1, because they are not at risk of experiencing **this particular** transition. Again, this is called forming the *risk set*

```
eclskk5<-eclskk5 %>% filter(is.na(pov_1)==F &
                           is.na(pov_2)==F &
                           is.na(pov_3)==F &
                           is.na(age_1)==F &
                           is.na(age_2)==F &
                           is.na(age_3)==F &
                           pov_1!=1)
```

Now we do the entire data set. To analyze data longitudinally, we need to reshape the data from the current “wide” format (repeated measures in columns) to a “long” format (repeated observations in rows). The `reshape()` function allows us to do this easily. It allows us to specify our repeated measures, time varying covariates as well as time-constant covariates.

```
e.long1 <- eclskk5 %>%
  #rename(wt = w4c4p_40, strata= w4c4p_4str, psu = w4c4p_4psu)%>%
  select(childid, male, race_rec, mlths, mgths, #time constant
         age_1, age_2, age_3, #t-varying variables
         pov_1, pov_2, pov_3,
         w6c6p_6psu, w6c6p_6str, w6c6p_20)%>%
  pivot_longer(cols = c(-childid, -male, -race_rec, -mlths, -mgths, -w6c6p_6psu, -w6c6p_6str, -w6c6p_20),
               names_to = c(".value", "wave"), #make wave variable and put t-v vars into
               names_sep = "_") %>% #all t-v variables have _ between name and time, like
  group_by(childid)%>%
  mutate(age_enter = age,
         age_exit = lead(age, 1, order_by=childid))%>%
  mutate(nexpov = dplyr::lead(pov, n=1, order_by = childid))%>%
  mutate(povtran = ifelse(nexpov == 1 & pov == 0, 1, 0))%>%
  filter(is.na(age_exit)==F)%>%
  ungroup()%>%
  filter(complete.cases(age_enter, age_exit, povtran,
                        mlths, mgths, race_rec, w6c6p_6psu))
```

Construct survey design and fit basic Cox model

Now we fit the Cox model using full survey design. In the ECLS-K, I use the longitudinal weight for waves 1-7, as well as the associated psu and strata id's for the longitudinal data from these waves from the parents of the child, since no data from the child themselves are used in the outcome.

```
options(survey.lonely.psu = "adjust")

des2<-svydesign(ids = ~w6c6p_6psu,
               strata = ~w6c6p_6str,
               weights=~w6c6p_20,
               data=e.long1,
               nest=T)

#Fit the model
fitl1<-svycoxph(Surv(time = age_enter, time2=age_exit, event = povtran)~mlths+race_rec, de
summary(fitl1)
```

Stratified 1 - level Cluster Sampling design (with replacement)

With (123) clusters.

```
svydesign(ids = ~w6c6p_6psu, strata = ~w6c6p_6str, weights = ~w6c6p_20,
         data = e.long1, nest = T)
```

Call:

```
svycoxph(formula = Surv(time = age_enter, time2 = age_exit, event = povtran) ~
         mlths + race_rec, design = des2)
```

n= 4084, number of events= 221

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)	
mlths	1.0034	2.7275	0.1715	0.1669	6.014	1.81e-09	***
race_recnhasian	-0.8670	0.4202	0.3735	0.3624	-2.393	0.0167	*
race_recnhblack	-0.1382	0.8710	0.2200	0.2475	-0.558	0.5767	
race_recnhwhite	-1.3062	0.2709	0.1586	0.2075	-6.296	3.06e-10	***
race_recother	-0.8036	0.4477	0.2761	0.1981	-4.056	5.00e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

	exp(coef)	exp(-coef)	lower .95	upper .95
mlths	2.7275	0.3666	1.9667	3.7826
race_recnhasian	0.4202	2.3797	0.2065	0.8549

race_recnhblack	0.8710	1.1482	0.5362	1.4147
race_recnhwhite	0.2709	3.6921	0.1804	0.4068
race_recother	0.4477	2.2335	0.3036	0.6602

```

Concordance= 0.695 (se = 0.024 )
Likelihood ratio test= NA on 5 df, p=NA
Wald test              = 115.7 on 5 df, p=<2e-16
Score (logrank) test = NA on 5 df, p=NA

```

(Note: the likelihood ratio and score tests assume independence of observations within a cluster, the Wald and robust score tests do not).

Model Residuals

There are several types of residuals for the Cox model, and they are used for different purposes.

First, we will extract the *Shoenfeld* residuals, which are useful for examining non-proportional hazards with respect to time. This means that the covariate effect could exhibit time-dependency.

First we extract the residuals from the model, then we fit a linear model to the residual and the observed (uncensored) failure times

WE DO NOT WANT TO SEE A SIGNIFICANT MODEL HERE!!!!

that would indicate dependence between the residual and outcome, or *non-proportionality*, similar to doing a test for heteroskedasticity in OLS

```

schoenresid<-resid(fitl1, type="schoenfeld")

fit.sr<-lm(schoenresid~des2$variables$age_enter[des2$variables$povtran==1])

fit.sr%>%
  broom::tidy()%>%
  filter(term != "(Intercept)")%>%
  select(response, estimate, statistic, p.value)

```

```

# A tibble: 5 x 4
  response      estimate statistic p.value
  <chr>          <dbl>      <dbl>   <dbl>

```

1	mlths	-0.0114	-0.318	0.751
2	race_recnhasian	-0.0140	-0.705	0.481
3	race_recnhblack	-0.0317	-1.25	0.212
4	race_recnhwhite	0.00812	0.221	0.825
5	race_recother	0.0312	1.47	0.142

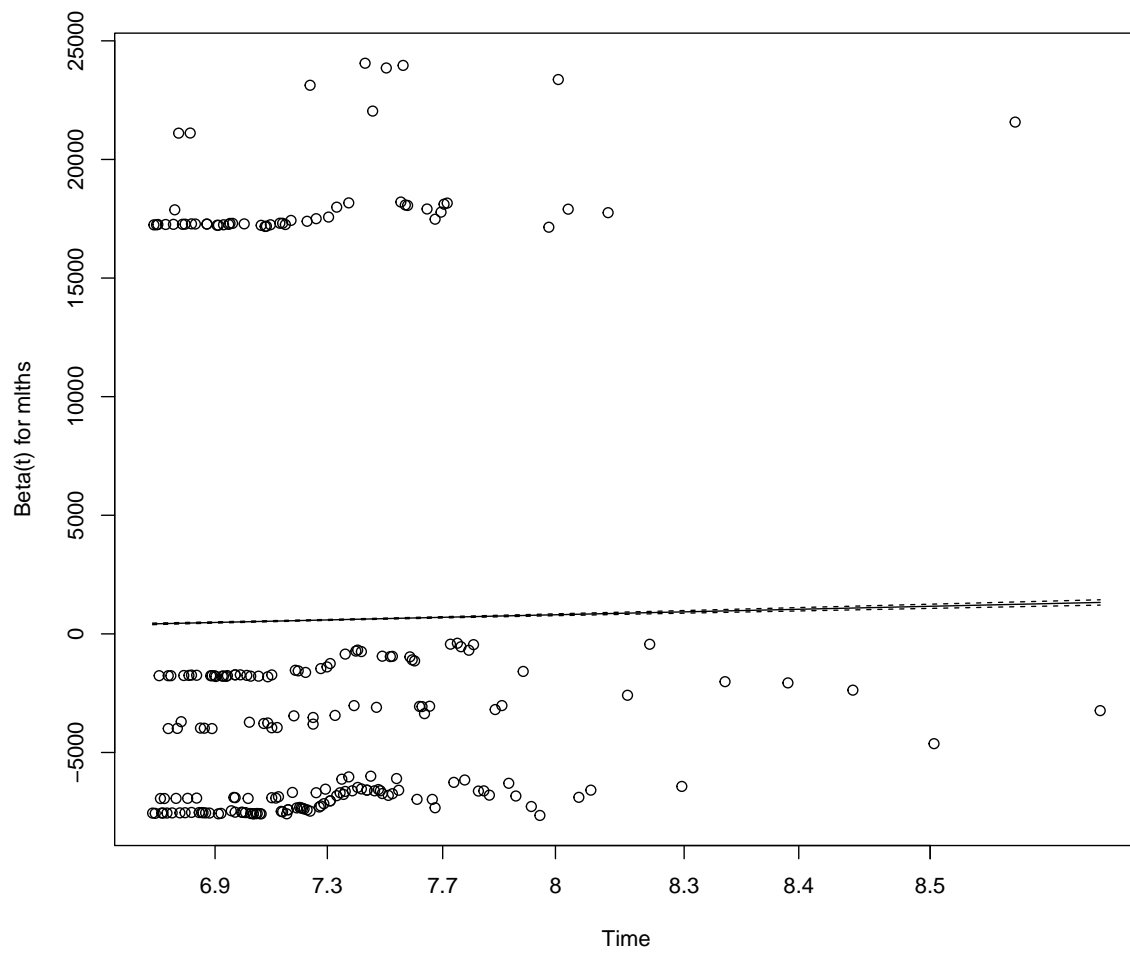
From these results, it appears that none of the variables are correlated with the timing of transition. This is what you want to see

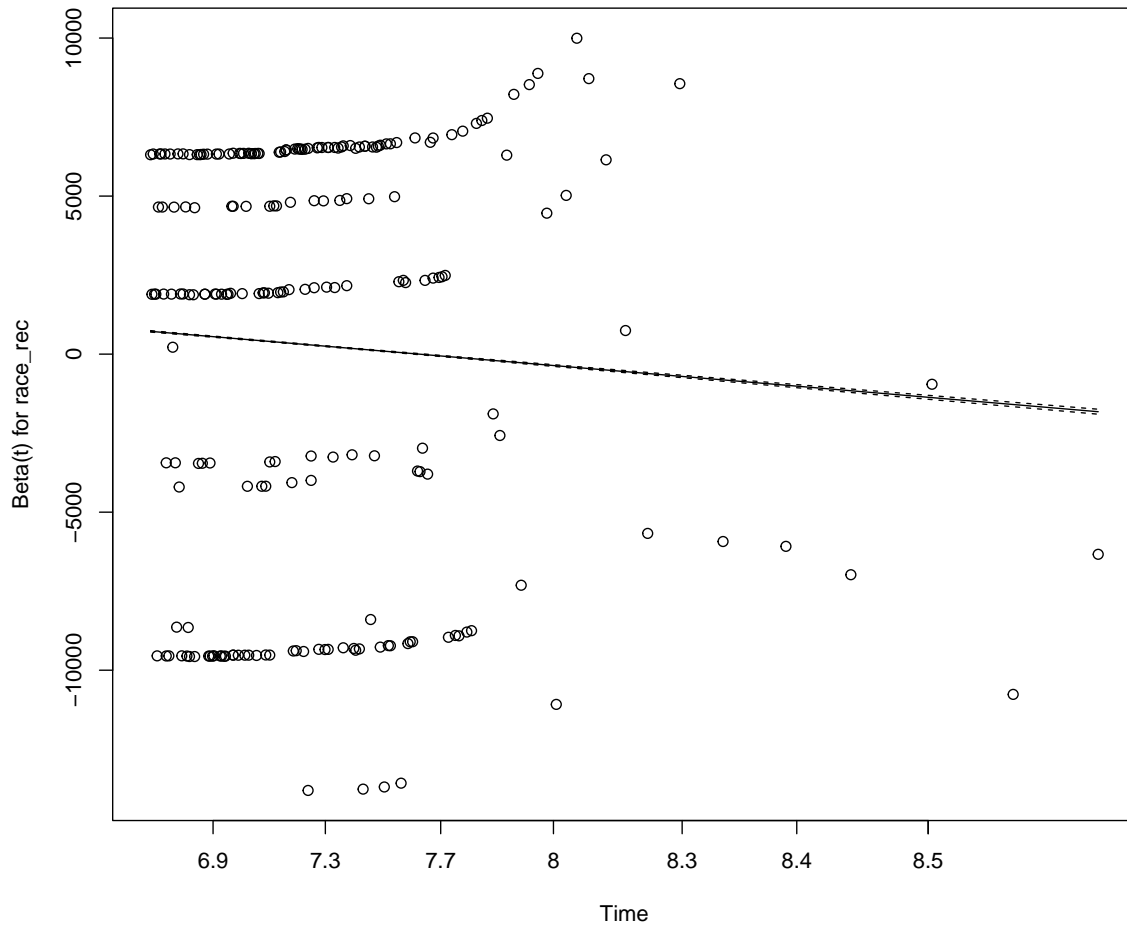
We can also get a formal test using weighted residuals in a nice pre-rolled form with a plot, ala [Grambsch and Therneau \(1994\)](#) :

```
fit.test<-cox.zph(fitl1)
fit.test
```

	chisq	df	p
mlths	0.000479	1	0.98
race_rec	0.000499	4	1.00
GLOBAL	0.000897	5	1.00

```
plot(fit.test, df=2)
```

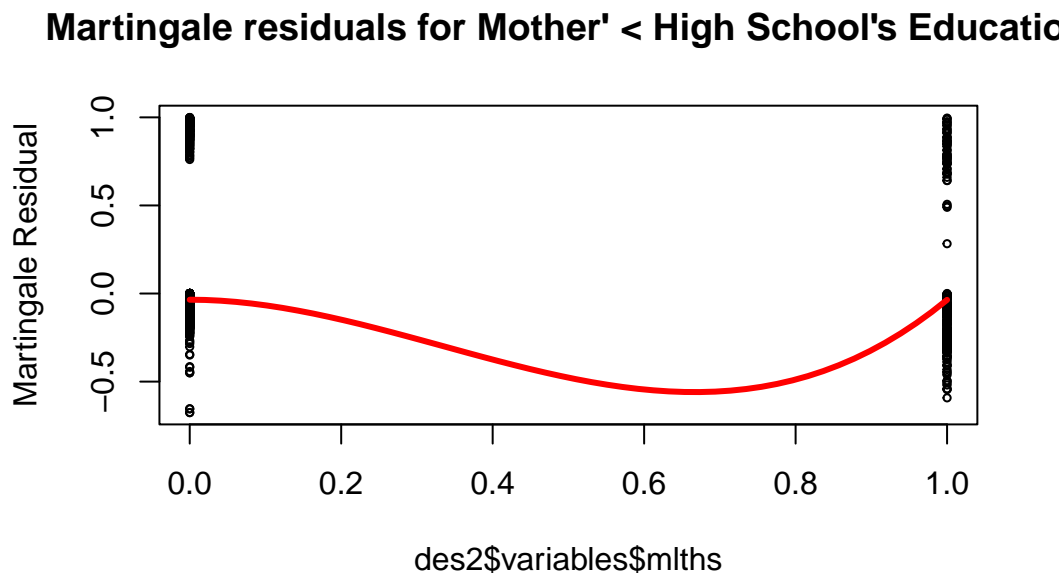


Here, we see the same result, with no significant relationship detected by the formal test. **This is what you want to see.**

Next we examine Martingale residuals. Martingale residuals are also useful for assessing the functional form of a covariate. A plot of the martingale residuals against the values of the covariate will indicate if the covariate is being modeled correctly, i.e. linearly in the Cox model. If a line fit to these residuals is a straight line, then the covariate has been modeled effectively, if it is curvilinear, you may need to enter the covariate as a quadratic, although this is not commonly a problem for dummy variables.

```
#extract Martingale residuals
res.mar<-resid(fitl1, type="martingale")

#plot vs maternal education
scatter.smooth(des2$variables$mlths, res.mar, degree = 2,
               span = 1, ylab="Martingale Residual",
               col=1, cex=.5, lpars=list(col = "red", lwd = 3))
title(main="Martingale residuals for Mother' < High School's Education")
```



Which shows nothing in the way of non-linearity in this case.

Stratification

Above, we observed evidence of non-proportional effects by education. There are a few standard ways of dealing with this in practice. The first is *stratification* of the model by the offending predictor. If one of the covariates exhibits non-proportionality we can re-specify the model so that each group will have its own baseline hazard rate. This is direct enough to do by using the `strata()` function within a model. This is of best use when a covariate is categorical, and not of direct importance for our model (i.e. a control variable).

```
fitl2<-svycoxph(Surv(time = age_enter, time2 = age_exit, event = povtran)~race_rec+strata(
                    design=des2)
summary(fitl2)
```

Stratified 1 - level Cluster Sampling design (with replacement)

With (123) clusters.

```
svydesign(ids = ~w6c6p_6psu, strata = ~w6c6p_6str, weights = ~w6c6p_20,
        data = e.long1, nest = T)
```

Call:

```
svycoxph(formula = Surv(time = age_enter, time2 = age_exit, event = povtran) ~
        race_rec + strata(mlths), design = des2)
```

n= 4084, number of events= 221

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)	
race_recnhasian	-0.8614	0.4226	0.3737	0.3643	-2.364	0.018059	*
race_recnhblack	-0.1583	0.8536	0.2207	0.2549	-0.621	0.534452	
race_recnhwhite	-1.3318	0.2640	0.1624	0.2204	-6.044	1.51e-09	***
race_recother	-0.7997	0.4495	0.2762	0.2057	-3.888	0.000101	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

	exp(coef)	exp(-coef)	lower .95	upper .95
race_recnhasian	0.4226	2.366	0.2069	0.8630
race_recnhblack	0.8536	1.172	0.5179	1.4066
race_recnhwhite	0.2640	3.788	0.1714	0.4066
race_recother	0.4495	2.225	0.3003	0.6726

Concordance= 0.659 (se = 0.026)

Likelihood ratio test= NA on 4 df, p=NA

Wald test = 50.52 on 4 df, p=3e-10

Score (logrank) test = NA on 4 df, p=NA

(Note: the likelihood ratio and score tests assume independence of observations within a cluster, the Wald and robust score tests do not).

Non-proportional effects with time

We can also include a time by covariate interaction term to model directly any time-dependence in the covariate effect. Different people say to do different things, some advocate for simply

interacting time with the covariate, others say use a nonlinear function of time, e.g. $\log(\text{time})$ * the covariate, others say use $\text{time}-1$ * covariate, which is called the “heavy side function”, according to Mills.

In this example, time is so limited that it doesn’t make sense to do this.

ANOVA like tests for factors

You can use the `regTermTest()` function in the `survey()` package to do omnibus tests for variation across a factor variable.

```
fit3<-svycoxph(Surv(time = age_enter, time2=age_exit, event = povtran)~mlths+race_rec,
               design=des2)
summary(fit3)
```

Stratified 1 - level Cluster Sampling design (with replacement)

With (123) clusters.

```
svydesign(ids = ~w6c6p_6psu, strata = ~w6c6p_6str, weights = ~w6c6p_20,
         data = e.long1, nest = T)
```

Call:

```
svycoxph(formula = Surv(time = age_enter, time2 = age_exit, event = povtran) ~
         mlths + race_rec, design = des2)
```

n= 4084, number of events= 221

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)	
mlths	1.0034	2.7275	0.1715	0.1669	6.014	1.81e-09	***
race_recnhasian	-0.8670	0.4202	0.3735	0.3624	-2.393	0.0167	*
race_recnhblack	-0.1382	0.8710	0.2200	0.2475	-0.558	0.5767	
race_recnhwhite	-1.3062	0.2709	0.1586	0.2075	-6.296	3.06e-10	***
race_recother	-0.8036	0.4477	0.2761	0.1981	-4.056	5.00e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

	exp(coef)	exp(-coef)	lower .95	upper .95
mlths	2.7275	0.3666	1.9667	3.7826
race_recnhasian	0.4202	2.3797	0.2065	0.8549
race_recnhblack	0.8710	1.1482	0.5362	1.4147
race_recnhwhite	0.2709	3.6921	0.1804	0.4068
race_recother	0.4477	2.2335	0.3036	0.6602

```

Concordance= 0.695 (se = 0.024 )
Likelihood ratio test= NA on 5 df, p=NA
Wald test              = 115.7 on 5 df, p=<2e-16
Score (logrank) test = NA on 5 df, p=NA

```

(Note: the likelihood ratio and score tests assume independence of observations within a cluster, the Wald and robust score tests do not).

```
regTermTest(fit3, ~mlths, method="LRT")
```

```

Working (Rao-Scott+F) LRT for mlths
in svycoxph(formula = Surv(time = age_enter, time2 = age_exit, event = povtran) ~
  mlths + race_rec, design = des2)
Working 2logLR = 22.79896 p= 1.0015e-05
df=1; denominator df= 72

```

```
#regTermTest(fit3, ~race_rec, method="LRT")
```

DHS data example

```

library(haven)
#load the data
dat<-read_dta("../data/ZAIR71FL.DTA")
dat<-zap_labels(dat)

```

In the DHS individual recode file, information on every live birth is collected using a retrospective birth history survey mechanism.

Since our outcome is time between first and second birth, we must select as our risk set, only women who have had a first birth.

The `bidx` variable indexes the birth history and if `bidx_01` is not missing, then the woman should be at risk of having a second birth (i.e. she has had a first birth, i.e. `bidx_01==1`).

I also select only non-twin births (`b0 == 0`).

The DHS provides the dates of when each child was born in Century Month Codes.

To get the interval for women who *actually had* a second birth, that is the difference between the CMC for the first birth `b3_01` and the second birth `b3_02`, but for women who had not had

a second birth by the time of the interview, the censored time between births is the difference between b3_01 and v008, the date of the interview.

We have 6124 women who are at risk of a second birth.

```
sub<-dat %>%
  filter(bidx_01==1&b0_01==0)%>%
  transmute(CASEID=caseid,
            int.cmc=v008,
            fbir.cmc=b3_01,
            sbir.cmc=b3_02,
            marr.cmc=v509,
            rural=v025,
            educ=v106,
            age = v012,
            agec=cut(v012, breaks = seq(15,50,5), include.lowest=T),
            partneredu=v701,
            partnerage=v730,
            weight=v005/1000000,
            psu=v021,
            strata=v022)%>%
  select(CASEID, int.cmc, fbir.cmc, sbir.cmc, marr.cmc, rural, educ, age, agec, partneredu)
mutate(agefb = (age - (int.cmc - fbir.cmc)/12))%>%
mutate(secbi = ifelse(is.na(sbir.cmc)==T,
                     int.cmc - fbir.cmc,
                     fbir.cmc - sbir.cmc),
       b2event = ifelse(is.na(sbir.cmc)==T,0,1))
```

Create covariates

Here, we create some predictor variables: Woman's education (secondary +, vs < secondary), Woman's age², Partner's education (> secondary school)

```
sub$educ.high<-ifelse(sub$educ %in% c(2,3), 1, 0)
sub$age2<-(sub$agefb)^2
sub$partnerhiedu<-ifelse(sub$partneredu<3,0,
                         ifelse(sub$partneredu%in%c(8,9),NA,1 ))

options(survey.lonely.psu = "adjust")

sub<- sub%>%
```

```

filter(complete.cases(secbi, partnerhiedu, educ.high, agefb))

des<-svydesign(ids=~psu, strata=~strata,
              data=sub[sub$secbi>0,],
              weight=~weight )

```

Fit the model

```

#use survey design

cox.s<-svycoxph(Surv(secbi,b2event)~educ.high+partnerhiedu+agefb+age2,
                design=des)

summary(cox.s)

```

Stratified 1 - level Cluster Sampling design (with replacement)
With (670) clusters.

```
svydesign(ids = ~psu, strata = ~strata, data = sub[sub$secbi >
0, ], weight = ~weight)
```

Call:

```
svycoxph(formula = Surv(secbi, b2event) ~ educ.high + partnerhiedu +
agefb + age2, design = des)
```

n= 2492, number of events= 1980

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)
educ.high	-0.2493914	0.7792749	0.0670396	0.0878240	-2.840	0.00452 **
partnerhiedu	0.0536853	1.0551525	0.0663283	0.0886441	0.606	0.54476
agefb	0.4995361	1.6479565	0.0348747	0.0423061	11.808	< 2e-16 ***
age2	-0.0079153	0.9921159	0.0005905	0.0007076	-11.187	< 2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

	exp(coef)	exp(-coef)	lower .95	upper .95
educ.high	0.7793	1.2832	0.6560	0.9256
partnerhiedu	1.0552	0.9477	0.8869	1.2554
agefb	1.6480	0.6068	1.5168	1.7904
age2	0.9921	1.0079	0.9907	0.9935

Concordance= 0.55 (se = 0.01)


```

Likelihood ratio test= NA on 4 df,  p=NA
Wald test              = 165.1 on 4 df,  p=<2e-16
Score (logrank) test = NA on 4 df,  p=NA

```

(Note: the likelihood ratio and score tests assume independence of observations within a cluster, the Wald and robust score tests do not).

```

#Schoenfeld test
fit.test<-cox.zph(cox.s)
fit.test

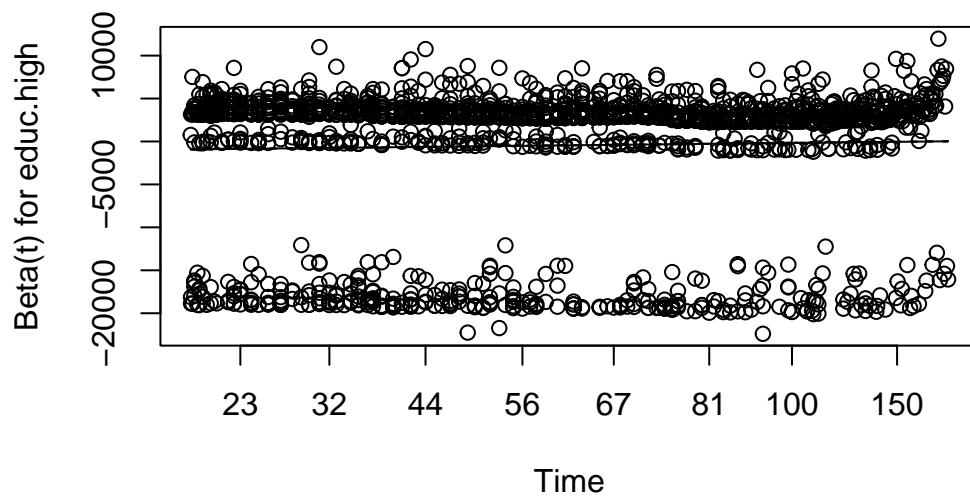
```

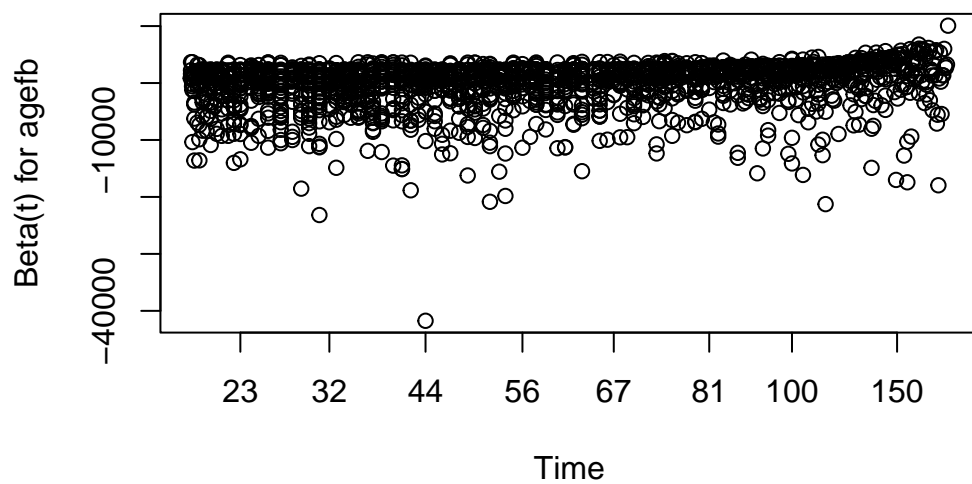
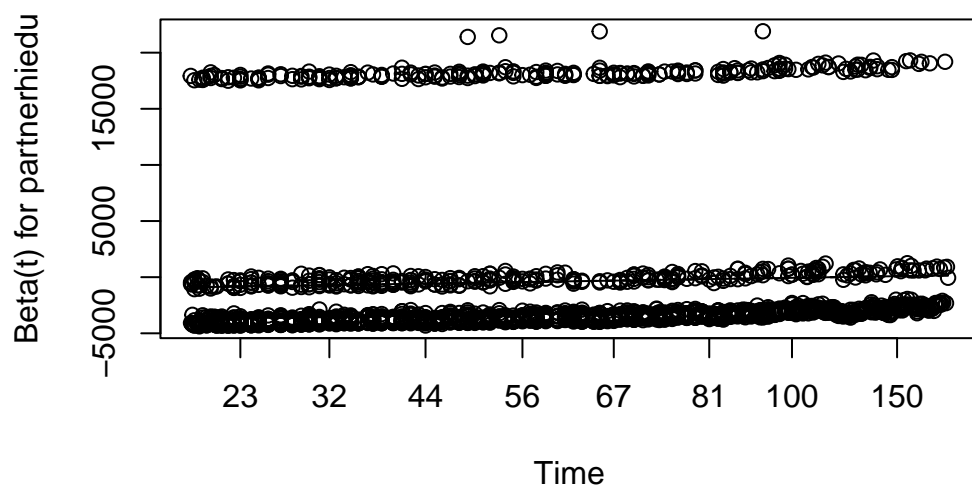
	chisq	df	p
educ.high	0.001026	1	0.97
partnerhiedu	0.000798	1	0.98
agefb	0.060697	1	0.81
age2	0.052766	1	0.82
GLOBAL	0.077515	4	1.00

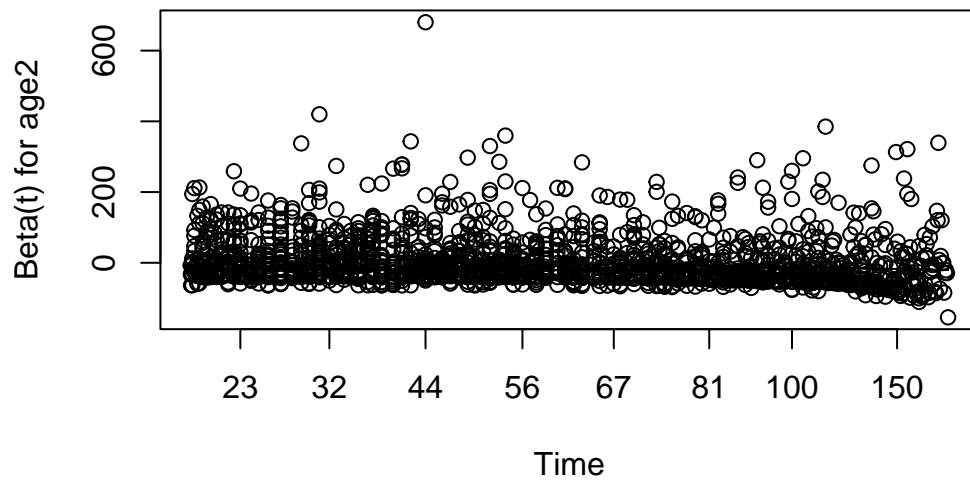
```

plot(fit.test, df=2)

```



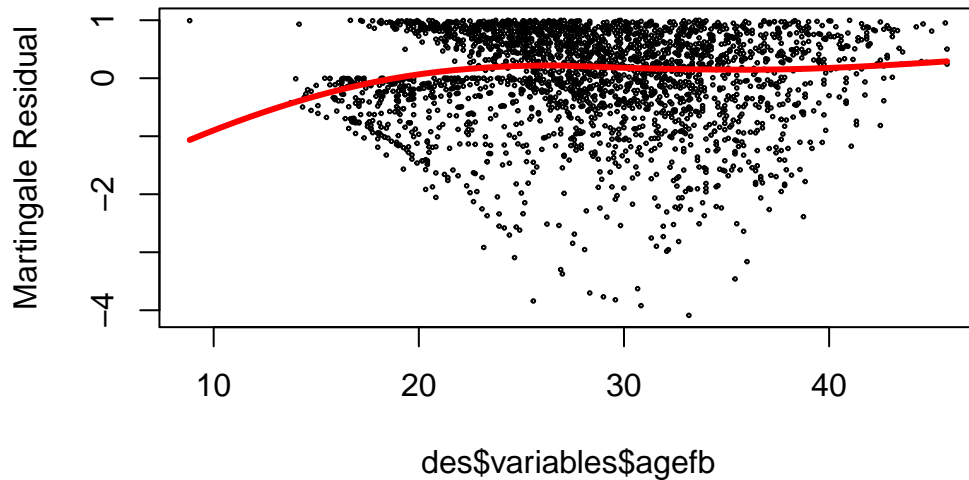




```
#martingale residuals
#extract Martingale residuals
res.mar<-resid(cox.s, type="martingale")

#plot vs maternal age
scatter.smooth(des$variables$agefb, res.mar,
              degree = 2,
              span = 1, ylab="Martingale Residual",
              col=1, cex=.25, lpars=list(col = "red",
                                         lwd = 3))
title(main="Martingale residuals for Mother Age' ")
```

Martingale residuals for Mother Age'



Non-proportional effects with time

We can also include a time by covariate interaction term to model directly any time-dependence in the covariate effect. Different people say to do different things, some advocate for simply interacting time with the covariate, others say use a nonlinear function of time, e.g. $\log(\text{time}) * \text{the covariate}$, others say use $\text{time}-1 * \text{covariate}$, which is called the “heavy side function”, according to Mills. Mills cites Allison, in saying that, to interpret the heavy side function, you go with the rule : “If β_2 is positive, then the effect of the covariate x increases over time, while if β_2 is negative, the effect of x decreases over time.”

```
sub.split<-survSplit(Surv(secbi, b2event)~.,
                    data= sub[sub$secbi>0,], cut=36, episode = "timegroup")
sub.split<-sub.split[order(sub.split$CASEID, sub.split$timegroup),]

sub.split$hv1<-sub.split$agefb*(1-sub.split$timegroup)
sub.split$hv2<-sub.split$agefb*(sub.split$timegroup)

head(sub.split, n=20)
```

	CASEID	int.cmc	fbir.cmc	sbir.cmc	marr.cmc	rural	educ	age	agec
1	1 13 2	1400	1204	NA	1333	1	1	41	(40,45]

2	1	13	2	1400	1204	NA	1333	1	1	41	(40,45]
3	1	36	2	1402	1351	NA	1372	1	2	27	(25,30]
4	1	36	2	1402	1351	NA	1372	1	2	27	(25,30]
5	2	29	2	1401	1088	NA	1240	2	2	45	(40,45]
6	2	29	2	1401	1088	NA	1240	2	2	45	(40,45]
7	2	29	3	1401	1286	1221	1383	2	1	36	(35,40]
8	2	29	3	1401	1286	1221	1383	2	1	36	(35,40]
9	2	34	4	1401	1371	1339	1391	2	2	26	(25,30]
10	2	35	2	1401	1146	1076	1145	2	1	47	(45,50]
11	2	35	2	1401	1146	1076	1145	2	1	47	(45,50]
12	2	59	2	1401	1168	1128	1132	2	1	47	(45,50]
13	2	59	2	1401	1168	1128	1132	2	1	47	(45,50]
14	2	65	2	1401	1275	1175	1164	2	1	39	(35,40]
15	2	65	2	1401	1275	1175	1164	2	1	39	(35,40]
16	2	79	1	1401	1301	1115	1252	2	3	48	(45,50]
17	2	79	1	1401	1301	1115	1252	2	3	48	(45,50]
18	2	96	11	1401	1401	1381	1345	2	2	23	(20,25]
19	3	2	2	1402	1386	1302	1132	1	2	39	(35,40]
20	3	2	2	1402	1386	1302	1132	1	2	39	(35,40]

	partneredu	partnerage	weight	psu	strata	agefb	educ.high	age2
1	1	44	1.396726	1	18	24.66667	0	608.4444
2	1	44	1.396726	1	18	24.66667	0	608.4444
3	2	31	1.396726	1	18	22.75000	1	517.5625
4	2	31	1.396726	1	18	22.75000	1	517.5625
5	0	48	1.206331	2	13	18.91667	1	357.8403
6	0	48	1.206331	2	13	18.91667	1	357.8403
7	2	52	1.206331	2	13	26.41667	0	697.8403
8	2	52	1.206331	2	13	26.41667	0	697.8403
9	2	33	1.206331	2	13	23.50000	1	552.2500
10	1	55	1.206331	2	13	25.75000	0	663.0625
11	1	55	1.206331	2	13	25.75000	0	663.0625
12	0	57	1.206331	2	13	27.58333	0	760.8403
13	0	57	1.206331	2	13	27.58333	0	760.8403
14	1	47	1.206331	2	13	28.50000	0	812.2500
15	1	47	1.206331	2	13	28.50000	0	812.2500
16	3	52	1.206331	2	13	39.66667	1	1573.4444
17	3	52	1.206331	2	13	39.66667	1	1573.4444
18	2	34	1.206331	2	13	23.00000	1	529.0000
19	1	42	0.560815	3	9	37.66667	1	1418.7778
20	1	42	0.560815	3	9	37.66667	1	1418.7778

	partnerhiedu	tstart	secbi	b2event	timegroup	hv1	hv2
1	0	0	36	0	1	0.00000	24.66667
2	0	36	196	0	2	-24.66667	49.33333

3	0	0	36	0	1	0.00000	22.75000
4	0	36	51	0	2	-22.75000	45.50000
5	0	0	36	0	1	0.00000	18.91667
6	0	36	313	0	2	-18.91667	37.83333
7	0	0	36	0	1	0.00000	26.41667
8	0	36	65	1	2	-26.41667	52.83333
9	0	0	32	1	1	0.00000	23.50000
10	0	0	36	0	1	0.00000	25.75000
11	0	36	70	1	2	-25.75000	51.50000
12	0	0	36	0	1	0.00000	27.58333
13	0	36	40	1	2	-27.58333	55.16667
14	0	0	36	0	1	0.00000	28.50000
15	0	36	100	1	2	-28.50000	57.00000
16	1	0	36	0	1	0.00000	39.66667
17	1	36	186	1	2	-39.66667	79.33333
18	0	0	20	1	1	0.00000	23.00000
19	0	0	36	0	1	0.00000	37.66667
20	0	36	84	1	2	-37.66667	75.33333

```
des3<-svydesign(ids=~psu, strata = ~strata ,
               weights=~weight, data=sub.split[is.na(sub.split$partnerhiedu)==F,])

cox.s2<-svycoxph(Surv(secbi,b2event)~educ.high+partnerhiedu+hv1+hv2,
                 design=des3)
summary(cox.s2)
```

Stratified 1 - level Cluster Sampling design (with replacement)

With (670) clusters.

```
svydesign(ids = ~psu, strata = ~strata, weights = ~weight, data = sub.split[is.na(sub.split$
F, ])
```

Call:

```
svycoxph(formula = Surv(secbi, b2event) ~ educ.high + partnerhiedu +
hv1 + hv2, design = des3)
```

n= 4320, number of events= 1980

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)
educ.high	-0.090737	0.913258	0.066556	0.091208	-0.995	0.320
partnerhiedu	0.066127	1.068362	0.066210	0.085337	0.775	0.438
hv1	0.150921	1.162905	0.007882	0.005713	26.415	<2e-16 ***

```

hv2          0.086805  1.090684  0.004594  0.004530 19.162  <2e-16 ***
---

```

```

Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

          exp(coef) exp(-coef) lower .95 upper .95
educ.high      0.9133      1.0950      0.7638      1.092
partnerhiedu    1.0684      0.9360      0.9038      1.263
hv1             1.1629      0.8599      1.1500      1.176
hv2             1.0907      0.9169      1.0810      1.100

```

```

Concordance= 0.651 (se = 0.008 )

```

```

Likelihood ratio test= NA on 4 df,   p=NA

```

```

Wald test          = 864 on 4 df,   p=<2e-16

```

```

Score (logrank) test = NA on 4 df,   p=NA

```

(Note: the likelihood ratio and score tests assume independence of observations within a cluster, the Wald and robust score tests do not).

So, for us β_2 in the heavyside function is positive, suggesting that the age effect increase over time

Aalen's additive regression model

An alternative model proposed by Odd Aalen in 1989 and 1993 describe a model that is inherently nonparametric and models the changes in relationships in a hazard model.

```

fita<-aareg(Surv(secbi,b2event)~educ.high+partnerhiedu+agefb+age2+cluster(strata),
            sub, weights = weight)

summary(fita)

```

```

          slope      coef se(coef) robust se      z      p
Intercept -0.055100 -2.85e-03 3.00e-04  2.47e-04 -11.60 7.09e-31
educ.high  -0.007290 -2.12e-04 6.65e-05  7.33e-05  -2.89 3.81e-03
partnerhiedu 0.000603 3.90e-05 6.44e-05  4.48e-05   0.87 3.84e-01
agefb        0.005790 2.51e-04 2.27e-05  2.27e-05  11.10 2.19e-28
age2        -0.000098 -3.95e-06 4.06e-07  4.07e-07  -9.71 2.80e-22

```

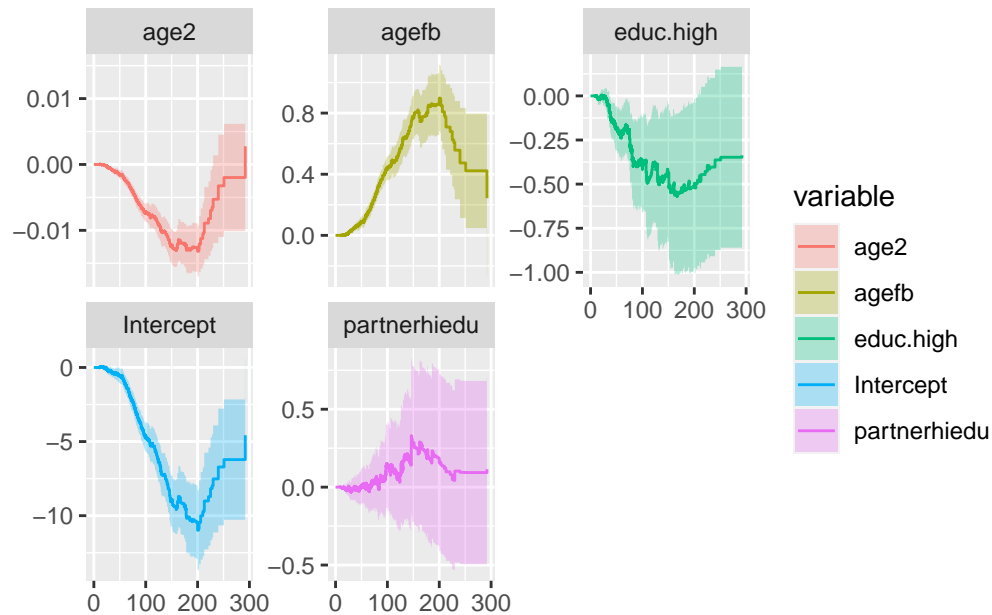
```

Chisq=369.83 on 4 df, p=<2e-16; test weights=aalen

```

```
library(ggfortify)
autoplot(fita)
```

Warning: `gather_()` was deprecated in tidyr 1.2.0.
Please use `gather()` instead.
This warning is displayed once every 8 hours.
Call `lifecycle::last_lifecycle_warnings()` to see where this warning was generated.



What is seen in the plots are the time-varying coefficients of the hazard model. For example the effect of `educ.high` is globally negative, suggesting higher education decreases the hazard, as we saw in the Cox model above. In the plot, the regression function initially decreases sharply but then plateaus, suggesting the education effect is really only time varying until about 100 months after the first birth.