

Biostat III Examination 2016 Answers

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Set-up

```
. global folder 5  
. set linesize 80
```

Commentary

In the following answers, the code and full Stata output are provided together with the answers. The full Stata output was not required in the given answers, but is given here to show how the answers were found.

Some brief comments are warranted on presentation. First, when the question asks for specific results, then those results should be presented separately in text, rather than only presenting the output from the statistical package. Second, the choice of non-proportional fonts makes it difficult to read output from the statistical package. Third, using colours in the graphics makes it difficult to discern which line is which in black-and-white printout. I suggest that using `scheme(s2mono)` would be useful for graphics in Stata.

Part 1

Question 1

We read in the dataset:

```
. import delimited "http://biostat3.net/download/exams/2016/$folder/incidence.c  
> sv", clear  
(6 vars, 360 obs)  
. egen agecat = cut(age), at(40, 50, 60, 70, 80, 90)
```

We then fit a Poisson regression with the number of lung cancer cases at the outcome (first argument), with the person-time of exposure as the `exposure` option. We include attained `age` as a linear, continuous effect in each model.

```
. poisson lc sex age, exposure(pt) nolog irr
```

```
Poisson regression                               Number of obs   =           360  
                                                LR chi2(2)      =           547.51  
                                                Prob > chi2     =           0.0000  
Log likelihood = -888.07465                    Pseudo R2      =           0.2356
```

```
-----  
      lc |           IRR   Std. Err.      z    P>|z|     [95% Conf. Interval]  
-----+-----  
      sex |    2.101776   .1927428     8.10  0.000     1.756011   2.515624  
      age |    1.095354   .0045096    22.12  0.000     1.086551   1.104229  
      _cons |    1.85e-06   5.02e-07   -48.72  0.000     1.09e-06   3.15e-06  
      ln(pt) |           1   (exposure)
```

```
-----
. poisson lc smoking age, exposure(pt) nolog irr
```

```
Poisson regression                               Number of obs   =       360
                                                LR chi2(2)      =      1303.83
                                                Prob > chi2     =       0.0000
Log likelihood = -509.91407                    Pseudo R2       =       0.5611
```

```
-----
          lc |          IRR   Std. Err.      z    P>|z|     [95% Conf. Interval]
-----+-----
    smoking |   15.63405   1.790159    24.01  0.000    12.49124   19.56759
         age |    1.100899   .0045774    23.12  0.000    1.091964   1.109907
        _cons |   4.70e-07   1.34e-07   -51.21  0.000    2.69e-07   8.21e-07
    ln(pt) |           1 (exposure)
```

```
-----
. poisson lc asbestos age, exposure(pt) nolog irr
```

```
Poisson regression                               Number of obs   =       360
                                                LR chi2(2)      =       586.93
                                                Prob > chi2     =       0.0000
Log likelihood = -868.36147                    Pseudo R2       =       0.2526
```

```
-----
          lc |          IRR   Std. Err.      z    P>|z|     [95% Conf. Interval]
-----+-----
    asbestos |    3.556892   .3804806    11.86  0.000    2.884149   4.386556
         age |    1.093873   .0044806    21.90  0.000    1.085126   1.10269
        _cons |   2.58e-06   6.76e-07   -49.18  0.000    1.55e-06   4.32e-06
    ln(pt) |           1 (exposure)
```

The age-adjusted incidence rate ratio for sex is 2.16 (95% confidence interval (CI): 1.80, 2.60). This association is highly significant ($p < 0.001$).

The age-adjusted incidence rate ratio for smoking is 18.45 (95% confidence interval (CI): 14.56, 23.37). This association is highly significant ($p < 0.001$).

The age-adjusted incidence rate ratio for asbestos is 3.68 (95% confidence interval (CI): 2.99, 4.53). This association is highly significant ($p < 0.001$).

We could have adjusted for attained age in several other ways, including quintiles or splines. To investigate this, we first use quintiles with sex:

```
. xtile ageQ5 = age, nquantiles(5)
. poisson lc sex i.ageQ5, exposure(pt) nolog irr base
```

```
Poisson regression                               Number of obs   =       360
                                                LR chi2(5)      =       527.17
                                                Prob > chi2     =       0.0000
Log likelihood = -898.24495                    Pseudo R2       =       0.2269
```

```
-----
          lc |          IRR   Std. Err.      z    P>|z|     [95% Conf. Interval]
-----+-----
         sex |    2.080201   .1907444     7.99  0.000    1.738017   2.489753
         |
    ageQ5 |
         1 |           1 (base)
         2 |    2.562157   .4319319     5.58  0.000    1.841234   3.565352
         3 |    6.721469   1.053021    12.16  0.000    4.944364   9.137301
         4 |   13.40016   2.123278    16.38  0.000     9.82281   18.28034
```

```

      5 |      20.4727   4.041424   15.29   0.000   13.90412   30.14443
      |
    _cons | .0000887   .0000132  -62.51   0.000   .0000662   .0001188
ln(pt) |           1 (exposure)
-----

```

This shows a very similar point estimate and standard errors to modelling attained age as a linear, continuous effect. We also investigate using restricted cubic splines:

```

. mkspline ageSpline = age, cubic nknots(4)
. poisson lc sex ageSpline*, exposure(pt) nolog irr base

```

```

Poisson regression                               Number of obs   =       360
                                                LR chi2(4)      =       564.98
                                                Prob > chi2     =       0.0000
Log likelihood = -879.33904                    Pseudo R2      =       0.2431
-----

```

```

      lc |      IRR   Std. Err.      z    P>|z|    [95% Conf. Interval]
-----+-----
    sex |  2.085053   .1911968     8.01   0.000    1.742059   2.495578
ageSpline1 |  1.112402   .0236588     5.01   0.000    1.066985   1.159752
ageSpline2 |  1.029289   .0634749     0.47   0.640    .912105   1.161528
ageSpline3 |  .8037657   .1399861    -1.25   0.210    .5713231   1.130778
    _cons |  7.62e-07   7.95e-07   -13.50   0.000    9.84e-08   5.89e-06
ln(pt) |           1 (exposure)
-----

```

Again, this shows a very similar point estimate and standard errors to modelling attained age as a linear, continuous effect. I accepted answers using any of quintiles, linear/continuous age, splines or similar functional forms.

In summary, lung cancer incidence is associated with age, sex, asbestos exposure and current smoking exposure.

Question 2

We now adjust for age, sex, smoking exposure and asbestos exposure in the same model.

```

. poisson lc age sex smoking asbestos, exposure(pt) nolog irr

```

```

Poisson regression                               Number of obs   =       360
                                                LR chi2(4)      =      1435.18
                                                Prob > chi2     =       0.0000
Log likelihood = -444.23989                    Pseudo R2      =       0.6176
-----

```

```

      lc |      IRR   Std. Err.      z    P>|z|    [95% Conf. Interval]
-----+-----
    age |  1.103361   .0046225    23.48   0.000    1.094338   1.112459
    sex |  1.475907   .1372634     4.19   0.000    1.229971   1.771019
 smoking | 15.13076   1.739477    23.63   0.000   12.07825   18.95472
 asbestos |  3.443032   .3718914    11.45   0.000    2.786124   4.254825
    _cons |  2.84e-07   8.41e-08   -50.98   0.000    1.59e-07   5.08e-07
ln(pt) |           1 (exposure)
-----

```

```

. est store ModelA

```

This shows clearly that each of attained age, sex, smoking and asbestos exposure are significantly associated with lung cancer incidence ($p < 0.001$ for all adjusted effects). The adjusted rate ratio (RR)

for age was 1.104 (95% CI: 1.095, 1.113) per year of age, indicating a rapid rise with increasing age. Males have higher rates of disease even after adjustment for other covariates (RR=1.45, 95% CI: 1.20, 1.74). Smoking is strongly associated with lung cancer incidence (RR=17.63, 95% CI: 13.90, 22.35). Finally, asbestos exposure has a rate ratio of 3.27 (95% CI: 2.64, 4.05).

Empirical evidence for confounding can be assessed in several ways. First, we can assess whether exposure to smoking and asbestos are associated:

```
. tab smoking asbestos [aw=pt], row
```

```
+-----+
| Key          |
|-----|
| frequency    |
| row percentage |
+-----+
```

smoking	asbestos		Total
	0	1	
0	252.38654	19.281283	271.66783
	92.90	7.10	100.00
1	81.849054	6.4831182	88.332172
	92.66	7.34	100.00
Total	334.2356	25.764401	360
	92.84	7.16	100.00

We see that the prevalence of exposure to asbestos is similar or slightly lower among never smokers (7.4%) and current smokers (8.0%). We are not able to undertake a formal statistical test with these weighted data.

Second, we can assess whether the estimated associations between lung cancer incidence and each of smoking and asbestos change after an adjustment for other covariates.

Comparing the linear age-adjusted model with the main effects model, we see that the rate ratio for asbestos changed from 3.68 to 3.42 (7% reduction), and the rate ratio for smoking changed from 18.45 to 17.63 (4% reduction). Again, there is limited evidence for confounding between smoking and asbestos.

Question 3

(a)

A regression model formula is

$$\log(\lambda(t|x)) = \beta_0 + \beta_1 \text{age} + \beta_2 I(\text{sex} = 1) + \beta_3 I(\text{smoking} = 1) + \beta_4 I(\text{asbestos} = 1) + \beta_5 I(\text{smoking} = 1 \ \& \ \text{asbestos} = 1)$$

where $\lambda(t|x)$ is the rate at attained age t given covariates x (including sex, smoking and asbestos), with coefficients $\beta_0, \beta_1, \beta_2, \beta_3, \beta_4$ and β_5 , and $I(\text{test})$ is 1 if the test is true and 0 if the test is false.

(b)

We now fit the interaction model:

```
. poisson lc age sex smoking##asbestos, exposure(pt) nolog irr
```

```
Poisson regression          Number of obs   =          360
                             LR chi2(5)           =       1447.96
                             Prob > chi2          =          0.0000
Log likelihood = -437.84669   Pseudo R2      =          0.6231
```

```
-----+-----
```

lc	IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
age	1.103094	.0046206	23.42	0.000	1.094075	1.112188
sex	1.473861	.1368159	4.18	0.000	1.228686	1.767959
1.smoking	19.3376	2.728971	20.99	0.000	14.66489	25.49919
1.asbestos	7.027045	1.516922	9.03	0.000	4.602826	10.72805
smoking# asbestos						
1 1	.4007745	.0999591	-3.67	0.000	.2458089	.653435
_cons	2.35e-07	7.16e-08	-50.17	0.000	1.30e-07	4.27e-07
ln(pt)	1	(exposure)				

```
-----+-----
```

```
. est store ModelB
. lrtest ModelA ModelB
```

```
Likelihood-ratio test                    LR chi2(1) =    12.79
(Assumption: ModelA nested in ModelB)    Prob > chi2 =    0.0003
```

Comparing Model A with Model B, we see that there is little evidence for a statistical interaction on a multiplicative scale. First, we note that the Wald test for the interaction term has a p-value of 0.18. Second, we see that the likelihood ratio test is also not significant, with $p = 0.19$.

(c)

From Model B, we can calculate the incidence rate for a males aged 62 years who has been exposed to asbestos and is a current smoker using several approaches. We can calculate the rate from the regression estimates, however we need to take account of the covariance terms to calculate the confidence interval, which is best done using tools provided by each statistical package. Using the `lincom` command:

```
. quietly poisson lc age sex smoking##asbestos, exposure(pt) nolog irr
. lincom sex + 1.smoking + 1.asbestos + 1.smoking#1.asbestos + 62*age + _cons,
> irr
```

```
( 1) 62*[lc]age + [lc]sex + [lc]1.smoking + [lc]1.asbestos +
      [lc]1.smoking#1.asbestos + [lc]_cons = 0
```

```
-----+-----
```

lc	IRR	Std. Err.	z	P> z	[95% Conf. Interval]	
(1)	.0082793	.0009574	-41.46	0.000	.0066003	.0103855

```
-----+-----
```

This shows that the incidence rate is 9.19 (95% CI: 7.46, 10.32) per 1000 person-years.

We can do the same analysis using the `margins` command:

```
. margins smoking##asbestos, predict(ir) at(age=62 sex=1)
```

```
Predictive margins                    Number of obs   =    360
Model VCE      : OIM
```

```
Expression   : Predicted incidence rate, predict(ir)
at           : age           =    62
              sex           =    1
```

```
-----+-----
```

		Delta-method				[95% Conf. Interval]	
		Margin	Std. Err.	z	P> z		
smoking							
	0	.0006102	.0000941	6.48	0.000	.0004257	.0007946
	1	.0056096	.0004958	11.31	0.000	.0046378	.0065813
asbestos							
	0	.0015459	.0000953	16.22	0.000	.0013592	.0017327
	1	.0046738	.0004901	9.54	0.000	.0037133	.0056344
smoking#							
asbestos							
	0 0	.000152	.0000209	7.27	0.000	.000111	.000193
	0 1	.0010683	.0001859	5.75	0.000	.000704	.0014326
	1 0	.0029398	.0001859	15.82	0.000	.0025756	.0033041
	1 1	.0082793	.0009574	8.65	0.000	.0064028	.0101558

Finally, we could also do this analysis with the `predict` command.

Part 2

Question 4

We read in the data using the following:

```
. display "Folder = $folder"
Folder = 5
. import delimited "http://biostat3.net/download/exams/2016/$folder/survival.csv", clear
(8 vars, 522 obs)
```

(a)

This question is equivalent to completing *Table 1* for a randomised controlled trial to assess whether randomisation led to balanced covariates. We use simple tests to assess whether treatment assignment varies substantially by age at diagnosis, sex, smoking exposure and asbestos exposure.

For age at diagnosis, we can use either a t-test or a non-parametric test:

```
. ttest age, by(tx)
```

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.	[95% Conf. Interval]	
0	252	63.37966	.6228296	9.887114	62.15302	64.6063
1	270	62.9975	.5749281	9.447033	61.86557	64.12943
combined	522	63.18199	.4225693	9.654575	62.35184	64.01214
diff		.3821607	.8462883		-1.280404	2.044725

```
diff = mean(0) - mean(1)
Ho: diff = 0
t = 0.4516
degrees of freedom = 520
```

```
Ha: diff < 0
Pr(T < t) = 0.6741
Ha: diff != 0
Pr(|T| > |t|) = 0.6518
Ha: diff > 0
Pr(T > t) = 0.3259
. ranksum age, by(tx)
```

Two-sample Wilcoxon rank-sum (Mann-Whitney) test

tx	obs	rank sum	expected
0	252	66577	65898
1	270	69926	70605
combined	522	136503	136503

unadjusted variance 2965410.00
 adjustment for ties 0.00

 adjusted variance 2965410.00

Ho: age(tx==0) = age(tx==1)
 z = 0.394
 Prob > |z| = 0.6934

We find no evidence that age differs by treatment modality ($p = 0.46$ for the t-test and $p = 0.61$ for the Wilcoxon test). For the other variables:

. tab tx sex, chi row

tx	sex		Total
	0	1	
0	92	160	252
	36.51	63.49	100.00
1	92	178	270
	34.07	65.93	100.00
Total	184	338	522
	35.25	64.75	100.00

Pearson chi2(1) = 0.3383 Pr = 0.561

. tab tx smoking, chi row

tx	smoking		Total
	0	1	
0	47	205	252
	18.65	81.35	100.00

1	46	224	270
	17.04	82.96	100.00
-----+			
Total	93	429	522
	17.82	82.18	100.00

Pearson chi2(1) = 0.2318 Pr = 0.630

. tab tx asbestos, chi row

```
+-----+
| Key      |
|-----|
| frequency|
| row percentage|
+-----+
```

tx	asbestos		Total
	0	1	
0	196	56	252
	77.78	22.22	100.00
1	215	55	270
	79.63	20.37	100.00
Total	411	111	522
	78.74	21.26	100.00

Pearson chi2(1) = 0.2670 Pr = 0.605

We find little evidence that randomisation varied by sex ($p = 0.09$), by smoking ($p = 0.21$) or by asbestos exposure ($p = 0.86$). We could check for potential confounding by sex in the survival analysis.

(b)

We `stset` the data using time since diagnosis as the primary time scale and then plot the Kaplan-Meier curves

```
. stset tsurv, failure(event) id(id)
```

```
          id: id
failure event: event != 0 & event < .
obs. time interval: (tsurv[_n-1], tsurv]
exit on or before: failure
```

```
-----
522 total observations
  0 exclusions
-----
```

```
522 observations remaining, representing
522 subjects
459 failures in single-failure-per-subject data
538.4558 total analysis time at risk and under observation
                                at risk from t =          0
                                earliest observed entry t =      0
                                last observed exit t =          5
```

```
. sts graph, by(tx) name(km1, replace) scheme(s2mono)
```

```
failure _d: event
```



```

analysis time _t: tsurv
id: id
. graph export exam_2016_km1.eps, name(km1) replace
(file exam_2016_km1.eps written in EPS format)
. * the following line is only needed on Linux
. !! convert -density 300 exam_2016_km1.eps exam_2016_km1_$folder.png
. sts test tx

```

```

failure _d: event
analysis time _t: tsurv
id: id

```

Log-rank test for equality of survivor functions

tx	Events observed	Events expected
0	221	238.29
1	238	220.71
Total	459	459.00

chi2(1) = 2.61

Pr>chi2 = 0.1059

. sts list, by(tx) at(1 2 3 4 5)

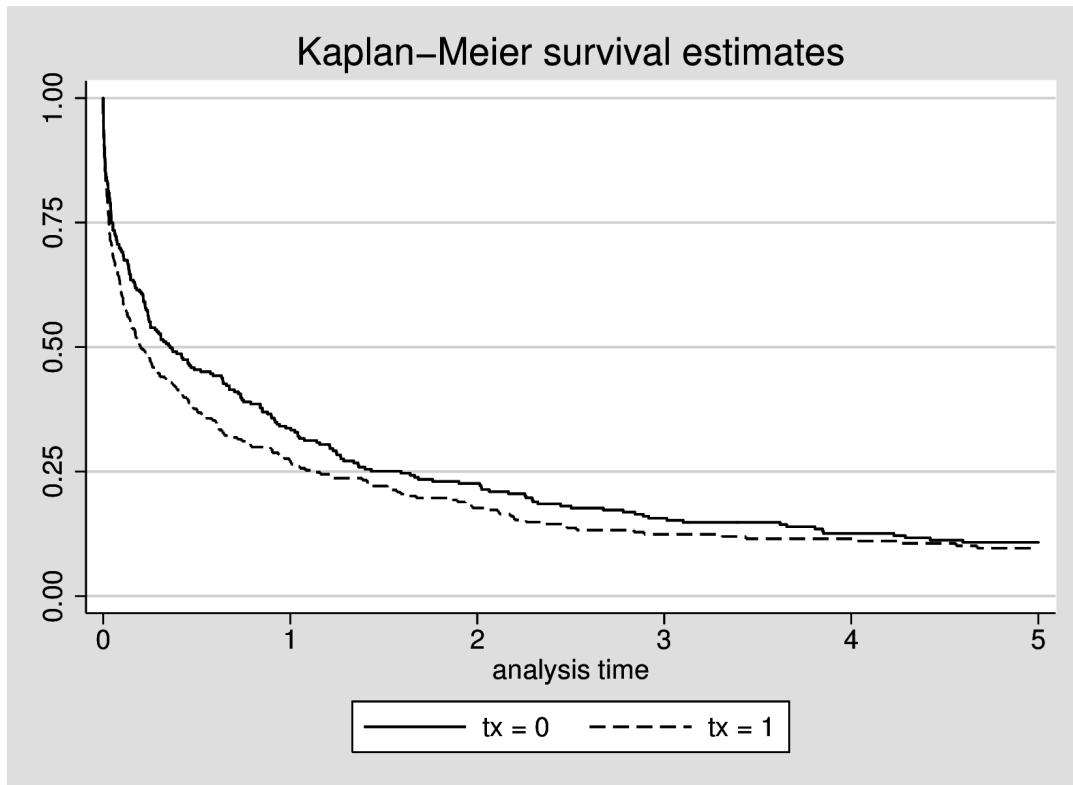
```

failure _d: event
analysis time _t: tsurv
id: id

```

	Time	Beg. Total	Fail	Survivor Function	Std. Error	[95% Conf. Int.]	
tx=0							
	1	83	166	0.3371	0.0299	0.2791	0.3960
	2	56	27	0.2261	0.0266	0.1761	0.2800
	3	39	17	0.1562	0.0232	0.1140	0.2045
	4	29	7	0.1258	0.0213	0.0877	0.1710
	5	24	4	0.1078	0.0201	0.0725	0.1510
tx=1							
	1	71	195	0.2721	0.0273	0.2200	0.3267
	2	45	24	0.1768	0.0237	0.1332	0.2257
	3	29	13	0.1239	0.0207	0.0870	0.1678
	4	26	2	0.1151	0.0201	0.0794	0.1580
	5	19	4	0.0963	0.0189	0.0633	0.1373

Note: survivor function is calculated over full data and evaluated at indicated times; it is not calculated from aggregates shown at left.



The Kaplan-Meier curves show that survival is poor for lung cancer patients, with fewer than 25% of patients surviving to 5 years. We also see that treatment with chemotherapy+radiotherapy leads to more deaths soon after diagnosis. It is unclear whether the rates are different after one year.

Although not specifically asked for, we also (i) used the log-rank test to compare the curves, finding strong evidence for a difference ($p = 0.0001$) and (ii) estimated survival to five years, where 9% (95% CI: 6, 13) survived for those on conventional treatment and 3% (95% CI: 1, 6) survived for those on chemotherapy+radiotherapy.

Question 5

Based on Question 4 (a), we first investigated whether age and sex were associated with survival and hence would be potential confounders:

```
. stcox tx sex age, nolog
```

```
      failure _d:  event
analysis time _t:  tsurv
           id:  id
```

```
Cox regression -- no ties
```

```
No. of subjects =          522          Number of obs =          522
No. of failures =          459
Time at risk    = 538.4557975
Log likelihood   = -2529.8439          LR chi2(3) =          3.64
                                          Prob > chi2 =          0.3035
```

	_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
	tx	1.154286	.1086023	1.53	0.127	.9599027 1.388033
	sex	.9927266	.0964916	-0.08	0.940	.8205293 1.201061
	age	.9950297	.0048949	-1.01	0.311	.9854821 1.00467

Adjusting for treatment modality, there is no evidence that either sex or age are associated with survival, with Wald test p-values of 0.75 and 0.25 for sex and age, respectively. Furthermore, fitting a Cox regression models with and without age and sex suggest that the effect of treatment modality is insensitive to inclusion of age and sex in the model. The hazard ratio for chemotherapy+radiotherapy compared with conventional therapy is 1.77 (95% CI: 1.47, 2.13), suggesting that the average hazard ratio for chemotherapy+radiotherapy is high over the five-year period.

For the time scale, we have initially used time since cancer diagnosis. There is a strong association between time since diagnosis and survival, suggesting that this is the best choice of primary time scale. Moreover, there is a suggestion of non-proportional hazards, with a higher rate ratio in the first year than for the later years. We could investigate using attained age as the primary time scale, but then we would need to finely model for the time since diagnosis, which would require modelling two time scales. For simplicity, we propose using time since diagnosis as the primary time scale.

Question 6

(i)

For an analysis of scaled Schoenfeld residuals, we use:

```
. estat phtest, detail
```

```
Test of proportional-hazards assumption
```

```
Time: Time
```

	rho	chi2	df	Prob>chi2
tx	-0.06370	1.84	1	0.1745
global test		1.84	1	0.1745

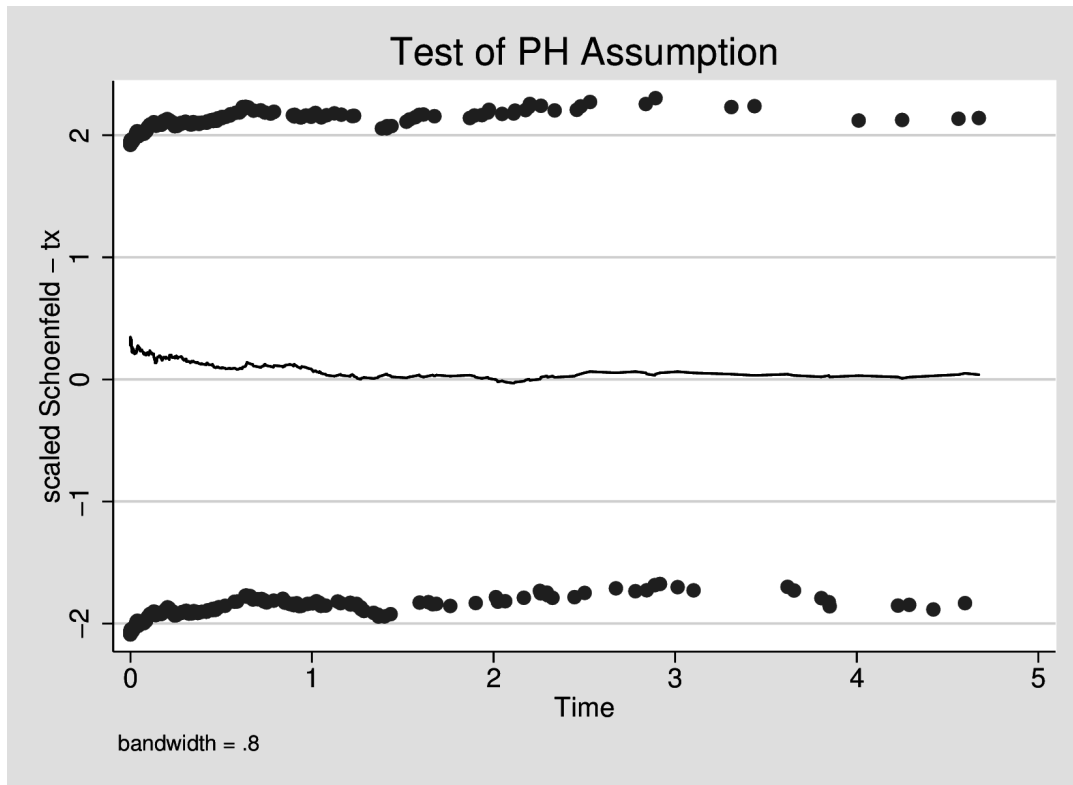
```
. estat phtest, plot(tx) name(phtest, replace) scheme(s2mono)
```

```
. graph export exam_2016_phtest.eps, name(phtest) replace
```

```
(file exam_2016_phtest.eps written in EPS format)
```

```
. * the following line is only needed on Linux
```

```
. !! convert -density 300 exam_2016_phtest.eps exam_2016_phtest_`$folder`.png
```



This shows that there is little evidence ($p = 0.14$) that the hazard ratio decreases with increasing time since diagnosis: the scaled residuals and linear time have a correlation of -0.07. From the plot of the scaled residuals and time, we see the running mean smoother dips early in the follow-up period and then is flat or very slightly declining. Given the number of events that are early in the period, we could also test using a log-transformation for time since diagnosis:

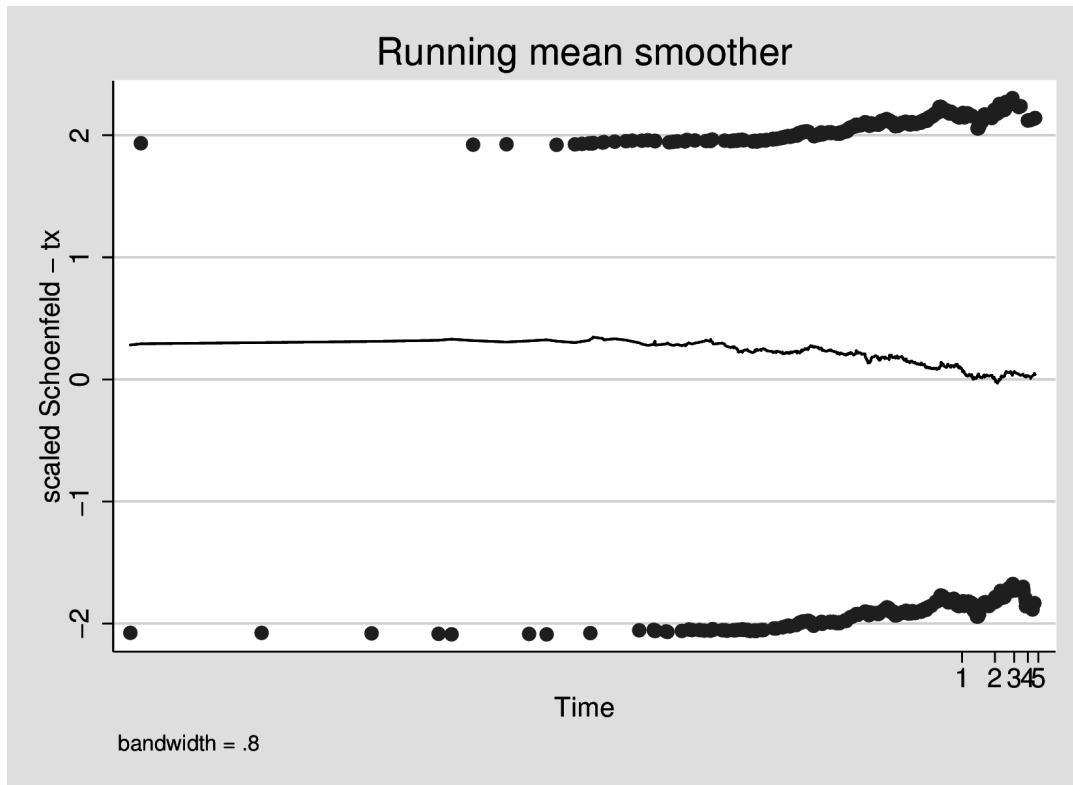
```
. estat phtest, detail log
```

```
Test of proportional-hazards assumption
```

```
Time: Log(t)
```

	rho	chi2	df	Prob>chi2
tx	-0.02676	0.33	1	0.5684
global test		0.33	1	0.5684

```
. estat phtest, log plot(tx) name(phtestlog, replace) scheme(s2mono)
. graph export exam_2016_phtestlog.eps, name(phtestlog) replace
(file exam_2016_phtestlog.eps written in EPS format)
. * the following line is only needed on Linux
. !! convert -density 300 exam_2016_phtestlog.eps exam_2016_phtestlog_$folder.p
> ng
```



Again, there is little evidence for non-proportionality ($p = 0.15$).

(ii)

We can test for piecewise-constant hazard ratios by splitting by time and fitting for an interaction. In the following, the "c" prefix indicates a continuous variable, while the "i" prefix indicates a factor variable.

```
. quietly import delimited "http://biostat3.net/download/exams/2016/$folder/sur
> vival.csv", clear
. quietly stset tsurv, fail(event) id(id)
. stsplit timeband, at(0, 1, max)
(152 observations (episodes) created)
. stcox sex i.tx##i.timeband, nolog

      failure _d:  event
analysis time _t:  tsurv
              id:  id
```

Cox regression -- no ties

```
No. of subjects =          522          Number of obs =          674
No. of failures =          459
Time at risk    = 538.4557975
Log likelihood  = -2529.6271          LR chi2(3) =          4.07
                                          Prob > chi2 =          0.2540
```

	_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
	sex	.9921265	.0964386	-0.08	0.935	.8200246 1.200348
	1.tx	1.234292	.1307633	1.99	0.047	1.002859 1.519133
	1.timeband	20.09028

```
tx#timeband |
      1 1 | .7588576 .1741348 -1.20 0.229 .4839889 1.189831
```

```
-----
. stcox tx sex c.tx#c.timeband, nolog
```

```
      failure _d: event
      analysis time _t: tsurv
              id: id
```

```
Cox regression -- no ties
```

```
No. of subjects =          522          Number of obs =          674
No. of failures =          459
Time at risk    = 538.4557975
Log likelihood   = -2529.6271          LR chi2(3) =          4.07
                                      Prob > chi2 =          0.2540
```

```
-----
      _t | Haz. Ratio   Std. Err.      z    P>|z|    [95% Conf. Interval]
-----+-----
      tx |  1.234292   .1307633     1.99   0.047    1.002859   1.519133
      sex |  .9921265   .0964386    -0.08   0.935    .8200246   1.200348
      |
      c.tx#|
      c.timeband | .7588576 .1741348 -1.20 0.229 .4839889 1.189831
-----
```

```
. stcox c.tx#i.timeband, nolog
```

```
      failure _d: event
      analysis time _t: tsurv
              id: id
```

```
Cox regression -- no ties
```

```
No. of subjects =          522          Number of obs =          674
No. of failures =          459
Time at risk    = 538.4557975
Log likelihood   = -2529.6304          LR chi2(2) =          4.06
                                      Prob > chi2 =          0.1311
```

```
-----
      _t | Haz. Ratio   Std. Err.      z    P>|z|    [95% Conf. Interval]
-----+-----
timeband#c.tx |
      0 |  1.233572   .1303855     1.99   0.047    1.002754   1.51752
      1 |  .9363188   .1906887    -0.32   0.747    .6281595   1.395654
-----
```

This model provides little or no evidence that the hazard ratio is time-dependent ($p = 0.57$). The hazard ratio in the first year is 1.83 (95% CI: 1.48, 2.25), while the hazard ratio after the first year is 1.60 (95% CI: 1.08, 2.39).

(iii)

We can re-fit the model in (ii) using Stata `stcox`'s `tv` and `te` options:

```
. stcox tx, nolog tv(c.tx) te(_t>=1)
```

```
      failure _d: event
```

```
analysis time _t: tsurv
id: id
```

Cox regression -- no ties

```
No. of subjects =          522          Number of obs =          674
No. of failures =          459
Time at risk   = 538.4557975
Log likelihood = -2529.6304          LR chi2(2) =          4.06
          Prob > chi2 =          0.1311
```

	_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
-----+-----						
main						
	tx	1.233572	.1303855	1.99	0.047	1.002754 1.51752
-----+-----						
tvc						
	tx	.7590309	.1741616	-1.20	0.230	.4841156 1.190063
-----+-----						

Note: variables in tvc equation interacted with _t>=1

Again, we find little evidence for a time-dependent hazard ratio ($p = 0.57$). We can model for a time-dependent hazard ratio that depends on time:

```
. stcox tx, nolog tvc(tx) texp(_t)
```

```
failure _d: event
analysis time _t: tsurv
id: id
```

Cox regression -- no ties

```
No. of subjects =          522          Number of obs =          674
No. of failures =          459
Time at risk   = 538.4557975
Log likelihood = -2529.4288          LR chi2(2) =          4.47
          Prob > chi2 =          0.1072
```

	_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
-----+-----						
main						
	tx	1.266262	.14277	2.09	0.036	1.015199 1.579413
-----+-----						
tvc						
	tx	.8718413	.0885579	-1.35	0.177	.7144569 1.063895
-----+-----						

Note: variables in tvc equation interacted with _t

The interpretation of this model is as follows: the hazard ratio at time 0 is 1.97 (95% CI: 1.57, 2.48); for each year since diagnosis, the rate tends to decrease by 1-0.84=16% (RR=0.84, 95% CI: 0.68, 1.05), although this trend is not significant ($p = 0.12$, as per the Schoenfeld test).

(iv)

Using `stpm2` with time-dependent hazard ratios, we use a low-dimensional natural spline for the time-dependent effect. We use a Wald test to check for time-dependence and plot the time-dependent hazard

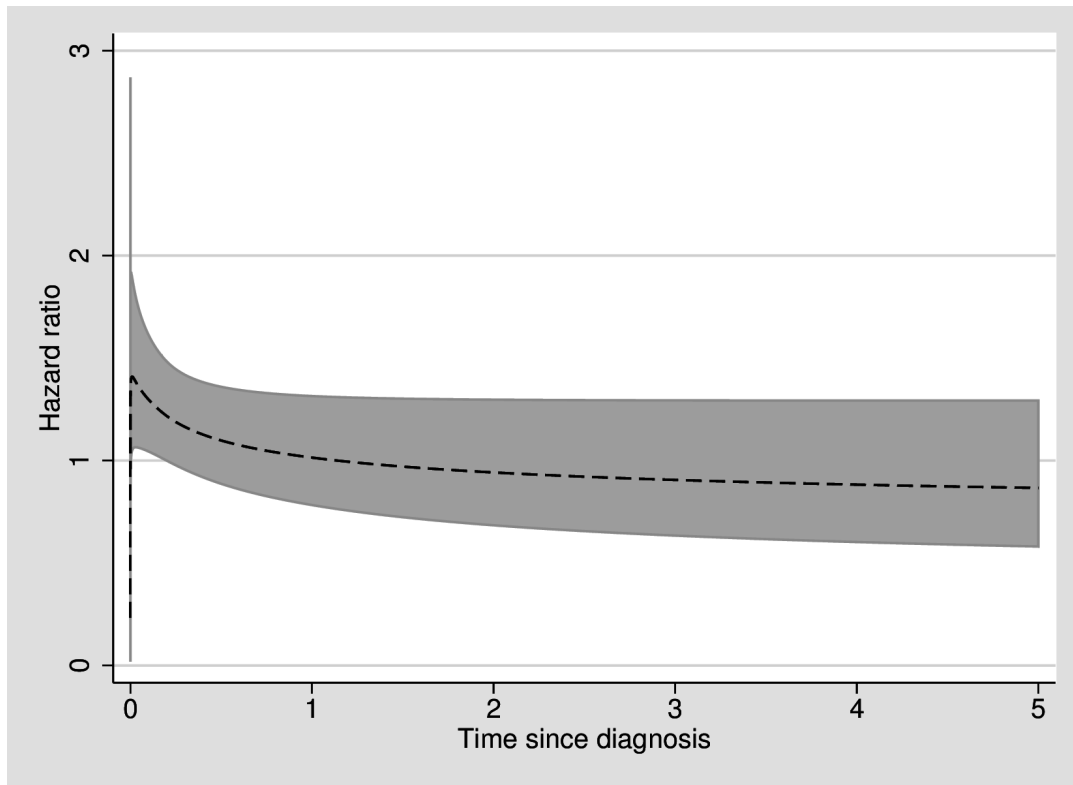
ratio:

```
. stpm2 tx, df(4) scale(hazard) nolog eform tvc(tx) dftvc(2)  
note: delayed entry models are being fitted
```

```
Log likelihood = -1195.3843          Number of obs   =          674
```

```
-----+-----  
          |      exp(b)   Std. Err.      z    P>|z|      [95% Conf. Interval]  
-----+-----  
xb          |  
    tx      |   1.184981   .1266655    1.59   0.112    .9610019    1.461162  
   _rcs1    |   3.020649   .2299325   14.52   0.000    2.601996    3.506662  
   _rcs2    |   .9115485   .0544966   -1.55   0.121    .8107575    1.024869  
   _rcs3    |   1.03003    .0282722    1.08   0.281    .9760816    1.08696  
   _rcs4    |   .9979048   .0169522   -0.12   0.902    .965226     1.03169  
  _rcs_tx1  |   1.074447   .1239423    0.62   0.534    .8570282    1.347023  
  _rcs_tx2  |   1.170552   .1053925    1.75   0.080    .9811864    1.396465  
   _cons    |   .5874006   .0462969   -6.75   0.000    .5033216    .6855249  
-----+-----
```

```
. test _rcs_tx1 _rcs_tx2  
  
( 1) [xb]_rcs_tx1 = 0  
( 2) [xb]_rcs_tx2 = 0  
  
      chi2( 2) =      3.51  
      Prob > chi2 =      0.1732  
. predict hr, hrnumerator(tx 1) ci  
. twoway (rarea hr_lci hr_uci _t if hr_uci<5, sort color(gs12)) (line hr _t if  
> hr_uci<5, sort), legend(off) xtitle("Time since diagnosis") ytitle("Hazard ra  
> tio") name(hr, replace) scheme(s2mono)  
. graph export exam_2016_hr.eps, name(hr) replace  
(file exam_2016_hr.eps written in EPS format)  
. * the following line is only needed on Linux  
. !! convert -density 300 exam_2016_hr.eps exam_2016_hr_$folder.png
```



We see that there is limited evidence for time-dependent hazards ($p = 0.37$ from the Wald test). We also see from the plot that the hazard ratio looks comparatively stable across the follow-up period.

Question 7

(a)

Advantages of using Poisson regression for Questions 5–6 include: (i) Poisson regression readily models for multiple time scales, where we could split on attained age and time since diagnosis and then model for main effects and interactions between those time scales and interactions between a time scale and other covariates; (ii) it is simpler to predict rates from Poisson regression, as the analysis is done on that scale.

Disadvantages of using Poisson regression include: (i) the need to split on the time scales, which may increase the size of the computational problem; (ii) the need to specify a functional form for the primary time scale using parametric functions, rather than using Cox regression's non-parametric formulation; (iii) crude time splitting will assume that rates are piece-wise constant, which may not be appropriate; (iv) risk calculations for Poisson regression require that the risk period involves constant rates or numerical integration.

(b)

Assuming that the follow-up time has been split for within one year of diagnosis and from one year of diagnosis, we can model the rate using:

$$\log(\lambda(t|\text{tx})) = \beta_0 + \beta_1 I(t < 1) + \beta_2 I(t \geq 1) + \beta_3 I(\text{tx} = 1) + \beta_4 I(\text{tx} = 1 \ \& \ t \geq 1)$$

A better formulation would be to include more time-splits for time since diagnosis. If we let time cuts be represented by t_j where $t_0 = 0$, then

$$\log(\lambda(t|\text{tx})) = \beta_0 + \sum_j \beta_j I(t_{j-1} < t \leq t_j) + \beta_{\text{tx}} I(\text{tx} = 1) + \beta_{\text{tx},t} I(\text{tx} = 1 \ \& \ t \geq 1)$$

We could also model using splines. Any similar formulation was accepted, including different formulations for the time-dependent hazard ratios.