

```

. use melanoma
. gen id=_n
. stset survtime , failure(status==1) id(id)
      id: id
      failure event: status == 1
      obs. time interval: (survtime[_n-1], survtime]
      exit on or before: failure
-----+-----
      205 total obs.
      0  exclusions
-----+-----
      205 obs. remaining, representing
      205 subjects
      57 failures in single failure-per-subject data
      441324 total analysis time at risk, at risk from t =      0
      earliest observed entry t =      0
      last observed exit t =      5565

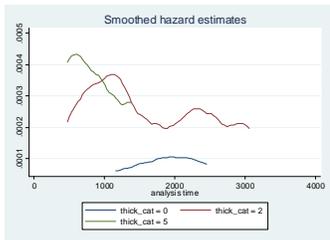
. // point is to learn about effect of the continuous variables
. egen thck_cat = cut(thck) , at(0, 2, 5, 20)
. egen age_cat = cut(age) , at(0, 35, 55, 65, 110)

```

```

. sts graph, by(thck_cat) hazard
      failure _d: status == 1
      analysis time _t: survtime
      id: id

```



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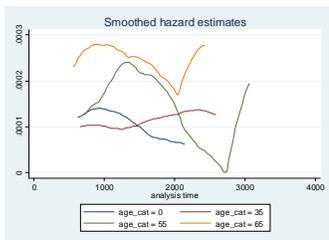
. // most info is binary
. replace thck_cat = thck >= 20
(% real changes made)

```

```

. sts graph, by(age_cat) hazard
      failure _d: status == 1
      analysis time _t: survtime
      id: id

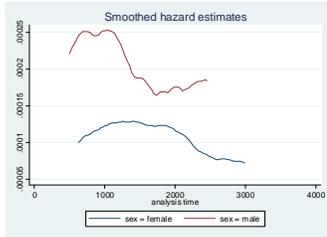
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```

. sts graph, by(sex) hazard
      failure_d: status == 1
      analysis_time_t: survtime
      id: id

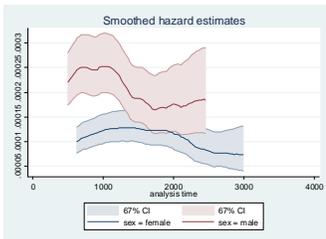
```



```

. sts graph, by(sex) hazard ci level(67)
      failure_d: status == 1
      analysis_time_t: survtime
      id: id

```



```

. stcox sex , tvc(sex) texp(log(survtime/1600)) nohr
      failure_d: status == 1
      analysis_time_t: survtime
      id: id

```

```

Iteration 0: log likelihood = -283.19925
Iteration 1: log likelihood = -279.41449
Iteration 2: log likelihood = -279.40855
Iteration 3: log likelihood = -279.40855
Refining estimates:
Iteration 0: log likelihood = -279.40855

```

Cox regression -- no ties

```

No. of subjects = 205           Number of obs = 1870
No. of failures = 57
Time at risk = 441324          LR chi2(2) = 7.59
Log likelihood = -279.40855    Prob > chi2 = 0.0225

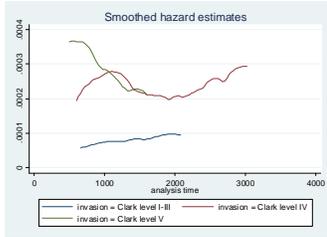
```

	_t	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
main	sex	.4545677	.3203576	1.42	0.156	-.1733217 1.082457
	tvc					
	sex	-.4606954	.3932363	-1.17	0.241	-1.231424 .3100337

Note: variables in tvc equation interacted with log(survtime/1600)

```
. sts graph, by(invasion) hazard
```

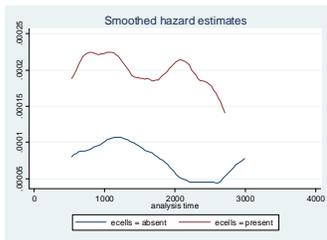
```
failure_d: status == 1  
analysis_time_t: survtime  
id: id
```



```
. // most info is binary  
. gen invas_cat = invasion > 0
```

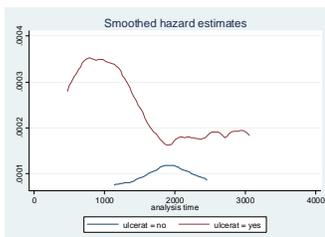
```
. sts graph, by(ecells) hazard
```

```
failure_d: status == 1  
analysis_time_t: survtime  
id: id
```



```
. sts graph, by(ulcerat) hazard
```

```
failure_d: status == 1  
analysis_time_t: survtime  
id: id
```



```

. replace age_cat = age > 35
(171 real changes made)

. // could use age as continuous, but most info is again binary
. // do not need xl here since all covars are binary 0,1
. stcox sex lnvas_cat ecelis ulcerat thck_cat age_cat

      failure_d: status == 1
analysis time _t: survtime
              id: id

No. of subjects =      205           Number of obs =      205
No. of failures =       57
Time at risk    =    441324

Log likelihood = -259.23961           LR chi2(6) =      47.92
                                      Prob > chi2 =     0.0000

```

_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
sex	1.683799	.473142	1.85	0.064	.9707453 2.92062
lnvas_cat	1.497389	.5872226	1.03	0.303	.6942526 3.229576
ecelis	1.688354	.5237427	1.69	0.091	.9192061 3.101087
ulcerat	2.370916	.7904118	2.59	0.010	1.233514 4.557097
thck_cat	2.096643	.9344657	1.66	0.097	.8751131 5.022331
age_cat	1.338263	.5257613	0.74	0.458	.6196287 2.890358

```

. // plausible that lnvas_cat and thck_cat are carrying same info
. estimates store full_mod

```

```

. stcox sex lnvas_cat ecel ulcer age_cat

      failure_d: status == 1
analysis time _t: survtime
              id: id

No. of subjects =      205           Number of obs =      205
No. of failures =       57
Time at risk    =    441324

Log likelihood = -259.79209           LR chi2(5) =      46.81
                                      Prob > chi2 =     0.0000

```

_t	Haz. Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
sex	1.581821	.4355338	1.67	0.096	.9221259 2.713468
ecelis	1.64861	.5094349	1.62	0.106	.8996822 3.020971
ulcerat	2.38814	.7973938	2.61	0.009	1.241212 4.594876
thck_cat	2.778232	.989867	2.87	0.004	1.381945 5.585299
age_cat	1.289179	.5045376	0.65	0.516	.5986551 2.776147

```

. lrtest full_mod

Likelihood-ratio test
(Assumption: _t nested in full_mod)           LR chi2(1) =      1.10
                                              Prob > chi2 =     0.2932

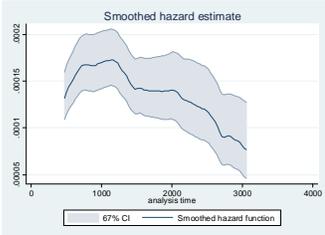
```

```

. // nature of baseline hazard
. sts graph , hazard ci level(67)

      failure_d: status == 1
analysis time _t: survtime
              id: id

```



```

// explore using Pan rgr
stgpl t_st_cat , every(250) after(0)
(1665 observations) (episodes) created

// do not use collapse without understanding it better
gen risk_time = _t - _t0

gen log_st = log(survtime/1500)

gen log_st_sq = log_st * log_st

poisson _d log_st log_st_sq sex ecol ulcer thck_cat age_cat , exp(risk_time)

Iteration 0: log likelihood = -295.01347
Iteration 1: log likelihood = -295.01347
Iteration 2: log likelihood = -295.01347

Poisson regression              Number of obs = 1870
                                LR chi2(6) = 49.73
                                Prob > chi2 = 0.0000
                                Pseudo R2 = 0.0777

Log likelihood = -295.01347

```

_d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
log_st	-.4692238	.3075451	-1.53	0.127	[-1.072001 , .1335536]
log_st_sq	-.3783439	.2278961	-1.66	0.097	[-.8250321 , .0683043]
sex	-.4300331	.2764531	-1.56	0.120	[-.1118049 , .9717172]
ecol	.5049301	.3087458	1.64	0.102	[-.1002005 , 1.110061]
ulcer	.9889528	.3344599	2.97	0.003	[.3203315 , 1.524074]
thck_cat	1.000505	.3561968	2.81	0.005	[.3023717 , 1.698638]
age_cat	.2378041	.3917115	0.61	0.543	[-.5290739 , 1.004682]
_cons	-10.62994	.4940477	-21.52	0.000	[-11.59825 , -9.661633]
risk_time	(exposure)				

```

// compare to coxreg
stcox sex ecol ulcer thck_cat age_cat , nhr

Failure _d: status == 1
analysis time _t: survtime
IG: id

No. of subjects = 205              Number of obs = 1870
No. of failures = 57
Time at risk = 441324              LR chi2(6) = 46.81
Log likelihood = -269.79209         Prob > chi2 = 0.0000

```

_t	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
sex	-.488577	.2753369	-1.77	0.078	[-.1010735 , .9882274]
ecol	.4999323	.3090388	1.62	0.106	[-.1067137 , 1.106578]
ulcer	.870149	.3338974	2.61	0.009	[.216088 , 1.524062]
thck_cat	1.021815	.3542938	2.87	0.004	[.3234918 , 1.720138]
age_cat	.2540055	.3913635	0.65	0.515	[-.5130529 , 1.021054]

```

// simplify baseline model
poisson _d log_st sex ecol ulcer thck_cat age_cat , exp(risk_time)

Iteration 0: log likelihood = -296.52572
Iteration 1: log likelihood = -296.52557
Iteration 2: log likelihood = -296.52557

Poisson regression              Number of obs = 1870
                                LR chi2(6) = 46.71
                                Prob > chi2 = 0.0000
                                Pseudo R2 = 0.0736

Log likelihood = -296.52557

```

_d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
log_st	-.0404921	.1668894	-0.36	0.717	[-.3875892 , .2666051]
sex	-.4421073	.2763889	-1.60	0.110	[-.090605 , .9838196]
ecol	.5110009	.3089322	1.65	0.098	[-.0944951 , 1.116487]
ulcer	.9920525	.3339436	2.97	0.003	[.327635 , 1.54657]
thck_cat	.9879986	.3539249	2.79	0.006	[.2743185 , 1.681679]
age_cat	.3387609	.3905463	0.87	0.414	[-.4466934 , 1.084219]
_cons	-10.50401	.4854094	-22.30	0.000	[-11.7754 , -9.872526]
risk_time	(exposure)				

```

. poisson _d sex ecal ul cer thi ck_cat age_cat , exp(risk_time)
Iteration 0: log likelihood = -296.59085
Iteration 1: log likelihood = -296.59083
Iteration 2: log likelihood = -296.59083

Poisson regression              Number of obs = 1870
                                LR chi2(3) = 46.55
                                Prob > chi2 = 0.0002
                                Pseudo R2 = 0.0728

Log likelihood = -296.59083

```

_d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
sex	-.4477186	.2758471	1.62	0.106	-.0929316 .9836692
ecal1s	.5133262	.3088862	1.66	0.097	-.0920438 1.118768
ulcerst	.9031567	.3316481	2.73	0.006	-.2550791 1.505194
thi ck_cat	.9662888	.3533995	2.73	0.006	-.2736385 1.658939
age_cat	.3364317	.3876971	0.87	0.386	-.4234406 1.095204
_cons	-10.82913	.4865467	-22.30	0.000	-11.78078 -9.877471
risk_time	(exposure)				

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. // constant hazard gives essentially correct estimates

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. // It was surprising that in the model before last that the log_st effect was so small
. // it is interesting in this regard that if we call all the covariables
. // the log_st effect becomes larger --- my interpretation of this is that when we ignore
. // the covariates there is some individual heterogeneity, and selection causes the
. // hazard to decrease, adjusting for the covariates removes this, so they may be
. // carrying most of the "heterogeneity"
. poisson _d log_st
Iteration 0: log likelihood = -255.08779
Iteration 1: log likelihood = -255.08779

Poisson regression              Number of obs = 1870
                                LR chi2(1) = 1.76
                                Prob > chi2 = 0.1849
                                Pseudo R2 = 0.0084

Log likelihood = -255.08779

```

_d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
log_st	-.2109464	.1640891	-1.35	0.176	-.5168167 .0949239
_cons	-3.558442	.1460562	-24.36	0.000	-3.844719 -3.272166