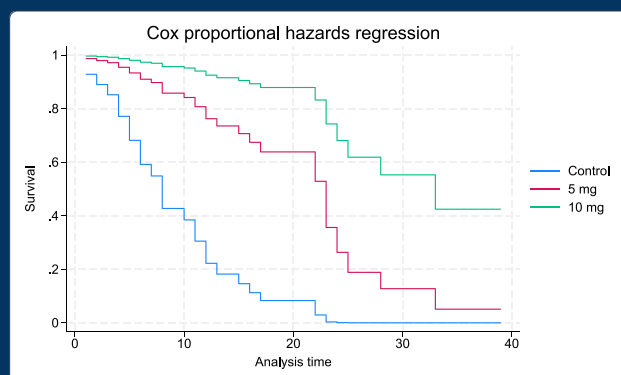


## Survival analysis

From Kaplan–Meier estimates of the survivor function to the Cox proportional hazards model, from competing-risks regression to multilevel survival models, Stata has everything you need to analyze your survival- or event-time data.



### • Survival-time data

- Single failure or multiple failures; right-, left-, and interval-censoring; left-truncation; gaps
- Support for complex survey designs

### • Life tables

- Tables and graphs with CIs
- Tests for equality of survivor functions
- Tests for trend

### • Graph survivor, cumulative hazard, and other functions

### • Cox proportional hazards model

- Stratified estimation
- Time-varying covariates **Updated**
- Shared frailty models
- Harrell's *C*, Somer's *D*, Gönen and Heller's *K*
- Tests for proportional hazards
- Goodness-of-fit plots **New**

### • Parametric survival models

- Weibull, exponential, Gompertz, lognormal, loglogistic, and generalized gamma
- Stratified models
- Individual or shared frailty
- Predictions of mean or median time to failure, survival probabilities, and hazards
- Goodness-of-fit plots **New**
- Bayesian estimation
- Finite mixture models

### • Competing risks model

- Fine and Gray proportional subhazards model
- Graph cumulative subhazard and cumulative incidence

### • Multilevel survival models

- Weibull, exponential, lognormal, loglogistic, and gamma
- Marginal predictions and marginal means

### • Structural equation models

- Weibull, exponential, lognormal, loglogistic, and gamma models
- Survival outcomes with other outcomes
- Path models, growth curve models, and more

### • Additive models of relative risk **New**

- Cox, parametric survival, interval-censored Cox, and interval-censored parametric survival models

### • Power analysis

- Log-rank test of survival curves, Cox models, exponential regression

### • Sample-size analysis for group sequential designs (GSDs) **New**

- Log-rank test of survival curves

### • Causal inference (treatment effects estimation)

- Regression adjustment, inverse-probability weighting (IPW), and doubly robust methods
- Average treatment effects (ATEs) and ATEs on the treated (ATETs)

### • Lasso and elastic net for Cox model **New**

- Select predictors via cross-validation, adaptive lasso, or BIC
- Penalized and postselection predictions
- Graph survivor and other functions

We begin by specifying that we have survival data using **stset**. Here **studytime** records the time of failure or censoring, and the variable **died** indicates whether the subject died or was censored.

```
Viewer - view st1.smcl
view st1.smcl X
+
Dialog ▾ Also see ▾ Jump to ▾
. . stset studytime, failure(died)

Survival-time data settings

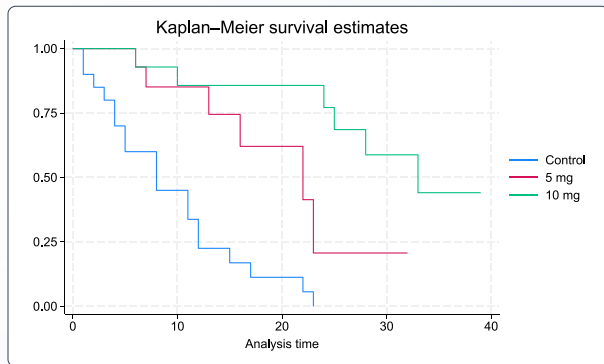
Failure event: died!=0 & died<.
Observed time interval: (0, studytime]
Exit on or before: failure

48 total observations
0 exclusions

48 observations remaining, representing
31 failures in single-record/single-failure data
744 total analysis time at risk and under observation
At risk from t = 0
Earliest observed entry t = 0
Last observed exit t = 39
CAP NUM INS
```

We are now ready to graph the survivor function for each level of our treatment,

```
. sts graph, by(dose)
```



or to fit a Cox proportional hazards model,

```
Viewer - view st2.smcl
view st2.smcl X
+
Dialog ▾ Also see ▾ Jump to ▾
. . stcox age i.dose

Failure _d: died
Analysis time _t: studytime

Iteration 0: Log likelihood = -99.911448
Iteration 1: Log likelihood = -82.331523
Iteration 2: Log likelihood = -81.676487
Iteration 3: Log likelihood = -81.652584
Iteration 4: Log likelihood = -81.652567
Refining estimates:
Iteration 0: Log likelihood = -81.652567

Cox regression with Breslow method for ties

No. of subjects = 48          Number of obs = 48
No. of failures = 31
Time at risk = 744          LR chi2(3) = 36.52
Log likelihood = -81.652567  Prob > chi2 = 0.0000

+-----+-----+-----+-----+-----+
| _t | Haz. ratio | Std. err. | z | P>|z| | [95% conf. interval] |
+-----+-----+-----+-----+-----+
| age | 1.118334 | .0409074 | 3.06 | 0.002 | 1.040963 | 1.201455 |
+-----+-----+-----+-----+-----+
| dose |          |          |          |          |          |          |
| 5 mg | .1805839 | .0892742 | -3.46 | 0.001 | .0685292 | .4758636 |
| 10 mg | .0520066 | .034103 | -4.51 | 0.000 | .0143843 | .1880305 |
+-----+-----+-----+-----+-----+
CAP NUM INS
```

and test for violations of the proportional-hazards assumption.

```
Viewer - view st3.smcl
view st3.smcl X
+
Dialog ▾ Also see ▾ Jump to ▾
. . estat phtest, detail

Test of proportional-hazards assumption

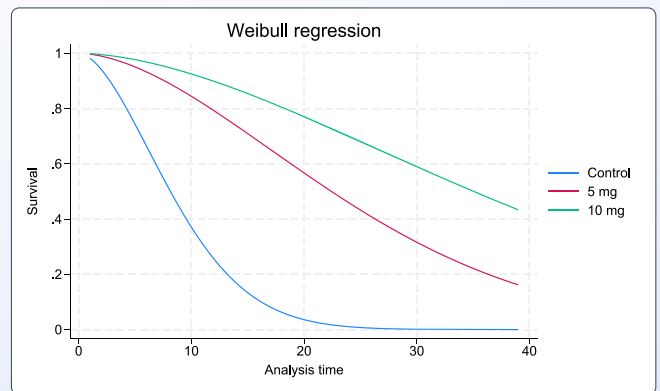
Time function: Analysis time

+-----+-----+-----+-----+
|          | rho | chi2 | df | Prob>chi2 |
+-----+-----+-----+-----+
| age | -0.06752 | 0.13 | 1 | 0.7199 |
| 1b.dose | . | . | 1 | . |
| 2.dose | 0.13698 | 0.50 | 1 | 0.4796 |
| 3.dose | -0.07380 | 0.16 | 1 | 0.6921 |
+-----+-----+-----+-----+
| Global test |          | 1.01 | 3 | 0.7994 |
+-----+-----+-----+-----+
CAP NUM INS
```

We can fit a Weibull model,

```
. streg age i.dose, distribution(weibull)
```

and plot the estimated survivor function for each dosage level,



or compute the marginal predictions of mean survival time for each dosage.

```
Viewer - view st4.smcl
view st4.smcl X
+
Dialog ▾ Also see ▾ Jump to ▾
. . margins dose, predict(mean time)

Predictive margins          Number of obs = 48
Model VCE: OIM

Expression: Predicted mean _t, predict(mean time)

+-----+-----+-----+-----+-----+
|          | Delta-method |          |          |          |          |
|          | Margin | std. err. | z | P>|z| | [95% conf. interval] |
+-----+-----+-----+-----+-----+
| dose |          |          |          |          |          |          |
| Control | 9.574003 | 1.336857 | 7.16 | 0.000 | 6.953812 | 12.19419 |
| 5 mg | 26.36447 | 6.992949 | 3.77 | 0.000 | 12.65854 | 40.0704 |
| 10 mg | 41.19108 | 10.31993 | 3.99 | 0.000 | 20.96438 | 61.41778 |
+-----+-----+-----+-----+-----+
CAP NUM INS
```

And that's just the beginning.