

Modeling long-term survival after surgery for esophageal cancer with *mlexp* command

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A patient's question

*“I got an esophagectomy a year ago.
What is my probability of surviving one more year?”*

A patient we met

The problem as a statistical model

We are interested in

$$f(y, d, x)$$

f : joint mixed density function

y : time to event

d : type of event (censored, cancer death, other deaths)

x : vector of patient's characteristics

$$f(y, d, x) = f_1(x)f_2(d|x)f_3(y|d, x)$$

The likelihood function

The likelihood function with an n -subject random sample is

$$l(\theta; y, d, x^T) = \prod_{i=1}^n f_1(x_i) f_2(d_i | x_i^T; \alpha^T) f_3(y_i | d_i, x_i^T; \beta^T)$$

$\theta = (\alpha^T, \beta^T)^T$ is the unknown parameter vector

The parameters in f_1 are of no inferential interest

The function f_2

The conditional probability of dying of a competing event is

$$f_2(d_i = 1 | x_i^T; \alpha^T) = 1 - \text{logit}^{-1}(x_i^T \alpha)$$
$$f_2(d_i = 2 | x_i^T; \alpha^T) = \text{logit}^{-1}(x_i^T \alpha)$$

The function f_3

The conditional density function of time to event is

$$f_3(y_i | d_i, x_i^T; \beta^T) = f_4(y_i | d_i, x_i^T; \beta^T)^{I(d_i \neq 0)} f_5(y_i | x_i^T; \beta^T)^{I(d_i = 0)}$$

The conditional probability of dying between y_i and $y_i + 1$ is

$$f_4(y_i | d_i, x_i^T; \beta^T) = S_6(y_i | d_i, x_i^T; \beta^T) - S_6(y_i + 1 | d_i, x_i^T; \beta^T)$$

S_6 is a parametric survival function

The conditional probability of being alive at y_i is

$$f_5(y_i | x_i^T; \theta^T) = \sum_{j=1}^2 S_6(y_i | d_i = j, x_i^T; \beta^T) f_2(d_i = j | x_i^T; \alpha^T)$$

The function S_6

We consider different functions

$$S_6(v_i) = 1 - [1 + \exp(-v_i)]^{-1} \quad \text{Loglogistic distribution}$$

$$S_6(v_i) = \exp[-\exp(v_i)] \quad \text{Weibull distribution}$$

$$S_6(v_i) = \exp[-\exp(v_i) + 1] \quad \text{Gompertz distribution}$$

$$\text{with } v_i = d_i c + x_i^T b + s_i^T e + d_i s_i^T q$$

s_i a vector of splines of $\log y_i$

We estimate a, b, c, e, q by maximizing the loglikelihood

The *mlexp* command

Maximum likelihood estimation of user-specified expressions

```
mlexp (lexp) [if] [in] [weight] [, options]
```

The loglikelihood function to maximize is

$$\log l(\theta^T) \propto \sum_{i=1}^n \log f_2(d_i | x_i^T; \alpha^T) + \log f_3(y_i | d_i, x_i^T; \beta^T)$$

Model without covariates

```
/* Log of time and time+1 */
gen y0 = log(y)
gen y1 = log(y+1/365.24)
gen cancer = death==2

/* Equation of parameters */
local a "{a}"
local v "{b}+{e}*y0+cancer*({c}+{q}*y0)"

/* Functions */
local Prob "invlogit(`a')"
local S60 "(1-invlogit(`v'))"
local S61 "`=substr(`S60',"y0',"y1",.)'"
local f5 "`=substr(`S61',"cancer',"1',.)')*`Prob'+
          (`=substr(`S61',"cancer',"0',.)')*(1-`Prob)'"
local f4 "`S60'-`S61'"
local log_f3 "(death!=0)*(log(`f4'))+(death==0)*log(`f5)'"
local log_f2 "(death==1)*log(1-`Prob')+(death==2)*log(`Prob)'"

mlexp (`log_f2'+`log_f3')
```

Model without covariates

Maximum likelihood estimation

Log likelihood = -5024.9791

Number of obs = 1,027

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
/a	.4837124	.1215812	3.98	0.000	.2454176	.7220073
/b	-2.515504	.1799259	-13.98	0.000	-2.868152	-2.162856
/e	.7411835	.0833531	8.89	0.000	.5778145	.9045525
/c	2.409246	.2436196	9.89	0.000	1.931761	2.886732
/q	.391697	.1070756	3.66	0.000	.1818327	.6015613

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The probability of dying of cancer is

$$f_2(d_i = 2 | x_i^T; \alpha^T) = [1 + \exp(-a)]^{-1}$$

```
di %3.2f 1/(1+exp(-_b[/a])) " ("
%3.2f 1/(1+exp(-_b[/a]+invnormal(.975)*_se[/a])) "; "
%3.2f 1/(1+exp(-_b[/a]-invnormal(.975)*_se[/a])) ") "
```

0.62 (0.56; 0.67)

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The parametric survival function is

$$S_6(y_i) = 1 - \left[1 + \exp \left(- (d_i c + x_i^T b + s_i^T e + d_i s_i^T q) \right) \right]^{-1}$$

Goodness of fit

Kaplan-Meier curve versus survival function f_5 from the model

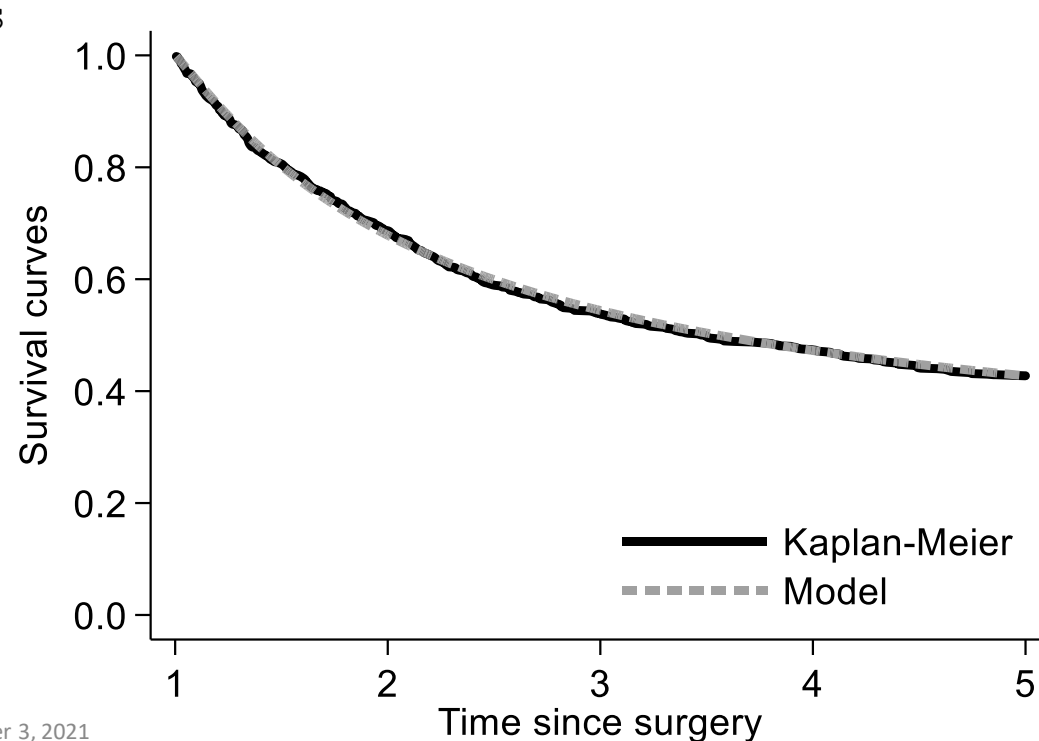
```
display "f5 = `f5'"
```

```
f5 = (1-invlogit({b}+{e}*y1+1*({c}+{q}*y1)))*invlogit({a})+  
      (1-invlogit({b}+{e}*y1+0*({c}+{q}*y1))*(1-invlogit({a}))
```

```
gen f5 = `=subinstr(subinstr(subinstr("`f5'", "{", "_b[/", ".), "}"  
      , "]" , .), "y1", "y0", .)'
```

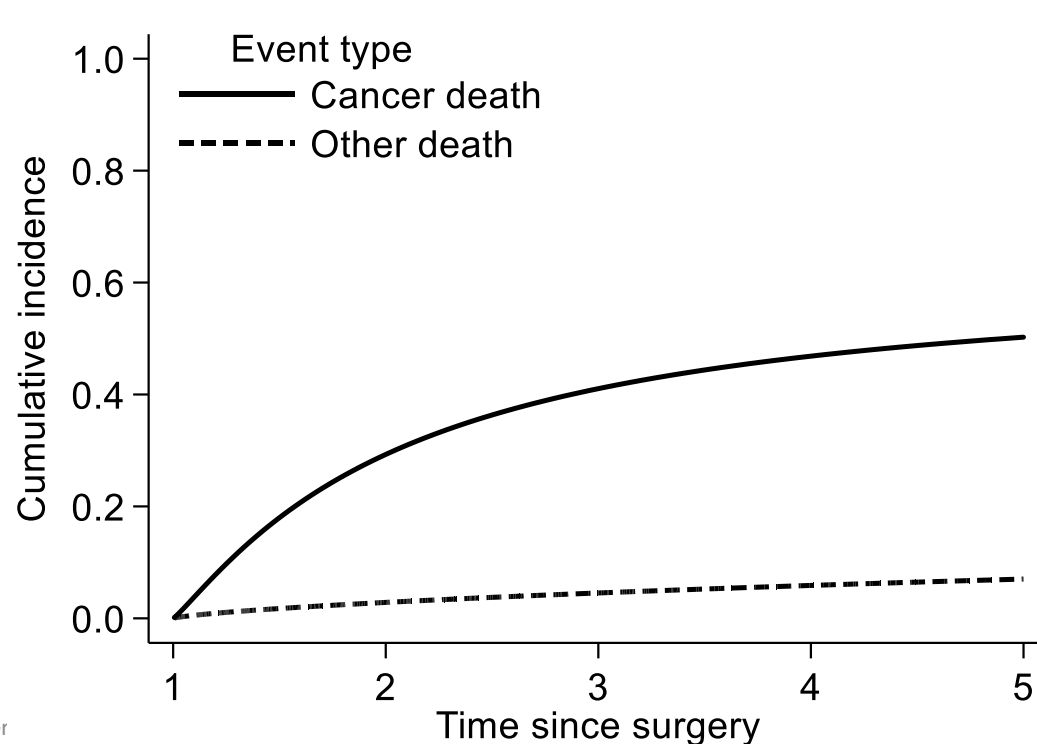
```
stset y, fail(death)
```

```
sts gen KM = s
```



Cumulative incidence functions

```
gen S6_cancer = `=subinstr(subinstr(subinstr("`S60'", "{", "_b[/", .)
    , "}", "]", .), "cancer", "1", .) '
gen S6_other = `=subinstr(subinstr(subinstr("`S60'", "{", "_b[/", .)
    , "}", "]", .), "cancer", "0", .) '
gen prob = `=subinstr(subinstr("`Prob'", "{", "_b[/", .), "}", "]", .) '
gen cif_cancer = (1-S6_cancer)*prob
gen cif_other = (1-S6_other)*(1-prob)
```



Model with covariates

```
local a "({a0}+{a1}*age+{a3}*resection+{a4}*stage2+
         {a5}*stage3+{a6}*stage4+{a7}*histology+{a9}*neoadjuvant) "
local b "({b0}+{b1}*age+{b2}*sex+{b3}*resection+{b5}*stage3+
         {b6}*stage4+{b7}*histology+{b8}*reoperation+{b11}*ed3) "
local v "`b'+{e0}*y0+cancer*({c0}+{q0}*y0) "

[... code omitted ...]
```

Model with covariates

Maximum likelihood estimation

Log likelihood = -4904.6917

Number of obs = 1,027

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
/a0	-1.262621	.2091682	-6.04	0.000	-1.672584	-.8526593
/a1	.2849556	.1065176	2.68	0.007	.076185	.4937263
/a3	1.623797	.5737225	2.83	0.005	.4993213	2.748272
/a4	1.569068	.21875	7.17	0.000	1.140326	1.99781
/a5	2.249153	.2830246	7.95	0.000	1.694435	2.803871
/a6	2.365552	.5775042	4.10	0.000	1.233665	3.497439
/a7	.5801036	.1954006	2.97	0.003	.1971255	.9630818
/a9	.4518313	.2148914	2.10	0.036	.030652	.8730106
/b0	-2.671793	.1901998	-14.05	0.000	-3.044577	-2.299008
/b1	.1492687	.0757875	1.97	0.049	.000728	.2978095
/b2	-.291121	.1557184	-1.87	0.062	-.5963235	.0140815
/b3	.6177573	.2126261	2.91	0.004	.2010178	1.034497
/b5	.5575737	.1523027	3.66	0.000	.2590658	.8560816
/b6	.9783877	.2963993	3.30	0.001	.3974556	1.55932
/b7	.2561525	.1421352	1.80	0.072	-.0224274	.5347323
/b8	.4617539	.2192609	2.11	0.035	.0320103	.8914974
/b11	-.3840374	.1957407	-1.96	0.050	-.7676821	-.0003928
/e0	.75694	.0846327	8.94	0.000	.5910629	.9228171
/c0	2.197287	.2313714	9.50	0.000	1.743807	2.650766
/q0	.43379	.1034186	4.19	0.000	.2310933	.6364866

Model with covariates

Maximum likelihood estimation

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The probability of dying of cancer is

$$f_2(d_i = 2 | x_i^T; \alpha^T) = [1 + \exp(-x_i^T a)]^{-1}$$

Model with covariates

Maximum likelihood estimation

Log likelihood = -4904.6917

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The parametric survival function is

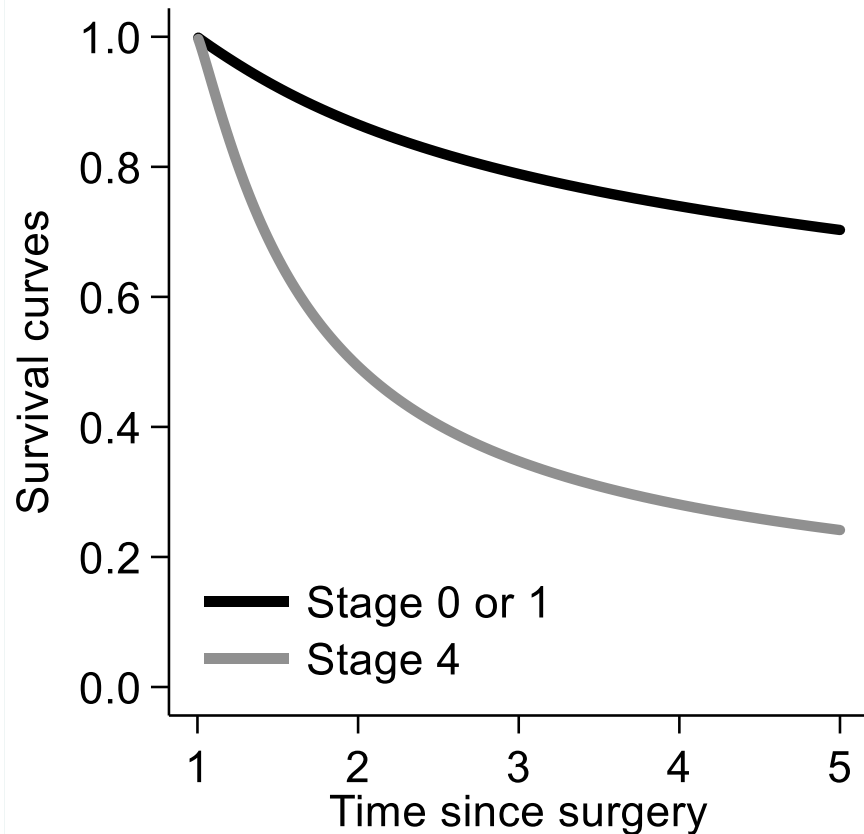
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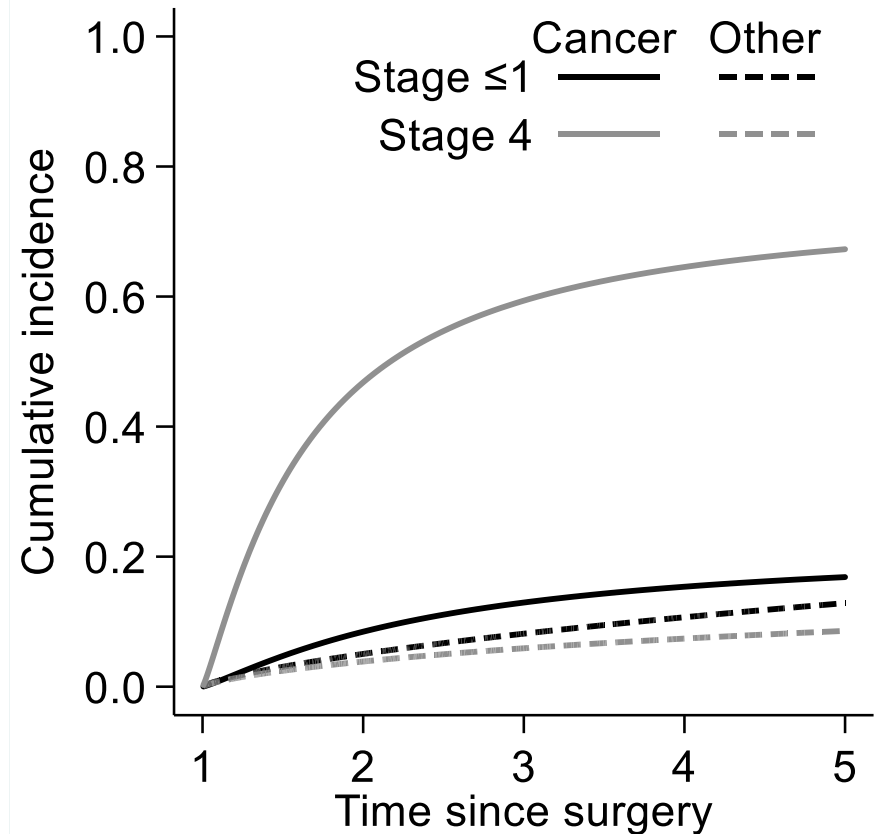
Individual predictions

A 65-year old male, adenocarcinoma, no reoperation, no neoadjuvant treatment, R0 resection, ≤ 12 years of education

Patient 1: tumor stage 1 or less



Patient 2: tumor stage 4



Individual predictions

A 65-year old male, adenocarcinoma, no reoperation, no neoadjuvant treatment, R0 resection, ≤ 12 years of education

Patient 1: tumor stage ≤ 1

Patient survived 1.25 years after esophageal surgery.

Probability of dying within the next 1.5 years

Any cause: 23.5%

Cancer: 17.1%

Years since surgery (≥ 1):

Years to prediction:

Age (18-90):

Sex: Man Woman

Years of Education: ≤ 12 >12

Tumor stage (0-4): 0 or 1 2 3 4

Resection margin: R0 R1-R2

Tumor histology: Squamous cell carcinoma Adenocarcinoma

Neoadjuvant therapy: No Yes

30-day reoperation: No Yes

Patient 2: tumor stage 4

Patient survived 1.25 years after esophageal surgery.

Probability of dying within the next 1.5 years

Any cause: 62.5%

Cancer: 58.5%

Years since surgery (≥ 1):

Years to prediction:

Age (18-90):

Sex: Man Woman

Years of Education: ≤ 12 >12

Tumor stage (0-4): 0 or 1 2 3 4

Resection margin: R0 R1-R2

Tumor histology: Squamous cell carcinoma Adenocarcinoma

Neoadjuvant therapy: No Yes

30-day reoperation: No Yes

Conclusions

Modeling requires understanding the problem at hand
mlexp is a flexible and computationally fast command
Few programming lines can code complex models