

# Example of cure-regression models - on lognormal

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## Abstract

The Gamel-Boag model is the mixture model fit with a lognormal failure density and a logistic cure-fraction link. This model seems to be the one used by Schmidt and Witte (1987). The purpose of this report is to show that the form of the lognormal of Spusto (2002) is the same as that of Schmidt and Witte (1987). The scale and shape parameters are the negative of each other.

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# 1 split-population description

A spilt-population model. Survival,  $S(t)$  where  $g(t)$  is time,  $t$ , itself or  $\tilde{t}$ . Please note that  $\tilde{t}$  is the scaled and shaped time:  $\tilde{t} = (\lambda t)^\gamma$  where the scale and shape are, respectively,  $\lambda = \exp(x_\lambda \beta_\lambda)$  and  $\gamma = \exp(x_\gamma \beta_\gamma)$ .  $\hat{\mu}_t$  &  $\hat{\sigma}$  below, respectively, refer to the mean time to fail in units of time,  $t$ , and the standard deviation or shape,  $(\gamma)$ . These apply to the two probabilistically equal alternative lognormal failure densities, equations 2 and 3.

## 1.1 mixture model

$$S(t) = \pi + (1 - \pi) (1 - F(g(t))) \quad (1)$$

## 1.2 fail density: lognormal, form Sposto

$$\hat{\mu}_t = \exp(-x_\lambda \beta_\lambda) \text{ & } \hat{\sigma} = \exp(-x_\gamma \beta_\gamma)$$

$$F(\tilde{t}) = \int_{-\infty}^{\ln(\tilde{t})} \frac{1}{\sqrt{2\pi}} e^{(-x^2/2)} dx \quad (2)$$

This form (Sposto, 2002) is really the same as the standard lognormal.

## 1.3 fail density: lognormal

$$\hat{\mu}_t = \exp(x_\lambda \beta_\lambda) \text{ & } \hat{\sigma} = \exp(x_\gamma \beta_\gamma)$$

$$F(t) = \int_{-\infty}^{\ln(t)} \frac{1}{(\gamma) \sqrt{2\pi}} e^{(-(x-(x_\lambda \beta_\lambda))^2/2\gamma^2)} dx \quad (3)$$

The standard lognormal fail density and is what is fit in Schmidt and Witte (1987) and also, for example, in Frankel and Longmate (2002) except that this form replaces the scale invariant  $\sigma$  with shape,  $(\gamma)$ .

## 1.4 fail density: exponential

$$F(t) = 1 - \exp\left(-\frac{t}{\lambda}\right) \quad (4)$$

## 1.5 cure-fraction link function:

logistic

$$\pi = \frac{\exp(x_\pi \beta_\pi)}{1 + \exp(x_\pi \beta_\pi)} \quad (5)$$

## 2 example: Schmidt & Witte model

### 2.1 sample data description

data description source at Stata Corp. Wooldridge et al. (1998) (does not include the gender (male) factor).

factor	description
tserved1	time served rounded to months (x100) median .12 (range: 0, 2.19)
age1	age in months (x1000) median .307 (range: .198, .933)
priors1	number of previous convictions (x10) median 0 (range: 0, 2.8)
white	=1 if not black (51%) n=744
felon	=1 if felony sentence (31%) n=454
alcohol	=1 if alcohol problems (21%) n=303
drugs	=1 if drug history (24%) n=349
property	=1 if property crime (25%) n=368

Table 1: factors

### 2.2 output

model featuring the fail density of equation 2 (cureregr Buxton (2013a)).

```
. cureregr tserved1 age1 priors1 white felon alcohol drugs property , dist(logn
> ormal) cl(mix) link(logist) scale() shape() nolog noshow
cf: logistic, kn: lognormal, model: mixture
cf_initial_coef: 0.4700 pi: 0.6154
No. of subjects = 1445
Number of obs      =      1445
LR chi2(8)        =     150.71
Prob > chi2       =     0.0000
Log likelihood = -3137.7484
```

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
cure_frac					
tserved1	-2.880944	.7033178	-4.10	0.000	-4.259422 -1.502467
age1	4.299798	.7287908	5.90	0.000	2.871395 5.728202
priors1	-2.011418	.4729662	-4.25	0.000	-2.938414 -1.084421
white	.6758835	.1436679	4.70	0.000	.3942995 .9574675
felon	1.00679	.2676913	3.76	0.000	.4821247 1.531455
alcohol	-.6741825	.1896951	-3.55	0.000	-1.045978 -.302387
drugs	-.4165466	.1662791	-2.51	0.012	-.7424477 -.0906455
property	-.5894715	.2427129	-2.43	0.015	-1.06518 -.113763
_cons	-.7904601	.2700485	-2.93	0.003	-1.319745 -.2611748
scale					
_cons	-3.211344	.0949789	-33.81	0.000	-3.397499 -3.025189
shape					
_cons	-.1818824	.0529575	-3.43	0.001	-.2856771 -.0780877

model featuring the fail density of equation 3 (cureregr1 Buxton (2013b)).

```
. cureregr1 tserved1 age1 priors1 white felon alcohol drugs property , dist(nor
> mallnt) cl(mix) link(logist) scale() shape() nolog noshow auxuse cformat(%6.4
> f) sformat(%3.2f) pformat(%5.4f)
cf: logistic, kn: lognormal, model: mixture
cf_initial_coef: 0.4700 pi: 0.6154
No. of subjects = 1445
Number of obs      =      1445
LR chi2(8)        =     150.71
Prob > chi2       =     0.0000
Log likelihood = -3137.7484
```

	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
cure_frac					
tserved1	-2.8809	0.7033	-4.10	0.0000	-4.2594    -1.5025
age1	4.2998	0.7288	5.90	0.0000	2.8714    5.7282
priors1	-2.0114	0.4730	-4.25	0.0000	-2.9384    -1.0844
white	0.6759	0.1437	4.70	0.0000	0.3943    0.9575
felon	1.0068	0.2677	3.76	0.0002	0.4821    1.5315
alcohol	-0.6742	0.1897	-3.55	0.0004	-1.0460    -0.3024
drugs	-0.4165	0.1663	-2.51	0.0122	-0.7424    -0.0906
property	-0.5895	0.2427	-2.43	0.0152	-1.0652    -0.1138
_cons	-0.7905	0.2700	-2.93	0.0034	-1.3197    -0.2612
mean_ln					
_cons	3.2113	0.0950	33.81	0.0000	3.0252    3.3975
shape					
_cons	0.1819	0.0530	3.43	0.0006	0.0781    0.2857
mean_t	24.8124	2.3567			20.5979    29.8893
stdev	1.1995	0.0635			1.0812    1.3307

Alas, the ancillary parameter estimates *mean\_t* and *stdev* only are displayed for the constant parameter even if more than one factor is included the respective *mean\_ln* and *shape* models.

### 2.3 model results

These results in tables 2 and 3 can be compared to those in Schmidt and Witte (1987). The results compare closely even though the factor, male, is not available as it is in Schmidt & Witte.

cox PH <sup>a</sup>			
	b	sig <sup>b</sup>	z
main			
tserved1	1.3650	***	8.10
age1	-3.4386	***	-6.88
priors1	0.8882	***	6.63
white	-0.4415	***	-5.01
felon	-0.5735	***	-3.97
alcohol	0.4280	***	4.08
drugs	0.2997	**	3.06
property	0.4190	**	3.03
lnL	-3813.672		

<sup>a</sup>re: proportional hazards model, *final specification*, in table 1 of Schmidt & Witte.

<sup>b</sup>\* < 0.05, \*\* < 0.01, \*\*\* < 0.001

Table 2: cox PH

factor	split lognormal model <b>t2a<sup>a</sup></b>			logit lognormal model <b>t2b<sup>b</sup></b>			logit/individual lognormal <b>t3<sup>c</sup></b>			logit/individual exponential <b>t4<sup>d</sup></b>		
	b	sig <sup>e</sup>	z	b	sig	z	b	sig	z	b	sig	z
<b>cure_frac</b>												
tserved1				-2.8809	***	-4.10	-1.5831	***	-3.80	-1.2902	**	-3.23
age1				4.2998	***	5.90	3.7469	***	4.99	3.8249	***	4.58
priors1				-2.0114	***	-4.25	-1.1582	***	-3.65	-2.0488	***	-3.88
white				0.6759	***	4.70	0.6695	***	4.32	0.7796	***	4.70
felon				1.0068	***	3.76	0.4735		1.67	0.3414		1.16
alcohol				-0.6742	***	-3.55	-0.4832	**	-2.59	-0.5197	**	-2.61
drugs				-0.4165	*	-2.51	-0.4226	*	-2.40	-0.3479	*	-2.02
property				-0.5895	*	-2.43	-0.2338		-0.92	-0.1196		-0.48
_cons	-0.8723	**	-3.09	-0.7905	**	-2.93	-0.8773	**	-3.14	-0.8387	**	-2.88
<b>mean_ln</b>												
tserved1	-1.9602	***	-5.98				-1.1552	***	-3.85			
age1	3.4541	***	6.07				1.1795		1.95			
priors1	-1.4350	***	-6.07				-0.6491	**	-2.87			
white	0.4729	***	3.92				-0.0072		-0.06			
felon	0.9364	***	4.75				0.6707	**	2.88			
alcohol	-0.6339	***	-4.31				-0.3029	*	-2.04			
drugs	-0.2957	*	-2.20				-0.0070		-0.05			
property	-0.6869	***	-3.69				-0.5596	**	-2.86			
_cons	3.3187	***	14.08	3.2113	***	33.81	3.1750	***	14.48			
<b>mean</b>	27.62			24.81			23.93					
<b>shape</b>												
_cons	0.3853	***	6.38	0.1819	***	3.43	0.1118	*	2.21			
<b>stdev</b>	1.47			1.20			1.12					
<b>scale</b>												
tserved1										-1.4549	***	-5.69
age1										1.2943		1.63
priors1										0.0662		0.28
white										-0.1685		-1.08
felon										0.7858	**	3.21
alcohol										-0.2227		-1.26
drugs										-0.1239		-0.80
property										-0.6842	***	-3.62
_cons										3.5309	***	12.74
<b>lnL</b>	-3145.759			-3137.748			-3121.393			-3138.226		

<sup>a</sup>re: split lognormal model in table 2 of Schmidt & Witte.

<sup>b</sup>re: logit lognormal model in table 2.

<sup>c</sup>re: logit/individual lognormal model in table 3 of Schmidt & Witte.

<sup>d</sup>re: logit/individual exponential model in table 3.

\* < 0.05, \*\* < 0.01, \*\*\* < 0.001

Table 3: split population models

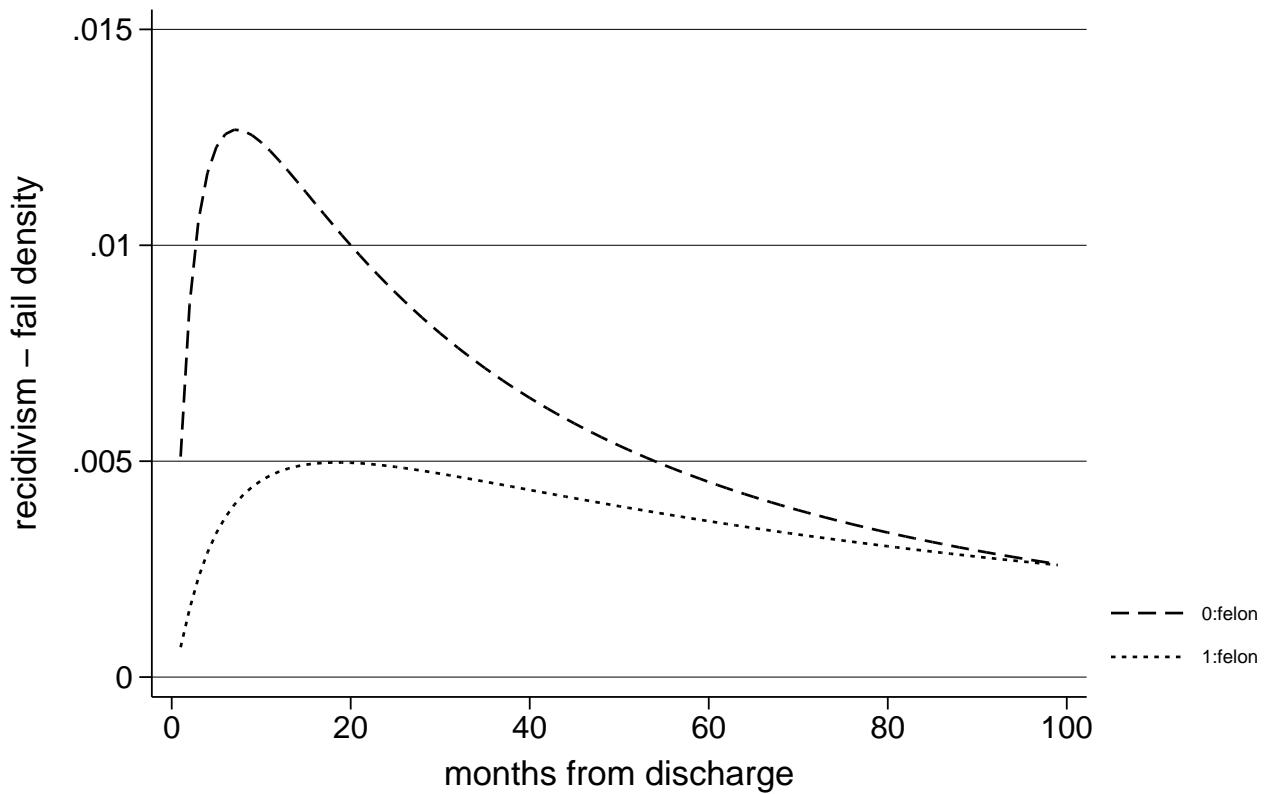


Figure 1: fail density split lognormal model

at median tserved1, age1, & priors1; alcohol, drugs, property, & white are ==0 except felon as noted. The average time to recidivism, among those reincarcerated, is about 25 months.

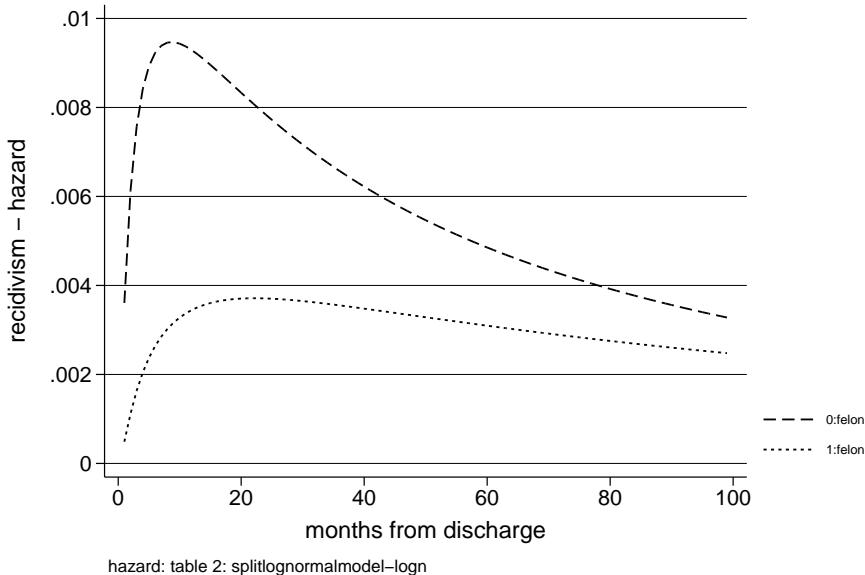


Figure 2: hazard split lognormal model  
at median tserved1, age1, & priors1; alcohol, drugs, property, & white are ==0 except felon as noted.

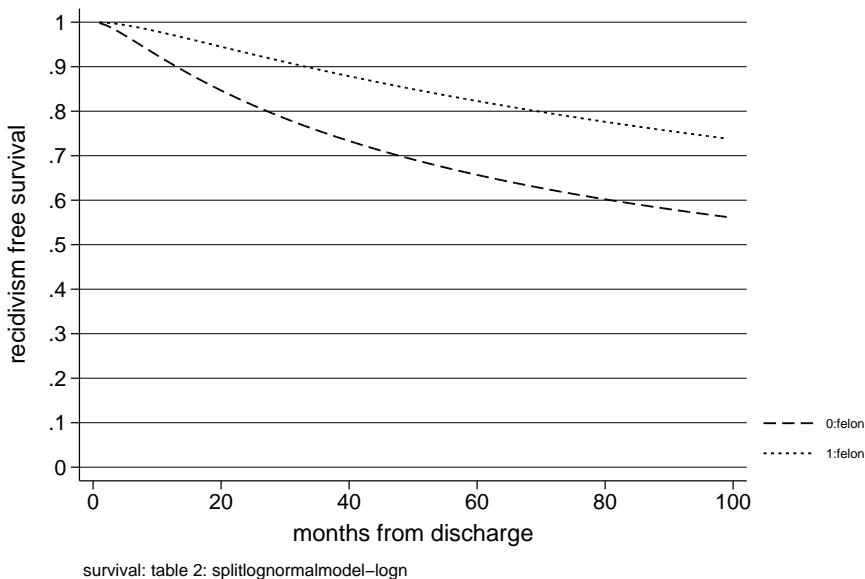


Figure 3: survival split lognormal model  
at median tserved1, age1, & priors1; alcohol, drugs, property, & white are ==0 except felon as noted.

## References

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