

Seminar at 國立清華大學統計所

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# Semiparametric inference for an accelerated failure time model with dependent truncation

Emura T\* & Wang W (2015), Semiparametric inference for an accelerated failure time model with dependent truncation, *Annals of the Institute of Statistical Mathematics*, DOI: 10.1007/s10463-015-0526-

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# Progress of the paper

- 2009-2010: Paper developed as my post-doc project under Prof. Weijing Wang
- **2011-2013:**  
Presented at 第20回南區統計研會, 國立中正大學  
Rejected by JRSSB, Biometrika, Sinica  
*Journal of Multivariate Analysis*  
(all do not recognize our idea of a new model)  
(I myself feel difficult to interpret the new method)
- Submit on 2013/10 → Now : the 2<sup>nd</sup> major revision at  
*Annals of the Institute of Statistical Mathematics (AISM)*

# Outlines

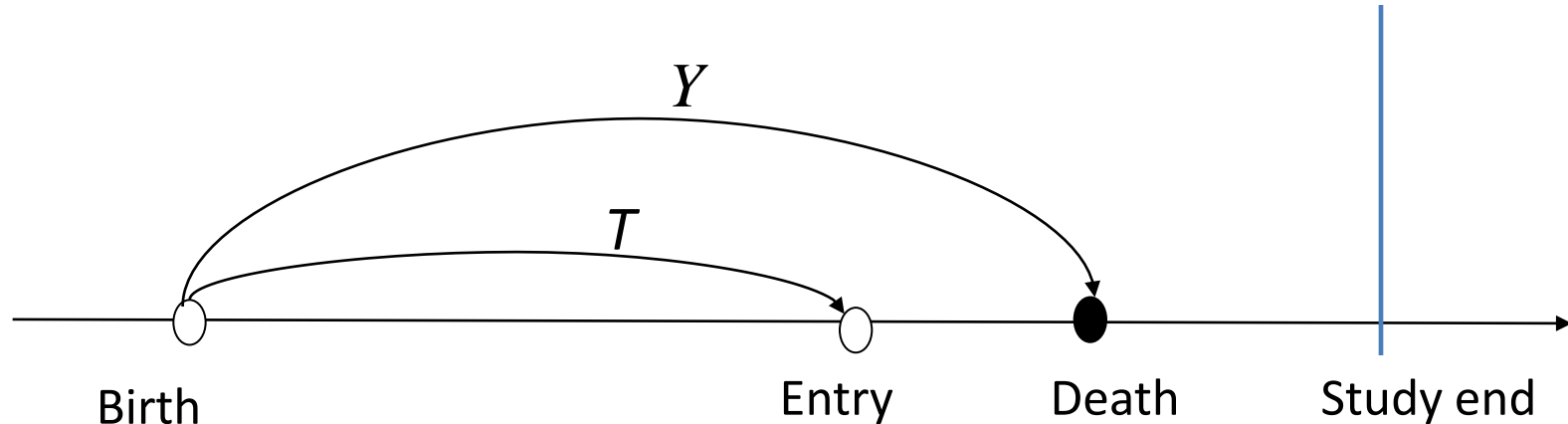
## Part I: Review

- Truncated data - Channing house data -
- Regression method - AFT model -

## Part II: Proposed method

- Proposed method
- Estimation procedure
- Asymptotic theory
- Simulation and data analysis
- Conclusion

# Left-truncated survival data



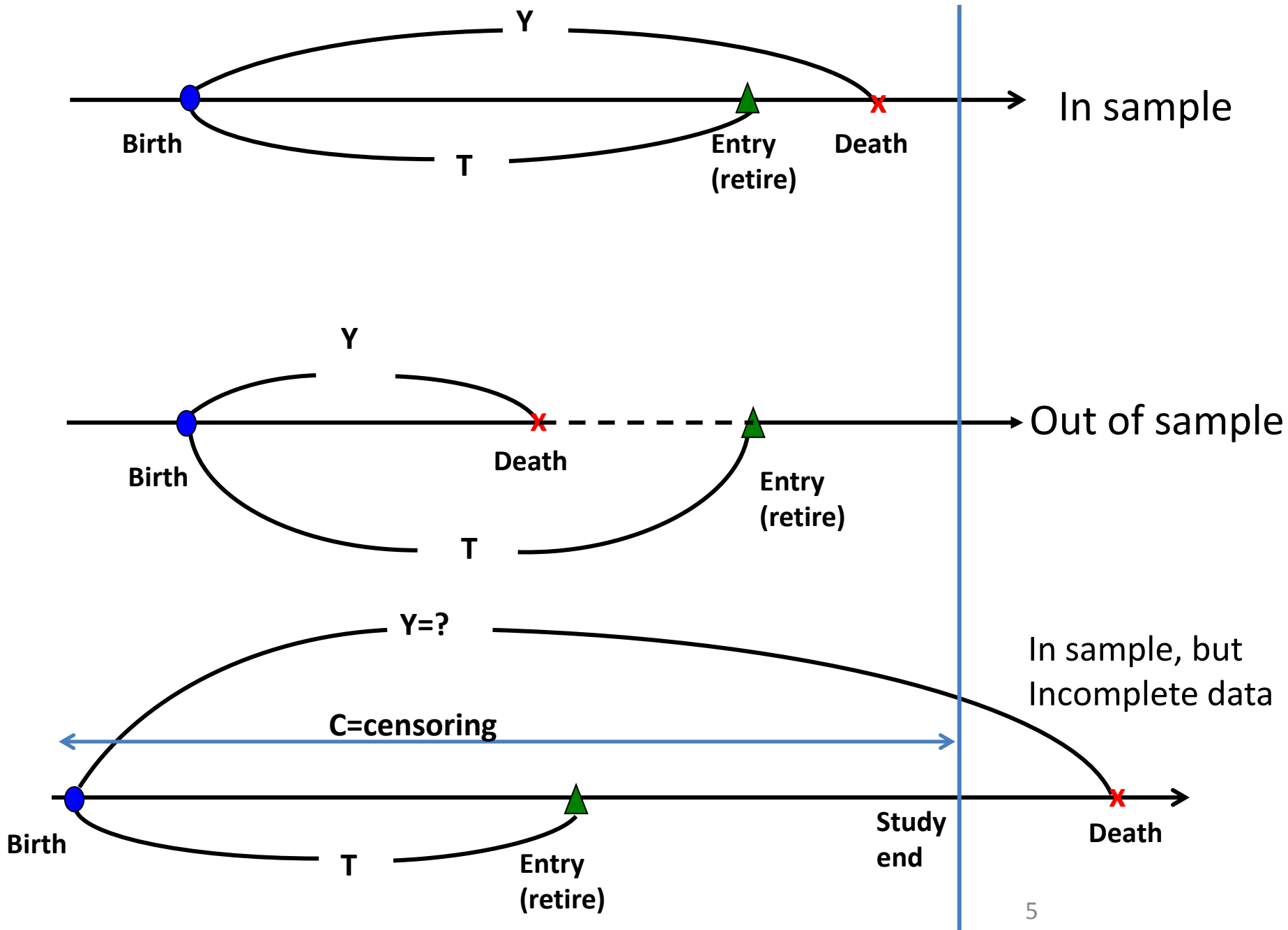
## Channing House data (Hyde, 1980)

Channing house is a retirement center in California

- 462 elderly residents (97 men + 365 women)
- Age at entry =  $T$
- Age at death =  $Y$  ( possibly right-censored )

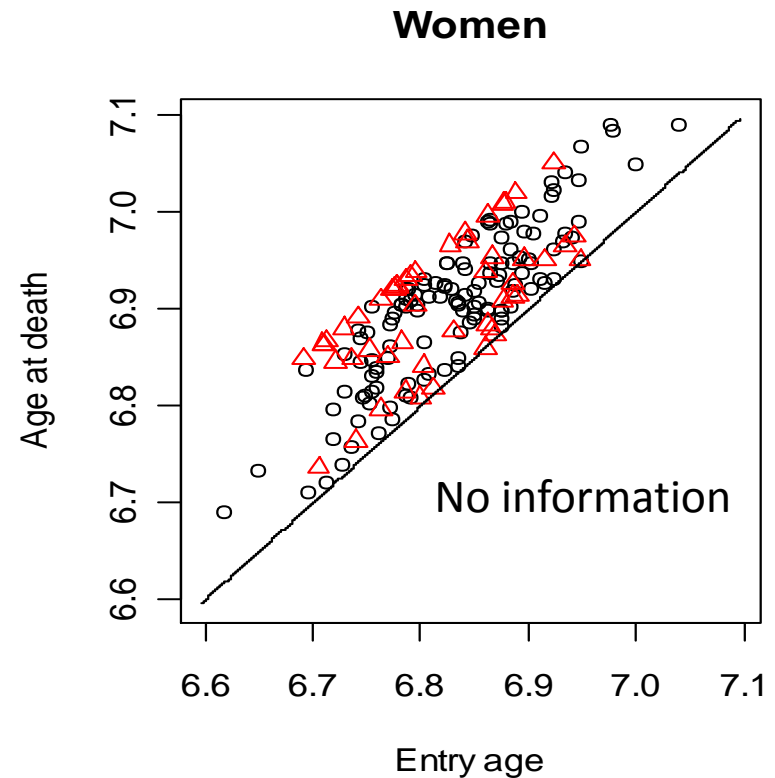
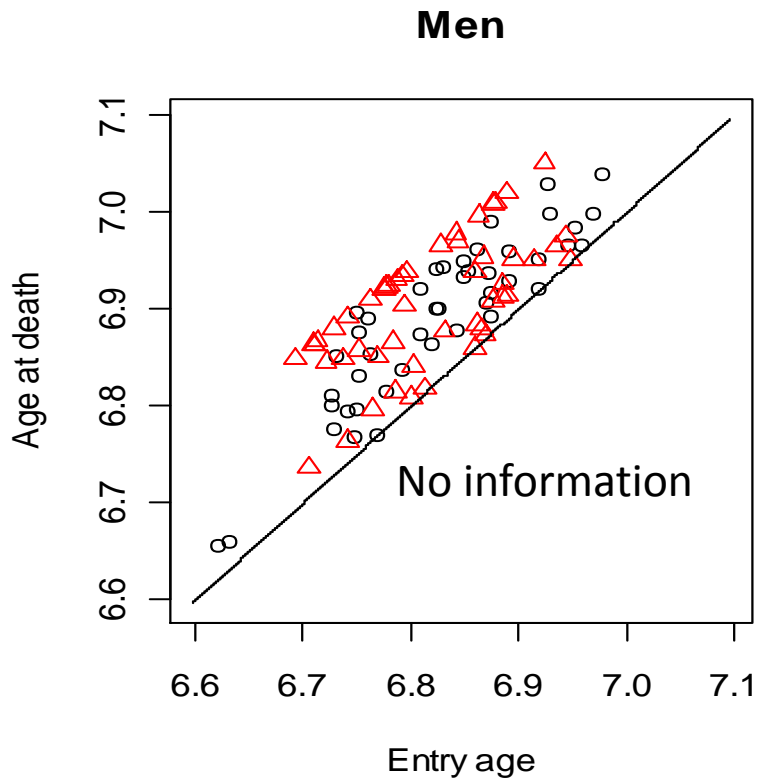
## Left-truncation criterion:

$$T \leq Y \quad \Rightarrow \quad \text{Sample is available}$$



# Left-truncation data

- Longer survivors have higher chance to be sampled
  - ➔ Observed lifetimes are longer than the random sampling from **the target population**
  - ➔ Ignoring left-truncation produces bias (often very serious bias)
- Many people ignore truncation or do not realize
  - The famous textbook for survival analysis Klein & Moeschberger (2003)
  - Answer to Exercise 3.7:** <--ignore left-truncation
- Truncation usually occurs when
  1. scale of survival is “**age**” (birth <---> death).
  2. the target is “**age** at disease”.



▲: Censored individual  
 ○: Died individual

- Hyde (1980) assumed:  
*knowing the person's entry age will provide no additional information about prospects for survival*
- That is,  $T \perp Y \mid \text{gender}$   
 (  $T$ : Age at entry;  $Y$ : Age at death )

- Existing methods for left-truncated and right-censored data

➔ Rely on *independent truncation assumption*

Hyde (1977, 1980)

➔ One-sample log-rank test

Wang, Jewell & Tsai (1986)

➔ Product-limit estimator

Lai and Ying (1991):

➔ Rank regression

He and Yang (1998):

➔ Estimating truncation probability

Su and Wang (2012):

➔ Joint models with longitudinal covariates



# Truncation data with covariates

## Left-truncated and right-censored data:

- $Y^*$  : Log-lifetime
- $T$  : Log-truncation time
- $C$  : Log-censoring time
- $\mathbf{X}$  :  $p$ -dimensional covariate

## Left-truncation:

A pair  $(T, Y, \Delta, \mathbf{X})$  is observed only when  $T \leq Y^*$ ,  
, where  $Y = Y^* \wedge C$ ,  $\Delta = I(Y^* \leq C)$

\*If  $T = -\infty$ , this is usual right-censored data with covariates  
→ fit Cox regression (1972) or AFT regression (Tsiatis 1990)

# Truncation data with covariates

## Observed data:

$\{(T_i, Y_i, \Delta_i, \mathbf{X}_i); (i = 1, \dots, n)\}$  subject to  $T_i \leq Y_i$

## AFT regression (Lai and Ying, 1991 AS)

Semiparametric accelerated failure time (AFT) model:

$$Y^* = \boldsymbol{\beta}'_0 \mathbf{X} + \varepsilon,$$

\*  $Y^*$  = Log - lifetime

\*  $\varepsilon$  = Error ( p.d.f of  $\varepsilon$  is unspecified )

\*  $\boldsymbol{\beta} \in R^p$

•  $X = 1$  (male),  $X = 0$  (female)

→  $\beta_0$  : The gender difference of mean lifetime  
in the scale of  $\log(\text{lifetime})$

# Lai & Ying's AFT regression

**Log-rank type estimating equation:**  $E_{\beta} \{ \mathbf{X}_i - E_{\beta} \mathbf{X}_i \} \approx 0$

$$\mathbf{U}_n(\boldsymbol{\beta}) = \sum_{i=1}^n \Delta_i \phi_i(\boldsymbol{\beta}) \left\{ \mathbf{X}_i - \frac{1}{R_i(\boldsymbol{\beta})} \sum_j \mathbf{X}_j I(e_j^T(\boldsymbol{\beta}) \leq e_i^Y(\boldsymbol{\beta}) \leq e_j^Y(\boldsymbol{\beta})) \right\},$$

where

$$e_i^T(\boldsymbol{\beta}) = T_i - \boldsymbol{\beta}'\mathbf{X}_i, \quad e_i^Y(\boldsymbol{\beta}) = Y_i - \boldsymbol{\beta}'\mathbf{X}_i: \quad \text{Residual}$$

$$R_i(\boldsymbol{\beta}) = \sum_j I(e_j^T(\boldsymbol{\beta}) \leq e_i^Y(\boldsymbol{\beta}) \leq e_j^Y(\boldsymbol{\beta})): \quad \text{Number at - risk}$$

**U-statistic form under the Gehan weight**  $\phi_i(\boldsymbol{\beta}) = R_i(\boldsymbol{\beta})$

$$\mathbf{U}_n^G(\boldsymbol{\beta}) = \sum_{i=1}^n \sum_{j=1}^n \Delta_i (\mathbf{X}_i - \mathbf{X}_j) I\{ e_j^T(\boldsymbol{\beta}) \leq e_i^Y(\boldsymbol{\beta}) \leq e_j^Y(\boldsymbol{\beta}) \}$$

$$\hat{\boldsymbol{\beta}}: \quad \mathbf{U}_n^G(\boldsymbol{\beta}) = \mathbf{0} \quad (\text{Lai \& Ying (1991) estimator})$$

# Lai & Ying's AFT regression

## Assumptions of Lai & Ying (1991):

$$\begin{cases} Y^* = \boldsymbol{\beta}'_0 \mathbf{X} + \varepsilon \\ (T, C, \mathbf{X}) \perp \varepsilon \quad \dots (A) \end{cases} \quad \leftarrow \text{Interpreted as "Independent truncation"}$$

## Why (A) is independent truncation?

By (A),  $Y^* - \boldsymbol{\beta}'_0 \mathbf{X} \perp T$ .

After adjusting for the effect of  $\mathbf{X}$ , the truncation variable  $T$  contains no information on survival  $Y^*$

**Motivating Example:** This model satisfy (A) only under  $\rho = 0$

$$\begin{bmatrix} Y^* \\ T \end{bmatrix} \sim N \left( \begin{bmatrix} \boldsymbol{\beta}'_0 \mathbf{X} \\ 0 \end{bmatrix}, \begin{bmatrix} 1 & \rho \\ \rho & 1 \end{bmatrix} \right), \quad C \sim N(0, 1)$$

# Independent truncation hold?

Testing quasi-independence

$$H_0 : \Pr(L = l, Y = y | L \leq Y) \propto dF_L(l)dF_Y(y)$$

**Available test statistics:**

1. [Chen et al. \(1996 JASA\)](#)

- Based on the conditional Pearson-correlation

2. [Tsai \(1990 Biometrika\)](#); [Martin & Betensky \(2005 JASA\)](#)

-Based on the conditional Kendall's tau

3. [Emura & Wang \(2010 JMVA\)](#) - Based on weighted-logrank test

(Optimal weight choice)

# Testing quasi-independence

$$H_0 : \Pr(L = l, Y = y | L \leq Y) \propto dF_L(l)dF_Y(y)$$

is rejected at 5 % level

*Table 4 of Emura and Wang (2010, JMVA).*

Tests of quasi-independence for the Channing House data.

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	Logrank test	Tsai test	Marting & Betensky test
	$L_{\rho=1}$		
P-value	0.048	0.043	0.040

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- Quasi-independence is questionable
  - Lai & Ying's AFT regression estimate may be biased (later shown via simulations)
- In Channing house data, the truncation (entry age) may be dependent on survival.

→ Motivate modeling for dependent truncation

Chaieb, Rivest, Abdous (2006 *Biometrika*)

Beaudoin and Lakhel-Chaieb (2008 *Stat. Med.*),

Emura and Wang (2010 *JMVA*)

Emura and Konno (2010 *Stat Papers*; 2012 *CSDA*)

Emura, Wang and Hung (2011 *Sinica*)

Emura and Wang (2012 *JMVA*)

Ding (2012 *Lifetime Data Analysis*)

Emura and Murotani (2014 *Test*, in revision)

(All the work is for iid case !)

## Part II: Proposed method



# Proposed method

**Proposed model (Semi-par AFT with dependent truncation):**

$$\begin{cases} Y^* = \boldsymbol{\beta}'_0 \mathbf{X} + \gamma_0 T + \varepsilon \\ (T, C, \mathbf{X}) \perp \varepsilon \end{cases} \quad \dots \quad (\text{B})$$

NOTE: Special case of  $\gamma_0 = 0$  ,  $\rightarrow$  Lai & Ying's AFT model

**Example: Bivariate normal model**

$$\begin{bmatrix} Y^* \\ T \end{bmatrix} \sim N \left( \begin{bmatrix} \boldsymbol{\beta}'_0 \mathbf{X} \\ \mu_L \end{bmatrix}, \begin{bmatrix} \sigma_{Y^*}^2 & \rho \sigma_{Y^*} \sigma_T \\ \rho \sigma_{Y^*} \sigma_T & \sigma_T^2 \end{bmatrix} \right), \quad C \sim N(1, 1)$$

$$\Rightarrow Y^* | \mathbf{X}, T \sim N \left( \boldsymbol{\beta}'_0 \mathbf{X} + \rho \frac{\sigma_{Y^*}}{\sigma_T} (T - \mu_L), \sigma_{Y^*}^2 (1 - \rho^2) \right)$$

This model satisfy (B) with

$$\gamma = \rho \frac{\sigma_{Y^*}}{\sigma_T}, \quad \varepsilon \sim N \left( -\rho \frac{\sigma_{Y^*}}{\sigma_T} \mu_L, \sigma_{Y^*}^2 (1 - \rho^2) \right)$$

# Proposed method

## Interpretation of our model:

$$Y^* = \boldsymbol{\beta}'_0 \mathbf{X} + \gamma_0 T + \varepsilon, \quad \varepsilon \sim f_\varepsilon$$

NOTE:

If  $\gamma_0 \neq 0$  then the Lai & Ying model do not hold in general

$$Y^* \neq \boldsymbol{\beta}'_0 \mathbf{X} + \varepsilon_0,$$

Indeed, the nonlinear model

$$f_{Y^*}(y | \mathbf{X}) = \int f_{Y^*}(y | t, \mathbf{X}) f_T(t | \mathbf{X}) dt = \int f_\varepsilon(y - \boldsymbol{\beta}'_0 \mathbf{X} - \gamma_0 t) f_T(t | \mathbf{X}) dt$$



Modeling this part:

Provocatively utilize truncation information  
in statistical modeling

# Estimation procedure

## Setting:

- Model : 
$$\begin{cases} Y^* = \boldsymbol{\beta}'_0 \mathbf{X} + \gamma_0 T + \varepsilon \\ (T, C, \mathbf{X}) \perp \varepsilon \end{cases} \quad \dots \quad (\text{B})$$

- Left-truncated & right-censored data :

$$\{(T_i, Y_i, \Delta_i, \mathbf{X}_i); (i = 1, \dots, n)\} \text{ subject to } T_i \leq Y_i$$

## Interest:

- 1) Joint estimation of  $(\boldsymbol{\beta}'_0, \gamma_0)$
- 2) Estimation of  $S_\varepsilon(t) = \Pr(\varepsilon > t)$

## Estimating equations for

- a)  $\boldsymbol{\beta}_0 \rightarrow$  Inverting the log-rank test statistics
- b)  $\gamma_0 \rightarrow$  Inverting the quasi-independence test statistics

# Estimation procedure

## Residual transformation:

$$\begin{cases} \varepsilon_i^Y(\boldsymbol{\beta}, \gamma) = Y_i - \boldsymbol{\beta}'\mathbf{X}_i - \gamma T_i & \text{residual log - survival} \\ \varepsilon_i^T(\boldsymbol{\beta}, \gamma) = T_i - \boldsymbol{\beta}'\mathbf{X}_i - \gamma T_i & \text{residual log - truncation} \end{cases}$$

## a) Log-rank estimating equation:

By assumption (B),  $H_0 : Y^* - \boldsymbol{\beta}'_0\mathbf{X} - \gamma_0 T \perp \mathbf{X}$  is true.

$$\begin{aligned} S_n^{\text{Logrank}}(\boldsymbol{\beta}, \gamma) \\ = - \sum_{i < j} (\mathbf{X}_i - \mathbf{X}_j) \text{sgn}\{(\varepsilon_i^Y(\boldsymbol{\beta}, \gamma) - \varepsilon_j^Y(\boldsymbol{\beta}, \gamma))\} I\{\tilde{\varepsilon}_{ij}^T(\boldsymbol{\beta}, \gamma) \leq \tilde{\varepsilon}_{ij}^Y(\boldsymbol{\beta}, \gamma)\} O_{ij}(\boldsymbol{\beta}, \gamma) \end{aligned}$$

## b) Quasi-independence estimating equation:

By assumption (B),  $H_0 : Y^* - \boldsymbol{\beta}'_0\mathbf{X} - \gamma_0 T \perp T - \boldsymbol{\beta}'_0\mathbf{X} - \gamma_0 T$  is true

$$\begin{aligned} S_n^{\text{Kendall}}(\boldsymbol{\beta}, \gamma) \\ = \sum_{i < j} \text{sgn}\{(\varepsilon_i^T(\boldsymbol{\beta}, \gamma) - \varepsilon_j^T(\boldsymbol{\beta}, \gamma))(\varepsilon_i^Y(\boldsymbol{\beta}, \gamma) - \varepsilon_j^Y(\boldsymbol{\beta}, \gamma))\} I\{\tilde{\varepsilon}_{ij}^T(\boldsymbol{\beta}, \gamma) \leq \tilde{\varepsilon}_{ij}^Y(\boldsymbol{\beta}, \gamma)\} O_{ij}(\boldsymbol{\beta}, \gamma). \end{aligned}$$

Martin & Betensky type statistic (2005 JASA)

# Estimation procedure

## Numerical solution :

$$\begin{pmatrix} \hat{\boldsymbol{\beta}} \\ \hat{\gamma} \end{pmatrix} : \begin{cases} \mathbf{0} = \mathbf{S}_n^{\text{Logrank}}(\boldsymbol{\beta}, \gamma) \\ 0 = S_n^{\text{Kendall}}(\boldsymbol{\beta}, \gamma) \end{cases} \quad \leftarrow \text{Non-monotonic step functions}$$

Use the simplex algorithm (Nelder and Mead, 1965) to find a minimum of

$$M(\boldsymbol{\beta}, \gamma) = \{ \|\mathbf{S}_n^{\text{Logrank}}(\boldsymbol{\beta}, \gamma)\|_2 + |S_n^{\text{Kendall}}(\boldsymbol{\beta}, \gamma)|^2 \} / n^2$$

Implementation available as “R optim routine”.

→ Not easy to choose good initial in automatic way.

(one of question raised by referees)

NOTE: Newton-Raphson, bisection method, and linear programming (Jin, Lin and Wei, 2003) do not work:

## Theorem 2 (Asymptotic normality) :

$$n^{1/2} (\hat{\boldsymbol{\beta}} - \boldsymbol{\beta}_0, \hat{\gamma} - \gamma_0) \rightarrow N(\mathbf{0}, \mathbf{A}_0^{-1} \mathbf{B}_0 \mathbf{A}_0^{-1})$$

Difficulty :  $\mathbf{A}_0 \equiv \partial \Phi(F; \boldsymbol{\beta}, \gamma) / \partial (\boldsymbol{\beta}, \gamma) |_{\boldsymbol{\beta}_0, \gamma_0}$

not differentiable if plug-in  $F = F_n$

## Kernel-based empirical variance estimator :

$$\hat{\mathbf{A}}_0 = [\hat{\mathbf{A}}_0^{(1)}(\hat{\boldsymbol{\beta}}, \hat{\gamma}; b_1), \dots, \hat{\mathbf{A}}_0^{(p+1)}(\hat{\boldsymbol{\beta}}, \hat{\gamma}; b_{p+1})]$$

$$\hat{\mathbf{A}}_0^{(k)}(\boldsymbol{\beta}, \gamma; b_k) = \int_{u \neq 0} \frac{1}{u} \Phi(F_n; \beta_1, \dots, \beta_k + u, \dots, \beta_p, \gamma) \frac{1}{b_k} K\left(\frac{u}{b_k}\right) du, \quad k = 1, \dots, p$$

$$\hat{\mathbf{B}}_0 = \sum_{j=1}^n \phi_{F_n}(T_j, Y_j, \delta_j, \mathbf{X}_j; \hat{\boldsymbol{\beta}}, \hat{\gamma}) \phi_{F_n}(T_j, Y_j, \delta_j, \mathbf{X}_j; \hat{\boldsymbol{\beta}}, \hat{\gamma})' / n$$

- Optimal bandwidth

$$\begin{aligned}
 MSE\{ \hat{\mathbf{A}}_0^{(k)}(\hat{\boldsymbol{\beta}}, \hat{\boldsymbol{\gamma}}; b_k) \} &= Var\{ \hat{\mathbf{A}}_0^{(k)}(\hat{\boldsymbol{\beta}}, \hat{\boldsymbol{\gamma}}; b_k) \} + Bias[\hat{\mathbf{A}}_0^{(k)}(\boldsymbol{\beta}, \boldsymbol{\gamma}; b_k)]^2 \\
 &\approx \frac{1}{nb_k} E\left[ \frac{\partial}{\partial \boldsymbol{\beta}_k} \phi_F\{ (T_j, Y_j, \delta_j, \mathbf{X}_j); \boldsymbol{\beta}_0, \boldsymbol{\gamma}_0 \} \right]^2 + \frac{b_k^4}{36} \left[ \frac{\partial^3}{\partial \boldsymbol{\beta}_k^3} \Phi(F; \boldsymbol{\beta}_0, \boldsymbol{\gamma}_0) \right]^2 \mu_2(K)^2,
 \end{aligned}$$

The bandwidth that minimizes the preceding expression becomes

$$b_k^{opt} = \left( 9E\left[ \frac{\partial}{\partial \boldsymbol{\beta}_k} \phi_F\{ (T_j, Y_j, \delta_j, \mathbf{X}_j); \boldsymbol{\beta}_0, \boldsymbol{\gamma}_0 \} \right]^2 \left[ \frac{\partial^3}{\partial \boldsymbol{\beta}_k^3} \Phi(F; \boldsymbol{\beta}_0, \boldsymbol{\gamma}_0) \right]^{-2} \mu_2(K)^{-2} \right)^{1/5} \frac{1}{n^{1/5}}$$

- In practice, use Silverman's reference bandwidth ([Sheather, 2004](#)) under the normal kernel

$$\hat{b}_k = 0.5 \min( S_k, IQR_k / 1.34 ) n^{-1/5}$$

# Estimation procedure

## Estimation of error distribution:

**Target:**  $S_\varepsilon(t) = \Pr(\varepsilon > t) = \prod_{u \leq t} \left\{ 1 - \frac{\Pr(\varepsilon = u)}{\Pr(\varepsilon \geq u)} \right\}$

$$\begin{cases} \varepsilon_i \approx \varepsilon_i^Y(\hat{\boldsymbol{\beta}}, \hat{\gamma}) = Y_i - \hat{\boldsymbol{\beta}}' \mathbf{X}_i - \hat{\gamma} T_i & \text{residual log-survival} \\ \varepsilon_i^T(\hat{\boldsymbol{\beta}}, \hat{\gamma}) = T_i - \hat{\boldsymbol{\beta}}' \mathbf{X}_i - \hat{\gamma} T_i & \text{residual log-truncation} \end{cases}$$

→  $\{ (\varepsilon_i^T(\hat{\boldsymbol{\beta}}, \hat{\gamma}), \varepsilon_i^Y(\hat{\boldsymbol{\beta}}, \hat{\gamma}), \Delta_i); i = 1, \dots, n \}$  subject to  $\varepsilon_i^T(\hat{\boldsymbol{\beta}}, \hat{\gamma}) \leq \varepsilon_i^Y(\hat{\boldsymbol{\beta}}, \hat{\gamma})$

: Left-truncated and right-censored data for  $\varepsilon$

## Product-limit estimator for with non-iid samples

$$\hat{S}_\varepsilon(t; \hat{\boldsymbol{\beta}}, \hat{\gamma}) = \prod_{u \leq t} \left\{ 1 - \frac{\sum_j I(\varepsilon_j^Y(\hat{\boldsymbol{\beta}}, \hat{\gamma}) = u, \Delta_j = 1)}{\sum_j I(\varepsilon_j^T(\hat{\boldsymbol{\beta}}, \hat{\gamma}) \leq u \leq \varepsilon_j^Y(\hat{\boldsymbol{\beta}}, \hat{\gamma}))} \right\}$$



**Theorem 3** *Under Assumptions I, II and V, the product-limit estimator*

*$\hat{S}_\varepsilon(t; \hat{\beta}, \hat{\gamma})$  converges in probability to  $S_\varepsilon(t)$ , uniformly over  $t \in [a, b]$ .*

- Asymptotic normality is remain to be done

# Simulation

**Model :**  $\begin{bmatrix} Y^* \\ T \end{bmatrix} \sim N\left(\begin{bmatrix} \beta_0 X \\ -1 \end{bmatrix}, \begin{bmatrix} 1 & \rho \\ \rho & 1 \end{bmatrix}\right), \quad C \sim N(1, 1), \quad X \sim U(0, 1)$

This induces the linear regression model

$$Y^* = \beta_0 X + \gamma_0 T + \varepsilon, \quad \varepsilon \sim N(\gamma_0, 1 - \gamma_0^2)$$

, where  $\rho = \gamma_0$  and  $S_\varepsilon(t) = 1 - \Phi\left\{\frac{t - \gamma_0}{(1 - \gamma_0^2)^{1/2}}\right\}$

**Parameter configurations:**

	1	2	3	4	5	6
$(\beta_0, \gamma_0)$	(0, -0.5)	(0, 0)	(0, 0.5)	(1, -0.5)	(1, 0)	(1, 0.5)
$\Pr(T \leq Y^*)$	0.72	0.76	0.84	0.80	0.85	0.84
$\Pr(C < Y^*   T \leq Y^*)$	0.30	0.27	0.23	0.41	0.39	0.34

# Simulation

## 1. Generate data

$(T_i, Y_i, \Delta_i, \mathbf{X}_i)$ , subject to  $T_i \leq Y_i$ , for  $i = 1, \dots, n$

**from model :**

$$Y^* = \beta_0 X + \gamma_0 T + \varepsilon, \quad \text{where } \rho = \gamma_0 \quad \text{and} \quad \varepsilon \sim N(\gamma_0, 1 - \gamma_0^2)$$

## 2. Estimate parameters

$$(\hat{\beta}, \hat{\gamma}) = \arg \min_{(\beta, \gamma)} M(\beta, \gamma):$$

using R optim with the true initial value  $(\beta_0, \gamma_0)$

$$\hat{S}_\varepsilon(t; \hat{\beta}, \hat{\gamma})$$

## 3. Variance estimation & confidence interval

$$[ \hat{\beta}_k - z_{\alpha/2} se(\hat{\beta}_k), \hat{\beta}_k + z_{\alpha/2} se(\hat{\beta}_k) ]$$

**Table 1** Simulation results for the proposed estimator  $(\hat{\beta}, \hat{\gamma})^\dagger$ .

		Estimation of $\beta_0$				Estimation of $\gamma_0$			
$(\beta_0, \gamma_0)$	$n$	Bias( $\hat{\beta}$ )	SD( $\hat{\beta}$ )	E{ $se(\hat{\beta})$ }	95%Cov	Bias( $\hat{\gamma}$ )	SD( $\hat{\gamma}$ )	E{ $se(\hat{\gamma})$ }	95%Cov
(0, -0.5)	150	0.003	0.333	0.352	0.926	-0.018	0.169	0.165	0.932
	300	-0.007	0.234	0.233	0.944	-0.002	0.112	0.112	0.944
(0, 0)	150	0.005	0.392	0.419	0.946	-0.008	0.173	0.169	0.942
	300	-0.002	0.275	0.287	0.954	0.002	0.113	0.116	0.946
(0, 0.5)	150	0.000	0.320	0.370	0.964	-0.001	0.124	0.132	0.968
	300	-0.006	0.238	0.253	0.958	0.000	0.085	0.087	0.962
(1, -0.5)	150	0.010	0.320	0.359	0.954	-0.021	0.163	0.155	0.950
	300	-0.003	0.234	0.246	0.952	-0.005	0.116	0.108	0.940
(1, 0)	150	0.041	0.411	0.430	0.954	-0.015	0.159	0.160	0.966
	300	0.012	0.284	0.289	0.950	-0.008	0.105	0.108	0.958
(1, 0.5)	150	0.012	0.331	0.355	0.954	0.007	0.115	0.127	0.970
	300	0.002	0.240	0.243	0.958	0.003	0.080	0.082	0.964

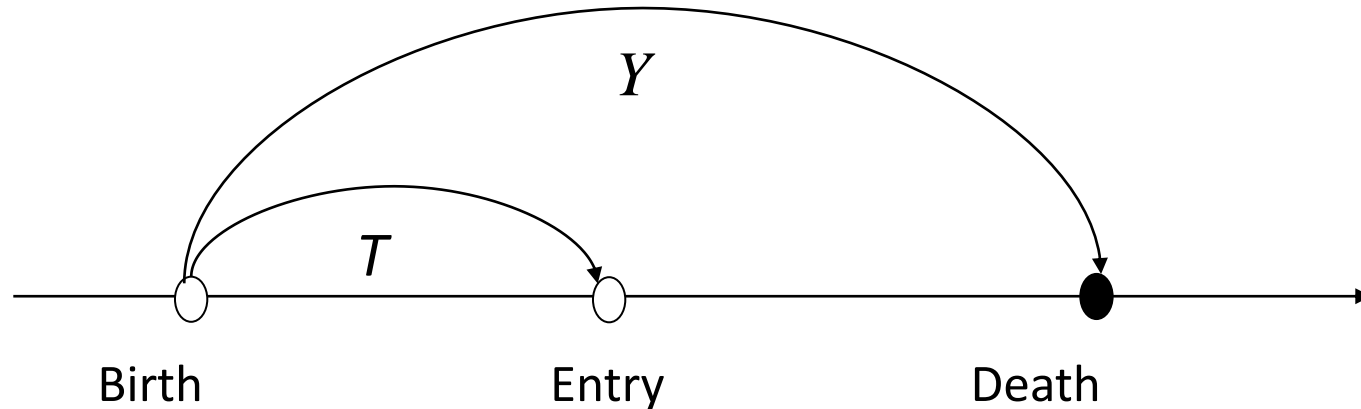
**Table 4** Simulation results for the proposed estimator  $\hat{S}_\varepsilon(t; \hat{\beta}, \hat{\gamma})^\dagger$ .

		$S_\varepsilon(t) = 0.25$		$S_\varepsilon(t) = 0.5$		$S_\varepsilon(t) = 0.75$	
$(\beta_0, \gamma_0)$	$n$	Bias	SD	Bias	SD	Bias	SD
(0, -0.5)	150	-0.076	0.090	-0.008	0.146	0.055	0.121
	300	-0.075	0.065	0.002	0.103	0.073	0.076
(0, 0)	150	0.006	0.109	0.000	0.146	-0.012	0.137
	300	0.003	0.072	0.003	0.098	-0.001	0.090
(0, 0.5)	150	0.044	0.100	0.004	0.129	-0.040	0.131
	300	0.041	0.076	0.000	0.098	-0.042	0.101
(1, -0.5)	150	-0.079	0.087	-0.010	0.149	0.056	0.120
	300	-0.078	0.063	0.000	0.109	0.071	0.083
(1, 0)	150	-0.005	0.102	-0.014	0.141	-0.020	0.131
	300	-0.002	0.071	-0.007	0.096	-0.009	0.089
(1, 0.5)	150	0.043	0.100	0.003	0.118	-0.039	0.113
	300	0.043	0.072	0.002	0.087	-0.040	0.082

**Table 3** Simulation results for comparing the proposed estimator with the estimator of  
Lai and Ying (1991)†.

		Proposed method			Lai & Ying (1991)			
$(\beta_0, \gamma_0)$	$n$	$E(\hat{\beta})$	$\text{Bias}(\hat{\beta})$	$SD(\hat{\beta})$	$E(\hat{\beta})$	$\text{Bias}(\hat{\beta})$	$SD(\hat{\beta})$	
Independent truncation	(-1, -0.5)	150	-0.966	0.034	0.327	-0.779	0.221	0.296
		300	-0.988	0.012	0.241	-0.795	0.205	0.212
	(-1, 0)	150	-1.007	-0.007	0.420	-1.026	-0.026	0.443
		300	-1.003	-0.003	0.322	-1.008	-0.008	0.307
	(-1, 0.5)	150	-1.002	-0.002	0.372	-1.554	-0.554	0.659
		300	-1.004	-0.004	0.292	-1.526	-0.526	0.441
Independent truncation	(1, -0.5)	150	1.010	0.010	0.320	0.829	-0.171	0.305
		300	0.997	-0.003	0.234	0.820	-0.180	0.217
	(1, 0)	150	1.041	0.041	0.411	1.027	0.027	0.411
		300	1.012	0.012	0.284	1.007	0.007	0.281
	(1, 0.5)	150	1.012	0.012	0.331	1.373	0.373	0.506
		300	1.002	0.002	0.240	1.363	0.363	0.345

# Data analysis



## Channing House data (Hyde, 1980)

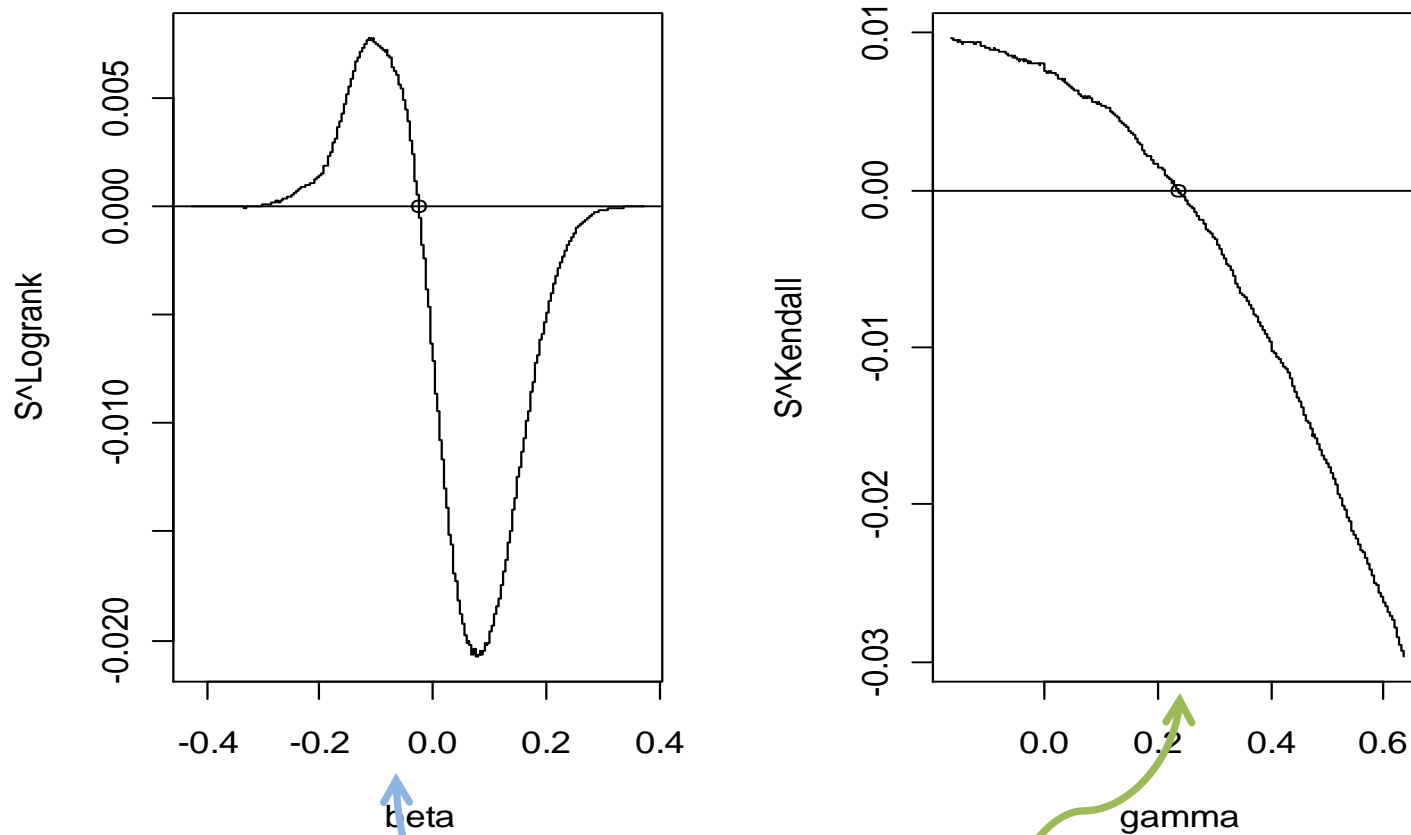
Available information for Individuals (n=462)

- $T$  : Entry age
- $Y^*$  : Age at death or censoring
- $X$ : Gender ( $X = 1$  for male;  $X = 0$  for female )

## Fit the proposed model:

$$Y^* = \beta_0 X + \gamma_0 T + \varepsilon, \quad \text{where } \varepsilon \text{ is unspecified}$$

# Data analysis



**Fig. 3.** Plots of  $S_n^{\text{Logrank}}(\beta, \hat{\gamma})$  and  $S_n^{\text{Kendall}}(\hat{\beta}, \gamma)$  based on the Channing house data.

The numerical solutions  $\hat{\beta} = -0.026$  and  $\hat{\gamma} = 0.236$  obtained from the grid search algorithm are indicated by “o”.



# Data analysis

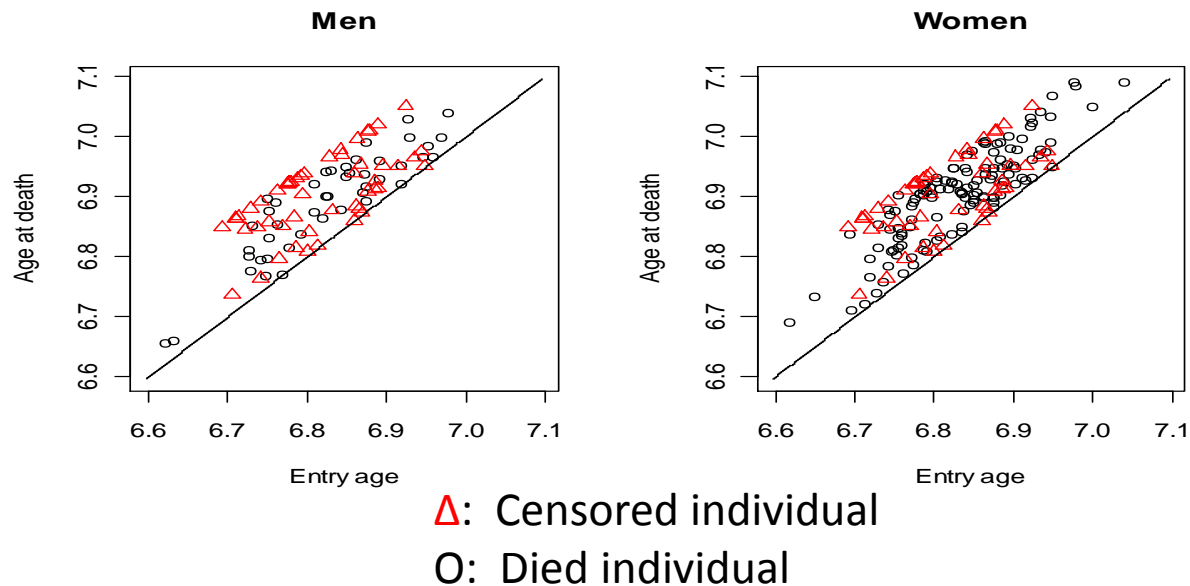
## Interpretation:

$$Y^* = -0.026 \times X + 0.236 \times T + \varepsilon$$

Age at death                      Gender                      Age at entry

$$\begin{cases} \hat{\beta} = -0.026 & \dots 95\% \text{ CI } (-0.067, 0.014) \\ \hat{\gamma} = 0.236 & \dots 95\% \text{ CI } (0.014, 0.459) \end{cases}$$

∴ Late entry to Channing house prolong the survival

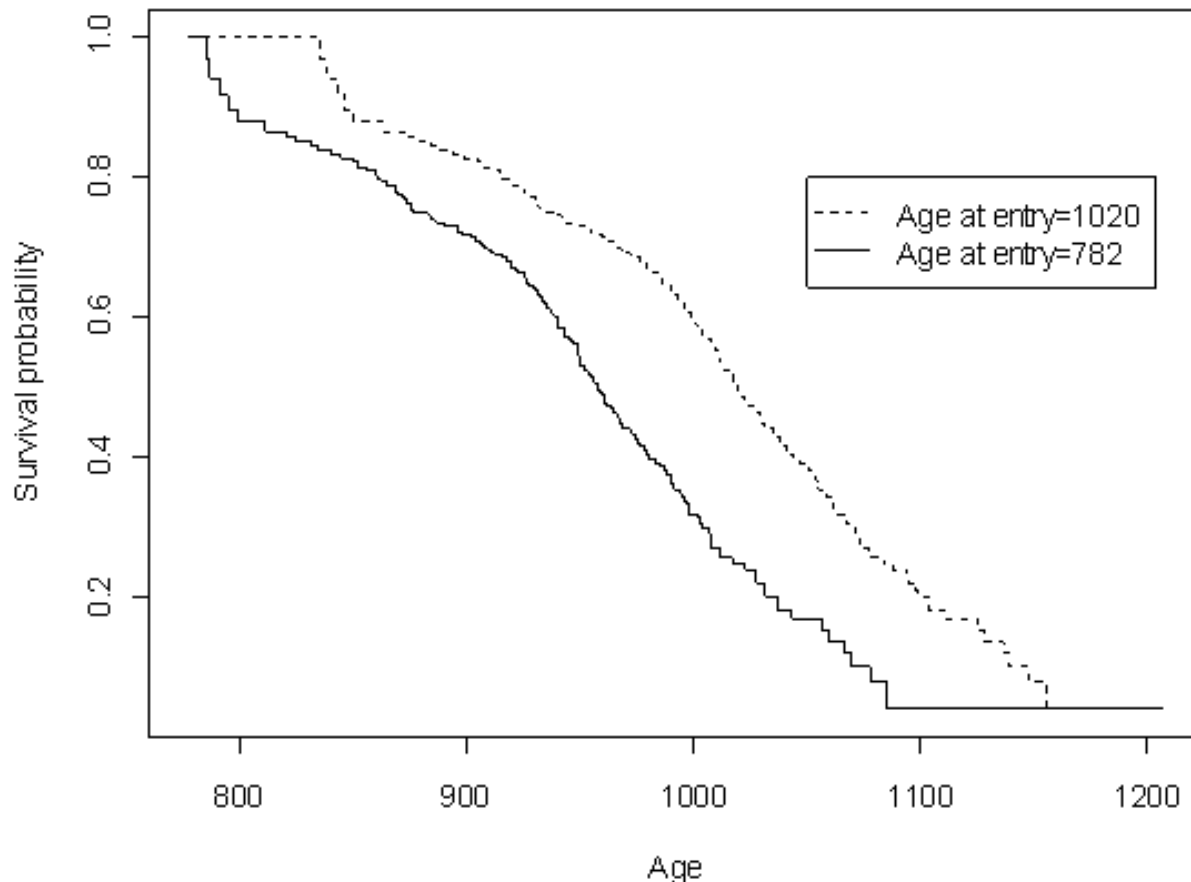


# Data analysis

## Subject specific survival :

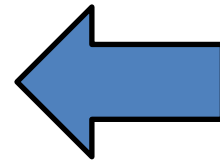
Survival for the two individual:

- ID#1: Entry age = 782 (month), sex = male
- ID#2: Entry age = 1020 (month), sex = male



# Conclusion

- We propose a semi-parametric AFT model which utilizes *both covariates and truncation variables* to model lifetimes
  - AFT of Lai & Ying (1991) can only utilize covariate as regressors
- We relax the independent truncation assumption in the Lai & Ying (1991)'s AFT method
- In Channing house data:
  - The entry age (truncation ) is informative for survival
  - Early entry → shorter survival
  - Late entry → Longer survival
  - Male → Shorter survival than female (non-significant)



Significant

**Thank you for your kind attention**