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**Nonparametric maximum likelihood estimation for
dependent truncation data based on copula
(this work is submitted to *Journal of Multivariate Analysis*)**

Takeshi Emura

Graduate Institute of Statistics, National Central University, Taiwan

Joint work with Weijing Wang

Institute of Statistics, National Chiao Tung University, Taiwan

Outlines

Part I: Literature Review & Motivation

- Truncation data:
- Traditional approaches
- Copula-based approaches

Part II: Proposed method

- Proposed method
- Simulation, Data analysis
- Extension to multiparameter copulas
(in progress)
- Conclusion

**This part is submitted to
*JMVA***

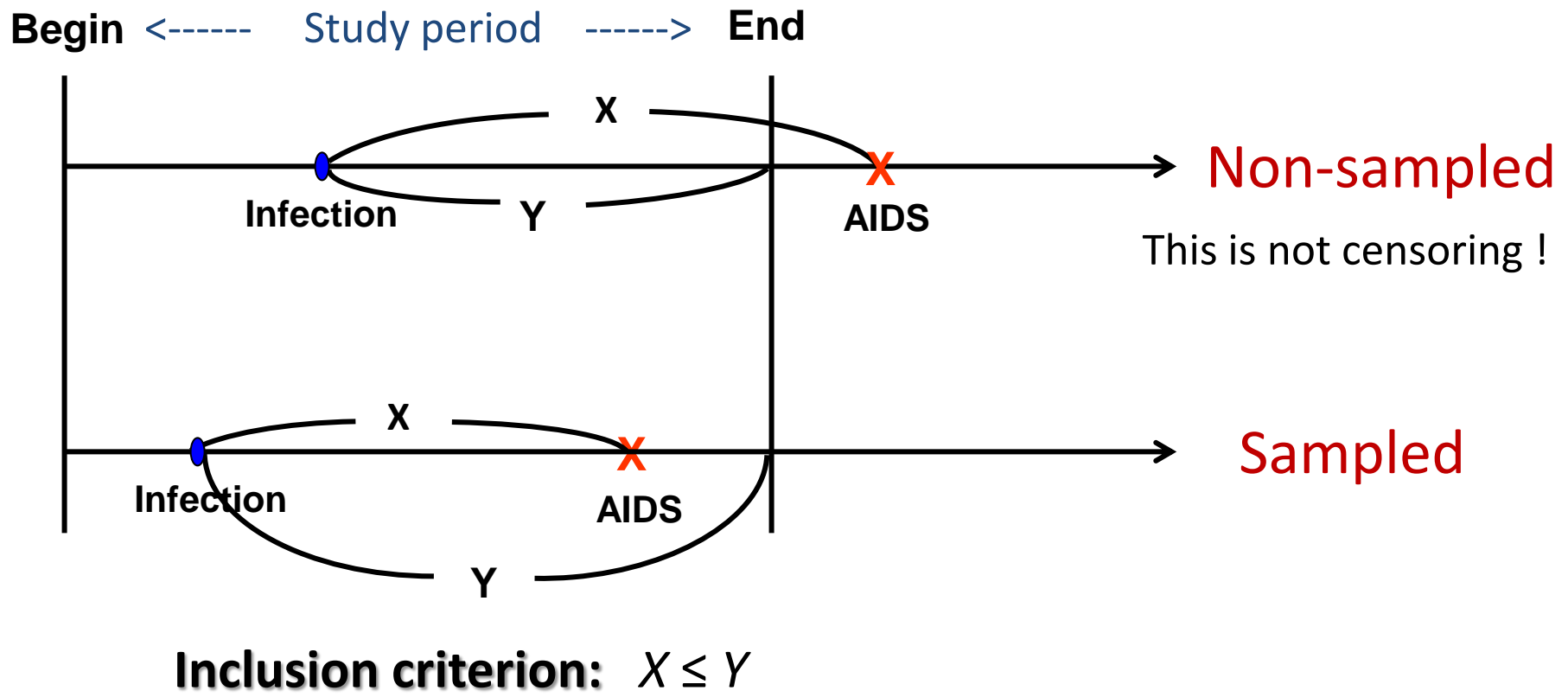
Part I

Part I: Literature Review & Motivation

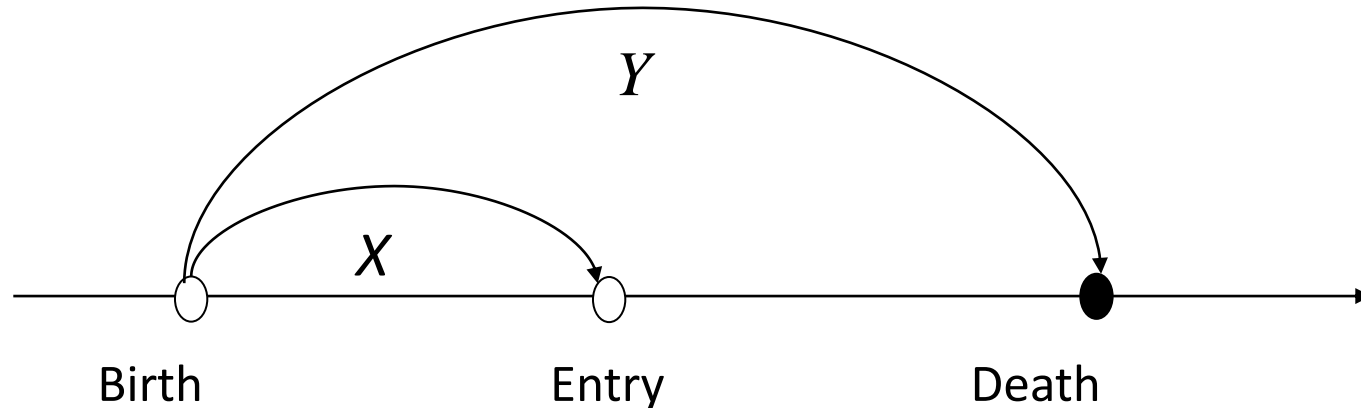
Truncation data

- Transfusion-related AIDS

(Lagakos et al., 1988; Kalbfleisch & Lawless, 1989)



Truncation data



Channing House data (Hyde, 1980)

Available information for Individuals (n=462)

- X: Entry age
- Y: Age at death or censoring (withdraw)
- Sex (97 man ; 365 women)

Inclusion criterion: $X \leq Y$

Truncation data

- Truncation data :

$$\{(X_j, Y_j); j = 1, \dots, n\}$$

$$\text{subject to } X_j \leq Y_j$$



i.i.d. from

$$\Pr(X \leq x, Y \leq y \mid X \leq Y),$$

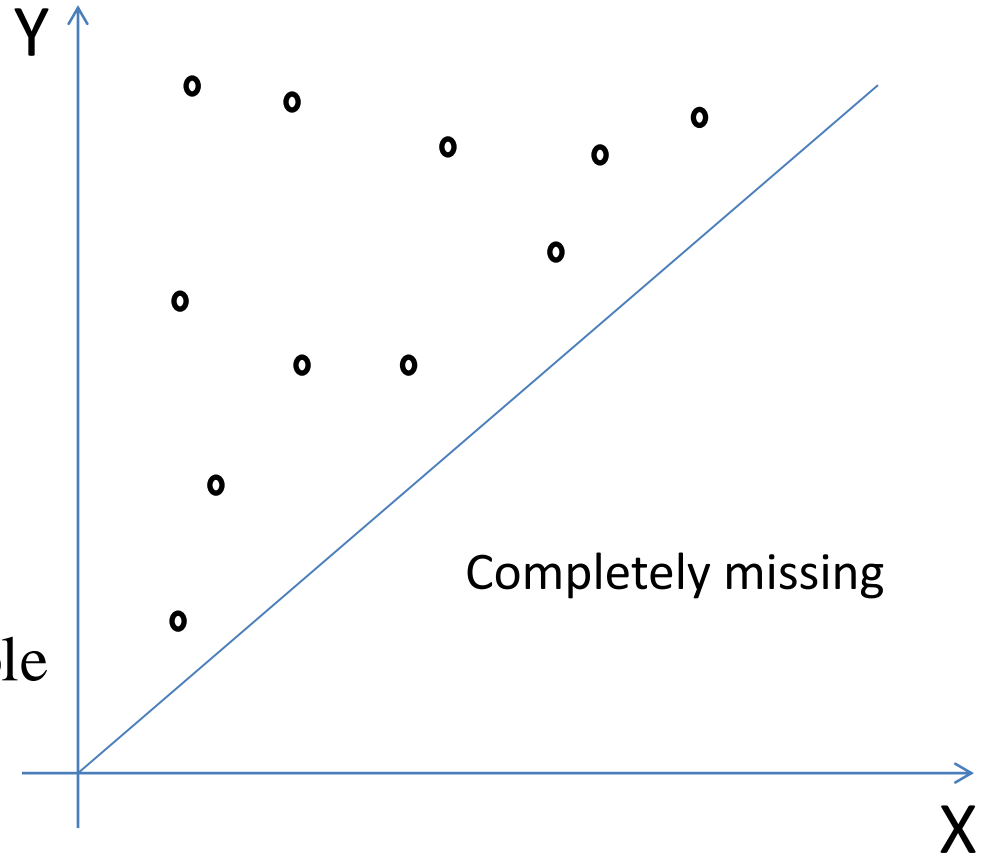
where (X, Y) is

the "*population*" random variable

Interest: Estimation of

$$F_X(x) = \Pr(X \leq x)$$

$$F_Y(y) = \Pr(Y \leq y)$$



Estimation under truncation

Traditional analysis

- Nonparametric estimator

$$\hat{F}_X(x) = \prod_{u>x} \left\{ 1 - \frac{\sum_{j=1}^n I(X_j = u)}{\sum_{j=1}^n I(X_j \leq u, Y_j \geq u)} \right\}, \quad \hat{F}_Y(y) = 1 - \prod_{u \leq y} \left\{ 1 - \frac{\sum_{j=1}^n I(Y_j = u)}{\sum_{j=1}^n I(X_j \leq u, Y_j \geq u)} \right\}$$

(Lynden-Bell, 1971; Lagakos et al., 1988)

- **Key assumption: Quasi-independence (Tsai, 1990):**

$$X \perp_Q Y : \Pr(X \leq x, Y \leq y | X \leq Y) \propto \int \int_{\substack{u \leq x, v \leq y \\ u \leq v}} dF_X(u) dF_Y(v)$$

*Quasi-independence assumption is testable

(Tsai, 1990; Chen et al., 1996; Martin & Betensky, 2005; Emura & Wang, 2010)

Estimation under truncation

Data : $\{(X_j, Y_j) (j = 1, \dots, n)\}$ subject to $X_j \leq Y_j$

1) Under $X \perp_Q Y$,

$$\hat{F}_X(x) = \prod_{u>x} \left\{ 1 - \frac{\sum_{j=1}^n I(X_j = u)}{\sum_{j=1}^n I(X_j \leq u, Y_j \geq u)} \right\}, \quad \hat{F}_Y(y) = 1 - \prod_{u \leq y} \left\{ 1 - \frac{\sum_{j=1}^n I(Y_j = u)}{\sum_{j=1}^n I(X_j \leq u, Y_j \geq u)} \right\}$$

2) Under Copula assumptions on (X, Y)

* weaker assumption than $X \perp Y$

(Chaieb et al., 2006, *Biometrika*; Beaudoin & Lakhel, 2008 *Stat. Med.*;
Emura, Wang & Hung, 2011, *Sinica*; Ding, 2011, Manuscript).

No explicit marginal estimators.

Focus of recent research: *Dependent Truncation*

(See also, Emura & Konno 2012: *Stat papers, CSDA*)

Copula

$$\Pr(X \leq x, Y \leq y) = C[\Pr(X \leq x), \Pr(Y \leq y)]$$

- **Example 1:** Independence copula

$$C[u, v] = uv$$

- **Example 2:** Frank copula (Genest, 1986; Frank, 1979)

$$C_\alpha[u, v] = \log_{\alpha^{-1}} \left\{ 1 + \frac{(\alpha^{-u} - 1)(\alpha^{-v} - 1)}{(\alpha^{-1} - 1)} \right\}, \quad \alpha > 0$$

$$\lim_{\alpha \rightarrow 1} C_\alpha[u, v] = uv$$

- **Example 3:** Normal copula

$$C_\rho[u, v] = \Phi_\rho[\Phi^{-1}(u), \Phi^{-1}(v)], \quad -1 < \rho < 1$$

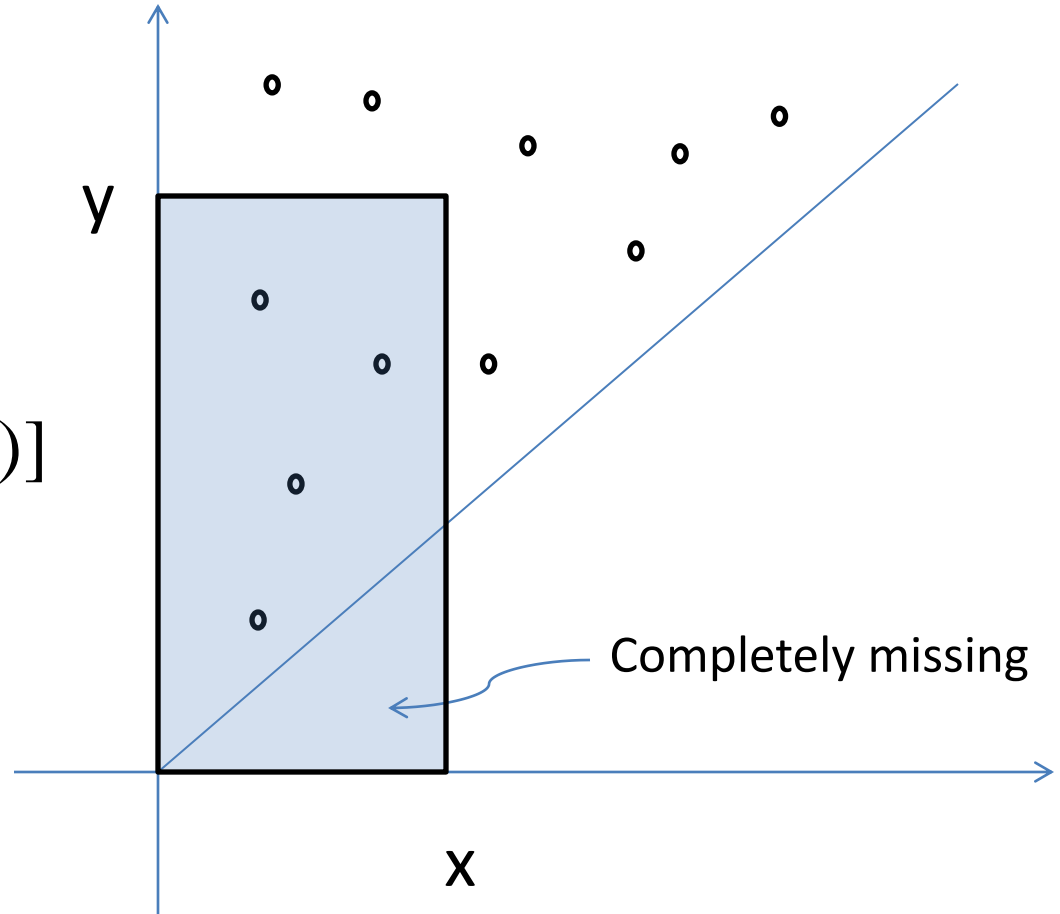
Φ_ρ : Joint CDF of standard bivariate normal

$$\lim_{\rho \rightarrow 0} C_\rho[u, v] = uv$$

Copula model

$$\Pr(X \leq x, Y \leq y) \\ = C[\Pr(X \leq x), \Pr(Y \leq y)]$$

The model is
unidentifiable



Copula model

$$\Pr(X \leq x, Y > y \mid X \leq Y)$$

$$= \frac{C_\alpha[F_X(x), S_Y(y)]}{c(\alpha, F_X, S_Y)}$$

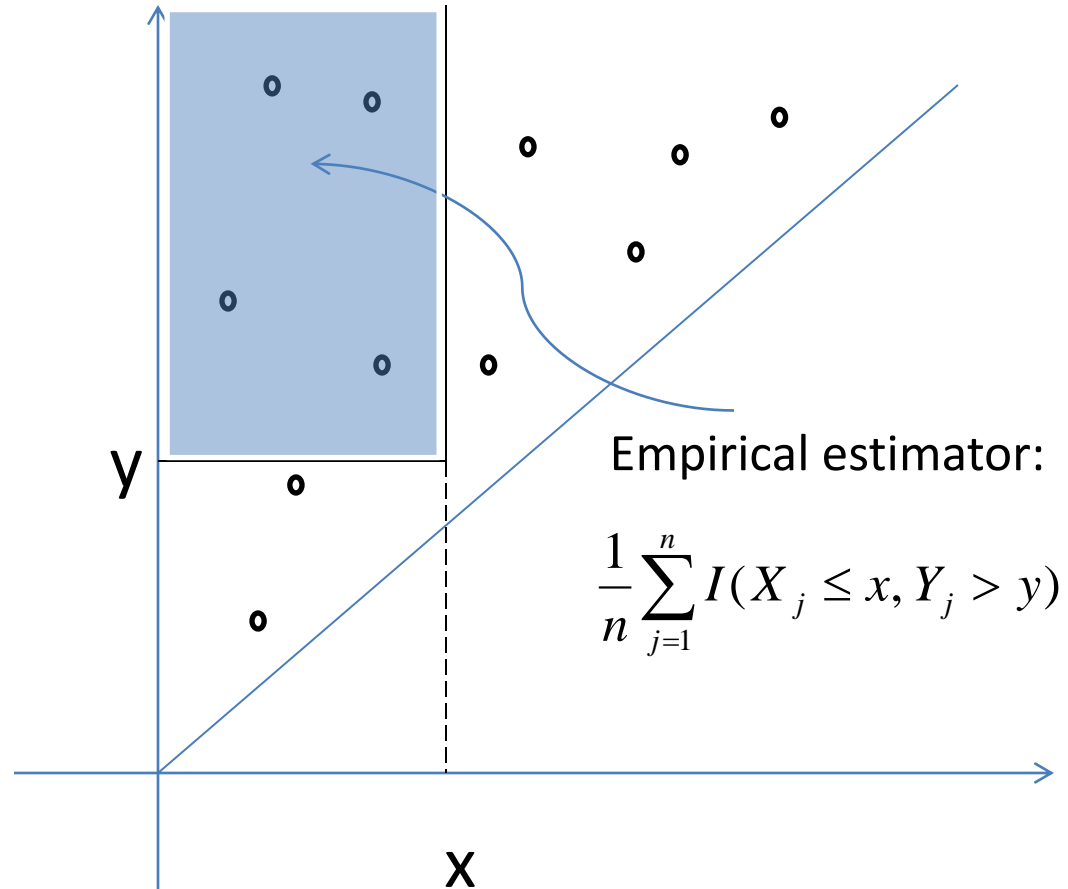
where

$$c(\alpha, F_X, S_Y) =$$

$$\iint_{x \leq y} \frac{\partial^2}{\partial x \partial y} C_\alpha[F_X(x), S_Y(y)] dx dy$$

- **Semi-survival copula**
(Chaieb et al., 2006, *Biometrika*)

- Quasi-independence: $C_\alpha[u, v] = uv$



Existing procedures

- **Archimedean family:** $C_\alpha[u, v] = \phi_\alpha^{-1}\{\phi_\alpha(u) + \phi_\alpha(v)\}$

Useful for solving moment equations:

[Chaieb et al. (2006) and Emura et al. (2011)]

$$\therefore \frac{1}{n} \sum_{j=1}^n I(X_j \leq t, Y_j > t) = \frac{C_\alpha[F_X(t), S_Y(t)]}{c(\alpha, F_X, S_Y)},$$

$$\Leftrightarrow \phi_\alpha \left(\frac{c(\alpha, F_X, S_Y)}{n} \sum_{j=1}^n I(X_j \leq t, Y_j > t) \right) = \phi_\alpha(F_X(t)) + \phi_\alpha(S_Y(t))$$

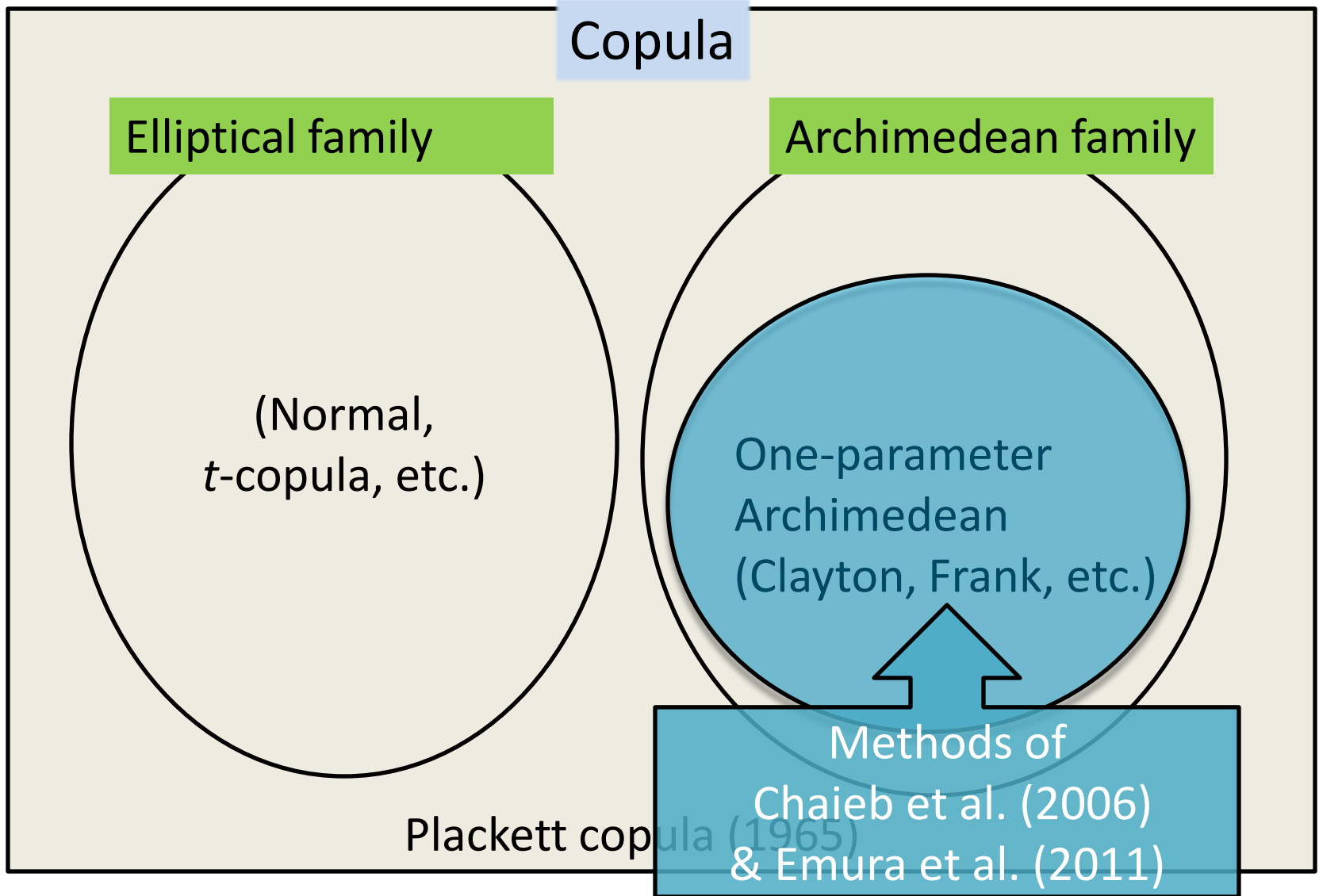
$$\Rightarrow F_X(t) = \phi_\alpha^{-1} \left\{ \phi_\alpha \left(\frac{c(\alpha, F_X, S_Y)}{n} \sum_{j=1}^n I(X_j \leq t, Y_j > t) \right) - \phi_\alpha(S_Y(t)) \right\}$$

$$S_Y(t) = \phi_\alpha^{-1} \left\{ \phi_\alpha \left(\frac{c(\alpha, F_X, S_Y)}{n} \sum_{j=1}^n I(X_j \leq t, Y_j > t) \right) - \phi_\alpha(F_X(t)) \right\}$$

Sequentially solve: $S_X(X_1) \equiv 1 \rightarrow F_X(X_1) \rightarrow S_X(X_2) \rightarrow \dots$

Assumption on "no ties": all $t \in (X_1, \dots, X_n, Y_1, \dots, Y_n)$ are different

Existing procedures



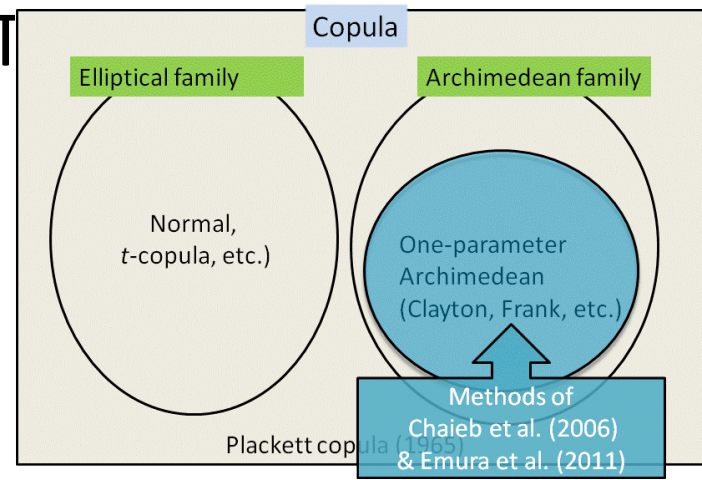
Part II: Proposed method

Proposed method

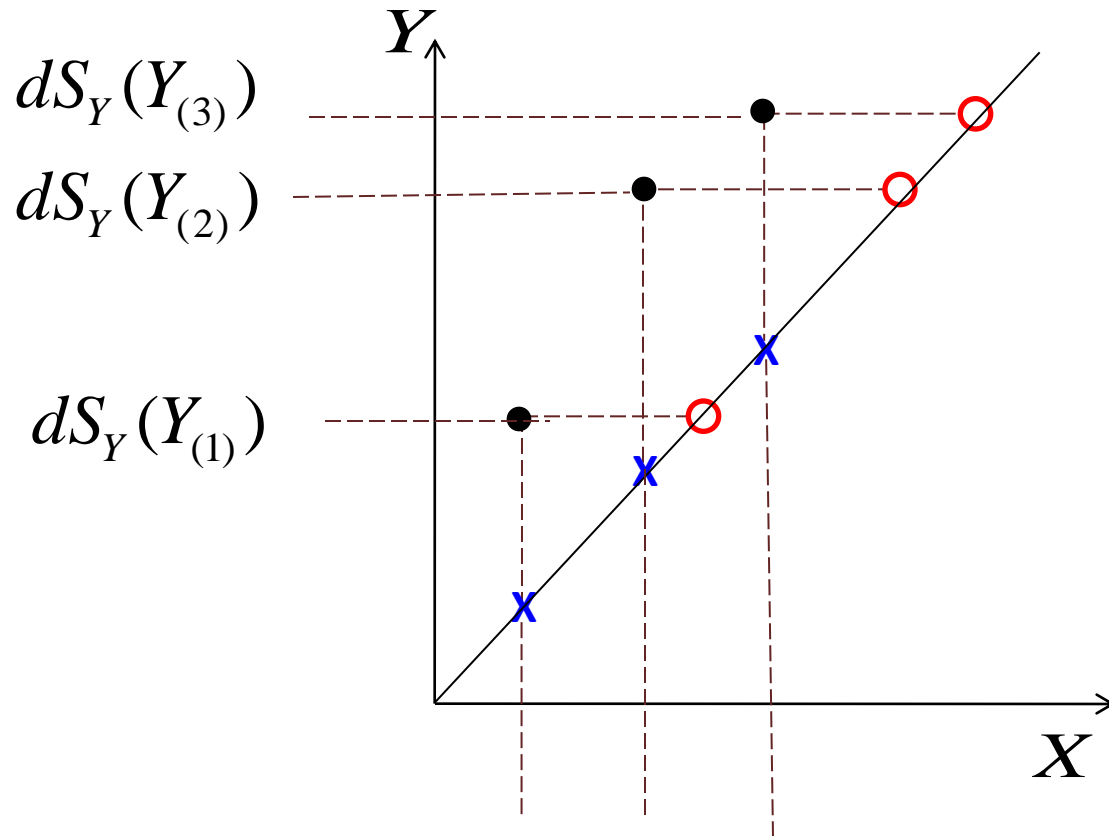
- The preceding two methods use moment-based estimating equations for (F_X, S_Y) under Archimedean copula family
- In this talk, we propose to get (\hat{F}_X, \hat{S}_Y) by the nonparametric maximum likelihood estimator (NPMLE; due to T

Advantage:

Potentially applicable for a broader class of copulas



NPMLE: due to Turnbull (1976)



Cumbersome constraints:

$$0 \leq dF_X(X_{(1)}) \leq 1$$

$$0 \leq dF_X(X_{(2)}) \leq 1$$

$$0 \leq dF_X(X_{(3)}) \leq 1$$

$$\sum_{i=1}^3 dF_X(X_{(i)}) = 1$$

$$dF_X(X_{(1)}) \quad dF_X(X_{(2)}) \quad dF_X(X_{(3)})$$

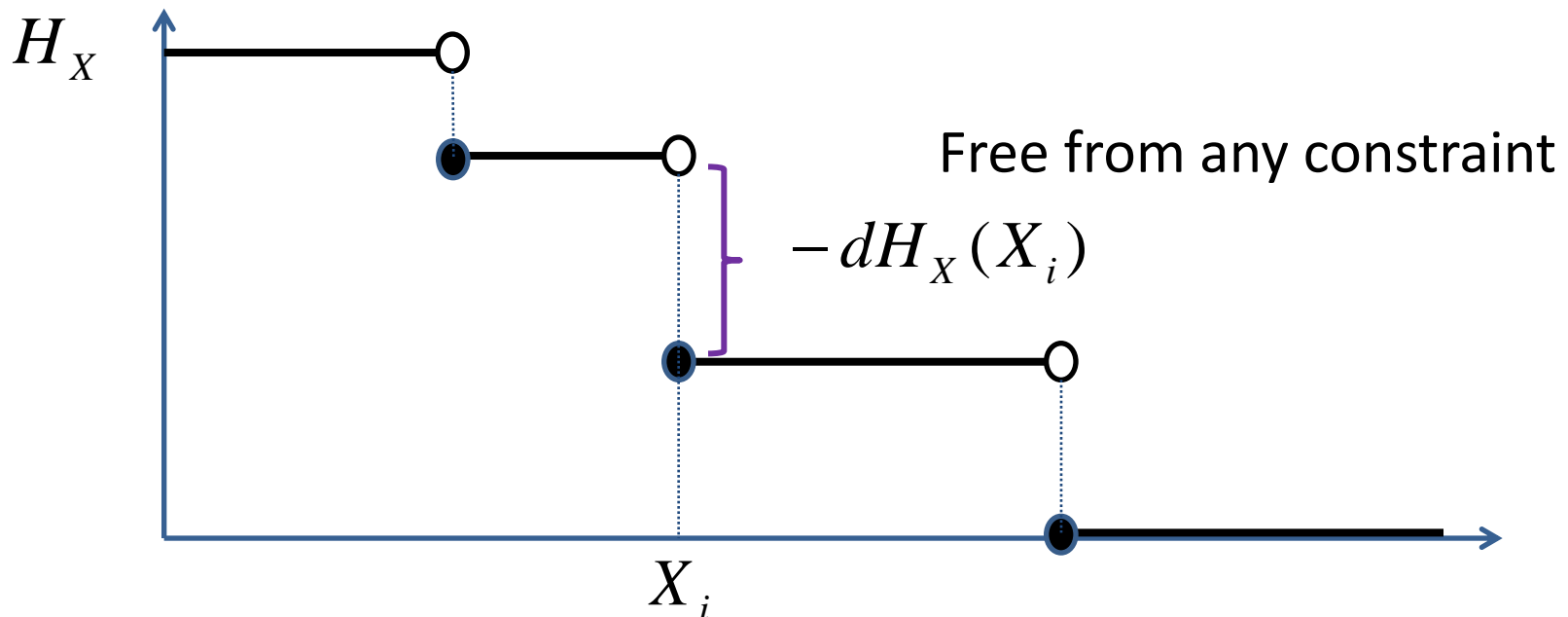
- Parameterize (F_X, S_Y) as follows:

$$F_X(x) = e^{-H_X(x)}, \quad S_Y(y) = e^{-\Lambda_Y(y)}$$

* $H_X(x)$: Reverse-time cumulative hazard

(Lagakos et al., 1988; Navaro & Ruiz, 1996)

* $\Lambda_Y(y)$: Cumulative hazard



Proposed method

- Semi-survival Copula (Chaieb et al., 2006)

$$\Pr(X \leq x, Y > y | X \leq Y) = \frac{C_\alpha [e^{-H_X(x)}, e^{-\Lambda_Y(y^-)}]}{c(\alpha, H_X, \Lambda_Y)},$$

where $c(\alpha, H_X, \Lambda_Y) = \iint_{x \leq y} -\frac{\partial^2}{\partial x \partial y} C_\alpha [e^{-H_X(x)}, e^{-\Lambda_Y(y^-)}] dx dy$

leading to the density

$$\Pr(X = x, Y = y | X \leq Y) = \frac{\eta_\alpha [H_X(x), \Lambda_Y(y^-)]}{c(\alpha, H_X, \Lambda_Y)} \{-dH_X(x)\} d\Lambda_Y(y),$$

where $\eta_\alpha [x, y] = e^{-x} e^{-y} \frac{\partial^2}{\partial u \partial u} C_\alpha [u, u] \Big|_{u=e^{-x}, v=e^{-y}}$

Proposed method

- Log-likelihood

$$l_n(\alpha, H_X, \Lambda_Y) =$$

$$\sum_{j=1}^n \log \eta_\alpha [H_X(X_j), \Lambda_Y(Y_j-)] + \log \{-dH_X(X_j)\} + \log d\Lambda_Y(Y_j) - \log c(\alpha, H_X, \Lambda_Y)$$

→ Maximize for $(2n+1)$ parameters

$$(\alpha, -dH_X(X_1), \dots, -dH_X(X_n), d\Lambda_Y(Y_1), \dots, d\Lambda_Y(Y_n))$$

Proposed method

- **Identifiability problem**

The maximum of $l_n(\alpha, H_X, \Lambda_Y)$ is not unique

(# of parameters = $2n+1$ > # of observed points = $2n$)

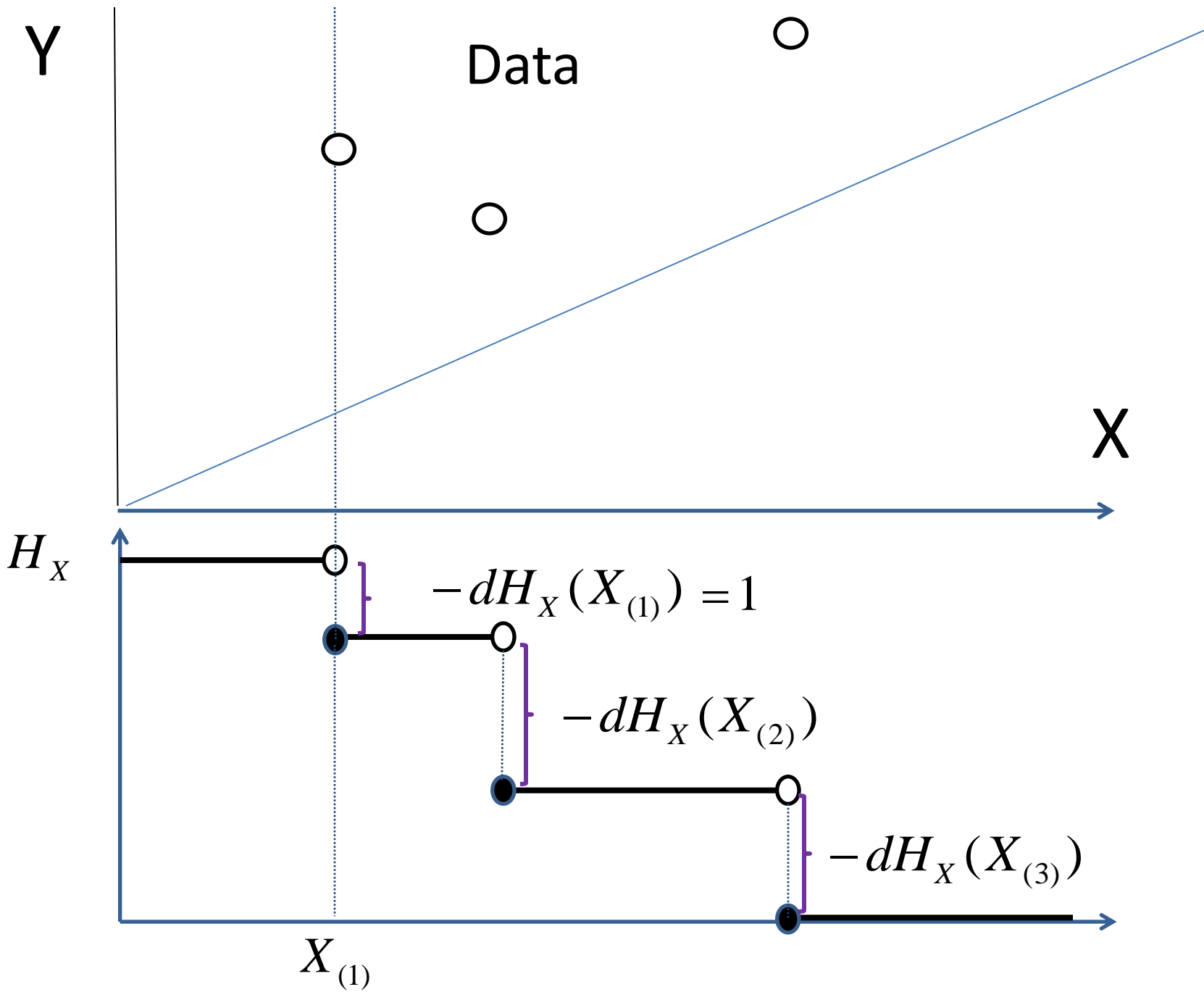
*analogy with the linear regression with $p > n$

- **Idea:** reduces to $2n-1$ parameters

$$(\alpha, -dH_X(X_1), \dots, -dH_X(X_n), d\Lambda_Y(Y_1), \dots, d\Lambda_Y(Y_n))$$



$$(\alpha, \underbrace{-dH_X(X_{(1)})}_{\equiv 1}, \dots, -dH_X(X_{(n)}), d\Lambda_Y(Y_{(1)}), \dots, \underbrace{d\Lambda_Y(Y_{(n)})}_{\equiv 1})$$



Proposed method

- $2n-1$ score equations

$$0 = \partial l_n(\alpha, H_X, \Lambda_Y) / \partial \alpha$$

$$0 = \partial l_n(\alpha, H_X, \Lambda_Y) / \partial \{-dH_X(X_{(j)})\}, \quad j = 2, \dots, n$$

$$0 = \partial l_n(\alpha, H_X, \Lambda_Y) / \partial d\Lambda_Y(Y_{(j)}), \quad j = 1, \dots, n-1$$

leading to a self-consistency (Turnbull, 1976)
type equations

$$H_X(x) = \int_x^\infty \frac{\sum_{j=1}^n I(X_j = u)}{\sum_{j=1}^n \Psi_j^{(1,0)}(u; \alpha, H_X, \Lambda_Y)}$$

$$\Lambda_Y(x) = \int_0^x \frac{\sum_{j=1}^n I(Y_j = u)}{\sum_{j=1}^n \Psi_j^{(0,1)}(u; \alpha, H_X, \Lambda_Y)}$$

Proposed method

- Quasi-Newton algorithm to maximize

$$l_n(\alpha, H_X, \Lambda_Y) =$$

$$\sum_{j=1}^n \log \eta_{\alpha}[H_X(X_j), \Lambda_Y(Y_j)] + \log\{-dH_X(X_j)\} + \log d\Lambda_Y(Y_j) - \log c(\alpha, H_X, \Lambda_Y)$$

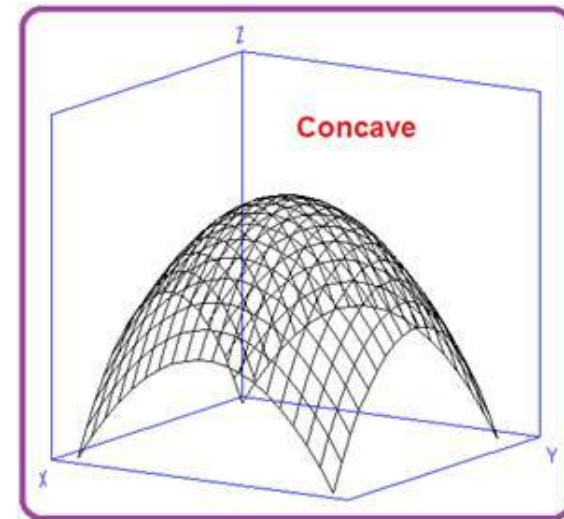
Requirements:

- twice differentiable with respect to α
- Unique maximum

Especially, one can apply “nlm” routine in R.

Convergence criteriat:

1. $2n-1$ scores are zero at $(\hat{\alpha}, \hat{H}_X, \hat{\Lambda}_Y)$
2. Hessian matrix at $(\hat{\alpha}, \hat{H}_X, \hat{\Lambda}_Y)$ is positive definite
(Hessian matrix are concave, it should be a global maxima)



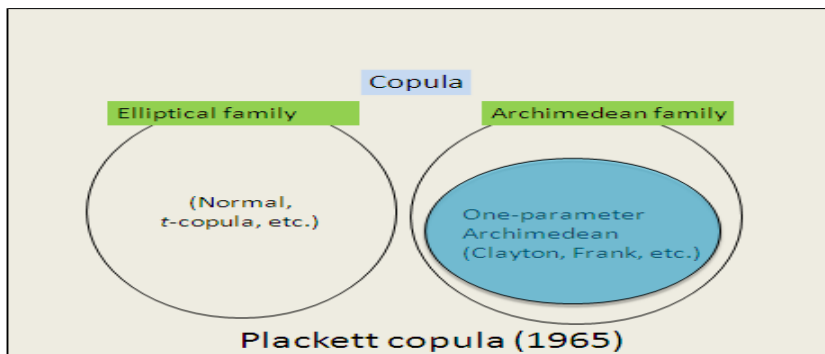
Proposed method

- The NPMLE $(\hat{\alpha}, \hat{H}_X, \hat{\Lambda}_Y)$ is **consistent & asymptotic normal**
- Observed Fisher information
= minus of the Hessian of $l_n(\alpha, H_X, \Lambda_Y)$

$$\hat{i}_n(\hat{\alpha}, \hat{H}_X, \hat{\Lambda}_Y) = \begin{bmatrix} \hat{i}_{n,11} & \hat{i}'_{n,12} \\ \hat{i}_{n,12} & \hat{i}_{n,22} \end{bmatrix}$$

- Consistent variance estimator, e.g.,

$$\hat{V}_n(\hat{\alpha}) \approx (\hat{i}_{n,11} - \hat{i}'_{n,12} \hat{i}_{n,22}^{-1} \hat{i}_{n,12})^{-1}$$



Simulation setting (I):

- **Plackett copula (not Archimedean family)**

$$C_{\alpha}[u, v] = \frac{1}{2(\alpha-1)} + \frac{u+v}{2} - \frac{[\{1 + (\alpha-1)(u+v)\}^2 - 4uv\alpha(\alpha-1)]^{1/2}}{2(\alpha-1)}$$

$$\alpha = 1/2.51, 1/5.11, 2.51, 5.11$$

(s.t. Spearman's rho = 0.25, 0.5, -0.25, -0.5)

- **Exponential margins**

$$H_X(x) = -\log(1 - e^{-1.5x})$$

$$\Lambda_Y(y) = 0.5y$$

- **Data generation:** $\Pr(X \leq x, Y > y) = C_{\alpha}[e^{-H_X(x)}, e^{-\Lambda_Y(y^-)}]$

If $X_j \leq Y_j$ then included in the sample. Otherwise truncated.

Repeat until we get n (=125 or 250) pair of (X_j, Y_j)

Positive dependence, 200 repetitions

<i>Parameter</i>		<i>Mean(Bias)</i>	<i>SE</i>	<i>SEE</i>	<i>95%Cov</i>
Spearman's $\rho = 0.25$ ($\alpha = 1/2.15$, $\Pr(X \leq Y) = 0.79$)					
$\log(\alpha) = -0.765$	$n = 125$	-0.778 (-0.013)	0.407	0.407	0.945
	$n = 250$	-0.697 (0.068)	0.311	0.296	0.965
$H_X(t) = 0.693$	$n = 125$	0.736 (0.043)	0.123	0.121	0.955
	$n = 250$	0.733 (0.040)	0.090	0.086	0.970
$\Lambda_Y(t) = 0.693$	$n = 125$	0.710 (0.017)	0.144	0.139	0.960
	$n = 250$	0.725 (0.032)	0.104	0.102	0.970
Spearman's $\rho = 0.50$ ($\alpha = 1/5.11$, $\Pr(X \leq Y) = 0.84$)					
$\log(\alpha) = -1.631$	$n = 125$	-1.642 (-0.011)	0.323	0.319	0.965
	$n = 250$	-1.652 (-0.021)	0.231	0.222	0.940
$H_X(t) = 0.693$	$n = 125$	0.726 (0.033)	0.101	0.092	0.910
	$n = 250$	0.716 (0.023)	0.067	0.064	0.920
$\Lambda_Y(t) = 0.693$	$n = 125$	0.704 (0.011)	0.110	0.102	0.960
	$n = 250$	0.701 (0.008)	0.068	0.069	0.950

Negative dependence, 200 repetitions

<i>Parameter</i>		<i>Mean(Bias)</i>	<i>SE</i>	<i>SEE</i>	<i>95%Cov</i>
Spearman's $\rho = -0.25$ ($\alpha = 2.15$, $\Pr(X \leq Y) = 0.72$)					
$\log(\alpha) = 0.765$	$n = 125$	0.859 (0.094)	0.598	0.554	0.960
	$n = 250$	0.717 (-0.048)	0.342	0.359	0.930
$H_X(t) = 0.693$	$n = 125$	0.809 (0.116)	0.313	0.244	0.960
	$n = 250$	0.717 (0.024)	0.139	0.138	0.935
$\Lambda_Y(t) = 0.693$	$n = 125$	0.793 (0.100)	0.363	0.267	0.960
	$n = 250$	0.699 (0.006)	0.139	0.137	0.930
Spearman's $\rho = -0.50$ ($\alpha = 5.11$, $\Pr(X \leq Y) = 0.70$)					
$\log(\alpha) = 1.631$	$n = 125$	1.758 (0.127)	0.818	0.598	0.915
	$n = 250$	1.708 (0.077)	0.534	0.386	0.955
$H_X(t) = 0.693$	$n = 125$	0.883 (0.190)	0.582	0.343	0.925
	$n = 250$	0.787 (0.094)	0.374	0.196	0.960
$\Lambda_Y(t) = 0.693$	$n = 125$	0.862 (0.169)	0.624	0.354	0.885
	$n = 250$	0.775 (0.082)	0.404	0.207	0.955

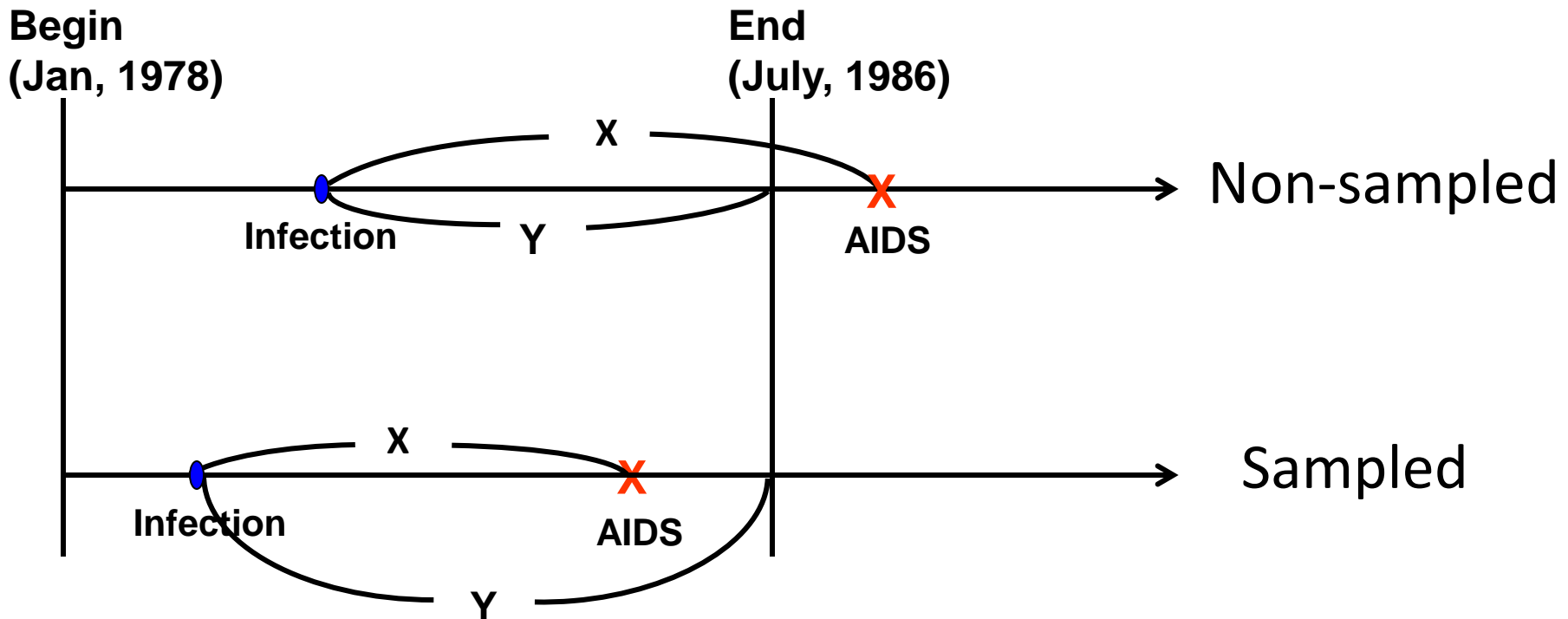
Data analysis

- **Transfusion-related AIDS (Kalbfleisch & Lawless, 1989, JASA)**

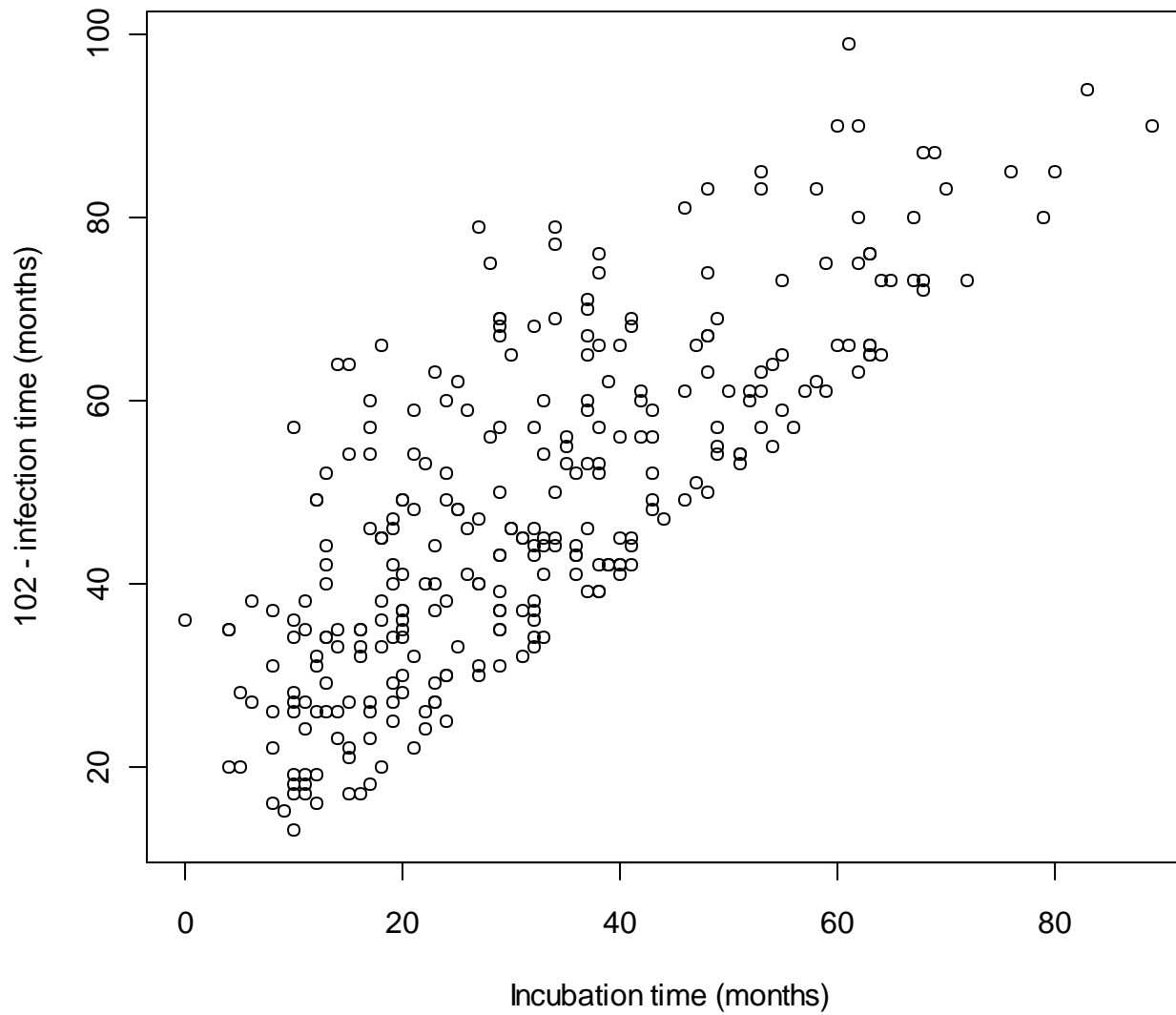
X : Time from infection to AIDS (month) ← Estimation

Y : 102 - time of infection (month)

n : sample size = 293



Transfusion-related AIDS data



Model selection

- $(K+1)$ candidate copulas

$$\begin{cases} C^{(0)}[u, v] = uv \\ C_{\alpha}^{(k)}[u, v], \quad k = 1, \dots, K \end{cases}$$

where $\lim_{\alpha \rightarrow 1} C_{\alpha}^{(k)}[u, v] = uv$

- Deviance: $C_{\alpha}^{(k)}$ vs. $C^{(0)}$

$$2\{l_n^{(k)}(\hat{\alpha}, \hat{H}_X, \hat{\Lambda}_Y) - l_n(1, \hat{H}_X^{\alpha=1}, \hat{\Lambda}_Y^{\alpha=1})\} \sim \chi_{df=1}^2$$

Step 1: Calculate deviances for K copulas

Step 2: Choose the copula with **smallest p-value** (<0.05)

Table 1: Analysis of the Transfusion-related AIDS data

$C_\alpha[u, v]$	Form ^(a)	$\hat{\alpha}$ (SE)	Kendall's τ on (X, Y)	95% CI for α	Deviance (p -value ^(b))
Clayton	Semi-survival	0.763 (0.033)	0.134	(0.701, 0.831)	19.028 (0.000)
	Regular	1.521 (0.172)	0.207	(1.218, 1.898)	8.568 (0.003)
	Survival	1.645 (0.233)	0.244	(1.246, 2.171)	5.228 (0.022)
Frank	Semi-survival	0.018 (0.014)	0.390	(0.004, 0.081)	10.828 (0.001)
Plackett	Semi-survival	0.189 (0.050)	0.356	(0.113, 0.316)	8.068 (0.005)
Normal	Semi-survival	-0.516 (0.083)	0.345	(-0.201, -0.831)	14.341(0.000)
t - (df=10)	Semi-survival	-0.520 (0.076)	0.350	(-0.234, -0.806)	9.559 (0.002)
t - (df=5)	Semi-survival	-0.507 (0.073)	0.344	(-0.223, -0.790)	3.959 (0.047)
Gumbel	Regular	1.459 (0.136)	0.315	(1.257, 1.821)	7.868 (0.005)
	Survival	1.340 (0.120)	0.254	(1.170, 1.678)	6.368 (0.012)
Two- parameter	Regular	$\hat{\alpha}$:1.521 (0.400) $\hat{\beta}$:1.000 ^(c)	0.207	α :(1.116, 3.348)	8.588 (0.003)
	Survival	$\hat{\alpha}$:1.344 (0.264) $\hat{\beta}$:1.235 (0.140)	0.309	α :(1.076, 2.551) β :(1.073, 1.756)	7.928 (0.019)

Smallest
P-value

(a) A Copula $C_\alpha[u, v]$ is used to model the distribution of (X, Y) in three different forms:

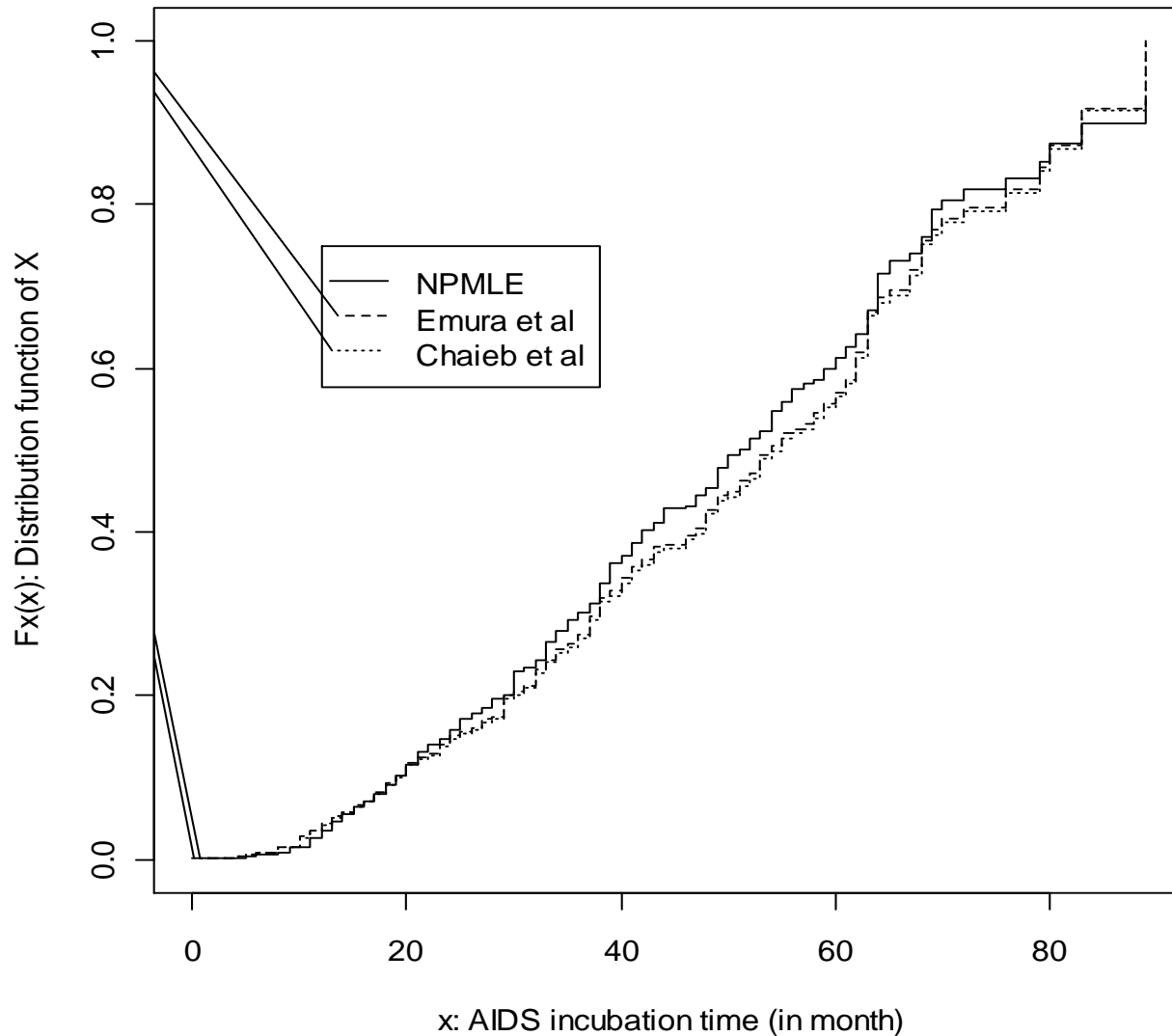
i) Semi-survival form: $\Pr(X \leq x, Y > y | X \leq Y) = C_\alpha[e^{-H_X(x)}, e^{-\Lambda_Y(y)}] / c,$

ii) Regular form: $\Pr(X \leq x, Y > y | X \leq Y) = (e^{-H_X(x)} - C_\alpha[e^{-H_X(x)}, 1 - e^{-\Lambda_Y(y)}]) / c,$

iii) Survival form: $\Pr(X \leq x, Y > y | X \leq Y) = (e^{-\Lambda_Y(y)} - C_\alpha[1 - e^{-H_X(x)}, e^{-\Lambda_Y(y)}]) / c.$

Under Clayton copula

$\hat{F}_X(x) = e^{-\hat{H}_X(x)}$: Time from infection to AIDS (month)



Extension to multiparameter copulas

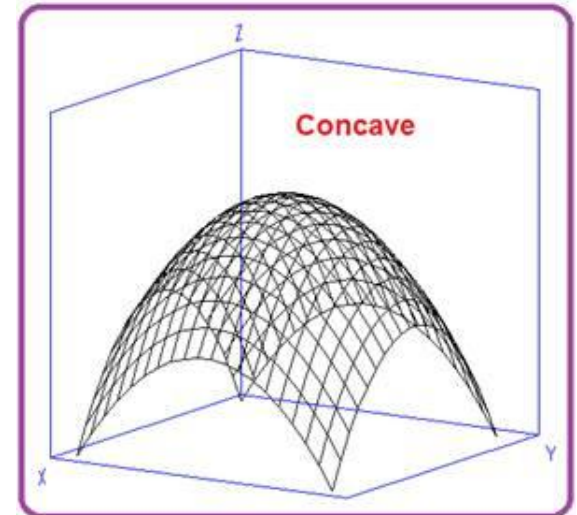
- In principle, quasi-Newton algorithm is applicable even when $\dim(\alpha) \geq 2$

$$l_n(\alpha, H_X, \Lambda_Y) =$$

$$\sum_{j=1}^n \log \eta_{\alpha}[H_X(X_j), \Lambda_Y(Y_j -)] + \log\{-dH_X(X_j)\} + \log d\Lambda_Y(Y_j) - \log c(\alpha, H_X, \Lambda_Y)$$

Requirements:

- Twice differentiable w.r.t α
- Unique maximum



Challenge: multiparameter copula

Simulation setting (II):

- **t -copula (multiparameter: $\alpha = (\rho, \nu)'$)**

$$C_{\rho, \nu}(u, v) = \Psi_{\rho} \{ \Psi^{-1}(u; \nu), \Psi^{-1}(v; \nu); \nu \},$$

$\Psi_{\rho}(\cdot; \nu)$: CDF of bivariate t with d.f. = ν

$\Psi(\cdot; \nu)$: CDF of t -distribution with d.f. = ν ,

$$\rho = -0.5, -0.25, 0.25, 0.5; \quad \nu = 5$$

- **Exponential margins**

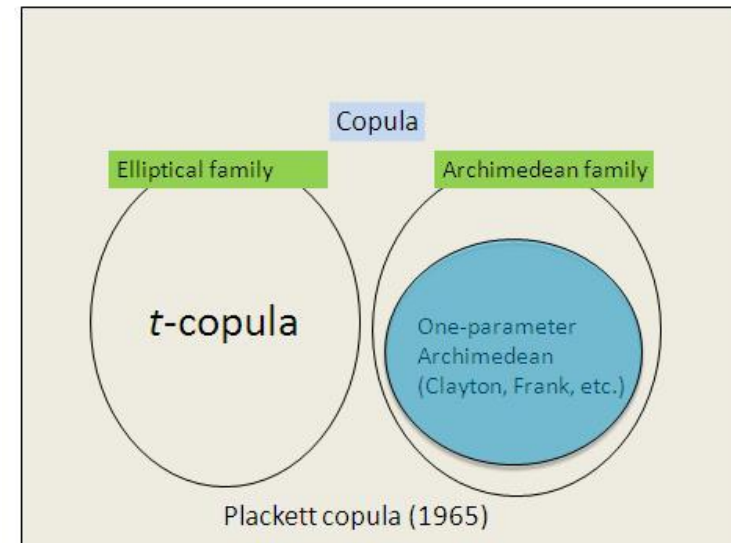
$$H_X(x) = -\log(1 - e^{-1.5x})$$

$$\Lambda_Y(y) = 0.5y$$

- **Data generation:** $\Pr(X \leq x, Y > y) = C_{\rho, \nu}[e^{-H_X(x)}, e^{-\Lambda_Y(y)}]$

If $X_j \leq Y_j$ then included in the sample. Otherwise truncated.

Repeat until we get n (=125 or 250) pair of (X_j, Y_j)



Challenge: multiparameter copula

- The maximum of $l_n(\rho, \nu, H_X, \Lambda_Y)$ may not be unique (?)

$$(\rho, \nu, -dH_X(X_1), \dots, -dH_X(X_n), d\Lambda_Y(Y_1), \dots, d\Lambda_Y(Y_n))$$

$$\# \text{ unknown} = 2n + 2$$



$$(\rho, \nu, \underbrace{-dH_X(X_{(1)}), \dots, -dH_X(X_{(n)})}_{\equiv 1}, d\Lambda_Y(Y_{(1)}), \dots, \underbrace{d\Lambda_Y(Y_{(n)})}_{\equiv 1})$$

$$\# \text{ unknown} = 2n$$

(Still # of parameters = # of observed points = $2n$)

Challenge: multiparameter copula

- Idea: Penalized NPMLE for $\dim(\alpha) \geq 2$

$$l_n(\alpha, H_X, \Lambda_Y) - \lambda \|\log(\alpha)\|^2,$$

where $\|\cdot\|$ is L_2 norm (Euclid norm),

and where $\lambda > 0$ is an arbitrary penalty parameter.

NOTE: $\lambda = \infty$:the NPMLE under independent

$\lambda = 0$:the NPMLE

- Choice of penalty parameter λ
As small as possible, e.g., $\lambda = 0.1$

Table 2: Simulation mean, standard error (SE), standard error estimate (SEE) and coverage rates of 95% confidence interval under the t-copula based on **10** runs., Penalty=0.1

Parameter	Mean	SE	SEE
Kendall's $\tau = 0.161$ ($\Pr(X \leq Y) = 0.79$)			
$\rho = -0.25$	-0.254	0.086	0.130
$\nu = 5$	6.150	5.307	5.175
$H_X(t) = 0.693$	0.676	0.095	0.098
$\Lambda_Y(t) = 0.693$	0.722	0.140	0.132
Kendall's $\tau = 0.333$ ($\Pr(X \leq Y) = 0.84$)			
$\rho = -0.50$	-0.510	0.073	0.102
$\nu = 5$	4.315	3.185	3.726
$H_X(t) = 0.693$	0.685	0.088	0.094
$\Lambda_Y(t) = 0.693$	0.734	0.152	0.122
Kendall's $\tau = -0.161$ ($\Pr(X \leq Y) = 0.73$)			
$\rho = 0.25$	0.189	0.122	0.144
$\nu = 5$	5.507	2.627	3.857
$H_X(t) = 0.693$	0.714	0.123	0.154
$\Lambda_Y(t) = 0.693$	0.737	0.284	0.195
Kendall's $\tau = -0.333$ ($\Pr(X \leq Y) = 0.72$)			
$\rho = 0.50$	0.466	0.146	0.132
$\nu = 5$	5.815	3.587	3.367
$H_X(t) = 0.693$	0.973	0.655	0.715
$\Lambda_Y(t) = 0.693$	0.962	0.720	0.707

Summary

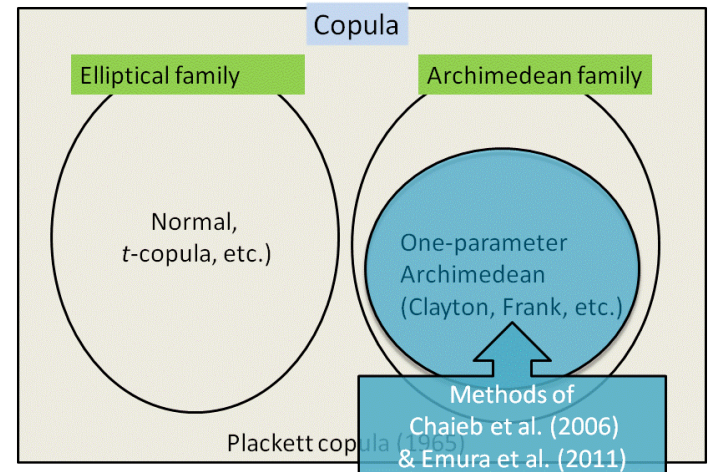
We proposed the NPMLE under semi-survival copula with reverse-time hazard modeling

Advantage of the proposed NPMLE

- Broader class of copula than existing methods

(Chaieb et al.(2006) & Emura et al. (2011))

- Equip Copula model selection via likelihood comparison



Disadvantage

- NPMLE is computationally very demanding (numerical maximization in high-dimensional space)

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Thank you for your kind attention