



Analyzing Survival Data with Competing Risks Using R

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경쟁위험자료란?
누적발생함수의 추정
비례위험 모형
좌절단 자료
시간변동 공변량 자료



OUTLINE

다변량 생존자료의 분류

자료 유형	순서 여부	
	Yes	No
동일한 유형	재발사건자료	군집자료
다른 유형	준경쟁위험자료	경쟁위험자료

Melanoma data

- ▶ Data on patients after operation for malignant melanoma collected at Odense University Hospital
- ▶ **cause**: 0: alive, 1: dead from melanoma, 2: dead from other cause
- ▶ **time**: survival time
- ▶ **ulcer**: ulceration (0: absent, 1: present)
- ▶ **thickness**: tumour thickness (mm)
- ▶ **sex**: 0: female, 1: male
- ▶ **age**

available at library(MASS)

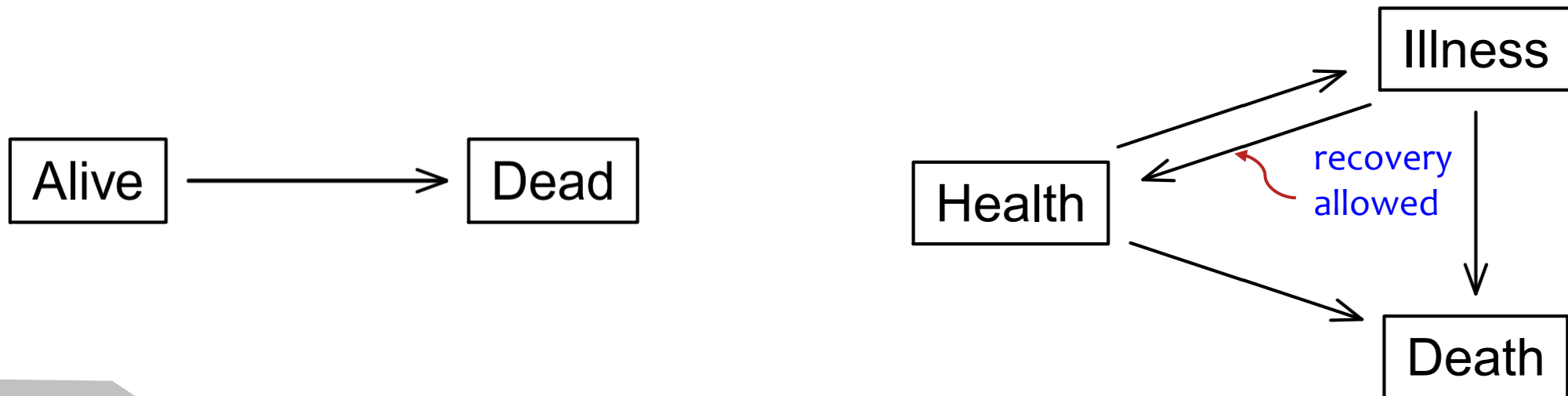
Hospital acquired pneumonia (HAP) in ICU data

- ▶ Assess the impact of HAP on ICU mortality
- ▶ id
- ▶ **start**: start of the observation time
- ▶ **stop**: failure time
- ▶ **status**: censoring status
- ▶ **event**: event type (2: death in ICU, 3: discharge alive)
- ▶ **pneu**: nosocomial pneumonia indicator (**time-varying**)
- ▶ adm.cens.exit: **administrative censoring times**
- ▶ age
- ▶ sex

available at library(kmi)

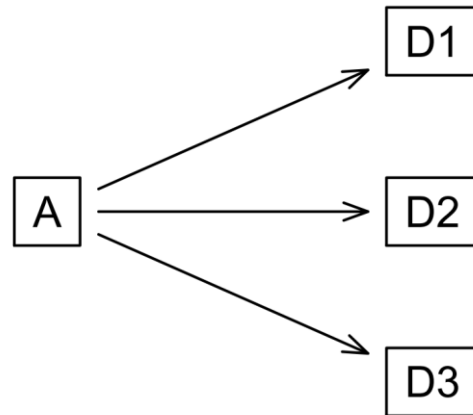
Multi-state model

- ▶ $K(\geq 2)$ 개의 상태를 가진 모형
- ▶ 생존자료 모형: 두 가지 상태(0: alive, 1: dead)를 가진 모형
 - ▶ 1: absorbing (흡수) state
- ▶ Illness-death model: 3 states (0: health, 1: illness, 2: death)



Competing risks model

- ▶ absorbing state가 여러 개인 multi-state model
- ▶ 한 사건의 발생은 다른 사건의 발생을 중도절단시킴
- ▶ Remarks
 - ▶ 어떤 사건의 발생이 다른 사건의 발생을 중도절단시키지만, 그 역은 성립하지 않는 경쟁위험을 준경쟁위험 (semi-competing risks) 이라고 함



Cause-specific hazard (CSH)

▶ $(T = \min(T_1, T_2), e \in \{1, 2\})$

▶ T_k : k -번째 원인에 의한 사건 발생 시간

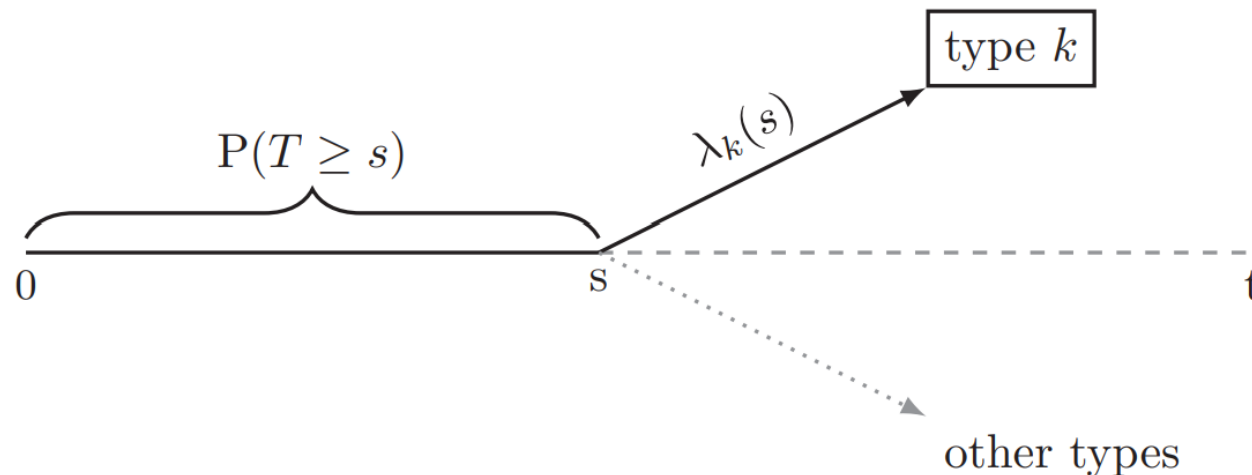
▶ $\lambda_k(t) = \lim_{\Delta t \downarrow 0} \frac{P(t \leq T < t + \Delta t, e = k | T \geq t)}{\Delta t}, k = 1, 2$: CSH

▶ $\lambda(t) = \sum_{k=1}^2 \lambda_k(t) = \lim_{\Delta t \downarrow 0} \frac{P(t \leq T < t + \Delta t | T \geq t)}{h}$: overall hazard

▶ $\Lambda_k(t) = \int_0^t \lambda_k(s) ds$: cumulative CSH (CCSH)

○ ○ ○ Cumulative incidence function (CIF) ○ ○ ○

- ▶ $F_k(t) = P(T \leq t, e = k) = \int_0^t S(s-) \lambda_k(s) ds$: CIF. t 이내에 k -번째 원인에 의해 사건이 발생할 확률
 - ▶ $S(t) = P(T > t) = \exp\left\{-\int_0^t \lambda(s) ds\right\}$: overall survival function



▶ Remarks refer to #8

▶ $\lambda_k(t) = \frac{F'_k(t)}{S(t)} \Rightarrow F'_k(t) = S(t)\lambda_k(t)$

▶ $F_1(t) + F_2(t) = 1 - S(t), \forall t$

▶ $F_k(\infty) = P(e = k) < 1$ 이므로 F_k 를 “subdistribution function”이라고 함

Observed data

- ▶ $\{(x_1, e_1 \delta_1), \dots, (x_n, e_n \delta_n)\}$
 - ▶ $x_i = \min(t_i, c_i)$, $\delta_i = I(t_i \leq c_i)$, $e_i \in \{1, 2\}$
 - ▶ c_i : 중도절단 시간 (only known under administrative censoring)
- ▶ $t_{(1)} < \dots < t_{(r)}$: unique ordered uncensored times
- ▶ d_{ik} : # of patients failing from cause k at $t_{(i)}$
- ▶ $d_i (= \sum_{k=1}^2 d_{ik})$: total # of patients failing from any cause at $t_{(i)}$
- ▶ n_i : # of patients at risk at $t_{(i)}$

Nelson-Aalen (NA) estimator

- ▶ $\hat{\lambda}_k(t_{(i)}) = \frac{d_{ik}}{n_i}$: estimated CSH
- ▶ $\hat{\Lambda}_k(t) = \sum_{t_{(i)} \leq t} \hat{\lambda}_k(t_{(i)})$: NA estimator

Aalen-Johansen (AJ) estimator

- ▶ $\hat{F}_k(t) = \sum_{m:t_{(m)} \leq t} \hat{S}(t_{(m)} -) \frac{d_{mk}}{n_m}$: estimated CIF
- ▶ $\hat{S}(t_{(m)} -) = \sum_{l:t_{(l)} < t_{(m)}} \left(1 - \frac{d_l}{n_l}\right)$: PL estimator of the probability of being free **of any event** just prior to $t_{(m)}$

Remarks on CIF estimation

- ▶ 만약 competing risks를 중도절단으로 간주한다면 $F_1(t)$ 를 **과대 추정**함
 - ▶ $S_1(t) = \exp\left\{-\int_0^t \lambda_1(s)ds\right\}$: probability of being free from **cause 1** just prior to t
 - ▶ $S_1(t) = \exp\left\{-\int_0^t \lambda_1(s)ds\right\} \geq \exp\left\{-\int_0^t (\lambda_1(s) + \lambda_2(s))ds\right\} = S(t)$
 $\Rightarrow \int_0^t S_1(s-) \lambda_1(s) ds \geq \int_0^t S(s-) \lambda_1(s) ds = F_1(t)$

Melanoma data

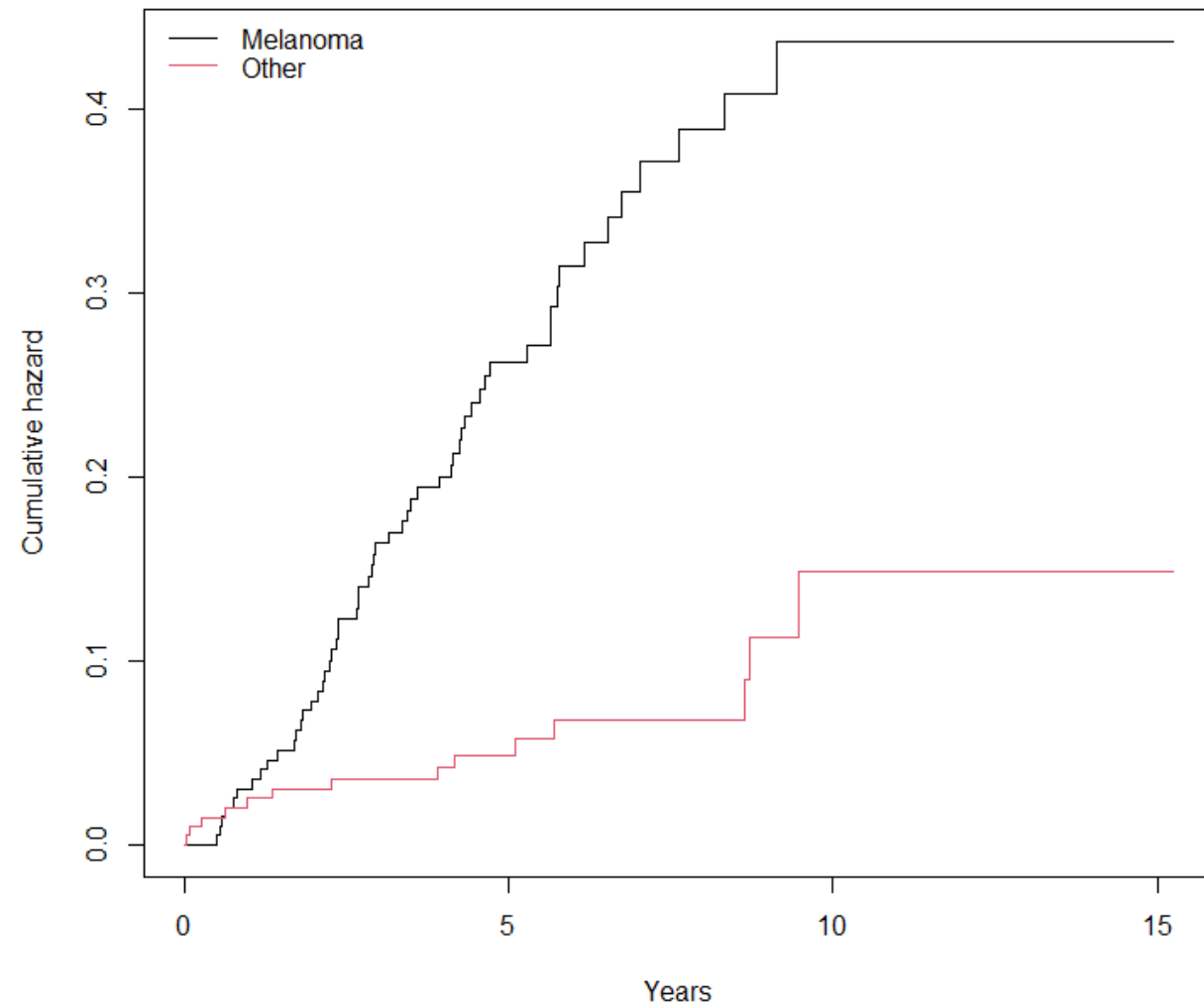
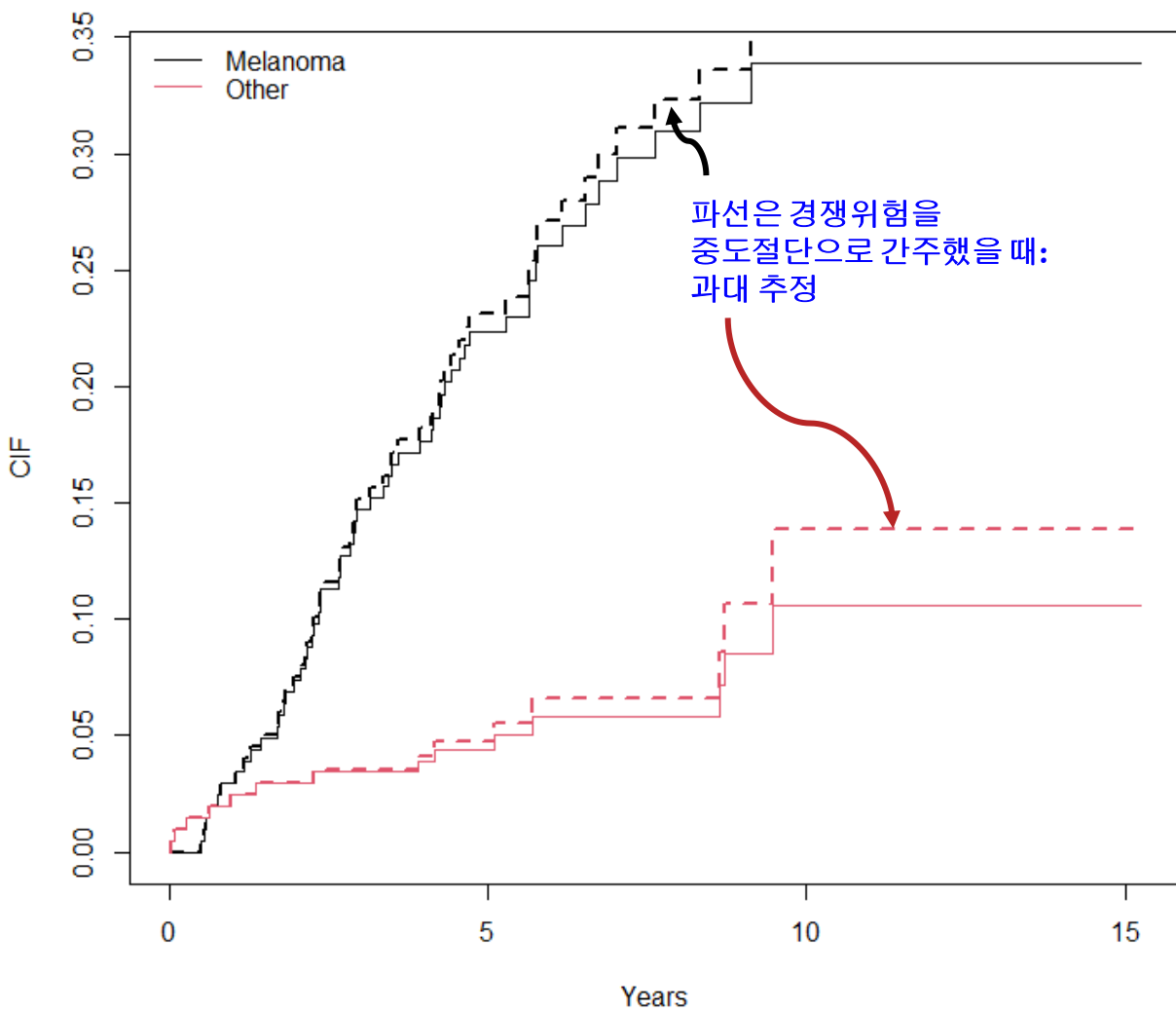
- ▶ 피부암 환자 256명을 대상으로 사망 원인을 조사:
- ▶ 중도절단: 134 (0), 피부암: 57 (1), 다른 원인: 14 (2)

```
> Melanoma[inx, c(1, 3:4, 6:8)]
      time  sex age thickness  ulcer cause
3 0.09582478 male 41    1.34 absent    0
5 0.50650240 male 52   12.08 present    1
1 0.02737851 male 76    6.76 present    2
```

Using R

- ▶ For NA: `survfit{survival}`, `mvna{mvna}`, ...
- ▶ For AJ: `survfit{survival}`, `cuminc{cmprsk}`, `etmCIF{etm}`, `Cuminc{mstate}`, ...

Estimated CCSH & CIF



Subdistribution hazard (SDH)

▶ $\lambda_k^*(t) \equiv -(\log(1 - F_k(t)))' \stackrel{\text{refer to \#10}}{=} \frac{F_k'(t)}{1 - F_k(t)}$: SDH. k -번째 subdistribution function에 대응하는 위험함수로 정의

▶ Remarks

▶ $\lambda_k^*(t) \stackrel{\text{refer to \#8}}{=} \lim_{\Delta t \downarrow 0} \frac{P(t \leq T_k < t + \Delta t | T \geq t \cup (T \leq t, e \neq k))}{\Delta t}$: SDH의 다른 정의

▶ 아직 사건이 발생하지 않은 환자 뿐만 아니라 이미 경쟁위험으로 사망한 환자도 위험 집합에 포함됨

Subdistribution hazard (SDH)

- ▶ $\Lambda_k^*(t) = \int_0^t \lambda_k^*(s) ds$: cumulative SDH (CSDH)
- ▶ $F_k(t) = 1 - \exp\left\{-\int_0^t \lambda_k^*(s) ds\right\}$: CIF의 다른 표현
- ▶ Remarks
 - ▶ $P(e = k) = F_k(\infty) = 1 - \exp(-\Lambda_k^*(\infty))$
 - ▶ $P(e = 1) + P(e = 2) = 1 \Leftrightarrow \exp(-\Lambda_1^*(\infty)) + \exp(-\Lambda_2^*(\infty)) = 1$
 - ▶ **SDHs cannot be independently modeled!**

CSH와 SDH의 관계

$$\blacktriangleright \lambda_1(t) = \left(1 + \frac{F_2(t)}{S(t)}\right) \lambda_1^*(t)$$

$$\blacktriangleright 1 - F_1(t) = 1 - P(T \leq t, e = 1) = P(T > t) + P(T \leq t, e = 2) \\ = S(t) + F_2(t)$$

- ▶ SDH는 CSH보다 **weighted down** 되고, 가중값은 시간에 따라 변하고 **competing events**에도 의존함

Estimation: SDH & CIF

- ▶ n_i^* : **estimated** # at risk at $t_{(i)} \Rightarrow n_i \leq n_i^*$
- ▶ $\hat{\lambda}_k^*(t_{(i)}) = \frac{d_{ik}}{n_i^*}$: estimated SDH
- ▶ $\hat{F}_k(t) = 1 - \prod_{i:t_{(i)} \leq t} (1 - \hat{\lambda}_k^*(t_{(i)}))$: estimated CIF (PL-form)

○○○ Contribution of individual l to n_i^* ○○○

- ▶ still at risk at $t_{(i)}$: 1
- ▶ competing event observed at $t_{(l)}$ before $t_{(i)}$:

$$P\{C \geq t_{(i)} | C \geq t_{(l)}\}$$

- ▶ $\Gamma(t) = P(C > t)$: censoring time distribution
- ▶ $\hat{\Gamma}(t)$: Kaplan-Meier estimate of $\Gamma(t)$ by reversing the role of T and C

- ▶ estimated as $\frac{\hat{\Gamma}(t_{(i)}^-)}{\hat{\Gamma}(t_{(l)}^-)}$



Using R



- ▶ For SDH: `survfit{survival}` along with `crprep{mstate}`
counting process format

Estimated weights

```
> maxT
```

```
[1] 9.138946
```

```
> EX
```

```
pid      time status2
3         3 0.09582478      0
5         5 0.50650240      1
114      114 5.70841889      2
```

pid=114 환자의 사망 이후 피부암으로 사망한 시점:

```
[1] 5.757700 5.771389 6.176591 6.537988 6.754278 7.022587 7.616701 8.328542 9.138946
```

```
> WT
```

	pid	Tstart	Tstop	status	weight.cens	count	failcode
117	3	0.000000	0.09582478	0	1.0000000	1	1
176	5	0.000000	0.50650240	1	1.0000000	1	1
528	114	0.000000	5.70841889	2	1.0000000	1	1
529	114	5.708419	5.75770021	2	0.9890110	2	1
530	114	5.757700	5.77138946	2	0.9778985	3	1
531	114	5.771389	6.17659138	2	0.8992170	4	1
532	114	6.176591	6.53798768	2	0.8536870	5	1
533	114	6.537988	6.75427789	2	0.7960055	6	1
534	114	6.754278	7.02258727	2	0.7374757	7	1
535	114	7.022587	7.61670089	2	0.6780018	8	1
536	114	7.616701	8.32854209	2	0.6295731	9	1
537	114	8.328542	9.13894593	2	0.4519325	10	1

```
> ipcw
```

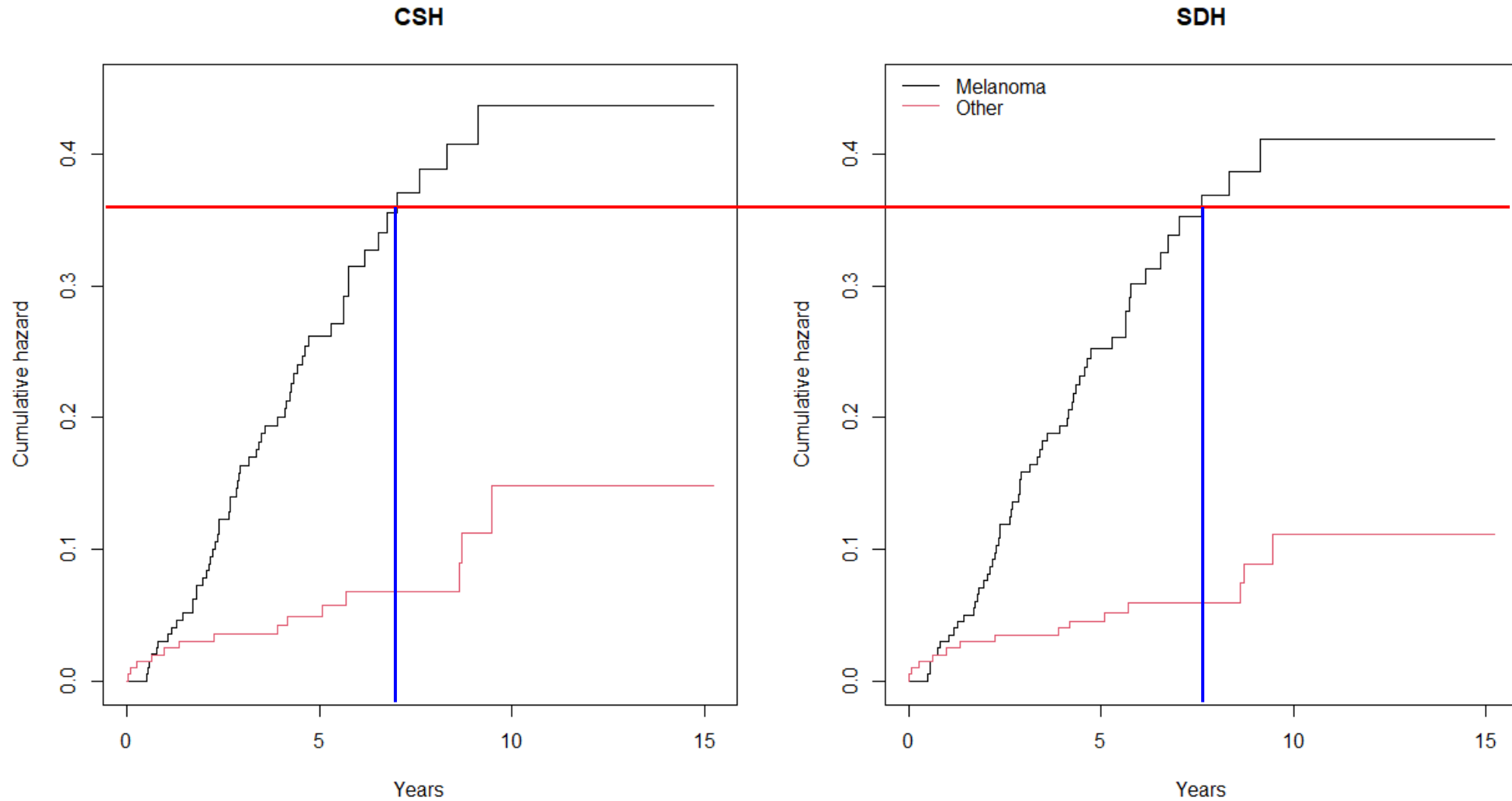
```
[1] 0.4519325
```


위험 집합 비교: n vs. n^*

```
> Comp[seq(1,194,15),]
```

	Time	nrisk.CS	nrisk.me1.SD	nrisk.oth.SD
[1,]	0.02737851	205	205.00000	205.00000
[2,]	1.44832307	189	194.99015	198.00000
[3,]	2.64750171	174	180.99015	197.00000
[4,]	4.12320329	159	166.94021	194.09573
[5,]	4.47364819	144	152.08521	177.05698
[6,]	4.90896646	125	131.69660	160.71688
[7,]	5.31690623	109	115.66296	139.06402
[8,]	5.64544832	94	100.66296	125.05327
[9,]	6.40383299	78	84.48061	97.98169
[10,]	7.03627652	62	67.14693	84.13023
[11,]	8.62970568	47	50.43076	70.86973
[12,]	9.26762491	32	32.00000	48.46820
[13,]	10.86379192	14	14.00000	14.00000

CCSH vs. CSDH



CIF 비교: AJ vs. FG

> Equiv[seq(1,194,15),]

	Time	CIF.me1.AJ	CIF.me1.FG	CIF.oth.AJ	CIF.oth.FG
[1,]	0.02737851	0.00000000	0.00000000	0.004878049	0.004878049
[2,]	1.44832307	0.04902198	0.04902198	0.029364888	0.029364888
[3,]	2.64750171	0.11765274	0.11765274	0.034267085	0.034267085
[4,]	4.12320329	0.18141214	0.18141214	0.039169283	0.039169283
[5,]	4.47364819	0.20689223	0.20689223	0.044197790	0.044197790
[6,]	4.90896646	0.22353960	0.22353960	0.044197790	0.044197790
[7,]	5.31690623	0.23013963	0.23013963	0.050456445	0.050456445
[8,]	5.64544832	0.24528498	0.24528498	0.050456445	0.050456445
[9,]	6.40383299	0.26936596	0.26936596	0.058111429	0.058111429
[10,]	7.03627652	0.29832966	0.29832966	0.058111429	0.058111429
[11,]	8.62970568	0.32177917	0.32177917	0.058111429	0.058111429
[12,]	9.26762491	0.33871751	0.33871751	0.085379085	0.085379085
[13,]	10.86379192	0.33871751	0.33871751	0.105947064	0.105947064

Proportional CSH models

- ▶ $\{(x_1, e_1 \delta_1, z_1), \dots, (x_n, e_n \delta_n, z_n)\}$: observed data
 - ▶ z_i : a vector of p covariates (risk factors)
 - ▶ eg: sex, treatment, age, tumour size, stage of cancer, blood measures, ... (**fixed**)
- ▶ Proportional CSH model:

$$\lambda_k(t|z_i) = \lambda_{0k}(t) \exp(\beta'_k z_i), k = 1, 2$$

CSH approach

- ▶ Use the partial likelihood principle

- ▶ $L(\beta_1, \beta_2) = \prod_{k=1}^2 L_k(\beta_k),$

$$L_k(\beta_k) = \prod_{i=1}^n \left\{ \frac{\exp\{\beta_k' z_i\}}{\sum_{j=1}^n I(x_j \geq x_i) \exp(\beta_k' z_j)} \right\}^{I(\delta_i e_i = k)}$$

- ▶ NPMLE: $\hat{\beta}_k$ as the solution of $\frac{\partial}{\partial \beta_k} \log L_k(\beta_k) = 0$

Using R

- ▶ Use `coxph{survival}`
- ▶ or `coxph{survival}` along with `mstep{mstate}` **stacked-long format**

```
Data:
  id from to trans Tstart      Tstop      time status age sex  ulcer
1   1   1  2     1     0 0.02737851 0.02737851     0  76 male present
2   1   1  3     2     0 0.02737851 0.02737851     1  76 male present
5   3   1  2     1     0 0.09582478 0.09582478     0  41 male absent
6   3   1  3     2     0 0.09582478 0.09582478     0  41 male absent
9   5   1  2     1     0 0.50650240 0.50650240     1  52 male present
10  5   1  3     2     0 0.50650240 0.50650240     0  52 male present

  thickness failcode
1         6.76 melanoma
2         6.76   Other
5         1.34 melanoma
6         1.34   Other
9        12.08 melanoma
10       12.08   Other
```

CS model: two analyses

call:

```
coxph(formula = Surv(time, cause == 1) ~ age + ulcer + thickness +  
sex, data = Melanoma)
```

	coef	exp(coef)	se(coef)	z	p
age	0.012198	1.012273	0.008297	1.470	0.14150
ulcerpresent	1.164479	3.204253	0.309751	3.759	0.00017
thickness	0.108945	1.115101	0.037734	2.887	0.00389
sexmale	0.432817	1.541594	0.267410	1.619	0.10554

Call:

```
coxph(formula = Surv(time, cause == 2) ~ age + ulcer + thickness +  
sex, data = Melanoma)
```

	coef	exp(coef)	se(coef)	z	p
age	0.07255	1.07525	0.02167	3.348	0.000813
ulcerpresent	0.10937	1.11557	0.59128	0.185	0.853256
thickness	0.04958	1.05083	0.08794	0.564	0.572891
sexmale	0.35801	1.43048	0.54859	0.653	0.514012

CS model: one analysis

Call:

```
coxph(formula = Surv(time, status) ~ age * strata(failcode) +  
      ulcer * strata(failcode) + thickness * strata(failcode) +  
      sex * strata(failcode), data = M.s)
```

	coef	exp(coef)	se(coef)	z	p
age	0.012198	1.012273	0.008297	1.470	0.14150
ulcerpresent	1.164479	3.204253	0.309751	3.759	0.00017
thickness	0.108945	1.115101	0.037734	2.887	0.00389
sexmale	0.432817	1.541594	0.267410	1.619	0.10554
age:strata(failcode)other	0.060354	1.062212	0.023203	2.601	0.00929
strata(failcode)other:ulcerpresent	-1.055112	0.348153	0.667501	-1.581	0.11395
strata(failcode)other:thickness	-0.059365	0.942363	0.095693	-0.620	0.53501
strata(failcode)other:sexmale	-0.074806	0.927923	0.610293	-0.123	0.90244

Likelihood ratio test=58.28 on 8 df, p=1.015e-09

n= 410, number of events= 71

SDH approach

- ▶ Modeling the CIF for failure from cause 1 conditional on the covariates, $F(t; z) = P(T \leq t, e = 1|z)$ [Fine & Gray, 1999]
- ▶ Proportional SDH model: $\lambda_1^*(t|z_i) = \lambda_{01}^*(t) \exp(\gamma_1' z_i)$
- ▶ $L_1(\gamma_1) =$

$$\prod_{i=1}^n \left\{ \frac{\exp\{\gamma_1' z_i\}}{\sum_{j=1}^n \left\{ I(x_j \geq x_i) + I(t_j < x_i < c_j, \delta_j e_j = 2) \frac{\hat{\Gamma}(x_i^-)}{\hat{\Gamma}(t_j^-)} \right\} \exp(\gamma_1' z_j)} \right\}^{I(\delta_i e_i = 1)}$$

1 under administrative censoring

Using R

- ▶ Use `crr{cmprsk}`
- ▶ or use `coxph{survival}` along with `crprep{mstate}`

```
> crp.r[crp.r$pid %in% c(3, 162, 171), ]
```

pid	Tstart	Tstop	status	weight.cens	age	sex	ulcer	thickness	count	failcode	
45	3	0.000000	0.09582478	0	1.0000000	41	male	absent	1.34	1	1
369	162	0.000000	8.71184120	2	1.0000000	49	male	absent	6.12	1	1
370	162	8.711841	9.13894593	2	0.8139535	49	male	absent	6.12	2	1
379	171	0.000000	9.13894593	1	1.0000000	60	female	present	3.22	1	1
416	3	0.000000	0.09582478	0	1.0000000	41	male	absent	1.34	1	2
909	162	0.000000	8.71184120	2	1.0000000	49	male	absent	6.12	1	2
918	171	0.000000	9.13894593	1	1.0000000	60	female	present	3.22	1	2
919	171	9.138946	9.46748802	1	0.8235294	60	female	present	3.22	2	2

FG model: two analyses

call:

```
crr(ftime = Melanoma$time, fstatus = Melanoma$cause, cov1 = covs,  
    failcode = 1, cencode = 0)
```

	coef	exp(coef)	se(coef)	z	p-value
age	0.00593	1.01	0.00929	0.638	0.5200
ulcer	1.12863	3.09	0.30344	3.719	0.0002
thickness	0.08999	1.09	0.03836	2.346	0.0190
sex	0.40503	1.50	0.27558	1.470	0.1400

call:

```
crr(ftime = Melanoma$time, fstatus = Melanoma$cause, cov1 = covs,  
    failcode = 2, cencode = 0)
```

	coef	exp(coef)	se(coef)	z	p-value
age	0.0570	1.059	0.0142	4.011	0.00006
ulcer	-0.1092	0.897	0.5866	-0.186	0.85000
thickness	0.0114	1.012	0.0851	0.135	0.89000
sex	0.2630	1.301	0.5923	0.444	0.66000

FG model: one analysis

```
call:
coxph(formula = Surv(Tstart, Tstop, status == failcode) ~ age *
      strata(failcode) + ulcer * strata(failcode) + thickness *
      strata(failcode) + sex * strata(failcode), data = crp.r,
      weights = weight.cens)
```

n= 953, number of events= 71

	coef	exp(coef)	se(coef)	robust se	z	Pr(> z)	
age	0.005928	1.005945	0.008026	0.009028	0.657	0.511465	
ulcerpresent	1.128630	3.091418	0.313844	0.300809	3.752	0.000175	***
thickness	0.089995	1.094168	0.038720	0.034614	2.600	0.009324	**
sexmale	0.405032	1.499350	0.268754	0.267154	1.516	0.129494	
age:strata(failcode)failcode=2	0.051030	1.052354	0.020666	0.015508	3.291	0.001000	***
strata(failcode)failcode=2:ulcerpresent	-1.237810	0.290019	0.668887	0.643621	-1.923	0.054455	.
strata(failcode)failcode=2:thickness	-0.078550	0.924456	0.093643	0.087838	-0.894	0.371184	
strata(failcode)failcode=2:sexmale	-0.142072	0.867559	0.610014	0.635984	-0.223	0.823232	

Left-truncated (late entry) data

- ▶ observed: $(x_i, e_i \delta_i, z_i) \rightarrow (x_i, e_i \delta_i, z_i, l_i)$, l_i : late entry
- ▶ For CSH: $I(x_i \geq t) \rightarrow I(x_i \geq t > l_i)$
- ▶ For SDH:

$$I(x_i \geq t) + I(t_i < t < c_i, \delta_i e_i = 2) \times \frac{\Gamma(t-)}{\Gamma(t_i-)} \rightarrow$$

$$I(x_i \geq t > l_i) + I(l_i \vee t_i < t < c_i, \delta_i e_i = 2) \times \frac{\Gamma(t-)}{\Gamma(t_i-)} \times \frac{\Phi(t-)}{\Phi(t_i-)}$$

- ▶ $\Phi(t) = P(L_i \leq t)$: estimated by reversing time

Time-varying covariates

- ▶ For CSH: `coxph{survival}` along with `tmerge{survival}`
counting process format
- ▶ For SDH: `cox.kmi{kmi}` with **imputed** data set [Ruan & Gray, 2008]

HAP in ICU data

	id	start	stop	status	pneu	adm.cens.exit	age	sex	outcome
131	23371	0	4	0	0	4	63.05753	F	0
132	23371	4	29	1	1	29	63.05753	F	2
159	30236	0	6	0	0	6	38.96164	M	0
160	30236	6	26	0	1	26	38.96164	M	0
265	1002890	0	4	1	0	358	76.77918	F	3
983	1021700	0	51	0	0	51	62.48219	F	0
993	1022004	0	7	0	0	7	2.09589	F	0
994	1022004	7	15	1	1	48	2.09589	F	3
1068	2001524	0	7	1	0	7	70.94475	F	2

pneumonia	event type			total
	censored	death in the unit	discharge alive	
yes	5	21	82	108
no	16	126	1063	1205
total	21	147	1145	1313

CSH model

```
coxph(formula = Surv(start, stop, outcome == 2) ~ pneu + sex +  
      age, data = icu.pneu)
```

	coef	exp(coef)	se(coef)	z	p
pneu1	-0.087804	0.915941	0.248345	-0.354	0.723673
sexM	-0.020834	0.979382	0.169716	-0.123	0.902299
age	0.019762	1.019958	0.005312	3.720	0.000199

FG model via imputation

```
cox.kmi(formula = Surv(start, stop, outcome == 2) ~ pneu + sex +  
      age, imp.data = imp.death)
```

Pooled estimates:

	coef	exp(coef)	se(coef)	t	p
pneu1	1.06374450	2.897199	0.24071329	4.4191348	9.909827e-06
sexM	0.11047993	1.116814	0.16978135	0.6507188	5.152280e-01
age	0.01827515	1.018443	0.00508217	3.5959344	3.232297e-04

FG model with complete censoring

```
coxph(formula = Surv(start, adm.cens.exit, outcome == 2) ~ pneu +  
      sex + age, data = icu.pneu)
```

	coef	exp(coef)	se(coef)	z	p
pneu1	1.064077	2.898162	0.240640	4.422	9.79e-06
sexM	0.112463	1.119031	0.169652	0.663	0.507390
age	0.018454	1.018625	0.005084	3.630	0.000284

References

- ▶ Beyersmann J, Allignol A, Schumacher M (2012). Competing risks and multistate models with R. Springer, New York.
- ▶ Geskus, RB (2015). Data analysis with competing risks and intermediate states. CRC press.



THANK YOU!!!