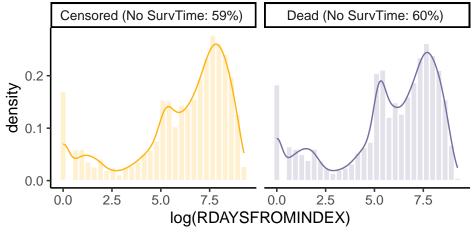
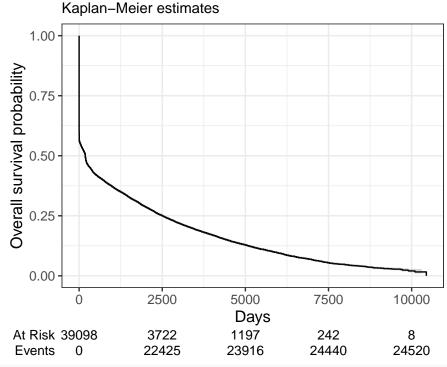
CS7

2023-04-10

```
### EDA
load("DataCleaned.RData")
data.subject <- data.impute %>% group by(RSUBJID) %>% filter(row number() == n()) %>%
  mutate(DEATH2 = ifelse(DEATH == 0, "Censored", "Dead")) %>%
  ungroup()
df.InstantDeath <- data.subject %>% group_by(DEATH2) %>%
  summarise(InstantDeath = mean(RDAYSFROMINDEX == 0)) %>%
  mutate(InstantDeath = str_c(round(InstantDeath*100, 0), "%"))
# Distribution of log-RDAYSFROMINDEX in the two groups
inner_join(data.subject, df.InstantDeath) %>%
  mutate(DEATH2 = str_c(DEATH2, " (No SurvTime: ", InstantDeath, ") ")) %>%
  ggplot() +
  geom_histogram(aes(log(RDAYSFROMINDEX), ..density.., fill = DEATH2),
                 color = "white", alpha = 0.2) +
  geom_density(aes(log(RDAYSFROMINDEX), color = DEATH2)) +
  scale_fill_manual(values = pal(2)) +
  scale_color_manual(values = pal(2)) +
  facet wrap(~DEATH2) +
  theme(legend.position = "none")
```





```
fig1 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ HXSMOKE, data = data.subject) %>%
  ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
      fill = "Smoke", color = "Smoke")
fig2 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ GENDER, data = data.subject) %>%
  ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
      fill = "Gender", color = "Gender")
fig3 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ DPPCI, data = data.subject) %>%
  ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
       fill = "DPPCI", color = "DPPCI")
fig4 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ ACS, data = data.subject) %>%
  ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
      fill = "ACS", color = "ACS")
fig5 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ HXCHF, data = data.subject) %%
  ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
      fill = "HXCHF", color = "HXCHF")
fig6 <- survfit2(Surv(RDAYSFROMINDEX + 0.1, DEATH) ~ DPCABG, data = data.subject) %>%
```

```
ggsurvfit() +
  add_confidence_interval() +
  labs(x = "Days", y = "Survival probability",
        fill = "DPCABG", color = "DPCABG")
ggarrange(fig1, fig2, fig3, fig4, fig5, fig6, nrow = 2, ncol = 3)
                                       1.00
                                                                           1.00
20.75 O.75 O.25 O.00
                                    20.75 Survival probability
                                                                        Survival probability
0.20.0
0.25
   0.00
                                       0.00
                                                                           0.00
                                                      5000
                                                            7500 10000
                                                                                          5000 7500 10000
            2500
                  5000
                       7500 10000
                                                2500
                                                                                     2500
                                                                                0
                  Days
                                                      Days
                                                                                           Days
           Smoke __ 0 __ 1
                                               Gender — 0 — 1
                                                                                DPPCI — 0 — 1 — 2
3.00 Survival probability 0.75 0.50 0.25 0.00
                                       1.00
                                                                           1.00
                                    Survival probability
                                                                        Survival probability
   0.00
                                       0.00
                                                                           0.00
                  5000 7500 10000
                                                                                          5000 7500 10000
                                                2500 5000 7500 10000
            2500
                                                                                     2500
        0
                                            Ö
                                                                                0
                  Days
                                                                                          Days
                                                      Days
  ACS 0 1 2 3 -
                                               HXCHF 0 1
                                                                              DPCABG — 0 — 1 — 2
# Cox regression model
surv_obj <- Surv(time = data.subject$RDAYSFROMINDEX+0.1, event = data.subject$DEATH,</pre>
                    type = "right")
model.cox <- coxph(surv_obj ~ .-RDAYSFROMINDEX-RSEQCATHNUM-RSUBJID-DEATH-DEATH2,</pre>
                       data = data.subject)
model.cox %>%
  tbl_regression(exp = TRUE)
```

Characteristic	HR	95% CI	p-value
AGE_G	1.01	1.01, 1.01	< 0.001
GENDER			
0			
1	1.07	1.03, 1.12	< 0.001
$RACE_G$			
1			
2	0.94	0.91, 0.98	0.002
3	0.88	0.82, 0.94	< 0.001
ACS			
0			
1	1.37	1.31, 1.43	< 0.001

Characteristic	HR	95% CI	p-value
$\frac{2}{2}$	1.00	0.95, 1.05	>0.9
3	1.67	1.56, 1.80	< 0.001
4	1.09	1.05, 1.12	< 0.001
CHFSEV	1.07	1.06, 1.09	< 0.001
DPCABG	1.01	1.00, 1.00	(0.001
0		_	
1	0.27	0.26, 0.29	< 0.001
2	0.88	0.84, 0.92	< 0.001
DPMI		,	
0			
1	0.63	0.60, 0.67	< 0.001
2	1.08	1.04, 1.13	< 0.001
DPPCI			
0	_		
1	0.38	,	< 0.001
2	0.85	0.78, 0.91	< 0.001
HXANGINA			
0			
1	0.77	0.74, 0.80	< 0.001
HXCEREB			
0			.0.001
1	1.09	1.05, 1.13	< 0.001
HXCHF			
0	0.74	0.71 0.79	<0.001
HXCOPD	0.74	0.71, 0.78	< 0.001
0			
1	1.16	1.11, 1.21	< 0.001
HXDIAB	1.10	1.11, 1.21	\0.001
0			
1	1.13	1.10, 1.16	< 0.001
HXHTN	1.10	1110, 1110	10.001
0		_	
1	0.95	0.93, 0.98	0.002
HXHYL		,	
0	_	_	
1	0.68	0.66, 0.70	< 0.001
HXMI			
0	_	_	
1			
HXSMOKE			
0	_		
1	1.02	0.99, 1.05	0.2
NUMPRMI	0.95	0.93, 0.97	< 0.001
DIASBP_R	0.99	0.99, 0.99	< 0.001
PULSE_R	1.00	1.00, 1.00	< 0.001
SYSBP_R	1.00	1.00, 1.00	0.3
CBRUITS			
0	1 19	1 00 1 17	ZO 001
1 HEICHT R	1.13	1.09, 1.17	< 0.001
HEIGHT_R S3	1.00	1.00, 1.00	0.8
DU			

Characteristic	HR	95% CI	p-value
0		_	
1	1.17	1.10, 1.23	< 0.001
WEIGHT R	1.0	0.99, 1.00	< 0.001
CREATININE R	1.01	1.01, 1.02	< 0.001
CATHAPPR —		,	
0		_	
1	0.51	0.44, 0.60	< 0.001
2	0.99	0.89, 1.09	0.8
3	0.91	0.82, 1.01	0.070
DIAGCATH			
0			
1	1.14	1.04, 1.24	0.003
INTVCATH			
0		_	
1	0.61	0.55, 0.67	< 0.001
CORDOM			
1	_	_	
2	0.87	0.82, 0.92	< 0.001
3	0.95	0.87, 1.03	0.2
LADST	1.00	1.00, 1.00	< 0.001
LCXST	1.00	1.00, 1.00	< 0.001
LMST	1.00	1.00, 1.00	< 0.001
LVEF_R	0.99	0.99, 0.99	< 0.001
NUMDZV	0.86	0.83, 0.88	< 0.001
PRXLADST	1.00	1.00, 1.00	0.067
RCAST	1.00	1.00, 1.00	< 0.001
DAYS2LKA	1.00	1.00, 1.00	< 0.001
DSCABG	1.24	1.20, 1.28	< 0.001
DSMI	1.22	1.15, 1.30	< 0.001
DSPCI	1.19	1.08, 1.32	< 0.001
DSSTROKE	1.16	1.11, 1.21	< 0.001