# Survival Analysis

2024.09.10

### Textbooks

Klein, J.P. and
 Moeschberger, M.L.
 (2003) Survival
 analysis: Techniques
 for censored and
 truncated data. 2nd
 Edition, Springer,
 New York.

Data sets: R package "Kmsurv"

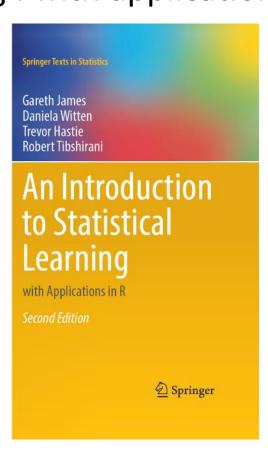


## Textbooks (Lasso, Tree based method)

1. Gareth James, Daniela Witten, Trevor Hastie, Robert Tibshirani. (2013). An introduction to statistical learning: with applications in R. New

York :Springer (Chapter 11)

https://www.statlearning.com/



## Website for the Class materials

https://github.com/lhchien-ndhu/113-1-Survival-Analysis.git

# Examples

## Right censored data

## 1.11 Times to Death for Patients with Cancer of the Tongue

A study was conducted on the effects of ploidy on the prognosis of patients with cancers of the mouth. Patients were selected who had a paraffin-embedded sample of the cancerous tissue taken at the time

of surgery. Follow-up survival data was obtained on each patient. The tissue samples were examined using a flow cytometer to determine if the tumor had an aneuploid (abnormal) or diploid (normal) DNA profile using a technique discussed in Sickle–Santanello et al. (1988). The data in Table 1.6 is on patients with cancer of the tongue. Times are in weeks.

The data is used in exercises.

tongue (KMsurv)

#### data from Section 1.11

#### Description

The tongue data frame has 80 rows and 3 columns.

#### Format

This data frame contains the following columns:

type

Tumor DNA profile (1=Aneuploid Tumor, 2=Diploid Tumor)

time

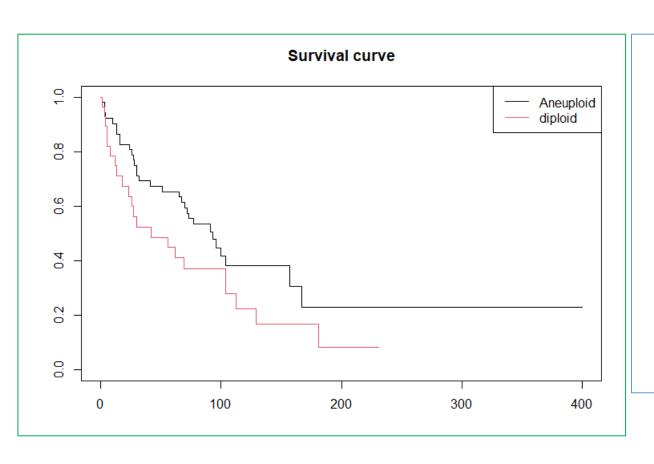
Time to death or on-study time, weeks

delta

Death indicator (0=alive, 1=dead)

> tongue

## Right censored data



tongue (KMsurv)

data from Section 1.11

Description

The tongue data frame has 80 rows and 3 columns.

Format

This data frame contains the following columns:

type

Tumor DNA profile (1=Aneuploid Tumor, 2=Diploid Tumor)

time

Time to death or on-study time, weeks

delta

Death indicator (0=alive, 1=dead)

## Left truncation data

## 1.16 Death Times of Elderly Residents of a Retirement Community

Channing House is a retirement center located in Palo Alto, California. Data on ages at death of 462 individuals (97 males and 365 females) who were in residence during the period January 1964 to July 1975 has been reported by Hyde (1980). A distinctive feature of these individuals was that all were covered by a health care program provided by the center which allowed for easy access to medical care without any additional financial burden to the resident. The age in months when members of the community died or left the center and the age when individuals entered the community is available on the authors' web site.

The life lengths in this data set are *left truncated* because an individual must survive to a sufficient age to enter the retirement community. Individuals who die at an early age are excluded from the study. Ignoring this left truncation leads to the problem of length-biased sampling. The concept of left truncation and the bias induced into the estimation process by ignoring it is discussed in section 3.4.

This data will be used in section 4.6 to illustrate how one estimates the conditional survival function for left-truncated data. The data is used in section 7.3 to illustrate the comparison of two samples (male and female), when there is left truncation and right censoring employing the log-rank test, and in Chapter 9 employing the Cox proportional hazards model.

channing (KMsurv)

#### data from Section 1.16

#### Description

The channing data frame has 462 rows and 6 columns.

#### Format

This data frame contains the following columns:

obs

Observation number

death

Death status (1=dead, 0=alive)

ageentry

Age of entry into retirement home, months

age

Age of death or left retirement home, months

time

Difference between the above two ages, months

gender

Gender (1=male, 2=female)

# > head(channing,10) obs death ageentry age time gender 1 1 1 1042 1172 130 2 2 2 1 921 1040 119 2 3 3 1 885 1003 118 2 4 4 1 901 1018 117 2 5 5 1 808 932 124 2 6 6 1 915 1004 89 2 7 7 1 901 1023 122 2 8 8 1 852 908 56 2 9 9 1 828 868 40 2 10 10 1 968 990 22 2

## Competing event

• 1.13 Bone Marrow Transplantation for Leukemia

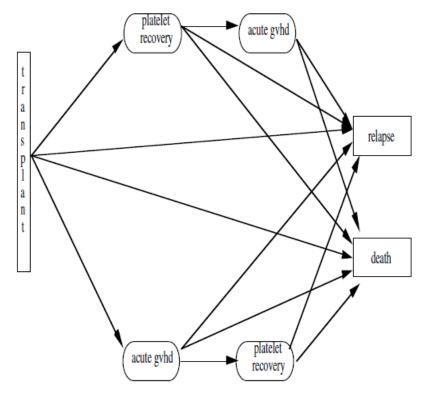


Figure 1.1 Recovery Process from a Bone Marrow Transplant