# Lecture twenty-two: Sample Size Determination

#### 1. The courses on design of clinical trials in the department

- (a) PQHS450: Clinical Trials and Intervention Studies.
- (b) EPBI446: Experimental Design for Biomedical Sciences

### 2. Software for Power Analysis and Sample Size Determination

- (a) Free one: http://stattools.crab.org (SWOG).
- (b) Commecial ones: nQuery Advisor; Power and Precision, etc.
- (c) SAS (version 9 and up): PROC POWER; PROC GLMPOWER.
- (d) Group sequential inference: EaSt @www.cytel.com.
- (e) https://www.trialdesign.org/ (Yuan & Lee from MD Anderson).

#### 3. The Aspects of the Design of a Clinical Trial

- (a) Primary/secondary questions
- (b) Study population: Eligibility, inclusion/exclusion criteria
- (c) Study designs:
  - i. Fundamental strategies in design: Randomization, blocking, stratification, balancing, and replication.
  - ii. Special designs:Latin square; factorial design; longitudinal studies with repeated measurements; cross-over trials, etc;
  - iii. Observational studies
- (d) Randomization
- (e) Blindness
- (f) Power and sample size
- (g) Design based on fixed sample size vs a (group) sequential design
- (h) Interim analysis and early stopping rules

## 4. Why power analysis and sample size (SS) calculation?

#### 5. SS requirement for a survival study with fixed SS

- (a) **Recall**: factors to determine the sample size of two group t-test of equal means: Significance level,  $\alpha$ , 1 or 2 sided test, means  $\mu_1$  and  $\mu_2$  (the difference between the two means), common standard error  $\sigma$ , power  $1 \beta$ .
- (b) Distinguishing between two treatment groups

Assuming PH model for the survival times

$$h_N(t) = \psi h_S(t),$$

where  $\psi$  is unknown hazard ratio. Let  $\theta = \log \psi$  be the log-hazard ratio, then

survival is longer under new treatment if  $\theta < 0$ 

survival is longer under standard treatment if  $\theta > 0$ 

no treatment difference if  $\theta = 0$ .

In order to test the null hypothesis  $H_0: \theta = 0$ , we have to choose a test statistic, for example, log-rank, test from Cox model, a test from parametric model (eg. exponential distribution), which are valid under certain assumptions.

(c) The required number of deaths:

Since we usually can not measure the acual survival times of censored observations, it is the number of acual deaths that is important, rather than the total number of subjects. What does the KM estimate look like if there is no event?

- i. The number of deaths depends on
  - A. The reference value of  $\theta$ :  $\theta_R$ , which reflects the magnitude of the treatment difference to be detected.
  - B. Type I error  $\alpha$ : the probability of rejecting  $H_0$  when  $H_0$  is true (false positive).
  - C. Type II error  $\beta$ : the probability of not rejecting  $H_0$  when  $H_0$  is false (false negative).
  - D.  $1 \beta$ : Power (probability to reject  $H_0$  when  $H_0$  is not true) to detect the difference.
  - E. The test statistic chosen

ii. The total required number of deaths for the two arms is

$$d = \frac{4(z_{\alpha/2} + z_{\beta})^2}{\theta_R^2},$$

where  $z_{\alpha/2}$ ,  $z_{\beta}$  are the upper  $\alpha/2$ , and upper  $\beta$  points, respectively of the standard normal distribution, N(0,1). If we define  $c(\alpha, \beta) = (z_{\alpha/2} + z_{\beta})^2$ , then

$$d = 4c(\alpha, \beta)/\theta_R^2$$
.

If the two treatment arms have unequal number of patients, say, with proportion  $\pi$ ,  $(1-\pi)$  in each group, then the formula becomes

$$d = \frac{c(\alpha, \beta)}{\pi (1 - \pi)\theta_R^2}.$$

- iii. The derivation of the required number of deaths (section 15.2.1)
  - A. based on log-rank test
  - B. Use several approximations
  - C. The estimate of the number of deaths based on the formula derived tends to be underestimated.
- iv. Example 15.1: Survival from chronic active hepatitis
  - A. The KM estimate of survival of patients under standard treatment:

Median = 3.3 years

$$S(2) = 0.7, S(4) = 0.45$$

$$S(5) = 0.41, S(6) = 0.25$$

- B. For similar population (eg. age, sex, race, disease status, ect), the expected survival rate at 5 years is 0.60 under the new treatment.
- C. Assume proportional hazard, then

$$S_N(t) = [S_S(t)]^{\psi},$$

thus,

$$\psi_R = \frac{\log(0.6)}{\log(0.41)} = 0.57,$$

and 
$$\theta_R = \log \psi_R = \log(0.57) = -0.563$$
.

D. given  $\alpha = 0.05$ ,  $\beta = 0.1$ ,  $c(\alpha, \beta) = 10.51$ . Therefore the required number of deaths is

$$d = \frac{4 \times 10.51}{0.563^2} = 133$$

- E.  $S_N(t)$  can be estimated under PH assumption
- (d) The required number of patients
  - i. The required number of patients depends on
    - A. The accrual period: say, length a
    - B. follow-up period: say length f
    - C. the total duration of a study is a + f
    - D. The death rate, p(death),
  - ii. Thus, the total required number of patients is

$$n = \frac{d}{p(death)},$$

iii. The death rate can be estimated by

$$p(death) = 1 - \frac{1}{6} \{ \bar{S}(f) + 4\bar{S}(0.5a + f) + \bar{S}(a + f) \},$$

where

$$\bar{S}(t) = \frac{S_S(t) + S_N(t)}{2},$$

The death rate formula is for equal number of patients in the two treatment arms.

- iv. The derivation of the probability of death p(death): section 15.3.1.
- v. Example 15.2: Survival from chronic active hepatitis (cont.)
  - A. Accrual period: 18 months (a = 18)
  - B. (subsequent) Fellow-up period: 24 months (f = 24)
  - C. The death rate:

$$p(death) = 1 - \frac{1}{6} \{ \bar{S}(24) + 4\bar{S}(33) + \bar{S}(42) \},$$

D. Based on estimated survival functions

$$\bar{S}(24) = \frac{S_S(24) + S_N(24)}{2} = \frac{0.7 + 0.82}{2} = 0.76,$$

$$\bar{S}(33) = \frac{S_S(33) + S_N(33)}{2} = \frac{0.57 + 0.73}{2} = 0.65,$$

$$\bar{S}(42) = \frac{S_S(42) + S_N(42)}{2} = \frac{0.45 + 0.63}{2} = 0.54,$$

- E. P(death) = 0.35, and the required number of patients is n = 133/0.35 = 380
- F. what if accrual period: 18 months (a=18) and (f=0), then P(death)=0.155, and the required number of patients is n=133/0.155=858
- vi. An approximate procedure
  - A. The average probability of survival beyond  $\tau$  is  $\frac{S_S(\tau)+S_N(\tau)}{2}$ .
  - B. The probability of death, in the period from the time origin to  $\tau$ , can be approximated by

$$1 - \frac{S_S(\tau) + S_N(\tau)}{2}.$$

C. The required number of patients is

$$n = \frac{2d}{2 - S_S(\tau) - S_N(\tau)},$$

- D. The choice of  $\tau$ : The average length of the follow-up: f+a/2.
- E. Example 15.3: