

생물학적 동등성시험에서 사용되는
통계방법에 대한 고찰

강 승 호

이화여자대학교 통계학과

* Bioavailability of a drug:

the rate and extent to which the active ingredient is absorbed and becomes available at the site of drug action

* brand-name drug vs generic drug

* ANDA

* AUC (Area Under the Curve)

C_0, C_1, \dots, C_k : plasma concentration obtained at time $0, t_1, \dots, t_k$

$$AUC(0-t_k) = \sum_{i=2}^k \frac{C_{i-1} + C_i}{2} (t_i - t_{i-1})$$

$$AUC(0-\infty) = AUC(0-t_k) + C_k/k_e$$

k_e : elimination rate

Assumption:

When two drugs products are equivalent in the rate and extent to site of drug action, it is assumed that they will be therapeutically equivalent.

* The comparison of the first moment (mean) of the distribution of the pharmacokinetic parameters (say, $AUC(0-\infty)$) for the two drug products refers to the comparison of "average bioavailability".

* Parallel Design vs Crossover Design

* intersubject variability

intrasubject variability

* Crossover design removes the intersubject variability and use only intra-subject variability.

* 2 x 2 Crossover Design

We wish to compare formulation A and B.

We make two sequences.

sequence 1 : A \rightarrow B

sequence 2 : B \rightarrow A

We assign $n_1 + n_2$ subjects randomly.

n_1 subjects are assigned to sequence 1

n_2 subjects are assigned to sequence 2

period1 washout period2

sequence 1	A	.	B
sequence 2	B	.	A

* carryover effect

* Statistical Model

$$Y_{ijk} = \mu + S_{ik} + P_j + F_{j,k} + C_{(j-1,k)} + e_{ijk}$$

where

μ : overall mean

S_{ik} : random effect of the i th subject
in the k th sequence

P_j : fixed effect of j th period

$F_{j,k}$: drug effect

$C_{(j-1,k)}$: carryover effect

e_{ijk} : random error

$$S_{ik} \sim N(0, \sigma_s^2)$$

$$e_{ijk} \sim N(0, \sigma_e^2)$$

S_{ik} and e_{ijk} are independent.

* 2 x 2 Crossover Design

sequence	period I	period II
1 (RT)	$\mu + P_1 + F_R$	$\mu + P_2 + F_T + C_R$
2 (TR)	$\mu + P_1 + F_T$	$\mu + P_2 + F_R + C_T$

* Test for Carryover Effects

Consider sum of observations in two periods.

$$U_{ik} = Y_{1k} + Y_{2k} \quad : \quad i = 1, \dots, n_k, \quad k = 1, 2$$

$$U_{i1} \sim N(2\mu + C_R, \sigma_u^2), \quad i = 1, \dots, n_1$$

$$U_{i2} \sim N(2\mu + C_T, \sigma_u^2), \quad i = 1, \dots, n_1$$

where

$$\sigma_u^2 = 2(2\sigma_s^2 + \sigma_e^2)$$

$$\text{Let } \overline{U}_{.k} = n_k^{-1} \sum_{i=1}^{n_k} U_{ik}, \quad k = 1, 2$$

$$\widehat{\sigma}_u^2 = \sum_{k=1}^2 \sum_{i=1}^{n_k} (U_{ik} - \overline{U}_{.k})^2$$

$$T_c = \frac{\overline{U}_{.2} - \overline{U}_{.1}}{\widehat{\sigma}_u \sqrt{n_1^{-1} + n_2^{-1}}}$$

Carryover effect exists if

$$|T_c| > t(\alpha/2, n_1 + n_2 - 2)$$

* Test for Average Bioequivalence

Responses such as AUC are usually positively skewed.

→ Not normal distribution

Take Log transformation!!!

Let $Y_{ijk}^* = \ln(Y_{ijk})$

$$\overline{Y_R^*} = 2^{-1}(\overline{Y_{.11}} + \overline{Y_{.22}})$$

$$\overline{Y_T^*} = 2^{-1}(\overline{Y_{.12}} + \overline{Y_{.21}})$$

The 90% confidence interval is (L_1, U_1) , where

$$L_1 = (\overline{Y_T^*} - \overline{Y_R^*})$$

$$- t(\alpha, n_1 + n_2 - 2) \widehat{\sigma}_d \sqrt{n_1^{-1} + n_2^{-1}}$$

$$U_1 = (\overline{Y_T^*} - \overline{Y_R^*})$$

$$+ t(\alpha, n_1 + n_2 - 2) \widehat{\sigma}_d \sqrt{n_1^{-1} + n_2^{-1}}$$

We conclude that two formulations are average bioequivalent if

$$L_1 > \ln(0.8) = -0.2231 \text{ and}$$

$$U_1 < \ln(1.25) = 0.2231$$

* Westlake's symmetric confidence interval

* Confidence interval based on Fieller's theorem

* Chow and Shao's confidence interval

— Interval Hypotheses Testing —

* Usual hypothesis testing

$$H_0: \mu_T = \mu_R \text{ vs } H_1: \mu_T \neq \mu_R$$

* In Bioequivalence

$$H_0: \mu_T \neq \mu_R \text{ vs } H_1: \mu_T = \mu_R$$

$$H_0: \mu_T - \mu_R \leq \theta_L \text{ or } \mu_T - \mu_R \geq \theta_U$$

$$\text{vs } H_1: \theta_L < \mu_T - \mu_R < \theta_U$$

where θ_L and θ_U are some clinically meaningful limits.

* The above hypotheses are decomposed into two sets of one-sided hypotheses.

$$H_{01}: \mu_T - \mu_R \leq \theta_L \text{ vs } H_{11}: \mu_T - \mu_R > \theta_L$$

and

$$H_{02}: \mu_T - \mu_R \geq \theta_U \text{ vs } H_{12}: \mu_T - \mu_R < \theta_U$$

If we reject H_{01} and H_{02} , then we conclude $\theta_L < \mu_T - \mu_R < \theta_U$

* Schuirman's Two One-Sided Procedure

We conclude that two formulations are bioequivalent if

$$\frac{(\overline{Y_T^*} - \overline{Y_R^*}) - \theta_L}{\widehat{\sigma}_d \sqrt{n_1^{-1} + n_2^{-1}}} > t(a, n_1 + n_2 - 2)$$

and

$$\frac{(\overline{Y_T^*} - \overline{Y_R^*}) - \theta_U}{\widehat{\sigma}_d \sqrt{n_1^{-1} + n_2^{-1}}} < -t(a, n_1 + n_2 - 2)$$

* Nonparametric Methods

When assumption of normality (for raw data or for log-transformed data) is violated, let

$$d_{ik} = 2^{-1}(Y_{2k} - Y_{1k}),$$

$$i = 1, \dots, n_k, \quad k = 1, 2$$

"period difference"

Let

$$b_{hik} = d_{ik} - \theta_h, \quad h = L, U$$

for subjects in sequence 1

$$b_{hik} = d_{ik}$$

for subjects in sequence 2

For a fixed $h = L, U$, b_{h1} and b_{h2} have the same distribution except for the mean difference.

We can construct Wilcoxon rank sum test based on the ranks of b_{h1} and

b_{h2} .

$R(b_{Lik}) = \text{rank of } b_{Lik} \text{ in the combined sample } b_{Lik}, \quad i = 1, \dots, n_k, \quad k = 1, 2$

$$W_L = R_L - 2^{-1} n_1 (n_1 + 1)$$

We reject H_{01} if $W_L > w(1 - \alpha)$

* Power of Schuirman's two one-sided t test procedure

* The conclusion of bioequivalence based on only average bioavailability may be misleading.

We need to investigate the assessment of equivalence in variabilities of bioavailability between two formulations.