A post hoc analysis of intra-subject coefficients of variation in pharmacokinetic measures to calculate optimal sample sizes for bioequivalence studies

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pISSN: 2289-0882 eISSN: 2383-5427 Because bioequivalence studies are performed using a crossover design, information on the intrasubject coefficient of variation (intra-CV) for pharmacokinetic measures is needed when determining the sample size. However, calculated intra-CVs based on bioequivalence results of identical generic drugs produce different estimates. In this study, we collected bioequivalence results using public resources from the Ministry of Food and Drug Safety (MFDS) and calculated the intra-CVs of various generics. For the generics with multiple bioequivalence results, pooled intra-CVs were calculated. The estimated intra-CVs of 142 bioequivalence studies were 14.7 \pm 8.2% for AUC and 21.7 \pm 8.8% for C_{max}. Intra-CVs of C_{max} were larger than those of area under the concentrationtime curve (AUC) in 129 studies (90.8%). For the 26 generics with multiple bioequivalence results, the coefficients of variation of intra-CVs between identical generics (mean \pm sd (min ~ max)) were 38.0 \pm 24.4% (1.9 ~ 105.3%) for AUC and 27.9 \pm 18.2 % (4.0 ~ 70.1%) for C_{max}. These results suggest that substantial variation exists among the bioequivalence results of identical generics. In this study, we presented the intra-CVs of various generics with their pooled intra-CVs. The estimated intra-CVs calculated in this study will provide useful information for planning future bioequivalence studies.

Introduction

One of the most important considerations in planning a bioequivalence study is the determination of the sample size and its associated power.[1-4] Statistically, power represents the probability the null hypothesis will be rejected when the alternative hypothesis is true.[5-7] Since the null hypothesis in bioequiva-

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lence studies is that the substances are bioinequivalent, the power of a bioequivalence study is the probability of proving bioequivalence when the products are in fact bioequivalent.[5, 7,8] Because finding the optimal sample size ensures adequate power, the sample size calculation is one of the most important steps in designing a bioequivalence study. Sample sizes that are too large increase the cost of the study and unnecessarily expose many subjects to the drug. In contrast, sample sizes that are too small increase the type 2 error and may result in study failure. According to the statistical guidelines of the U.S. FDA and EMA, 80% or 90% power is recommended for bioequivalence studies.[9]

The determination of the sample size requires information on the intra-subject coefficient of variation (intra-CV) of pharmacokinetic measures. However, the calculated intra-CVs of identical generics vary considerably among studies. For example, the reported intra-CVs of metformin's maximum concentration (C_{max}) were 12.1% and 24.8% in two different bioequivalence studies.[10] These results suggested that choosing a sample size based on a single bioequivalence result can be insufficient to achieve adequate power for planning a trial.

The Ministry of Food and Drug Safety (MFDS) of Korea has released the results of bioequivalence studies to the public since January 2014.[11] These data include information for power and sample size calculations in bioequivalence studies (i.e., 90% confidence intervals for the area under the concentration-time curve (AUC) and C_{max} , and sample sizes). These data also show that there has been considerable variability in the sample sizes for bioequivalence studies on the same generic drugs.

To aid in designing bioequivalence studies, this study aimed to investigate appropriate sample sizes by analyzing the intra-CV of AUC and C_{max} from 142 bioequivalence results of 58 generic drugs obtained from public resources provided by the MFDS of Korea.

Methods

Study data

The data for the analysis were obtained from the public bioequivalence results database on the Ministry of Food and Drug Safety's (MFDS) homepage (http://www.mfds.go.kr/).[11] A total of 183 bioequivalence study results published from Jan 2015 to Nov 2015 were considered for analysis. Among 183 bioequivalence results, 41 results from fixed-dose combinationdrugs were excluded to avoid statistical complications. The 142 analyzed bioequivalence studies were performed with a standard two period, two sequence crossover design involving fasting, healthy male volunteers.

Statistical analysis

Using the PowerTOST package (ver. 1.2-08) in the R statistical program (ver. 3.1.3), the intra-CV, post-hoc power and appropriate sample size needed for bioequivalence studies to attain more than 80% and 90% power were calculated with the equations below:[2, 8, 12-15]. For sample size calculation, the larger of the two intra-CVs from AUC or Cmax was used.

Point estimate (PE) based on a confidence interval (CI)

$$= \sqrt{CI_{low}} * CI_{high}$$

Margin of error on the log scale = $2^* \left(\frac{\Delta_{GL}}{\sqrt{\frac{1}{n_1} + \frac{1}{n_2}} * t_{1^- 2a,n! + n2^-}}\right)^2$

(t: t-values of the student t distribution; α : probability of type 1 error; n1 and n2: sample sizes of each group)

Intra-CV (%) = $100^{*}\sqrt{e^{MSE}-1}$

Sample size (N) $\geq \frac{2^{*LN(1+CV^2)*(Z_{\lfloor \underline{B}^{+Z_{-a}})^2}}{In(1.25)^2}}{In(1.25)^2}$

(α : probability of type 1 error; β : probability of type 2 error; CV = Intra-CV)

For generics with multiple bioequivalence results, pooled CVs weighted by sample size were calculated using the equations below, and these were used for calculating the optimal sample size:[4,14]

Pooled CV = $\sqrt{e^{\sum \ln(Intra-CV2+1)df \sum df} - 1}$

Confidence limit of pooled intra-CV = $\sqrt{e^{\sum \ln(Intra-CV2+1)df}/\chi^2}$ a.Edf -1

(df: degrees of freedom; 89 $\chi^2_{a, Edf}$: critical value of chi square estimates)

Results

Basic characteristics of bioequivalence studies analyzed

In total, 142 bioequivalence study results from 58 generics were evaluated in this study. Fifty-five generics were enteral formulations (i.e., 4 extended release formulations and 51 immediate release formulations), and 3 generics were topical formulations.

Intra-coefficients of variation for pharmacokinetic measures and sample size

The intra-CV of C_{max} was larger than that of AUC in 129 studies (90.8%), and this was consistent with previous reports that considered C_{max} the cornerstone for bioequivalence approval. [16] The estimated intra-CV (mean ± sd (min ~ max)) for C_{max} was 21.7 ± 8.8% (5.4 ~ 54.0%), and that for AUC was 14.7 ± 8.2% (3.2 ~ 56.4%) (Table 1).

The average total sample size (mean±sd) to obtain greater than 80% power was 26±20. In 44 out of 58 of the generics evaluated, the optimal sample sizes were larger than the minimal sample size for bioequivalence studies requested by the MFDS (n=12). For 14 (24.1%) generics, the estimated intra-CV of AUC and/ or C_{max} was larger than 30%, the threshold for classifying a drug as 'Highly Variable Drugs'. The estimated sample sizes of these 14 generics with estimated intra-CVs larger than 30% were 58.3±22.8 (min=42, max=120), far larger than the average estimated sample sizes of the 45 generics with intra-CVs of less than 30% (16.8±6.5, min=4, max=32). For the 26 generics with multiple bioequivalence results, substantial variations between the products of identical generics were found. The coefficient of variation (%) in intra-CV estimates between the products of identical generics ranged from 4.0% to 70.1% with respect to C_{max} and 1.9% to 105.3% for AUC.



Table 1. Weighted mean of intra-subject coefficient of variation (pooled intra-CV) and sample size for bioequivalence studies of 58 generics

Generics (Number of studies)	Pooled intra-CV (90% cor	nfidence interval) or intra-CV*	Sample size for bioequiva- lence study [#]	
	AUC	C _{max}	80% power	90% powe
Octylonium bromide*	56.4	40.6	120	164
Clopidogrel bisulfate*	39.1	44.9	82	110
Lansoprazole*	34.5	42.6	74	100
Naltrexone HCI*	28.8	40.9	68	92
Carvedilol*	20.7	36.6	56	76
Ranitidine HCI (2)	21.5 (19.0~24.0)	34.2 (30.0~38.4)	50	66
Desmopressin acetate*			46	
	32.7	26.9		62
Levetiracetam*	13.6	32.0	44	60
Esomeprazol mag. dihy. (2)	29.6 (26.6~32.6)	31.7 (28.4~35.0)	44	58
Atorvastatin ca. hyd.*	16.4 (15.7~17.1)	31.6 (30.2~33.0)	44	58
Pentoxifylline ER*	27.6	31.5	44	58
Telmisartan*	16.0	31.5	44	58
Celecoxib (8)	15.7 (14.8~16.6)	30.8 (29.1~32.5)	42	56
Linezolid (3)	10.2 (9.2~11.2)	26.6 (24.0~29.2)	32	42
Irbesartan (2)	18.0 (16.1~19.9)	25.4 (22.6~28.2)	30	38
Ecabet sodium*	22.8	24.6	28	36
Amlodipine besylate capsule*	9.1			
		24.0	26	36
Pramipexole HClpatch*	23.8	19.0	26	34
Duloxetine HCI*	19.3	23.0	24	32
Olmesartan medoxomil (3)	16.2 (14.4~18.0)	22.7 (20.1~25.3)	24	32
Atomoxetine HCI (3)	9.8 (8.9~10.7)	22.4 (20.2~24.6)	24	30
Entecavir (2)	12.4 (10.8~14.0)	22.3 (19.3~25.3)	24	30
Fentanyl patch*	14.6	22.0	22	30
Rosuvastatin ca. (22)	17.0 (164~17.6)	21.2 (20.5~21.9)	22	28
Aripiprazole (2)	9.8 (8.4~11.2)	21.1 (18.0~24.2)	22	28
Duloxetine hydrochloride capsule (8)	17.0 (16.0~18.0)	20.6 (19.7~21.5)	20	26
Buspirone HCI*	10.0	20.6	20	26
Donepezil HCI (3)	14.7 (13.0~16.4)	20.3 (18.0~22.6)	20	26
Mirtazapine (3)	8.1 (7.3~8.9)	20.1 (18.0~22.2)	20	26
Nizatidine*	7.5	20.0	20	26
Rivastigmine patch (6)	17.8 (16.6~19.0)	19.7 (18.4~21.0)	18	24
Sitagliptin phosphate hyd (5)	6.7 (6.2~7.2)	18.3 (16.8~19.8)	16	22
Topiramate (3)	7.5 (6.7~8.3)	18.3 (16.2~20.4)	16	22
Choline alfoscerate*	17.8	18.2	16	22
Tramadol HCI (2)	11.4 (10.0~12.8)	18.0 (15.7~20.3)	16	22
Mosapride citrate hydrate*	17.8	18.0	16	22
Risperidone (3)	16.1 (14.4~17.8)	17.8 (16.0~19.6)	16	20
Fluoxetine HCl capsule (2)	7.7 (6.6~8.8)	17.7 (15.2~20.2)	16	20
Propiverine HCI*	17.7	17.0	16	20
Bicalutamide*	14.4	16.7	14	18
Paroxetine HCI hydrate*	13.0	16.3	14	18
Lafutidine (2)	16.3 (14.1~18.5)	13.4 (11.6~15.2)	14	18
Imatinib mesylate (2)	12.8 (11.2~14.4)	16.1 (14.1~18.1)	14	18
Sitagliptin phosphate*	5.7	16.1	14	18
Levofloxacin hydrate*	13.3	15.3	12	16
Moxifloxacin hyd (2)	10.9 (9.5~12.3	15.3 (13.3~17.3)	12	16
Metformin HCI ER*	10.3	14.3	12	14
Pramipexole HCI mono. (3)	10.4 (9.4~11.4)	13.9 (12.6~15.2)	10	14
Gabapentin capsule*	10.2	13.2	10	12
Tadalafil*	13.1	12.6	10	12
Oxycodone HCI ER*	11.2	12.5	10	12
Escitalopram oxalate (3)	11.2 (9.8~12.6)	12.5 (11.0~14.0)	10	12
Meloxicam*	10.2	12.4	10	12
Irsogladine maleate (3)	8.1 (7.6~8.6)	12.4 (11.2~13.6)	10	12
Solifenacin succinate*	11.7	11.9	8	12
S-amlodipine besylate*	6.7	6.6	6	6
Memantine HCI*	5.2	5.4	4	6

ER, Extended release, mag, magnesium; dihy,dihydrate; hyd,hydrate; ca.,calcium; HCl,hydrochloride; dehy, dehydrate; mono, monohydrate. Sample sizes for bioequivalence studies of various generics were calculated based on the higher of the intra-CVs (i.e., either from AUC or C_{max}). * When the number of studies is 1. # Total sample size for 2X2 cross-over bioequivalence study.

Discussion

In the present study, we calculated the intra-CVs of various generics and evaluated the extent of inter-study variability. Large variations were observed for the estimated intra-CVs of pharmacokinetic measures between the study results of identical generics. Intra-CV is probably affected by drug's intrinsic factors such as absolute oral bioavailability and acidity.[17] However, extrinsic factors can substantially contribute to the variation in Intra-CV of same substance. The reason could be variability in drug concentration analysis, hospital site, protocol deviation, and manufacturing. Our results suggest that pooling of intra-CVs from multiple bioequivalence results will produce more reliable estimates of intra-CVs for designing bioequivalence studies. In this study, we present the pooled CV and its upper 80% confidence limit for 26 generics with multiple bioequivalence results (Table 1). The estimated intra-CV values and the information on inter-study variability will provide useful information for future planning of bioequivalence studies for the generics analyzed. To validate our results, we compared our data to other ethnic groups in 3 highly replicated generic drugs. Intra-CVs of C_{max} were 21.2% for rosuvastatin in Indonesian, [18] 29.0% for celecoxib in Taiwan[19] and 20.2% for duloxetine in Thai subjects.[20] All of them were quite similar to our results.

Our study has some limitations regarding the estimation of intra-CVs for reference drugs because we only analyzed 2x2 crossover studies. To estimate true intra-CVs of reference drugs, 2x3 or 2x4 replicative designs that allow replicative administration of reference products are needed.[21] In addition, all of the generic drugs we analyzed were successfully bioequivalent with their reference drugs, which may lead to biased results. However, our study results can be interpreted as reasonable approximations for the values of the true intra-CVs because we calculated pooled CVs from multiple studies.

In conclusion, we estimated the intra-CVs of various generics and the optimal sample sizes for bioequivalence studies. Our study results will be useful for planning future bioequivalence studies.

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Conflicts of interests

-Authors: None of the authors have any conflicts of interest to declare. -Reviewers: Nothing to declare

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