Review Article

Theme: Facilitating Oral Product Development and Reducing Regulatory Burden through Novel Approaches to Assess Bioavailability/Bioequivalence Guest Editors: James Polli, Jack Cook, Barbara Davit, and Paul Dickinson

Implementation of a Reference-Scaled Average Bioequivalence Approach for Highly Variable Generic Drug Products by the US Food and Drug Administration

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Abstract. Highly variable (HV) drugs are defined as those for which within-subject variability (%CV) in bioequivalence (BE) measures is 30% or greater. Because of this high variability, studies designed to show whether generic HV drugs are bioequivalent to their corresponding HV reference drugs may need to enroll large numbers of subjects even when the products have no significant mean differences. To avoid unnecessary human testing, the US Food and Drug Administration's Office of Generic Drugs developed a reference-scaled average bioequivalence (RSABE) approach, whereby the BE acceptance limits are scaled to the variability of the reference product. For an acceptable RSABE study, an HV generic drug product must meet the scaled BE limit and a point estimate constraint. The approach has been implemented successfully. To date, the RSABE approach has supported four full approvals and one tentative approval of HV generic drug products.

KEY WORDS: bioequivalence; generic drugs; highly variable drugs; reference-scaled average bioequivalence; US Food and Drug Administration.

INTRODUCTION

Acceptable bioequivalence (BE) between a generic drug product and its corresponding reference product is among the criteria required by the US Food and Drug Administration (FDA) for marketing approval of a new generic drug. A systemically active generic drug is considered to be bioequivalent to the reference-listed drug if the rate and extent of absorption of the two products do not show a significant difference (1). Drug rate and extent of absorption are typically assessed by conducting in vivo BE studies in human subjects in which generic and reference drug plasma pharmacokinetic (PK) profiles are characterized and compared. The PK data are used to obtain peak drug plasma concentration (C_{max}) as the BE measure of rate of absorption and area under the drug plasma concentration versus time profile (AUC) as the BE measure of extent of absorption. Most BE studies enroll healthy normal subjects who receive single doses of the generic (test) or reference product via a two-way crossover design. For the most part, the US FDA asks generic drug applicants to statistically compare generic and reference AUC and C_{max} values using the two one-sided tests (TOST) procedure (2). Under the TOST, the 90% confidence interval (CI) around the geometric mean ratio (GMR) of the test and reference values of C_{max} and AUC is required to fit within BE limits, set from 80% to 125% (3). The width of the 90% CI depends upon the number of subjects in the study and the variability of the BE measure.

Highly variable (HV) drugs are defined as drugs in which the within-subject variability (defined as the %CV) in one or more of the BE measures is 30% or greater (4,5). A survey of

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generic drug products reviewed at the FDA from 2003 to 2005 suggested that about 20% of the generic drugs evaluated for marketing approval are HV due to their drug substance dispositional characteristics (6). Determining the BE of HV generic drugs is challenging because the high within-subject variability means that large numbers of subjects may be needed to show BE. Figure 1 shows the results of two hypothetical BE studies with GMRs near 1.0. Product A meets the BE limits because within-subject variability is relatively low. Product B fails to meet the BE limits because of high within-subject variability. Thus, although product B, with a GMR near 1.0, appears to be well-designed to perform the same as the reference product *in vivo*, it will be necessary to increase the number of study subjects—perhaps dramatically—in order for product B to meet the BE limits.

Several factors influence the sample size needed to meet the regulatory criteria for acceptable BE. First, each onesided test is carried out at the 5% level of significance, which corresponds to a 90% CI (7). The 5% level of significance represents the type I error rate (α) , which is the probability of incorrectly deeming as bioequivalent two formulations whose true (population) GMR fails to meet the BE limits. The second factor influencing sample size is study power, defined as the likelihood or chance of correctly demonstrating BE when it, in fact, exists (8,9). A third factor influencing sample size is the test/reference BE measure ratios. If the true test/ reference ratio differs from unity, the overall power to show BE is reduced at any given sample size, resulting in an increase in the number of study subjects needed. Other factors influencing sample size include the study design and the expected within-subject variability. For example, a replicate four-way crossover BE study design, in which each subject receives the test and reference products twice, requires fewer subjects than a two-way crossover BE study design. As within-subject variability increases, the number of



Fig. 1. The 80–125% BE limits are represented along the *x*-axis as two "goal posts." The BE limits are compared to the hypothetical 90% CIs of the test/reference BE measure GMRs for two drugs, a drug with normal variability (*Drug A*) and an HV drug (*Drug B*). The 90% CIs of the two drugs are represented by *colored bars*. For drug A (normal variability), the 90% CI (*green bar*) meets the BE limits. For drug B (HV), the 90% CI (*red bar*) fails to meet the acceptance limits. As the width of the CI is influenced by the number of study subjects, in the hypothetical case of drug B, it is likely that the study would have met the BE limits if more subjects had been used

subjects needed in a crossover design will also increase, assuming that all other factors remain constant. Thus, BE study sample size is calculated based on a type I error rate of 5% per test, the desired study power, and the best estimates of test/reference ratios and within-subject variability. Table I shows how these factors influence the number of subjects needed to provide an 80% chance of an acceptable BE study.

As shown in Table I, the number of study subjects needed to show BE increases dramatically for HV drugs. The FDA observed in a survey of generic drug product BE studies reviewed from 2003 to 2005 that studies of HV drugs generally used more subjects than studies of lower variability drugs (6).

SHOULD "ONE-SIZE-FITS-ALL" BE LIMITS APPLY TO HV DRUGS

One important observation is that clinical data strongly support a conclusion that HV drugs have wide therapeutic indices. Otherwise, there would have been significant safety issues and lack of efficacy during the pivotal safety and efficacy clinical trials required for initial FDA marketing approval (10). In other words, the reference product, when dosed on different occasions, was safe and efficacious, despite high PK variability. The majority of HV drugs appear to fall into Biopharmaceutics Classification System class II or IV (low aqueous solubility-high intestinal permeability and low aqueous solubility-low intestinal permeability, respectively) (11,12). Dispositional characteristics of HV drugs include extensive presystemic metabolism, low bioavailability, high acid lability, and high lipophilicity (6). Consequently, plasma concentrations of these HV drugs are often very low. In such situations, it may not be possible to accurately characterize PK profiles, with the result that the within-subject variability of BE measures can exceed 30% (13). As the FDA discourages unnecessary human testing, these observations raise questions about whether large numbers of subjects

Table I. The Number of Study Subjects Required to Show BE with 80% Power is a Function of Within-Subject Variability and GMR (Sample Size Estimations are for the Case $\sigma_{WT} = \sigma_{WR}$ and $\sigma_D = 0$)

Within- subject %CV	GMR (%)	Sample size for a two-way crossover study	Sample size for a four-way fully replicated crossover	
15	100	10	6	
	105	12	8	
	110	20	12	
30	100	32	18	
	105	38	20	
	110	68	36	
45	100	66	34	
	105	80	42	
	110	142	72	
60	100	108	56	
	105	132	66	
	110	236	118	
75	100	156	80	
	105	190	96	
	110	340	172	

should be used in BE studies of drug products for which high variability does not appear to impact safety and efficacy. It should be stressed that this concern pertains to high PK variability due to drug substance dispositional characteristics and has nothing to do with the formulation performance assessment that is the key question in BE comparisons.

An additional concern is that the large sample sizes needed for BE studies of HV drugs may serve to deter the development of new generic products (14,15). For example, a generic product line may be abandoned because of a high failure rate obtained in *in vivo* BE studies or it may be necessary to repeat the *in vivo* studies several times until the outcome of an acceptable BE study is achieved (16). Not only does this situation lead to unnecessary human testing, but it can also serve to increase the prices of generic drugs, conflicting with the principle that generic drugs should be relatively low-cost because they are not subject to extensive and unnecessary clinical testing.

A final concern is that an HV reference product may not be shown to be bioequivalent to itself in a crossover study using a relatively modest number of subjects (e.g., 18–40) (13). This concern is supported by data from *in vivo* studies of brand-name (reference drug product) formulations of chlorpromazine and verapamil, both of which were characterized by high within-subject PK variability exceeding 30% (17,18). Such findings raise the question of whether the 80–125% BE limits are too stringent for HV drug products.

EVOLUTION OF A NEW BE APPROACH FOR HV GENERIC DRUGS AT THE FDA

The issues surrounding BE evaluation of HV drugs and proposals for modifying the BE approach for such products were discussed over many years within the pharmaceutical sciences community, in the literature, and at various national and international venues (17-21). In 2004, FDA considered seriously whether an alternative BE approach to the "one-size-fits-all" paradigm should be developed for generic HV drugs. In April of that year, FDA's Office of Generic Drugs (OGD) brought the issue of optimizing BE study designs for HV drugs before its Pharmaceutical Science and Clinical Pharmacology Advisory Committee (hereafter Advisory Committee). Speakers from the FDA, academia, and industry speakers presented the various issues surrounding BE evaluation of HV drugs to the Advisory Committee (22). Among the proposals debated were whether to change the BE limits, for example, using a GMR constraint of 80-125% or expanding the 90% CI limits to 70-143% (23) versus whether to use reference product within-subject PK variability to scale the BE limits (16). Following deliberations, the Advisory Committee made several recommendations (22).

- There is a need to understand where the variability originated. Prior knowledge of all biostudies may help set more appropriate specifications to make decisions.
- There is an undue reliance on the use of 80–125% BE limits for all drug products. As such, a paradigm shift for HV drugs is in order.

• The use of reference scaling should be explored. A limit on the GMR should be used along with reference scaling.

FDA's OGD agreed to initiate a series of simulation studies to determine whether such an approach was feasible and, if so, how to optimize the approach. Subsequently, the OGD formed an interdisciplinary working group (hereafter Working Group) to investigate the feasibility of applying a reference-scaled average bioequivalence (RSABE) approach to generic HV drugs. The Working Group conducted simulations to investigate expanding BE limits as a function of reference product within-subject variability. Based on the simulation study results, the Working Group members presented a tentative proposal for an RSABE method to the Advisory Committee for Pharmaceutical Science and Clinical Pharmacology in April of 2006. Speakers from academia and from the FDA discussed the advantages and limitations of the following approaches: (a) implementing several different scaling methods (24-27); (b) recommending a minimum number of subjects for RSABE (25,26); and (c) including a constraint on the GMR of the BE study (25-28). The Advisory Committee's recommendations at the meeting's conclusion are summarized below (29).

- The Advisory Committee was not asked to comment on the merits of RSABE as it had previously advocated investigating the approach in April 2004.
- The Advisory Committee agreed that these studies should enroll a minimum number of subjects. The vote was split as to whether this number should be 24 or 36.
- The majority of Advisory Committee members either voted against a GMR constraint or abstained from voting on this issue.

FDA's OGD considered the Advisory Committee's recommendations in finalizing the RSABE approach for HV drugs (30).

FDA'S RECOMMENDATIONS FOR RSABE STUDIES OF HV DRUGS

Throughout this section, the three basic approaches used for determining if two products are bioequivalent will be illustrated using a series of inequalities based on population parameters.

Average Bioequivalence

The usual manner of statistically analyzing BE study data is by the average bioequivalence (ABE) approach. The acceptance of BE is stated if the difference between the logarithmic means is between preset regulatory limits, as shown below:

$$(\mu_T - \mu_R)^2 \le \theta_A^2$$

where $\mu_{\rm T}$ is the population average response of the logtransformed measure for the test (T) formulation, $\mu_{\rm R}$ is the population average response of the log-transformed measure for the reference (R) formulation, and θ_A is equal to ln(1.25).

As $-\ln(1.25) = \ln(0.8)$, the BE limits are:

$$\ln(0.8) \le (\mu_T - \mu_R) \le \ln(1.25)$$

Individual Bioequivalence

For a time, the FDA worked toward implementing an individual bioequivalence (IBE) approach for studies submitted to New Drug Applications (NDAs) and Abbreviated New Drug Applications (ANDAs, for generic drugs). It was argued that requiring drug products to meet an IBE rather than an ABE criterion would improve formulation switchability (31,32). The proposed criterion for acceptable IBE included the (a) comparison of test and reference means; (b) comparison of within-subject variances; (c) assessment of subject-by-formulation interaction; and (d) ability to scale the BE limits if within-subject variability following the administration of the reference product exceeded a predetermined value (15). An additional justification offered for implementing IBE was that the ability to scale BE limits would address the concerns associated with BE studies of HV drugs (33–35).

Under IBE, the inequality used to determine if two products are bioequivalent is as follows:

$$\frac{\left(\mu_T - \mu_R\right)^2 + \sigma_D^2 + \left(\sigma_{WT}^2 - \sigma_{WR}^2\right)}{\sigma_{WR}^2} \le \theta_I$$

where σ_D^2 is the population subject-by-formulation interaction variance components, σ_{WT}^2 is the population withinsubject variance of the test formulation, σ_{WR}^2 is the population within-subject variance of the reference formulation, and θ_I is the BE limit for IBE.

From 1999 to 2001, at the FDA's request, the pharmaceutical industry applied the IBE study design and analysis to NDAs and ANDAs for modified-release drug products (36). The IBE was used to evaluate the BE of modified-release drug products because it was thought that, due to the relative complexity of modified-release formulations, the likelihood was greatest of detecting possible subject-by-formulation interactions with these types of drug products. However, analysis of these data failed to detect the presence of clinically significant subject-by-formulation interactions (37). The FDA concluded that expansion of the IBE approach to all BE studies submitted for marketing approval was not warranted and subsequently halted its implementation (38). Notably, during a discussion of IBE at its November 29, 2001 meeting, the Advisory Committee for Pharmaceutical Science and Clinical Pharmacology suggested that the FDA consider applying reference scaling to ABE studies of HV drugs (39).

Reference-Scaled Average Bioequivalence

The RSABE for both AUC and C_{max} is evaluated as shown below:

$$\frac{(\mu_T - \mu_R)^2}{\sigma_{WR}^2} \le \theta_S$$

where σ_{WR}^2 is the population within-subject variance of the reference formulation, $\theta_S = \frac{(\ln(1.25))^2}{\sigma_{W0}^2}$ is the BE limit, and σ_{W0}^2 is a predetermined constant set by the regulatory agency.

Under this model, the implied limits (which represent FDA's desired consumer risk model) on $\mu_{\rm T} - \mu_{\rm B}$ are:

$$-\left[\ln(1.25)\frac{\sigma_{WR}}{\sigma_{W0}}\right] \le \mu_T - \mu_R \le \ln(1.25)\frac{\sigma_{WR}}{\sigma_{W0}}$$

If $\sigma_{WR} = \sigma_{W0}$, the implied limits are equal to the standard unscaled BE limits of $\pm \ln(1.25)$ (0.80 to 1.25). If $\sigma_{WR} > \sigma_{W0}$, the implied limits are wider than the standard limits. If $\sigma_{WR} < \sigma_{W0}$, the implied limits are narrower than the standard limits.

The FDA recommends a mixed scaling approach, as the Agency has determined that it is acceptable for the implied limits to be wider than the standard limits only when σ_{WR} is large (as for HV drugs). The mixed scaling model is as shown below.

T and R are considered BE if:

$$\frac{(\mu_T - \mu_R)^2}{\sigma_{W0}^2} \le \frac{(\ln(1.25))^2}{\sigma_{W0}^2} \text{ if } \sigma_{WR} \le \sigma_{W0}$$

and if:

$$\frac{(\mu_T - \mu_R)^2}{\sigma_{WR}^2} \le \frac{(\ln(1.25))^2}{\sigma_{W0}^2} if \, \sigma_{WR} > \sigma_{W0}$$

FDA sets the value of σ_{W0} at 0.25 (30,40). Under this mixed scaling model and with σ_{W0} =0.25, the implied limits on $\mu_{T}-\mu_{R}$ are as depicted in Fig. 2.

Direct implementation of FDA's desired consumer risk model is impossible because σ_{WR} is a characteristic of the entire population and thus not directly measured in any particular study. Therefore, FDA has proposed an implementation algorithm (described in the "FDA Draft Guidance for



Fig. 2. Implied BE limits are plotted as a function of the population reference product within-subject variability of the BE measure. When $\sigma_{W0} \le 0.25$, for an acceptable BE study, the 90% CI of the BE measure test/reference GMRs must fall within the 80–125% limits. When $\sigma_{W0} > 0.25$, the implied BE limits scale as reference product within-subject variability increases. The slope of this portion of the curve is determined by the value of σ_{W0}



Fig. 3. Decision tree showing the process by which FDA's OGD decide whether it is suitable to use RSABE or unscaled ABE. The first condition to be met is that the study protocol must state *a priori* that RSABE will be the method of statistical analysis

Industry, Bioequivalence Recommendations for Progesterone Oral Capsules") and discussed in more detail later in this paper. In the FDA-recommended implementation algorithm for mixed scaling studies, the observed within-subject variability of the reference s_{WR} (determined in the BE study) is compared to a cutoff value of 0.294, above which reference scaling is used. This implementation reduces the type I error (defined relative to FDA's desired consumer risk model) when the within-subject variability is near σ_{W0} .

FDA RECOMMENDATIONS FOR DESIGNING RSABE STUDIES

The reference product is administered twice in order to determine its within-subject variability (30). As such, the BE study can use either a partial replicate (three-way crossover: RTR, RRT, or TRR) or full replicate (four-way crossover: RTRT or TRTR) design, but should enroll a minimum of 24 subjects (40,41). The FDA recommends an s_{WR} cutoff value of 0.294, at or above which reference scaling is permitted and below which the unscaled limits of 0.8 to 1.25 are applied (30,40). The selection of 0.294 as the variation at which use of reference scaling of the limits is consistent with the general understanding that drugs may be considered HV if the within-subject coefficient of variation (%CV) observed in the study

is $\geq 30\%$ and, as such, is determined by using the conversion formula of $s^2 = \ln(CV^2 + 1)$.

FDA determined that there are advantages to choosing an s_{WR} cutoff somewhat larger than σ_{W0} . The Working Group conducted a series of simulations of BE study results (1,000,000 per condition) to compare the effects of applying different values to σ_{W0} . The study results showed that using a σ_{W0} of 0.25 (1) increased the study power compared to ABE, without causing relatively large numbers of studies to pass when the GMR \geq 1.2 and (2) resulted in a lower inflation of type I error compared to a σ_{W0} of 0.294 (42,43).

The FDA recommends a secondary point estimate constraint criterion of 0.8 to 1.25 on the GMR (30,40). As it is possible that, using RSABE, two products can be shown to be bioequivalent but have an estimated GMR outside of the 0.8 to 1.25 range, it is thought that use of the secondary point estimate constraint will improve the confidence of clinicians and patients (15,28). Several simulation studies investigating the relationship between FDA's recommended RSABE approach and study power have shown that, when within-subject variability exceeds 50-60%, the point estimate constraint becomes the dominant regulatory criterion rather than the scaling (43-45). It has also been shown that applying the point estimate constraint can increase the sample size needed to show BE for drugs with within-subject variability >50-60% (45). Thus, using a point estimate constraint results in a situation where the benefits of using RSABE are reduced when applied to drugs with very high within-subject variability.

However, it has also been argued that, without a point estimate constraint in effect, the permissiveness of the RSABE approach can become excessively high (42). Another argument favoring incorporating a point estimate constraint is that it has a long history of successful application by regulatory agencies as a BE study acceptance criterion (15). Using a point estimate constraint on the GMR as a criterion for study outcome acceptance is used by the Therapeutic Products Directorate of Health Canada for C_{max} in all BE studies (46) and, until 2003, by the FDA for both AUC and C_{max} in BE studies that were conducted under fed conditions.

APPLICATION OF RSABE TO REGULATORY REVIEW OF ANDAS FOR GENERIC HV DRUGS

Using the RSABE approach recommended by the FDA, two products are bioequivalent when the 95% upper confidence bound for $\frac{(\mu_T - \mu_R)^2}{\sigma_{WR}^2}$ is $\leq \theta_S$ or, equivalently, a 95% upper confidence bound for $(\mu_T - \mu_R)^2 - \theta_S \sigma_{WR}^2$ is ≤ 0 . In

Table II. Summary BE Statistics for Two HV Generic Drugs, Analyzed Using RSABE or Unscaled ABE, Depending upon the Value of s_{WR}

east-square	geometric means	, point estimates	and 90%	confider	nce interval	s	
T/R ratio	Lower 90% CL	Upper 90% CL	s_{WR}^2	s _{wR}	Criteria bound	Method used	Outcome
1.17	93.91	132.91	0.5723	0.7565	-0.2892	Scaled/PE	Pass
1.11	94.64	124.91	0.3452	0.5875	-0.1767	Scaled/PE	Pass
1.19	100.47	133.26	0.3768	0.6138	-0.1684	Scaled/PE	Pass
0.99	89.82	109.53	0.07425	0.2725	-0.03914	Unscaled	Pass
1.02	92.57	111.77	0.06561	0.2561	-0.03126	Unscaled	Pass
0.98	85.70	110.54	0.1069	0.3270	-0.05294	Scaled	Pass
	east-square T/R ratio 1.17 1.11 1.19 0.99 1.02 0.98	east-square geometric means T/R ratio Lower 90% CL 1.17 93.91 1.11 94.64 1.19 100.47 0.99 89.82 1.02 92.57 0.98 85.70	east-square geometric means, point estimates T/R ratio Lower 90% CL Upper 90% CL 1.17 93.91 132.91 1.11 94.64 124.91 1.19 100.47 133.26 0.99 89.82 109.53 1.02 92.57 111.77 0.98 85.70 110.54	east-square geometric means, point estimates and 90% T/R ratioT/R ratioLower 90% CLUpper 90% CL s_{WR}^2 1.1793.91132.910.57231.1194.64124.910.34521.19100.47133.260.37680.9989.82109.530.074251.0292.57111.770.065610.9885.70110.540.1069	east-square geometric means, point estimates and 90% confiderT/R ratioLower 90% CLUpper 90% CL s_{WR}^2 s_{WR} 1.1793.91132.910.57230.75651.1194.64124.910.34520.58751.19100.47133.260.37680.61380.9989.82109.530.074250.27251.0292.57111.770.065610.25610.9885.70110.540.10690.3270	east-square geometric means, point estimates and 90% confidence interval T/R ratioconfidence interval s_{WR}^2 1.1793.91132.910.57230.7565-0.28921.1194.64124.910.34520.5875-0.17671.19100.47133.260.37680.6138-0.16840.9989.82109.530.074250.2725-0.039141.0292.57111.770.065610.2561-0.031260.9885.70110.540.10690.3270-0.05294	east-square geometric means, point estimates and 90% confidence intervalsT/R ratioLower 90% CLUpper 90% CL s^2_{WR} s_{WR} Criteria boundMethod used1.1793.91132.910.57230.7565 -0.2892 Scaled/PE1.1194.64124.910.34520.5875 -0.1767 Scaled/PE1.19100.47133.260.37680.6138 -0.1684 Scaled/PE0.9989.82109.530.074250.2725 -0.03914 Unscaled1.0292.57111.770.065610.2561 -0.03126 Unscaled0.9885.70110.540.10690.3270 -0.05294 Scaled/PE



Fig. 4. Outcome of the review of 64 RSABE studies reviewed at the OGD. All but two of the studies met the BE acceptance criteria. For 36 of these studies, there were no other deficiencies related to the BE portion of the ANDA submission. For 26 studies, other miscellaneous deficiencies unrelated to RSABE analysis (most often related to the bioanalytical method used) were present in the BE portion of the submission. One study did not meet the point estimate constraint and thus the study did not meet the RSABE acceptance criteria. Another study could not be analyzed using RSABE because the s_{WR} for C_{max} was <0.294. However, when the TOST was applied, the 90% CI for C_{max} did not fall within the 80–125 limits; thus, this study did not meet the ABE acceptance criteria

addition, the GMR of the two products should fall between 0.8 and 1.25.

Steps for Conducting an RSABE Analysis

It should be noted that we do not know the values of the above population parameters and can never know their values. What we can and do in actuality is calculate CIs around the BE metrics, log(AUC) or $log(C_{max})$.

FDA posted a Guidance for Industry providing stepby-step instructions on how to statistically analyze BE study data using RSABE (40). The intention to use RSABE for an HV drug should be stated *a priori* in the study protocol. The first step in the analysis is to determine s_{WR} , the within-subject standard deviation (SD) of the reference product estimated from the study, for the BE measures AUC and C_{max} . If $s_{WR} < 0.294$, then the TOST procedure should be used to determine ABE for the BE measure. If $s_{WR} \ge 0.294$, then RSABE can be used for the BE measure. Figure 3 illustrates the decision-making process that the FDA uses when determining whether it is suitable to use RSABE for a generic drug.

Once it has been decided that RSABE can be applied to a BE measure, the next step is to determine the 95% upper confidence bound for $(\mu_T - \mu_R)^2 - \theta_S \sigma_{WR}^2$ based on *Howe's Approximation I* (47). The test and reference product are concluded to be bioequivalent if:

- (a) the 95% upper confidence bound for $(\mu_T \mu_R)^2 \theta_S \sigma_{WR}^2$ is ≤ 0 AND
- (b) the test/reference GMR in the study falls within [0.8, 1.25].

Case Studies

Two examples, provided in Table II, show how FDA applies RSABE to studies of HV generic drugs. Both cases are of approved generic drugs, for which the BE studies used three-way partial replicate study designs. As FDA scientific reviewers conduct their own independent calculations in performing BE analysis, the data in Table II are from FDA's calculations, performed using SAS® Version 9.2.

As shown in Table II, in the case of drug A, the first step of the analysis determined that values of s_{WR} exceeded the regulatory cutoff of 0.294 for AUC_{0-t}, AUC_∞, and C_{max} . Thus, the reviewer proceeded to use RSABE for all three BE measures. Table II shows that the calculated 95% upper confidence bound values for all three parameters were <0. Therefore, the product met the first BE acceptance criterion. In addition, the point estimates for all three BE measures are within 0.8 and 1.25. Thus, in this study, drug A met the two RSABE acceptance criteria and was deemed bioequivalent to its corresponding reference.

The analysis of drug B, shown in Table II, was conducted in a different manner because it was not suitable to use RSABE for all BE measures. In the first part of the analysis, $s_{\rm WR}$ for $C_{\rm max}$ exceeded 0.294 and values of $s_{\rm WR}$ for AUC_{0-t} and AUC_{∞} were <0.294. Thus, ABE with the TOST procedure was used for AUC, for which the 90% CIs met the 80-125% limits. RSABE analysis was only applied to $C_{\rm max}$, which met both acceptance criteria, as the 95% upper confidence bound was <0 and the test/reference GMR point estimate was within 0.8 and 1.25. The case of drug B illustrates the important points that (a) both ABE and RSABE analysis can be used for data from the same study depending on BE measure variability and (b) there is no penalty if the applicant uses a three-way or four-way study design with the intent of using RSABE but s_{WR} fails to make the 0.294 cutoff.

Statistics on BE Study Designs in Recent ANDA Submissions of Generic HV Drugs

Since 2007, the FDA has evaluated 46 ANDAs containing 64 studies in which RSABE was applied. Figure 4 displays the statistics on these studies. Sixty-two of these studies met the RSABE acceptance criteria, although some studies were considered "incomplete" for other reasons, many due to questions relating to the bioanalytical method (48). Only two studies were unacceptable because they did not meet the acceptance criteria. In the first instance, as one BE measure, s_{WR} , was <0.294, the RSABE analysis could not be applied. Thus, it was necessary to apply ABE and the TOST procedure, but the study did not meet the acceptance limits because the upper confidence limit exceeded 125%. In the second instance, the study data met the

Table III. Number of Failed BE Studies Reviewed at FDA's OGD Since the "All Bioequivalence Studies" Rule Became Effective in July 2009

	В	E Studies	ANDAs		
Description	Number	Percent of total	Number	Percent of total	
Within-subject variability of AUC and $C_{\text{max}} \ge 30\%$	92	45	45	37	
Within-subject variability of AUC and $C_{\text{max}} < 30\%$	113	55	76	63	
Totals	205	100	121	100	

Table IV. Changes Made to the Study Design or Formulation toAchieve Successful Pivotal BE Studies of HV Drugs after the InitialBE Studies Failed to Meet the Acceptance Criteria

Change made	No. of ANDAs	Percent
Increase in sample size in	41	91
Changed to three-way study design and RSABE	3	7
Reformulated Total	1 45	2 100

criteria for applying the RSABE analysis, and the 90% upper confidence bound for all three BE measures was <0, but the GMRs of all BE measures exceeded 1.25. Interestingly, the applicant conducted a second successful study on this product which met the BE limits using RSABE analysis without reformulating or increasing the number of subjects.

As of this writing, FDA has fully approved four HV generic drug products for which RSABE was used. In addition, one HV generic drug product for which RSABE was used was tentatively approved (because a patent related to the reference product has not yet expired).

One of the major reasons offered to justify moving away from the "one-size-fits-all" BE limits for HV drugs was concern that BE studies would be unnecessarily repeated because of underpowering. Until the "All Bioequivalence Studies" Final Rule became effective in 2009, it was not possible for the FDA to assess the prevalence of failed and repeated BE studies on HV drugs because generic applicants were not required to submit failed BE studies (49). As a result, FDA's assessment of the scope of BE problems in regulatory submissions of HV drugs was possibly biased because the Agency did not see the results of failed BE studies. As applicants are now required to submit to ANDAs all BE studies conducted on the same drug product formulation as the one intended for marketing (50,51), it is hoped that such submissions will provide valuable information about the development of HV drug formulations.

To test the hypotheses that a high percentage of BE studies fail because they are of (a) HV drugs and (b) do not enroll enough subjects, we surveyed all failed BE studies of generic drugs reviewed from the time that the Final Rule went into effect until the time of this writing. As shown in Table III, 205 failed BE studies were reviewed during this time period. Of these, 92 of these failed BE studies were of HV drugs, representing 45% of the total. This appears to be a disproportionately high percentage of failed BE studies, as HV drugs are thought to represent about 20% of drug products submitted to FDA in ANDAs.

In addition, we surveyed the reasons why these 92 BE studies submitted in 45 ANDAs failed. The results confirm that a high percentage of studies of HV drugs appear to fail the BE acceptance criteria due to underpowering. As shown in Table IV, for all but one of 45 different ANDAs, generic products that initially failed one or more two-way crossover studies with TOST analysis met the BE limits in subsequent studies when power was increased, either by enrolling more subjects or by changing to an RSABE design.

RSABE APPROACHES FOR HV DRUGS IN OTHER JURISDICTIONS

In August 2010, the European Medicines Agency (EMA) issued a new Guideline on the Investigation of Bioequivalence, recommending a scaling approach for BE assessment of HV drugs (15,42,52,53). EMA recommends that BE limits scale with variability only for C_{max} and only up to a maximum within-subject variability value of 50% (where %CV is defined as $100\sqrt{e^{s_{WR}^2} - 1}$), after which they remain constant at the static limits of 69.84% to 143.19%. In BE

Timeline of Events in FDA's Development of the RSABE Approach



Fig. 5. This timeline summarizes the evolution of the RSABE approach at the FDA. The FDA first proposed reference scaling as a component of IBE analysis, although the FDA subsequently decided not to continue implementing IBE. Three years later, in 2004, the FDA began developing RSABE using the scaling approach from IBE. The first ANDA using RSABE was submitted in 2007. The first tentative approval of an HV drug for which RSABE was successfully applied was in 2009. The first full approval of an HV drug supported by acceptable RSABE was in 2011. From 2004 to 2010, FDA scientists published several peer-reviewed scientific articles on RSABE and made a number of presentations at national and international venues. The FDA has also posted a Draft Guidance for Industry with a step-by-step description of how to perform RSABE

studies, C_{max} is usually more variable than AUC (15,54,55). Either a three-way or four-way crossover design, in which the reference product is administered twice, can be used in the BE study, in order to determine the reference product withinsubject variability for scaling. The applicant should demonstrate that the within-subject variability of the reference product C_{max} is >30%. The extent of the widening of the acceptance limits is defined based upon the within-subject variability seen in the BE study using scaled average BE according to $[U, L] = \exp[\pm k \times s_{WR}]$, where U is the upper limit of the acceptance range, L is the lower limit of the acceptance range, k is the regulatory constant set to 0.760, and s_{WR} is the within-subject SD of the log-transformed values of C_{max} of the reference product (15,42,52). Using these values, the regulatory limits range from 80% to 125% when the within-subject %CV is 30% to 69.84-143.19% when the within-subject variability is 50%. EMA also recommends a point estimate constraint on the C_{max} GMR if scaling is used.

The South African Medicines Control Council suggests that the RSABE may be used for AUC and C_{max} if specified *a priori* in the protocol and justified by sound scientific principles; applicants also have the option of applying static BE limits with a wider acceptance interval for HV C_{max} (56,57).

CONTROVERSIES

One of the more controversial aspects of the RSABE approach is the selection of the value of σ_{W0} , the variability at which the BE limits can potentially begin to scale. The FDA sets σ_{W0} at 0.25, but also proposes an implementation algorithm that does not permit BE limits to scale until the observed within-subject variability reaches the cutoff value of 0.294 for s_{WR} . Endrenyi and Tothfalusi (44) argue that the FDA's approach to RSABE is discontinuous and conducted simulations claiming that, when σ_{W0} ranges from 0.246 to 0.294, consumer risk is substantially larger than 5% and the regulatory uncertainty about a decision on acceptance or rejection is enhanced. This claim is based on a different definition of consumer risk than FDA uses. Endrenyi and Tothfalusi define consumer risk relative to BE limits of 80-125% (44). FDA's position is that a true population variance of $\sigma_{WR} \ge 0.25$ justifies using BE limits that are wider than 80– 125%. The use of $s_{WR} \ge 0.294$ as part of FDA's implementation method affects the producer risk-there will be a higher probability of failing a product that is acceptable according to FDA's consumer risk model.

Results of simulations conducted by members of the HV Drug Working Group support the position that using a cutoff value of 0.294 for s_{WR} maintains an acceptable type I error rate relative to FDA's desired consumer risk model. This can be illustrated by using a value of 0.29399 for σ_{WR} . In this case, the acceptable limits on $\mu_T - \mu_R$ are $\pm \ln(1.25) \times 0.29399/0.25 =$ ± 0.262408 . Results of simulated three-way partial replicate studies of 36 subjects result in 46,434 conclusions of BE out of 1,000,000 simulated studies or an estimated type I error rate of 0.046434 relative to FDA's desired consumer risk model. It is important to carefully note the distinction between σ_{WR} and s_{WR} in the analysis of the scaling method.

Finally, likelihood of regulatory uncertainty in evaluating RSABE study results when within-subject variability is near

30% is substantially reduced by the evaluation approach used by BE review scientists in FDA's OGD. Although the RSABE analysis is triggered when $s_{WR} \ge 0.294$ in a threeway or four-way replicated study, review scientists are encouraged to evaluate the appropriateness of reference scaling in the context of other available information on the PK variability of a given drug. The three-level review process for all BE submissions in the OGD (reviewer, team leader, and division director) also encourages critical evaluation of data and consistent policy application.

FUTURE DIRECTIONS

Lack of harmonization in setting the regulatory constant and cutoff for RSABE could potentially be resolved by using the approach for all generic products, as was formerly proposed for IBE. Such an approach could also be more favorable to ensuring drug product switchability. The FDA recently proposed to adapt RSABE for BE studies of narrow therapeutic index (NTI) drugs (58). Since NTI drugs are characterized by low within-subject variability, this would effectively result in narrower BE limits.

CONCLUSIONS

Figure 5 provides a timeline showing how RSABE evolved at the FDA, from inception to application to ANDAs for generic HV drugs. FDA implemented RSABE for generic HV wide therapeutic index drugs to ease regulatory burden. In the RSABE approach recommended by the FDA, the within-subject variability of the reference product BE measure is determined in a three-way or four-way replicated study which should use at least 24 subjects. The implied BE limits scale to reference within-subject variability once $\sigma_{\rm WR} = \sigma_{\rm W0} = 0.25$ or greater. However, to preserve an acceptable (<5%) type I error rate, applicants cannot apply reference scaling to calculate BE limits until the reference product within-subject SD in the BE study is at least 0.294. Acceptance criteria are that the 95% upper confidence bound for the GMR test/reference BE measure ratio is ≤ 0 and the point estimate of the BE measure GMR is within 0.8 to 1.25. The approach has been applied successfully to ANDAs, with 64 RSABE studies submitted and reviewed since 2007. To date, FDA has granted four full marketing approvals and one tentative approval for several diverse HV generic drug products based on the RSABE analysis. The RSABE is not a "one-size-fits-all" approach, but rather a system that selfadjusts to the characteristics of the reference. As a result, RSABE could be used to potentially set appropriately strict BE limits for all products.

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