

# Consumer risk in SABE

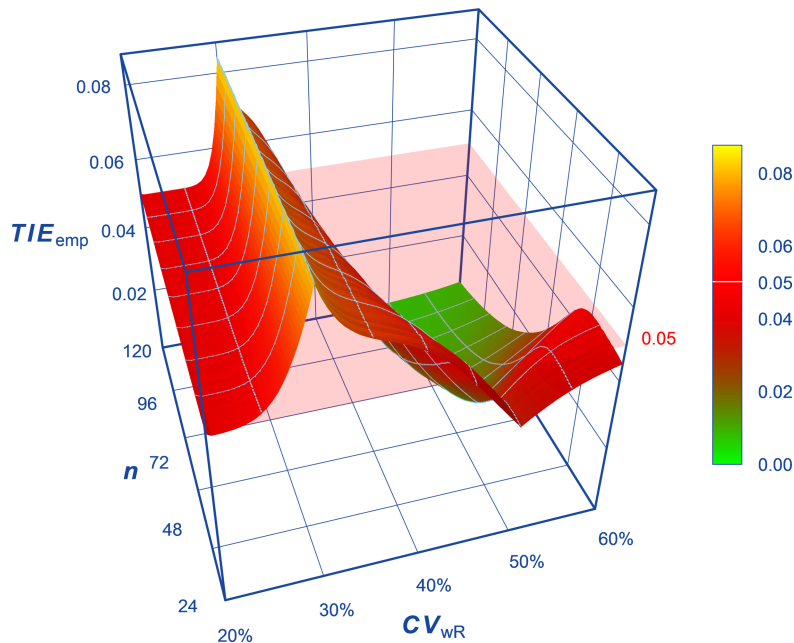
## Empiric Type I Error assessed at $\exp(\pm\theta_s \cdot s_{wR})$

weighted impact factor 3.37 (1.24 – 6.45)

1. Tóthfalusi L, Endrényi L, García-Arieta A. Evaluation of bioequivalence for highly variable drugs with scaled average bioequivalence. *Clin Pharmacokinet.* 2009; 48(11): 725–43.
2. Haidar SH, Makhlof F, Schuirmann DJ, Hyslop T, Davit B, Conner D, Yu LX. Evaluation of a Scaling Approach for the Bioequivalence of Highly Variable Drugs. *AAPS J.* 2008; 10(3): 450–4.
3. Endrényi L, Tóthfalusi L. Regulatory and Study Conditions for the Determination of Bioequivalence of Highly Variable Drugs. *J Pharm Pharmaceut Sci.* 2009; 12(1): 138–49.
4. Karalis V, Symillides M, Macheras P. On the leveling-off properties of the new bioequivalence limits for highly variable drugs of the EMA guideline. *Europ J Pharm Sci.* 2011; 44(4): 497–505.
5. Wonnemann M, Frömke C, Koch A. Inflation of the Type I Error: Investigations on Regulatory Recommendations for Bioequivalence of Highly Variable Drugs. *Pharm Res.* 2015; 32(1): 135–43.
6. Muñoz J, Alcaide D, Ocaña J. Consumer's risk in the EMA and FDA regulatory approaches for bioequivalence in highly variable drugs. *Stat Med.* 2016; 35(12): 1933–43.
7. Labes D, Schütz H. Inflation of Type I Error in the Evaluation of Scaled Average Bioequivalence, and a Method for its Control. *Pharm Res.* 2016; 33(11): 2805–14.
8. Tóthfalusi L, Endrényi L. An Exact Procedure for the Evaluation of Reference-Scaled Average Bioequivalence. *AAPS J.* 2016; 18(2): 476–89.
9. Tóthfalusi L, Endrényi L. Algorithms for Evaluating Reference Scaled Average Bioequivalence: Power, Bias, and Consumer Risk. *Stat Med.* 2017; 36(27): 4378–90.
10. Molins E, Cobo E, Ocaña J. Two-Stage Designs Versus European Scaled Average Designs in Bioequivalence Studies for Highly Variable Drugs: Which to Choose? *Stat Med.* 2017; 36(30): 4777–88.
11. Endrényi L, Tóthfalusi L. Bioequivalence for highly variable drugs: regulatory agreements, disagreements, and harmonization. *J Pharmacokin Pharmacodyn.* 2019; 46(2): 117–26.
12. Deng Y, Zhou XH. Methods to control the empirical type I error rate in average bioequivalence tests for highly variable drugs. *Stat Methods Med Res.* 2019; 29(6): 1650–67.
13. Ocaña J, Muñoz J. Controlling type I error in the reference-scaled bioequivalence evaluation of highly variable drugs. *Pharm Stat.* 2019; 18(5): 583–99.
14. Schütz H, Labes D, Wolfsegger MJ. Critical Remarks on Reference-Scaled Average Bioequivalence. *J Pharm Pharmaceut Sci.* 2022; 25: 285–96.

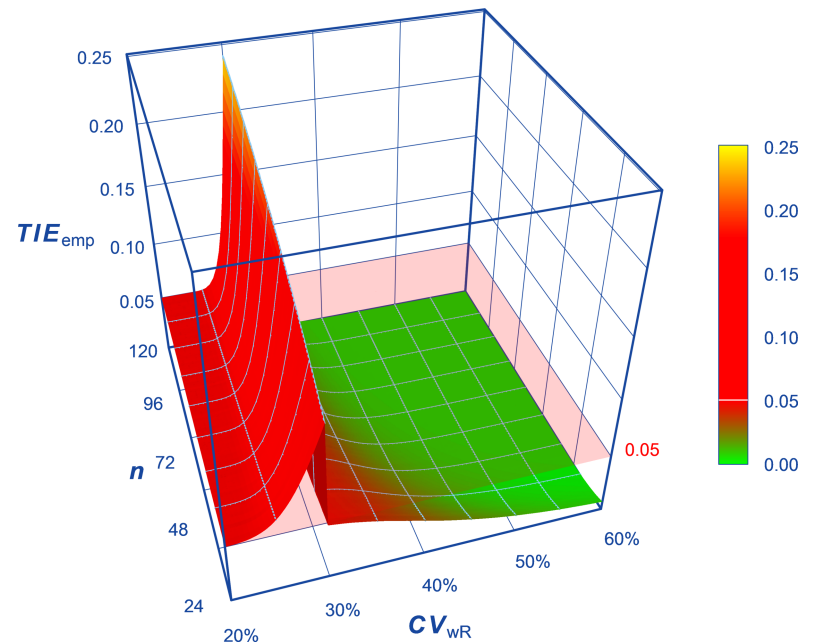
# Consumer risk in SABE

## ABEL (EMA and others)



$TIE_{emp}$  at  $CV_{wR}$  30%;  $n$  24: **0.0804**,  $n$  120: **0.0838**

## RSABE (FDA 'implied limits')



$TIE_{emp}$  at  $CV_{wR}$  30%;  $n$  24: **0.1335**,  $n$  120: **0.2418**

2-sequence 4-period full replicate design,  $CV_{wT} = CV_{wR}$

# Consumer risk in the FDA's RSABE

## Haidar *et al.* (2008), Section 'Results and Discussion'

» Furthermore, a  $\sigma_{w0}$  of 0.25 results in a lower inflation of Type I error compared to a  $\sigma_{w0}$  value of 0.294. Type I error, defined as the risk of concluding two products are bioequivalent when in fact they are not, is 0.05 (or 5%) for average BE. **It is undesirable for any new method to significantly deviate from this value.** «

- 100 runs of  $10^6$  simulations with random seeds ( $CV_{wT} = CV_{wR} = 30\%$ , partial replicate design,  $n = 36$ ), passing studies with  $GMR = 1.25$ :
  - minimum 13.15%
  - median 13.24%
  - maximum 13.90%
- Does  $\approx 13\%$  significantly deviate from 5%?  
**Yes, it does.**
- Does RSABE as implemented show such an undesirable property?  
**Yes, indeed.**

# The FDA's 'desired consumer risk model'

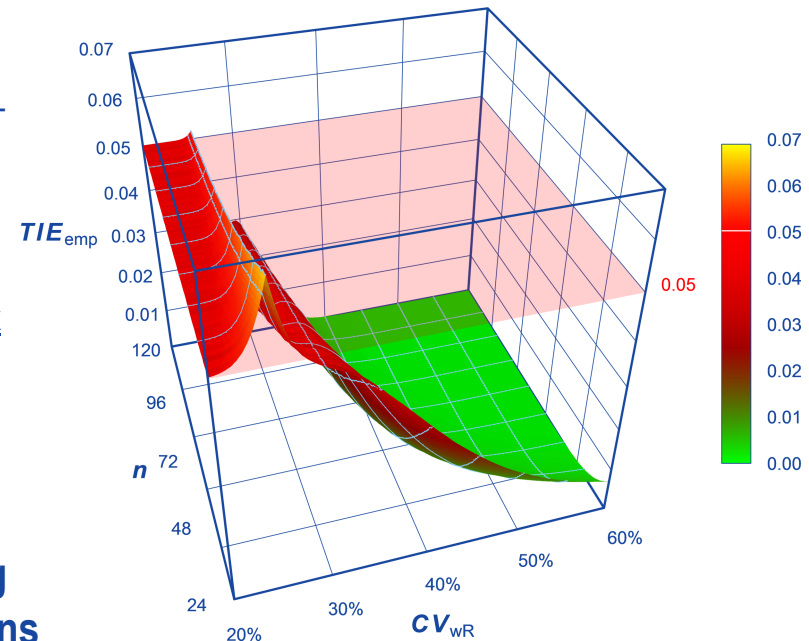
## Empiric Type I Error assessed at

- 0.8000 or 1.2500 if  $s_{WR} \leq 0.25$
- $\exp(\pm\theta_s \cdot s_{WR})$  if  $s_{WR} > 0.25$

Davit *et al.* Implementation of a Reference-Scaled Average Bioequivalence Approach for Highly Variable Generic Drug Products by the US Food and Drug Administration. AAPS J. 2012; 14(4): 915–24.  
<https://doi.org/10.1208/s12248-012-9406-x>

## Section 'Controversies'

» 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. «



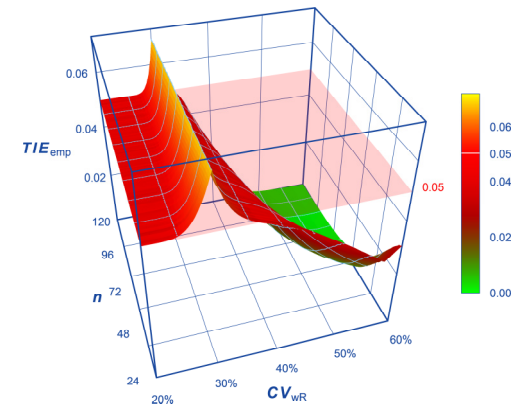
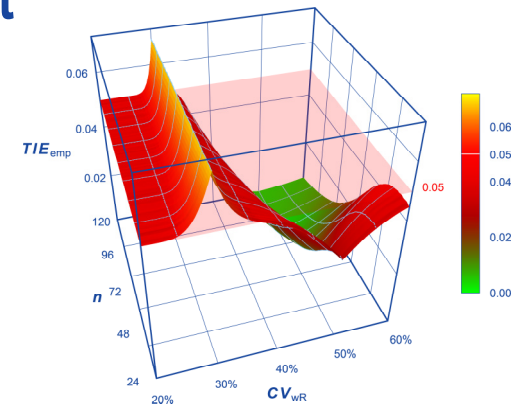
$TIE_{emp}$  at  $CV_{WR} \approx 25.396\%$  ( $s_{WR} 0.25$ );  
 n 24: 0.0663, n 120: 0.0501

# The FDA's proposal for harmonization

The EMA and Health Canada should implement Howe's approximation<sup>1</sup> while keeping their current regulatory conditions<sup>2</sup>

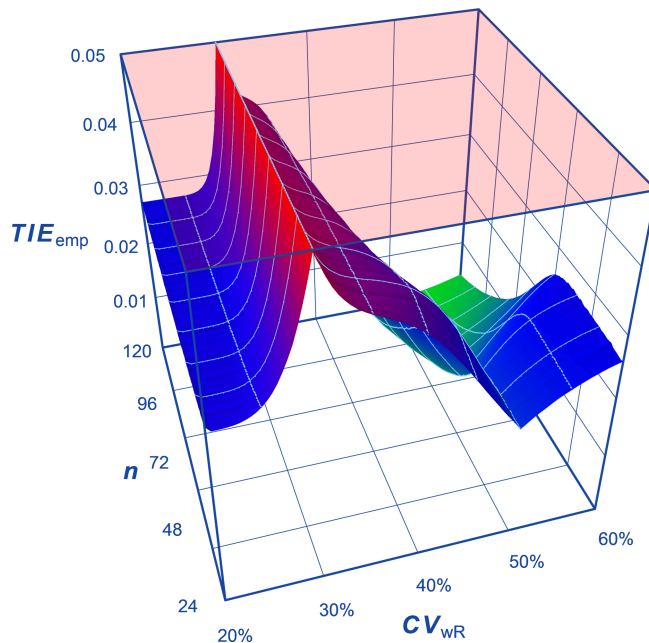
- Regulatory constant  $k = 0.760$
- Cap of scaling
  - EMA  $CV_{wR} = 50\%$        $\max TIE_{emp} \mathbf{0.0686}$
  - HC  $CV_{wR} \approx 57.382\%$        $\max TIE_{emp} \mathbf{0.0690}$

1. Howe W.G. Approximate Confidence Limits on the Mean of  $X + Y$  Where  $X$  and  $Y$  Are Two Tabled Independent Random Variables. *J Am Stat Assoc.* 1974; 69(347): 789–94. <https://doi.org/10.2307/2286019>
2. Muñoz J, Alcaide D, Ocaña J. Consumer's risk in the EMA and FDA regulatory approaches for bioequivalence in highly variable drugs. *Stat Med.* 2016; 35(12): 1933–43. <https://doi.org/10.1002/sim.6834>



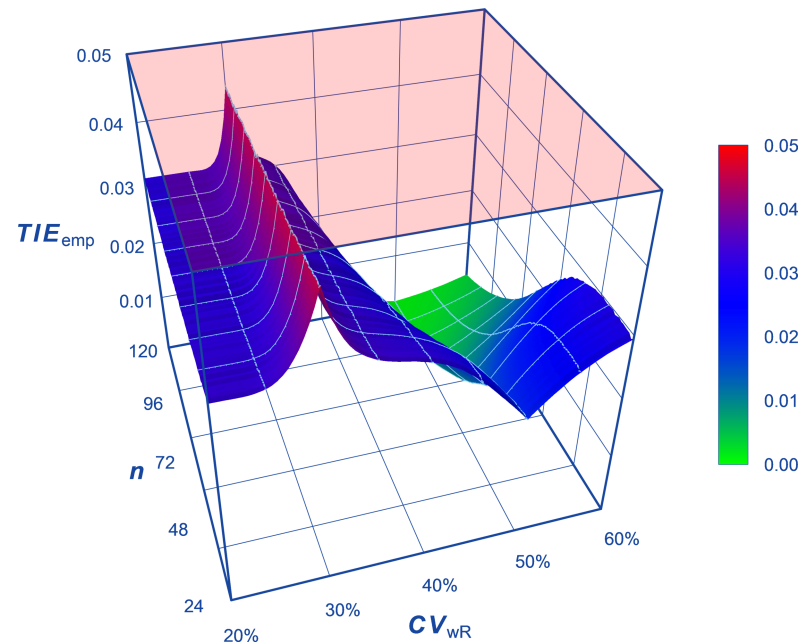
# Alternative: Iteratively adjusted $\alpha$

Molins *et al.* (2017)



$TIE_{emp}$  at  $CV_{wR}$  30%: 0.0500 ✓

Ocaña *et al.* (2019)



$TIE_{emp}$  at  $CV_{wR}$  30%;  $n$  24: 0.0430,  $n$  120: 0.0456 ✓

2-sequence 4-period full replicate design,  $CV_{wT} = CV_{wR}$  (evaluation for the EMA's ABEL)