

Univariate Bioequivalence

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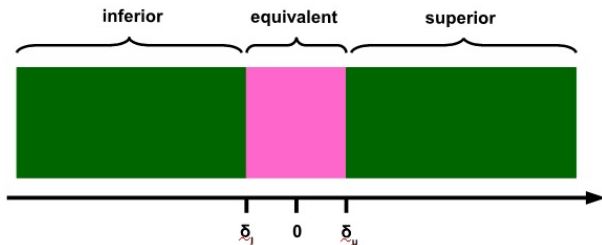


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What is bioequivalence?

Equivalence is not equality



Pharmacological question:

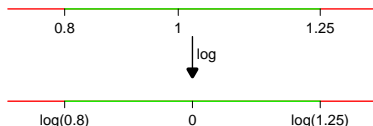
Is test product T equivalent to reference product R in terms of key PK parameters (e.g. AUC_{0-t} , $AUC_{0-\infty}$, C_{max})?

Statistical question:

Is the log-ratio of PK parameters μ^T / μ^R for test and reference within a pre-defined acceptable range $[\Delta_l, \Delta_u]$?

Testing average bioequivalence

A common choice is $\Delta_L = \log(0.8)$ and $\Delta_U = \log(1.25)$ so that the resulting interval $[-0.223, 0.223]$ is symmetric around 0



... but why logarithms?

PK parameters like AUC and C_{max} are typically assumed to be log-normal and therefore logarithmised

Parameter for analysis: $\theta = \log\left(\frac{\mu^T}{\mu^R}\right) = \log(\mu^T) - \log(\mu^R)$

θ is then assumed to be normal with variance σ^2

Two one-sided tests (TOST)

Standard procedure for average bioequivalence: two one-sided t -tests, each at level α (Schuirmann 1987)

1) Test $H_{01}: \theta \leq \Delta_l$ vs. $H_{A1}: \theta > \Delta_l$ at level α

2) Test $H_{02}: \theta \geq \Delta_u$ vs. $H_{A2}: \theta < \Delta_u$ also at level α

3) Conclude bioequivalence if both H_{01} and H_{02} are rejected

$$H_0 \equiv H_{01} \cup H_{02}$$

$$H_A \equiv H_{A1} \cap H_{A2}: -\Delta < \theta < \Delta$$

This procedure controls the type I error rate at level α due to the intersection-union principle (Berger 1982)

A slightly peculiar relationship

The TOST procedure at level α gives the same test decisions as a confidence interval inclusion approach using. . .

1) . . . either a two-sided $100(1 - 2\alpha)\%$ interval around $\hat{\theta}$

$$\left[\hat{\theta} - SE(\hat{\theta})t_{1-\alpha,\nu}, \hat{\theta} + SE(\hat{\theta})t_{1-\alpha,\nu} \right]$$

where $SE(\hat{\theta})$ is the standard error of $\hat{\theta}$, and $t_{1-\alpha,\nu}$ the $100(1 - \alpha)\%$ quantile of Student's t -distribution with ν degrees of freedom

2) . . . or an “expanded” two-sided $100(1 - \alpha)\%$ interval around $\hat{\theta}$

$$\left[\min \left(0, \hat{\theta} - SE(\hat{\theta})t_{1-\alpha,\nu} \right), \max \left(0, \hat{\theta} + SE(\hat{\theta})t_{1-\alpha,\nu} \right) \right]$$

In either case bioequivalence may be concluded if and only if the interval lies entirely within $[\Delta_L, \Delta_U]$ (Berger & Hsu 1996)

Case study

Single-dose ticlopidine hydrochloride (250mg active ingredient) administered as a tablet of commercial reference product Tiklid (R) or test formulation (T)



2 x 2 x 2 crossover study with 24 healthy male volunteers randomised to sequence RT or TR (Marzo *et al.* 2002)

AUC_{0-t} , $AUC_{0-\infty}$, C_{max} data in R package `jocre` (Pallmann 2017)

Case study

Exercises:

1. Calculate the mean and standard error of the difference (T vs. R) of the logarithms of AUC_{0-t} .
2. Test whether formulations T and R are bioequivalent regarding AUC_{0-t} . Use the TOST procedure at $\alpha = 0.05$ with the conventional [80%, 125%] bioequivalence range.
3. Compute the corresponding 90% confidence interval as well as the 95% “expanded” interval.
4. Translate the point estimate and interval boundaries back to the original scale.

Case study: code & results

```
### Dataset
```

```
library(jocre)
data(marzo)
head(marzo, 5)
```

```
## Volunteer Sequence Cmax_T Cmax_R AUC_T AUC_R AUCinf_T AUCinf_R
## 1 1 TR 784.3 878.2 2021.7 2665.2 2131.4 3030.1
## 2 2 RT 304.2 211.7 901.7 685.9 1107.9 798.6
## 3 3 TR 307.3 259.6 741.4 654.0 806.2 696.1
## 4 4 TR 156.7 307.8 475.6 753.7 509.9 833.7
## 5 5 RT 745.6 1036.2 2521.4 2781.9 2784.0 3015.6
```

```
### Differences of logarithms
```

```
marzo$logdiffAUC <- log(marzo$AUC_T) - log(marzo$AUC_R)
head(marzo, 5)
```

```
## Volunteer Sequence Cmax_T Cmax_R AUC_T AUC_R AUCinf_T AUCinf_R
## 1 1 TR 784.3 878.2 2021.7 2665.2 2131.4 3030.1
## 2 2 RT 304.2 211.7 901.7 685.9 1107.9 798.6
## 3 3 TR 307.3 259.6 741.4 654.0 806.2 696.1
## 4 4 TR 156.7 307.8 475.6 753.7 509.9 833.7
## 5 5 RT 745.6 1036.2 2521.4 2781.9 2784.0 3015.6
## logdiffAUC
## 1 -0.27634036
## 2 0.27355003
## 3 0.12543294
## 4 -0.46041725
## 5 -0.09831984
```


Case study: code & results

```
### Mean & SE  
mean(marzo$logdiffAUC)  
  
## [1] -0.08027212  
  
exp(mean(marzo$logdiffAUC))  
  
## [1] 0.9228652  
  
sd(marzo$logdiffAUC) / sqrt(nrow(marzo))  
  
## [1] 0.05876836
```

```
### TOST p-values  
t.test(x=marzo$logdiffAUC, alternative="less", mu=log(1.25))$p.value  
  
## [1] 1.55846e-05  
  
t.test(x=marzo$logdiffAUC, alternative="greater", mu=log(0.8))$p.value  
  
## [1] 0.01162843
```

Case study: code & results

```
### 90% confidence interval

t.test(x=marzo$logdiffAUC, conf.level=0.9)$conf.int

## [1] -0.1809935  0.0204493
## attr(,"conf.level")
## [1] 0.9

exp(t.test(x=marzo$logdiffAUC, conf.level=0.9)$conf.int)

## [1] 0.8344408  1.0206598
## attr(,"conf.level")
## [1] 0.9

### 95% "expanded" confidence interval: same as 90% interval
```

Literature

Berger RL (1982) Multiparameter hypothesis testing and acceptance sampling. *Technometrics*, **24**(4), 295–300.

Berger RL, Hsu JC (1996) Bioequivalence trials, intersection-union tests and equivalence confidence sets. *Statistical Science*, **11**(4), 283–319.

Jaki T, Pallmann P, Wolfsegger MJ (2013) Estimation in AB/BA crossover trials with application to bioequivalence studies with incomplete and complete data designs. *Statistics in Medicine*, **32**(30), 5469–5483.

Marzo A, Dal Bo L, Rusca A, Zini P (2002) Bioequivalence of ticlopidine hydrochloride administered in single dose to healthy volunteers. *Pharmacological Research*, **46**(5), 401–407.

Pallmann P (2017) `jocre`: joint confidence regions. R package version 0.3.3.
<https://cran.r-project.org/package=jocre>

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