### FY 2020 Generic Drug Regulatory Science Initiatives Public Workshop

Breakout 3: In Vitro Bioequivalence Methods Pina D'Angelo May 4, 2020

# **IVPT Data Analysis: Raw Data**

### 1. Missing Data:

'Skin sections (diffusion cells) that are discontinued from the study based upon criteria specified in the study protocol may be replaced with new skin sections....' (Acyclovir, FDA draft guidance 2016).

- The purpose of replacing missing data as per the guidance is to ensure balanced data, but does this introduce bias?
- Are there mathematical methods to account for missing data instead of replacement, i.e., a weighted average?

## **IVPT Data Analysis: Raw Data**

#### 2. Zero Values:

Logarithmic transformation of zero values are non existent ( $-\infty$ ) and as such, a profile with zeros will result in that data being excluded from the inferential statistical analyses.

- Initial research suggests that imputing  $\frac{1}{2}$  LOQ generally appears to <u>over</u> estimate  $s_{WR}$  and that treating BLOQ values as missing generally appears to <u>under</u> estimate  $s_{WR}$  leading to an <u>increase</u> type-I error.
- Imputation with the LOQ seemed to behave well, however, more research is needed.
- Other, more sophisticated methods of imputation would be interesting to research.

# **IVPT Data Analysis: Outliers**

- 1. There are multiple methods of outlier detection.
  - Which is most suited for IVPT?
- The recommended statistical methodology to evaluate BE for IVPT includes a mixed criterion that uses s<sub>WR</sub> as a cutoff point.
  - Falsely declaring a value to be outlying may lead to the removal of pertinent data, which may in turn affect BE conclusions.
  - BUT not excluding data that is legitimately outlying may lead to over estimating  $s_{WR}$  and perhaps concluding that BE criteria is met using the RSABE method when in fact the product may not be highly variable.
  - Conducting sensitivity analyses to examine the influence of the removal of outlying value(s) may be warranted.

# **IVPT Data Analysis: ABE Modelling**

 When s<sub>WR</sub> ≤ 0.294, <u>ABE</u> analyses are in order. To declare T and R bioequivalent, the (1-2α)\*100% confidence interval :

$$exp(\overline{I} \pm t_{(n-1),\infty} * \sqrt{S_I^2/n})$$

must be contained within the limits [1/m, m]. (Acyclovir FDA draft guidance)

- Initial research suggests that the statistical method of analyses from the Acyclovir FDA draft guidance may be more powerful in concluding BE but also suggests an <u>increase</u> in type I error compared to using ANOVA suited for replicate data.
- Further research would be warranted in this area.