

Virtual bioequivalence workflow

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FDA grant U01FD006549: Virtual bioequivalence (VBE) workflow

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- ▶ Scientific officer:
 - ▶ Eleftheria Tsakalozou
- ▶ Objective: VBE assessment of reference and test formulations

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- Reference: in vitro dissolution profile + PK data
- Test: in vitro dissolution profile



Build API oral model with reference IV and solution/IR data



Identify relevant physiologic parameters that affect PK using sensitivity and uncertainty analysis



Identify posterior distributions and correlations for relevant physiologic parameters and formulation properties based on individual PK data for reference using “digital twins”



Generate a virtual population that captures the measured variability



Simulate individual plasma concentration vs time profiles for reference and test



Use non-compartmental analysis to get AUC, C_{max} and T_{max} and complete BE statistics appropriate for study design

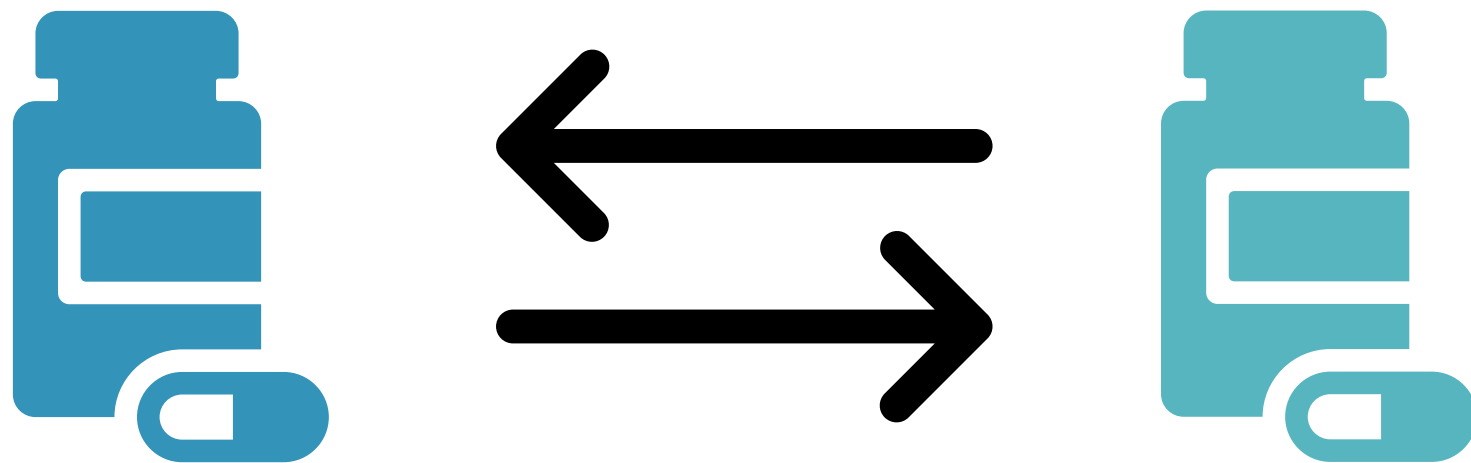


Assess VBE by analyzing the probability of demonstrating BE as a function of number of participants and sampling



Applications

- ▶ Predicting establishment of bioequivalence for a test formulation
- ▶ Predicting a “dissolution safe space”
- ▶ Optimizing clinical trial design
- ▶ Risk assessment
 - ▶ Generic drug development
 - ▶ Pre-/post-approval formulation changes
 - ▶ Scale-up and post-approval changes



Virtual bioequivalence workflow

Requirements for workflow

- ▶ PK-Sim model for molecule (.pkml)
 - ▶ Dissolution profiles for reference and test formulations
 - ▶ Sensitive model parameters as informed by local sensitivity analysis in PK-Sim, previously assessed and capable of capturing IIV
- ▶ Observed plasma concentration-time profiles for study population
- ▶ Study population demographics
(species, population, sex, weight, height, age, BMI, gestational age)

Building an oral PK-Sim model

- ▶ Physicochemical properties of the drug
- ▶ Known physiological processes
- ▶ IV data informs systemic parameters of disposition
- ▶ Solution/IR data informs system parameters of absorption
- ▶ We are assuming that the plasma concentration-time profiles are sensitive to formulation input (e.g. dissolution profile)
- ▶ At the end of this step, we have...
 - ▶ PBPK model that describes systemic disposition and absorption of API



Case study: Bupropion

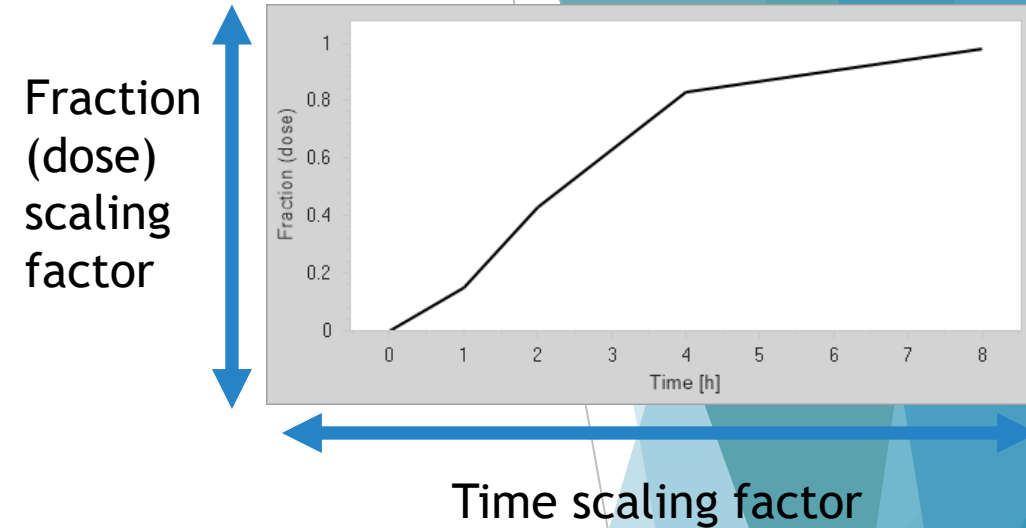
- ▶ Note: Bupropion is just a pilot model used to demonstrate a crossover virtual bioequivalent assessment
- ▶ Bupropion PK-Sim models:
 - ▶ Reference: SR 150 mg
 - ▶ Test: XL 150 mg
 - ▶ Sensitive parameters:
 - ▶ Enzyme concentration, Dissolution scaling factors
- ▶ Plasma concentration-time profiles for 32 individuals
- ▶ Study population demographics (sex, population, weight, height)

Capturing posterior distributions - Non-parametric optimal design (NPOD)

- ▶ Study individuals are recreated as “digital twins” in PK-Sim using demographic data available
- ▶ Parameter ranges are specified for sensitive parameters to be fit
- ▶ Input: reference formulation model, observed data, parameter ranges, study individuals (error function, initial support points)
- ▶ Output:
 - ▶ Posterior distributions (support points and weights)
 - ▶ Capable of capturing non-normal distributions, e.g. poor and extensive metabolizers
 - ▶ Correlation matrix between parameters and population demographics (weight, height)

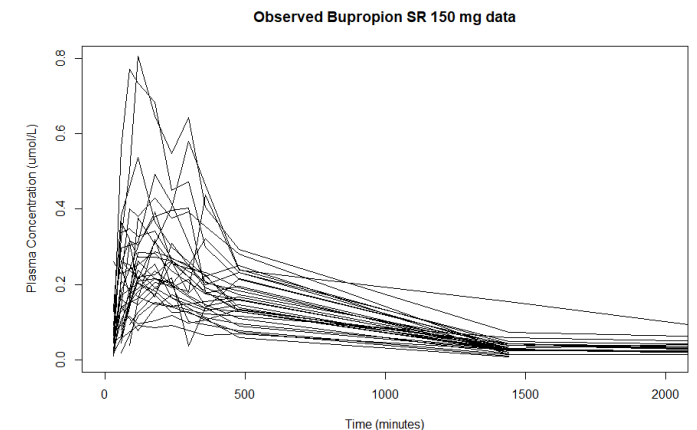
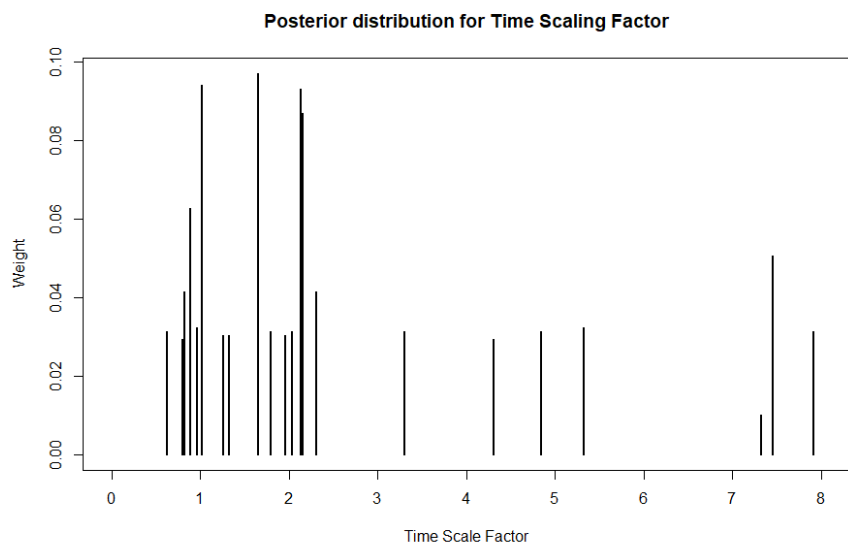
Establishing an in-vitro-in-vivo relationship (IVIVR)

- ▶ Mechanistic:
Formulation parameters (e.g. particle size distribution)
- ▶ In the absence of *a priori* knowledge:
Dissolution scaling factors
- ▶ Reference data (formulation input and plasma concentration-time profiles) is used to calibrate the IVIVR
- ▶ We assume that the IVIVR applies to test formulations
 - ▶ Ideally, the test formulation is similar in excipients and release mechanisms compared to the reference formulation



Case study: Bupropion

- Reference bupropion model: SR 150 mg
- Parameter ranges are specified for sensitive parameters to be fit
 - Time scaling factor on dissolution profile (i.e. x-axis) - [0,10]
 - Fraction (dose) scaling factor on dissolution profile (i.e. y-axis) - [0,1000]
 - Enzyme concentration - [0,1000]



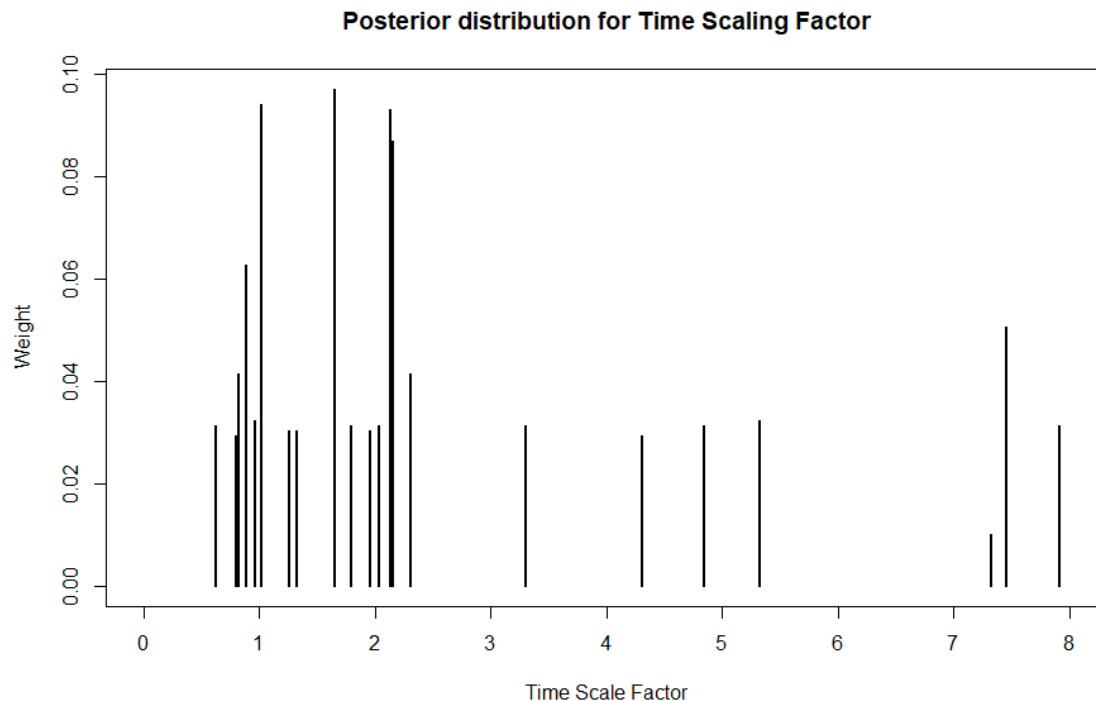
ID	Sex	Population	Weight (kg)	Height (cm)
1	Female	European	72.7	167.64
2	Female	European	84.4	162.56
3	Male	BlackAme	73.4	172.72
4	Male	MexicanA	79.8	167.64
5	Female	European	91.1	181.61
6	Female	Asian	74	172.72
7	Male	Asian	77.7	170.18
8	Female	European	73.5	167.64
9	Male	European	80.6	187.96
10	Male	European	94.7	170.18

```
> corr_matrix
      theta1      theta2      theta3      weight      Height
theta1  1.00000000  0.5655141 -0.64178739 -0.30277260 -0.09624631
theta2  0.56551415  1.0000000 -0.39004246 -0.23190258 -0.13329034
theta3 -0.64178739 -0.3900425  1.00000000  0.06232417  0.13168306
weight -0.30277260 -0.2319026  0.06232417  1.00000000  0.38472801
Height -0.09624631 -0.1332903  0.13168306  0.38472801  1.00000000
```



Case study: Bupropion

- ▶ A reference population of 1000 individuals is generated that maintains the captured correlations from the non-parametric population algorithm



	theta1	theta2	theta3	weight	height
1	1.2622520	137.60412	813.9782	107.93443	19.36761
2	0.8930129	43.19603	820.1831	61.03642	17.51424
3	1.6527643	47.60304	678.9087	99.33016	17.54653
4	2.1600338	62.21984	820.1831	86.89934	17.39655
5	2.0332647	110.45191	821.0817	55.76694	17.20736
6	0.8022292	55.45703	841.1291	78.72036	17.00976
7	2.3100450	100.05055	820.1831	62.96131	16.68466
8	0.8930129	59.12914	946.9103	61.52944	18.16568
9	4.3044085	58.06531	933.7260	55.49252	17.55697
10	1.9635224	43.19603	933.7260	57.89853	18.83575

```
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theta3 -0.64178739 -0.3900425  1.00000000  0.06232417  0.13168306  
weight -0.30277260 -0.2319026  0.06232417  1.00000000  0.38472801  
Height -0.09624631 -0.1332903  0.13168306  0.38472801  1.00000000
```

Generate virtual population and PK profiles

- ▶ OSPsuite is used to generate a virtual population
- ▶ The virtual population parameters are updated based on the reference population
- ▶ PK-Sim is used to generate the PK profiles for both the reference and test formulations



Case study: Bupropion

- ▶ A virtual population of 1000 individuals is generated
- ▶ Virtual individuals are compared to reference population and the appropriate parameter values are updated

	theta1	theta2	theta3	weight	height
1	1.2622520	137.60412	813.9782	107.93443	19.36761
2	0.8930129	43.19603	820.1831	61.03642	17.51424
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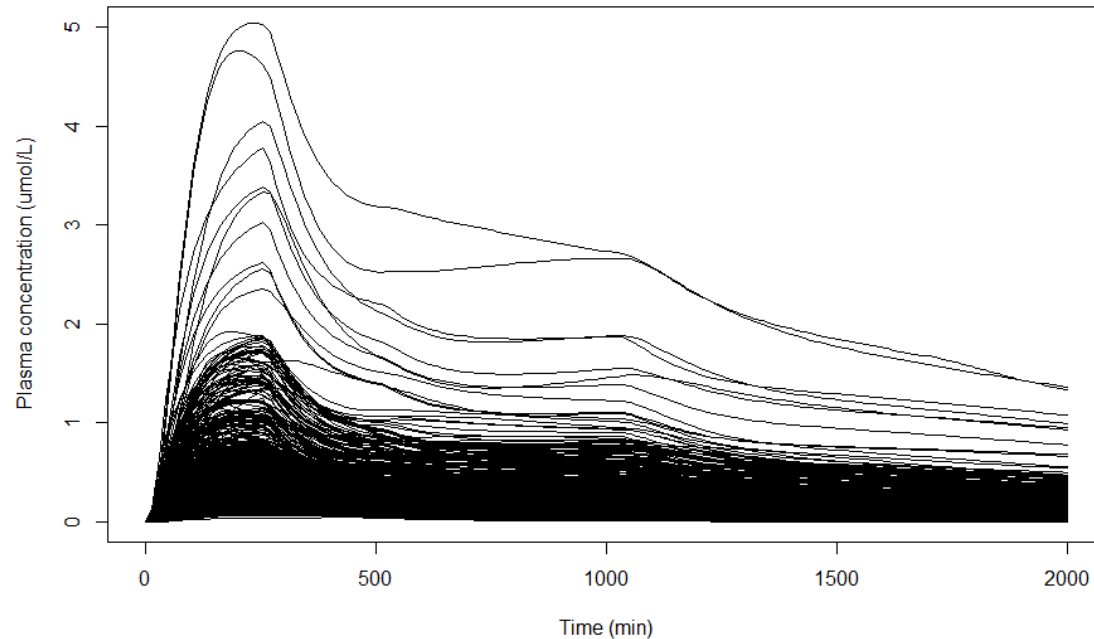
IndividualId	Gender	Population	Organism Weight	Organism Height	Applications PO	Applications PO 15	Liver and Intestina
0	MALE	European_ICRP_2002	69.67197648	17.34863337	1.652764258	43.1960325	966.1631612
1	MALE	European_ICRP_2002	80.66367394	17.31971653	0.893012886	55.45703163	946.9102584
2	MALE	European_ICRP_2002	11.6418776	7.739270429	4.304408454	137.6041153	678.9087472
3	MALE	European_ICRP_2002	73.25415791	17.61428916	0.802229246	43.1960325	986.9442911
4	MALE	European_ICRP_2002	86.73495227	17.28527499	2.1600338	62.21983888	820.1831479
5	MALE	European_ICRP_2002	67.65264677	17.44606382	2.138830071	100.0505509	373.7770325
6	MALE	European_ICRP_2002	63.98918339	16.50535396	7.454465816	110.4519128	886.9854164
7	MALE	European_ICRP_2002	13.79027881	9.18601242	4.304408454	137.6041153	678.9087472
8	MALE	European_ICRP_2002	62.59547048	16.41364181	7.454465816	82.09933596	670.7724672
9	MALE	European_ICRP_2002	61.88576613	17.19057429	4.839407277	82.09933596	813.9781987
10	MALE	European_ICRP_2002	55.92556148	15.07635357	1.963522405	41.58242253	820.1118292



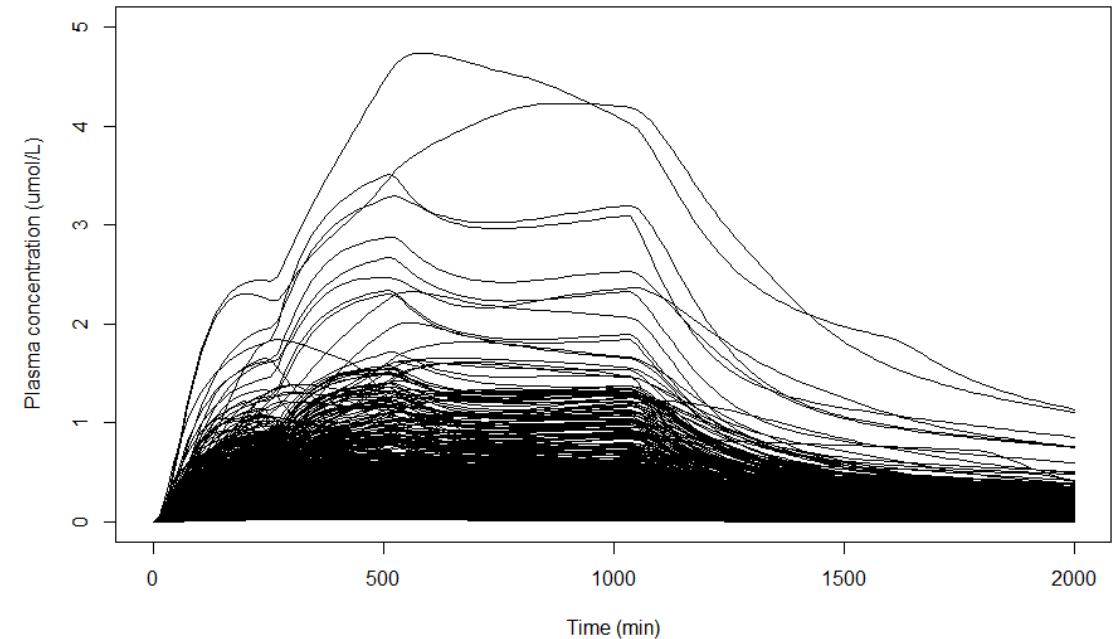
Case study: Bupropion

- ▶ Plasma concentration-time profiles are generated for the virtual population for the reference formulation and for the test formulation
- ▶ We assume that the IVIVR applies to test formulations

Bupropion SR 150 mg



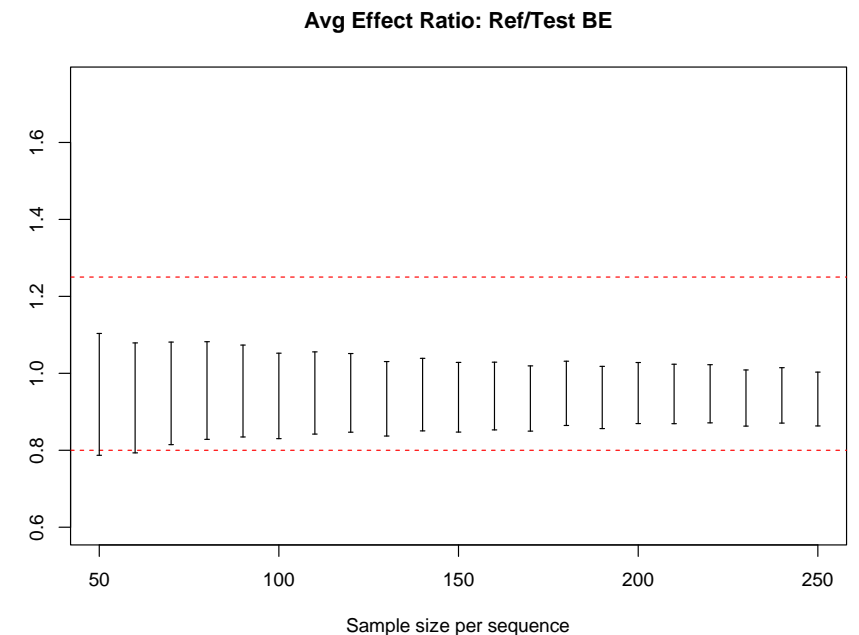
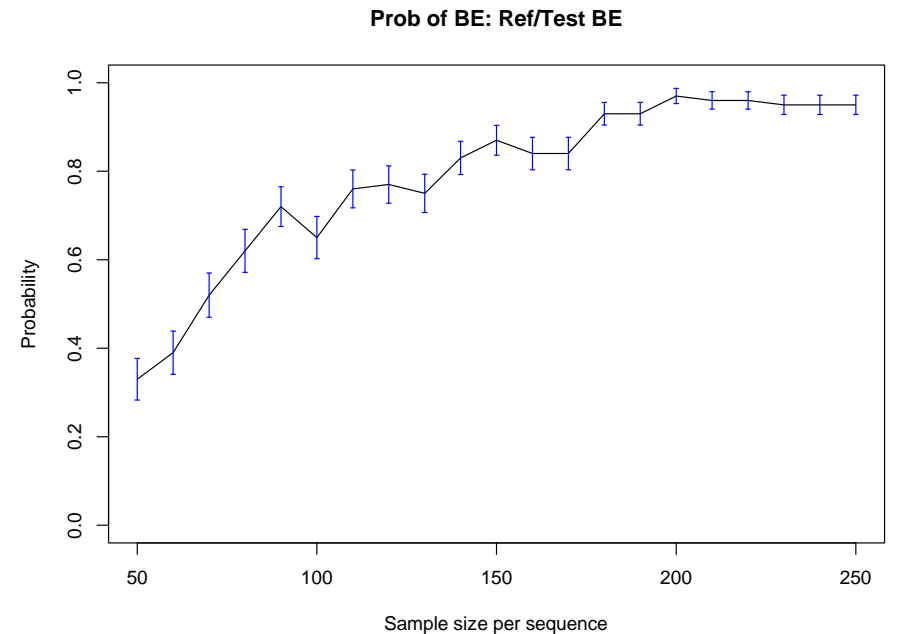
Bupropion ER 150 mg



Clinical trial simulator (CTS)

- *in progress*

- ▶ Input: Simulated plasma concentration-time profiles for two formulations (reference and test)
- ▶ CTS calculates the summary statistics (AUC, C_{max}, etc.)
- ▶ Virtual bioequivalence would then be evaluated from these summary statistics
- ▶ Multiple scenarios possible:
 - ▶ Parallel, i.e. different populations, different formulations
 - ▶ Crossover, no replication (AB)
 - ▶ Crossover with partial or full replication (e.g. TTR, TTRR) with IOV



Inter-occasion variability (IOV)

- *in progress*

▶ Mechanistic approach

- ▶ Identify physiological parameters that are affected by IOV for the molecule of interest
 - ▶ Use NPOD to identify posterior distributions and correlations for these parameters (in addition to other fitted parameters)
 - ▶ Sample from these distributions during the generation of the virtual populations
- ▶ IOV is incorporated when the virtual populations are generated, i.e. before plasma concentration-time profiles are simulated

Applications

- ▶ Predicting establishment of bioequivalence for a test formulation
- ▶ Creating a “dissolution safe space”
- ▶ Optimizing clinical trial design
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Deliverables related to FDA grant

- ▶ Virtual bioequivalence workflow - R package
 - ▶ PK-Sim
 - ▶ OSP Suite R toolbox (ospsuite, will be discussed in Session 4) (<https://github.com/Open-Systems-Pharmacology/OSPSuite-R>)
 - ▶ Non-parametric optimal design (NPOD)
 - ▶ Clinical trial simulator (CTS)
- ▶ Mechanistic absorption models (will be discussed in Session 4)
 - ▶ Dermal model (<https://github.com/Open-Systems-Pharmacology/Skin-permeation-model>)
 - ▶ Inhalation model (<https://github.com/Open-Systems-Pharmacology/Inhalation-model>)
- ▶ The VBE workflow will be available on the Open Systems Pharmacology Github page (<https://github.com/Open-Systems-Pharmacology>)
 - ▶ Current goal: August 2021

Acknowledgements

- ▶ Andrea Edginton
(University of Waterloo)
 - ▶ Cindy Hoi Ting Yeung
 - ▶ Dagmar Hajducek
 - ▶ Abdullah Hamadeh
- ▶ Jörg Lippert (Bayer)
 - ▶ André Dallman
 - ▶ Juri Solodenko
 - ▶ Rolf Burghaus
- ▶ Michael Neely
(Children's Hospital Los Angeles)
 - ▶ Julian Oltalvaro
 - ▶ Walter Yamada
 - ▶ Alona Kryshchenko (CSUCI)
 - ▶ Jay Bartroff (USC)
- ▶ Eleftheria Tsakalozou (FDA)