

Virtual bioequivalence workflow

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FDA grant U01FD006549: Virtual bioequivalence (VBE) workflow

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 - ▶ Andrea Edginton - University of Waterloo
 - ▶ Jörg Lippert - Bayer
- ▶ Scientific officer:
 - ▶ Eleftheria Tsakalozou
- ▶ Objective: Develop platform within OSP framework for VBE assessment

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 - ▶ Reference: in vitro dissolution profile + individual 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 analyses



Identify posterior distributions based on individual PK data 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

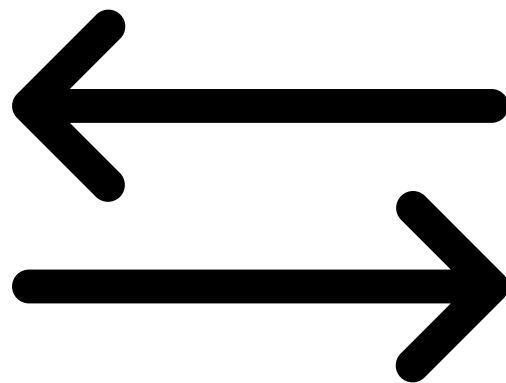


Optimize clinical trial design (e.g number of participants and sampling)



Applications

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



Virtual bioequivalence workflow

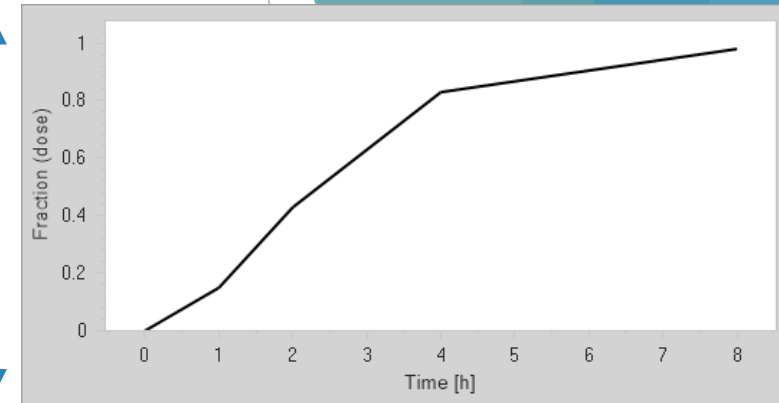
Building an oral PK-Sim model

- ▶ Physicochemical properties of the drug
- ▶ Known physiological processes
- ▶ IV data informs systemic parameters of disposition
- ▶ Solution/IR data informs systemic 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

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

Fraction
(dose)
scaling
factor



Time scaling factor



Case study: Bupropion

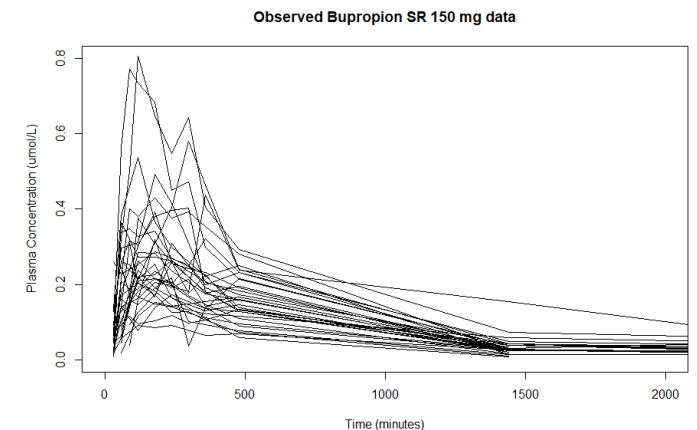
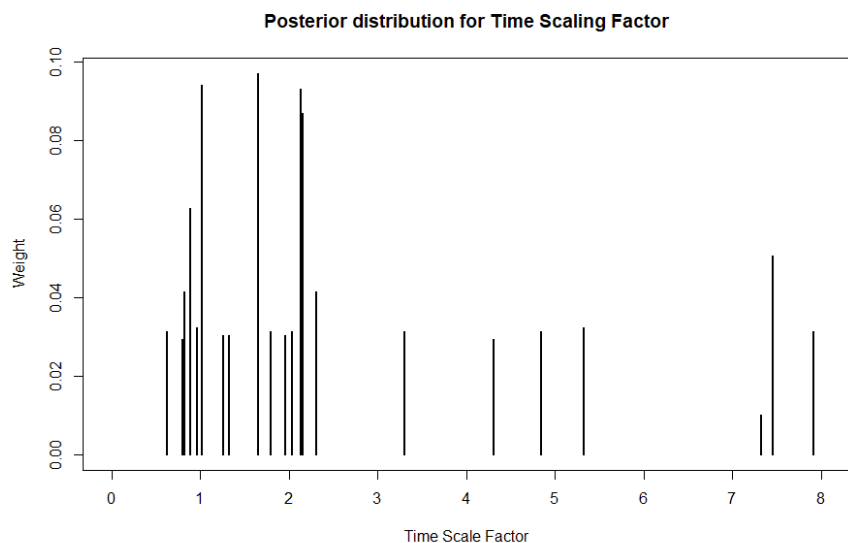
- ▶ Note: Bupropion is just a pilot model used to demonstrate a crossover virtual bioequivalence assessment
- ▶ Bupropion PK-Sim models:
 - ▶ Reference: SR 150 mg
 - ▶ Test: XL 150 mg
 - ▶ Sensitive parameters:
 - ▶ Enzyme reference concentration, Dissolution scaling factors
- ▶ Plasma concentration-time 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: API oral model, observed data, parameter ranges, study individuals
- ▶ 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)

Case study: Bupropion

- ▶ Reference formulation: 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 reference concentration - [0,1000]



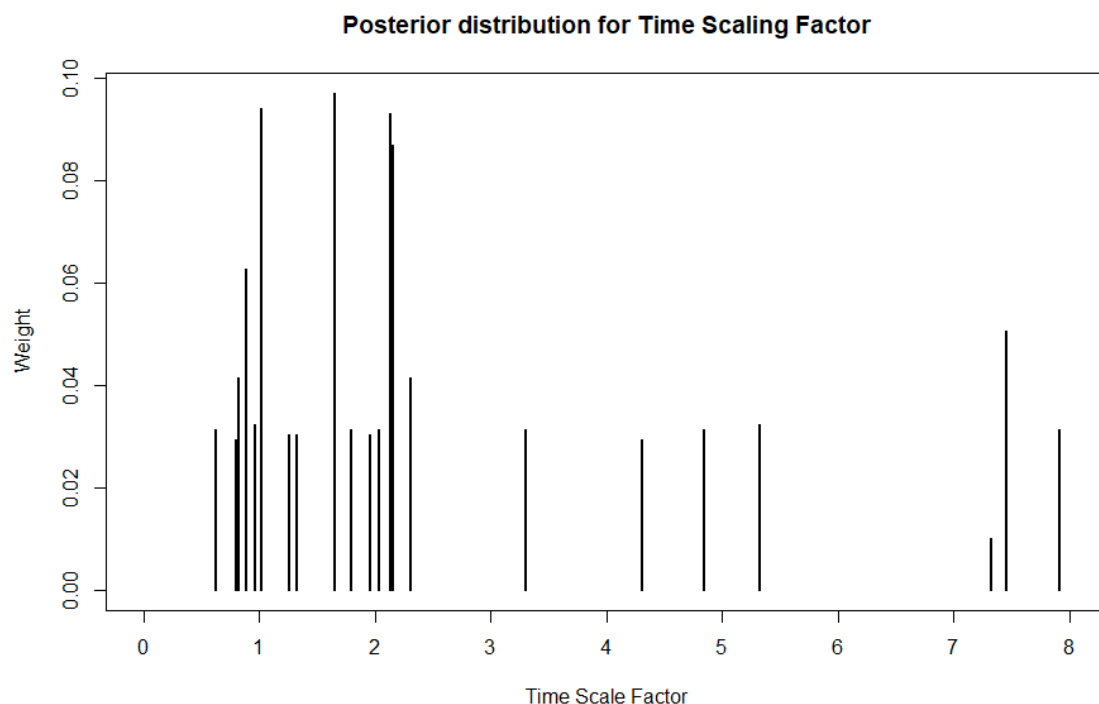
ID	Sex	Populatio	Weight (k	Height (cm)
1	Female	European	72.7	167.64
2	Female	European	84.4	162.56
3	Male	BlackAme	73.4	172.72
4	Male	MexicanA	79.8	167.64
5	Female	European	91.1	181.61
6	Female	Asian	74	172.72
7	Male	Asian	77.7	170.18
8	Female	European	73.5	167.64
9	Male	European	80.6	187.96
10	Male	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 table of 1000 parameter sets 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

```
> corr_matrix
```

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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

- ▶ OSP suite R toolbox is used to generate a virtual population
- ▶ The virtual population is updated based on the reference table
- ▶ 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 table and the appropriate parameter values are updated

	theta1	theta2	theta3	weight	height
1	1.2622520	137.60412	813.9782	107.93443	19.36761
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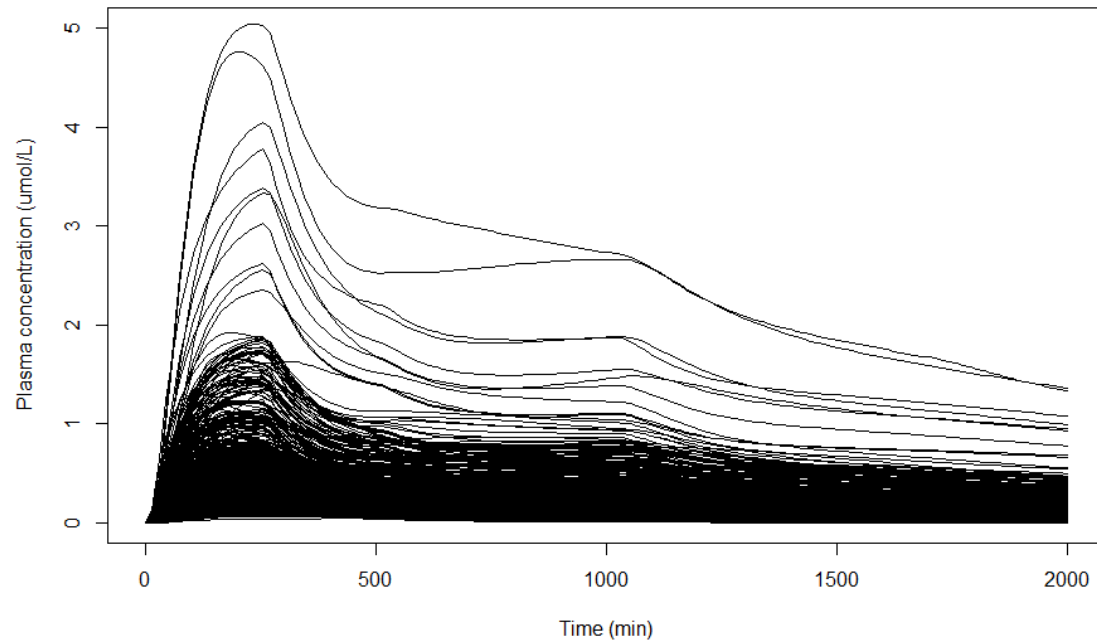
IndividualId	Gender	Population	Organism Weight	Organism Height	Applications PO	Applications PO 15	Liver and Intestina
0	MALE	European_ICRP_2002	69.67197648	17.34863337	1.652764258	43.1960325	966.1631612
1	MALE	European_ICRP_2002	80.66367394	17.31971653	0.893012886	55.45703163	946.9102584
2	MALE	European_ICRP_2002	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



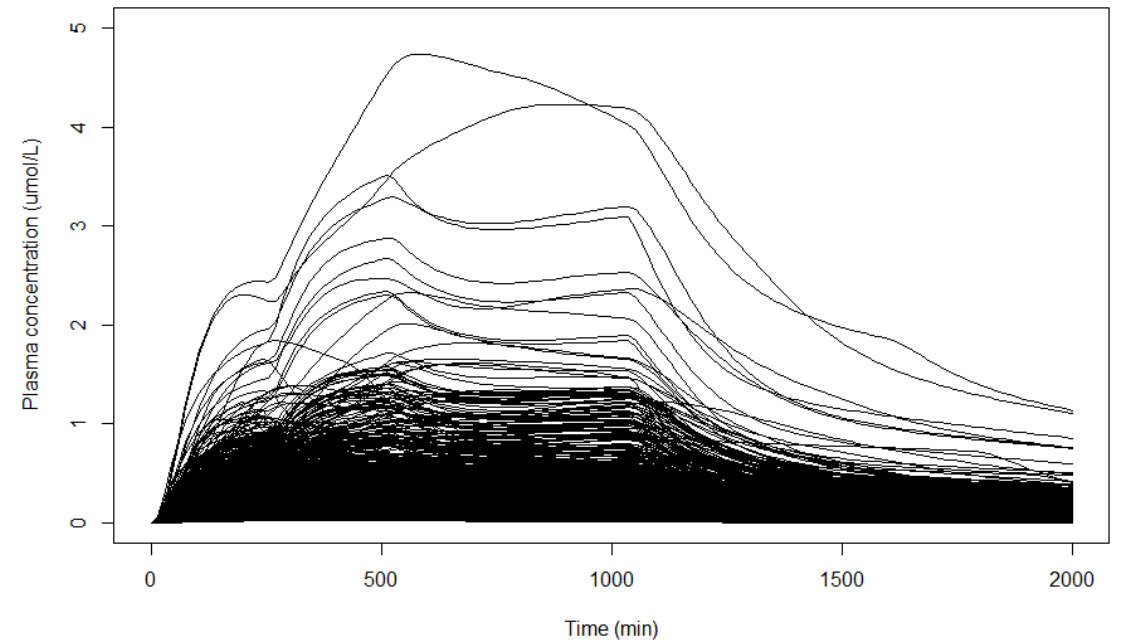
Case study: Bupropion

- ▶ 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



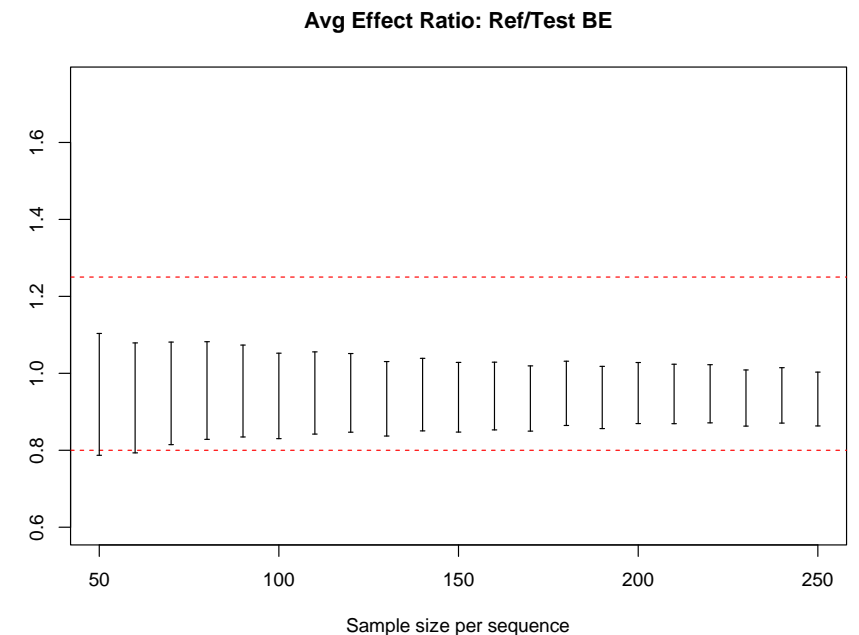
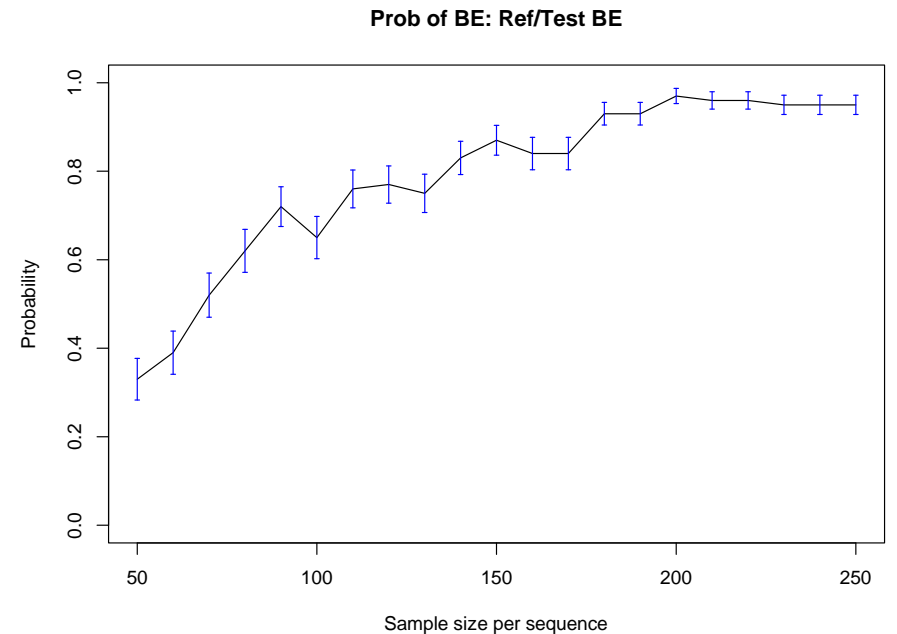
Bupropion ER 150 mg



Clinical trial simulator (CTS)

- *in progress*

- ▶ Input: Simulated plasma concentration-time profiles for two formulations (reference and test)
- ▶ CTS calculates the summary statistics (AUC, Cmax, 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 given API
 - ▶ 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”
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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
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 - ▶ Cindy Hoi Ting Yeung
 - ▶ Dagmar Hajducek
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 - ▶ Jay Bartroff (USC)
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