



## Reproducibility and Industrialization of the *in vitro* Hollow-Fiber System (HFS-TB)

Debra Hanna, Executive Director, Critical Path to TB Drug Regimens  
20 March 2017

## Critical Path to TB Drug Regimens Methodologies Landscape

- Methods landscape
- Academic approach to method development **versus**
- Methodologies designed as drug development tools

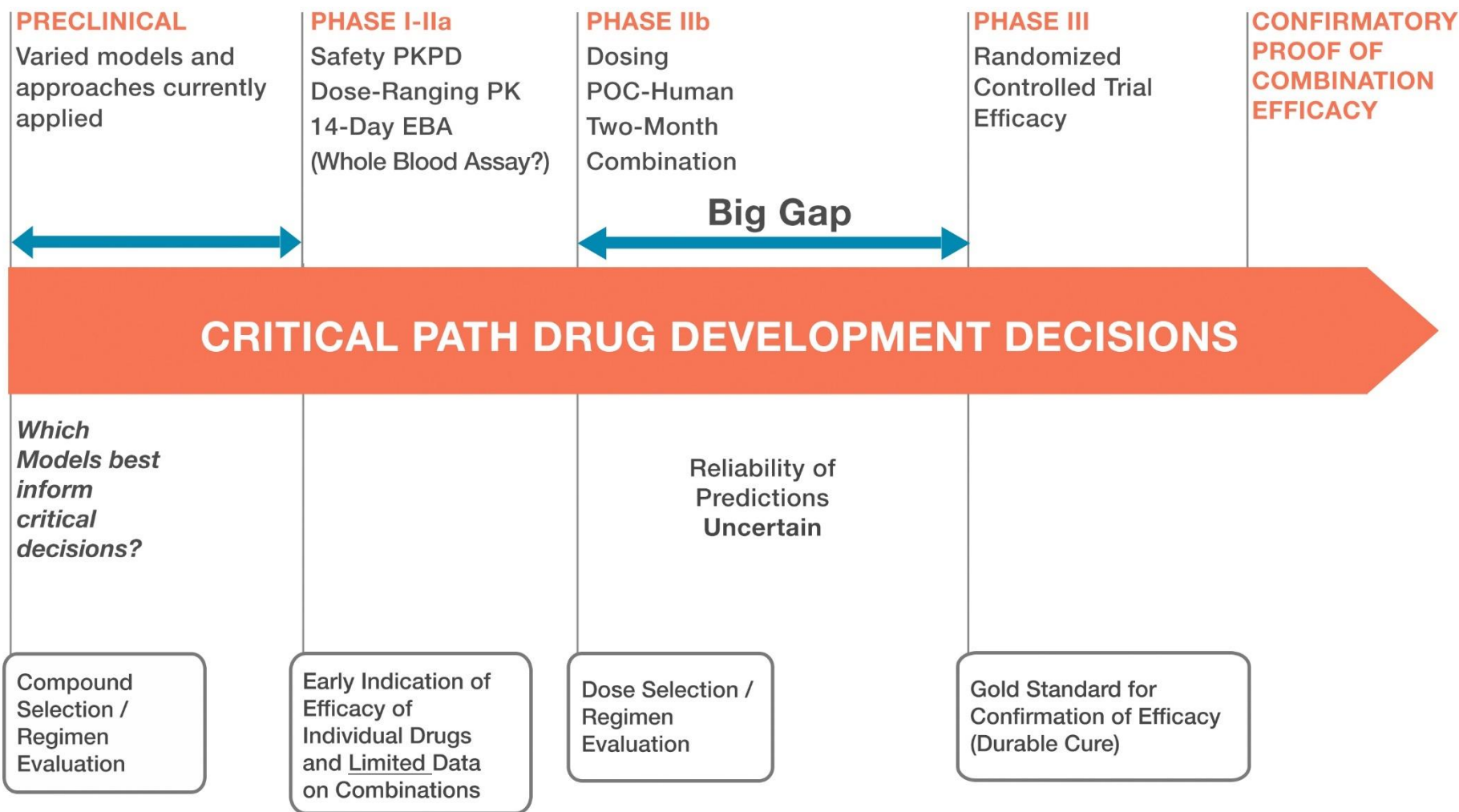
## *In vitro* HFS-TB Model

- HFS-TB model system
- CPTR multi-stage assessment evidence-based approach
- Predictive accuracy assessment leading to EMA qualification for use

## Reproducibility Assessment, Industrialization, Implementation

- Design and approach
- Summary results of experiments
- New work

# Current TB Regimen Development Risk of Late-Stage Attrition



## PRECLINICAL

### IN VITRO

- Static drug concent.
- HFS-TB

## PHASE I-IIa

- Safety PKPD
- Dose-Ranging PK
- 14-Day EBA

## PHASE IIb

- Dosing
- POC-human

## PHASE III

- Randomized
- Controlled Trial
- Efficacy

## CONFIRMATORY PROOF OF COMBINATION EFFICACY

Big Gap

# CRITICAL PATH DRUG DEVELOPMENT DECISIONS

### IN VIVO

- Murine models
- Guinea pig
- Rabbit
- Non-human
- Primate

PBPK  
Modeling

Accurate  
IVIVE  
Extrapolation

Accurate  
PKPD  
Translation

Early Indication of  
Efficacy of Individual  
Drugs and Data on  
Combinations

Dose Selection /  
Regimen Evaluation

Quantitative  
Assessment of Liquid  
Culture Biomarker

Population PKPD

Model-Based  
Meta-Analysis  
Phase III Trials

Systems  
Pharmacology/  
Mechanism-Based  
Models

PENULTIMATE TB CLINICAL TRIAL  
SIMULATION TOOL

Increase Reliability of  
Predictions for Dose  
Selection and  
Efficacy Outcomes

## Mission

- Evidence-based evaluation of innovative drug development tools to address preclinical to clinical translation
- Focus on *in vitro* methodologies supporting efficacy and safety toxicology assessment
- Submission for regulatory endorsement

## Goal

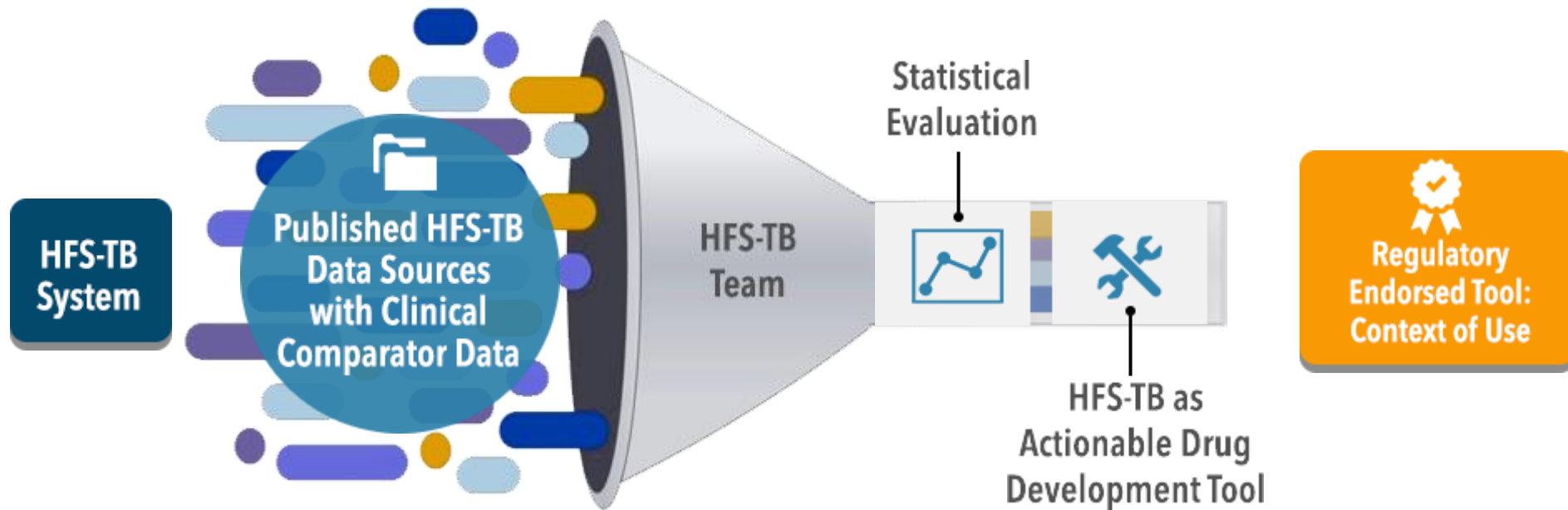
- Follow EMA and FDA Guidance on novel methodology and DDT qualification
- Gather all relevant published and unpublished data sources or aggregation
- Assess clinical translation of innovative preclinical novel methodologies/DDTs to test new TB drug candidates and regimens

## HFS-TB Evidence

- Significantly more quantitative HFS-TB PKPD data available than for any *in vivo* methodology for TB
- Supported thorough assessment of predictive accuracy for clinical outcomes



# Supportive Data Aggregation



# Regulatory Interactions for HFS-TB



FEBRUARY 20, 2013

LOI submission

FEBRUARY 27, 2013

LOI discussion

OCTOBER 16, 2013

VXDS document  
submission

NOVEMBER 15, 2013

VXDS meeting

FEBRUARY 4, 2014

Submission of  
comments to FDA  
draft guidance



EUROPEAN MEDICINES AGENCY  
SCIENCE MEDICINES HEALTH

Briefing document  
submission (for  
qualification  
opinion)

FEBRUARY 24, 2014

SAWP meeting

MAY 6, 2014

Draft  
qualification  
opinion

NOVEMBER 18, 2014

Public comment  
period

NOVEMBER 18, 2014 –  
JANUARY 9, 2015

Final  
qualification  
opinion

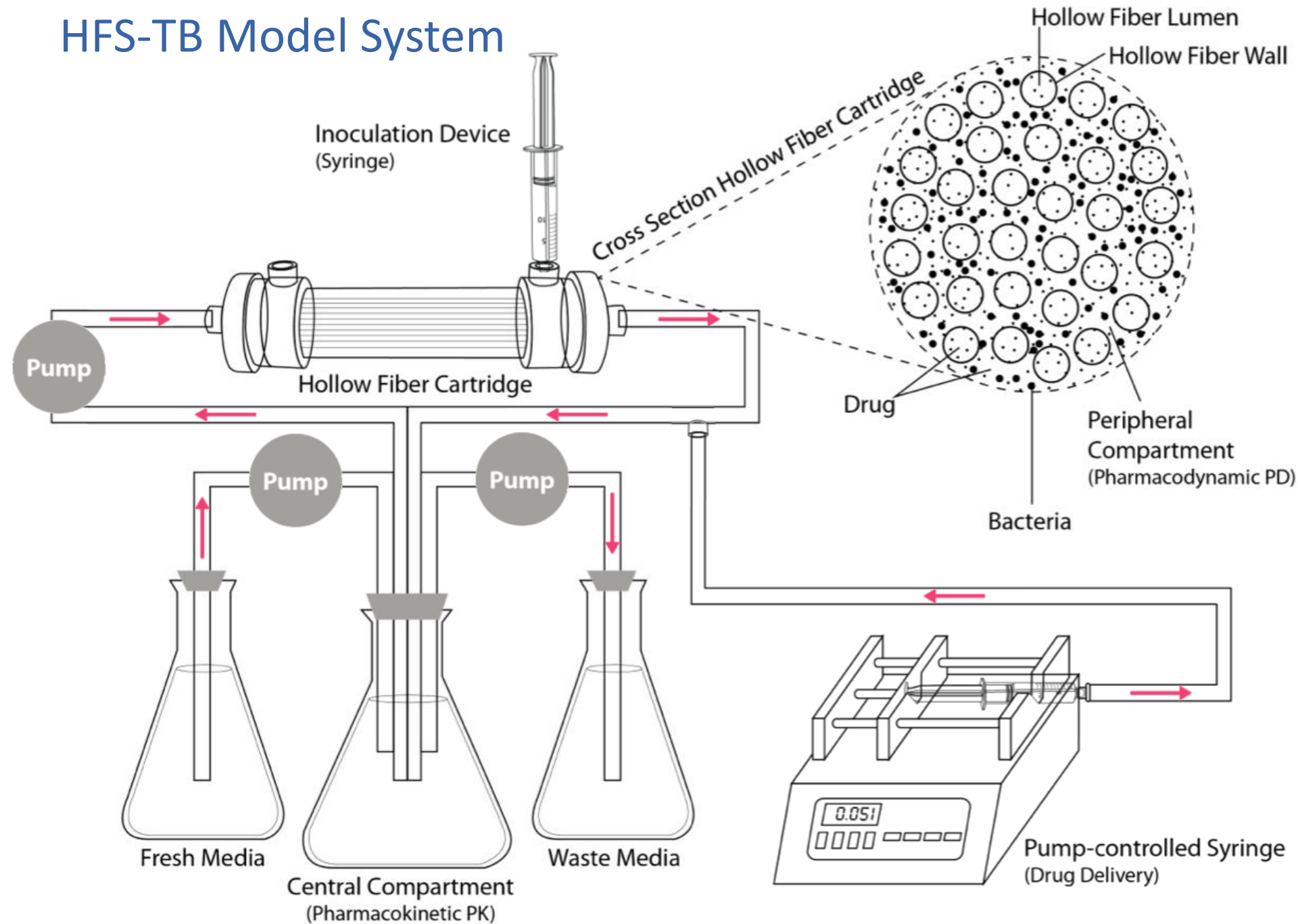
JANUARY 26, 2015

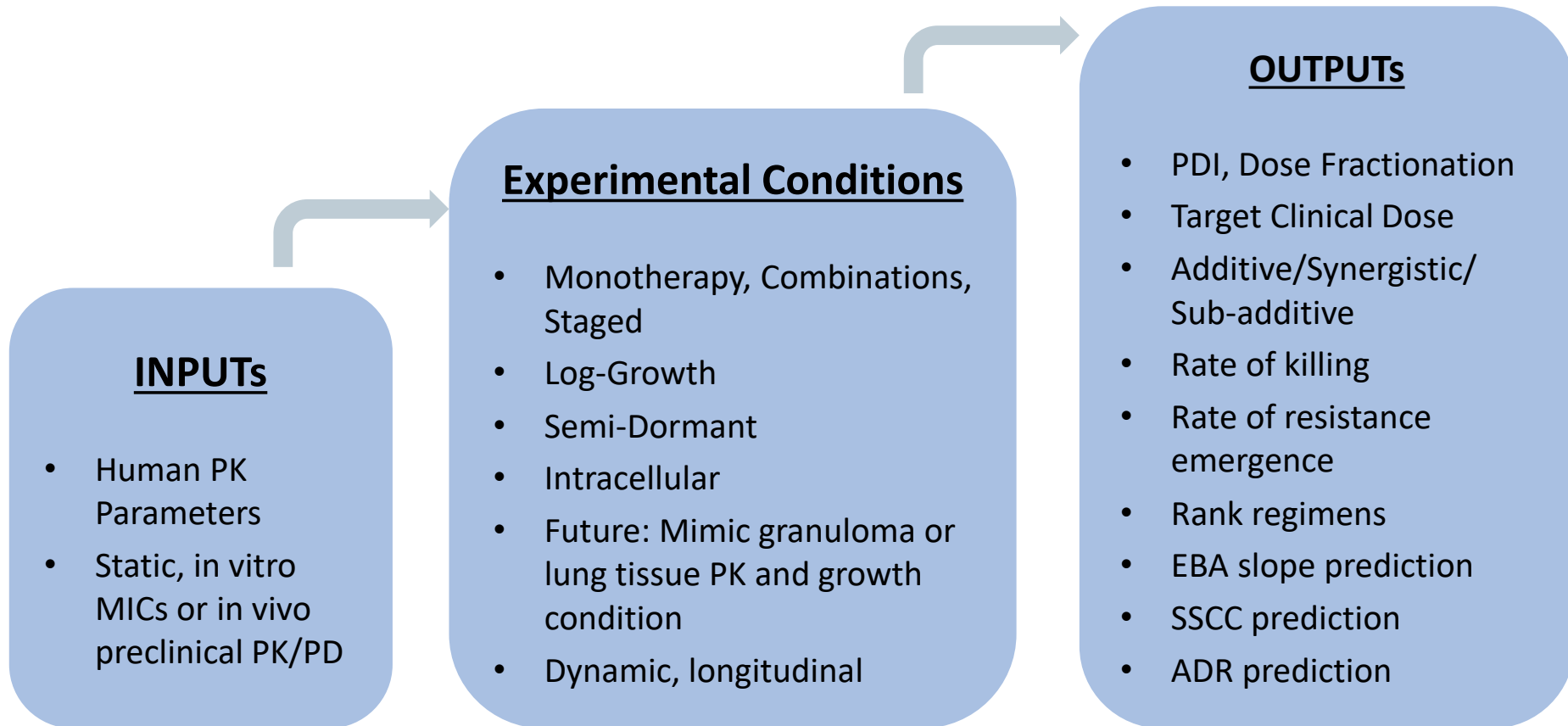


- *HFS-TB qualified for use in drug development programs as additional and complementary tool*
- *HFS-TB can be used in regulatory submissions, esp. for informed design and interpretation of clinical studies*
- *HFS-TB is recommended to be useful as follows:*
  - ✓ To provide preliminary proof of concept for developing a specific drug or combination to treat tuberculosis
  - ✓ To select the pharmacodynamic target (e.g.  $T_{>MIC}$ , AUC/MIC)
  - ✓ To provide data to support PK/PD analyses leading to initial dose selection for non-clinical and clinical studies
  - ✓ To assist in confirming dose regimens for later clinical trials taking into account human PK data and exposure-response relationships



# HFS-TB Model System





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## Objectives

- Characterize signal-to-noise in HFS-TB system under all growth conditions
- Jan 2014-Feb2016

## Design

162 Experiments

- 6 Treatment arms
  - Positive control = HRZE
  - REMox1 = MRZE
  - REMox2= HRZM
  - High Dose MRZ
  - H 3 days = MRZ
  - Untreated Control
- 3 Growth Conditions
  - Log
  - Semi-dormant
  - Intracellular
- 3 Teams
  - Leader + 4 laboratorians
- All Expts in Triplicate

## Experimental Questions

- Variability in PK measurement
  - Intra and inter experiment
  - Ability to achieve targeted AUC and Cmax
- Variability in kill rates
  - Across growth phases

# Results: HFS-TB REMox Reproducibility

## System Reliably Achieves Targets for Cmax and AUC

Drug	PK parameter	Target	Observed Mean $\pm$ SD	%CV	MAPE (%)	% Accuracy (95% CI)	% Bias (95% CI)
INH	Peak (mcg/mL)	6.80	6.65 $\pm$ 0.29	4.03	3.87	96.10 (95.40-96.90)	2.19 (1.32 to 3.06)
INH	AUC <sub>0-24</sub> (mcg*hr/mL)	24	25.80 $\pm$ 1.30	5.05	8.07	91.90 (90.90-93.00)	-7.58 (-8.75 to -6.38)
RIF	Peak (mcg/mL)	6.0	6.1 $\pm$ 0.11	1.75	1.50	98.00 (98.00-99.00)	-1.17 (-1.58 to -0.80)
RIF	AUC <sub>0-24</sub> (mcg*hr/mL)	22	25.00 $\pm$ 1.30	5.45	12.00	88.00 (87.00-89.00)	-11.81 (-13.19 to -10.45)
Hi RIF	Peak (mcg/mL)	18.0	18.00 $\pm$ 0.30	1.65	1.8	98.0 (98.00-99.00)	1.67 (-2.17 to -1.22)
Hi RIF	AUC <sub>0-24</sub> (mcg*hr/mL)	66.0	70.00 $\pm$ 2.90	4.11	6.50	93.00 (92.00-95.00)	-6.36 (-7.27 to -5.00)
PZA	Peak (mcg/mL)	54	50.0 $\pm$ 0.11	3.48	7.35	92.60 (91.90-93.40)	7.94 (7.18 to 8.72)
PZA	AUC <sub>0-24</sub> (mcg*hr/mL)	390	430 $\pm$ 29.5	6.85	10.80	89.20 (87.70-90.70)	-9.53 (-11.95 to -8.62)
Hi PZA	Peak (mcg/mL)	108	99.50 $\pm$ 2.88	2.89	7.85	92.10 (91.40-92.90)	7.85 (7.13 to 8.58)
Hi PZA	AUC <sub>0-24</sub> (mcg*hr/mL)	780	815.0 $\pm$ 56.90	6.98	6.08	93.90 (92.30-95.50)	-4.43 (-6.42 to -2.44)

# Results: HFS-TB REMox Reproducibility

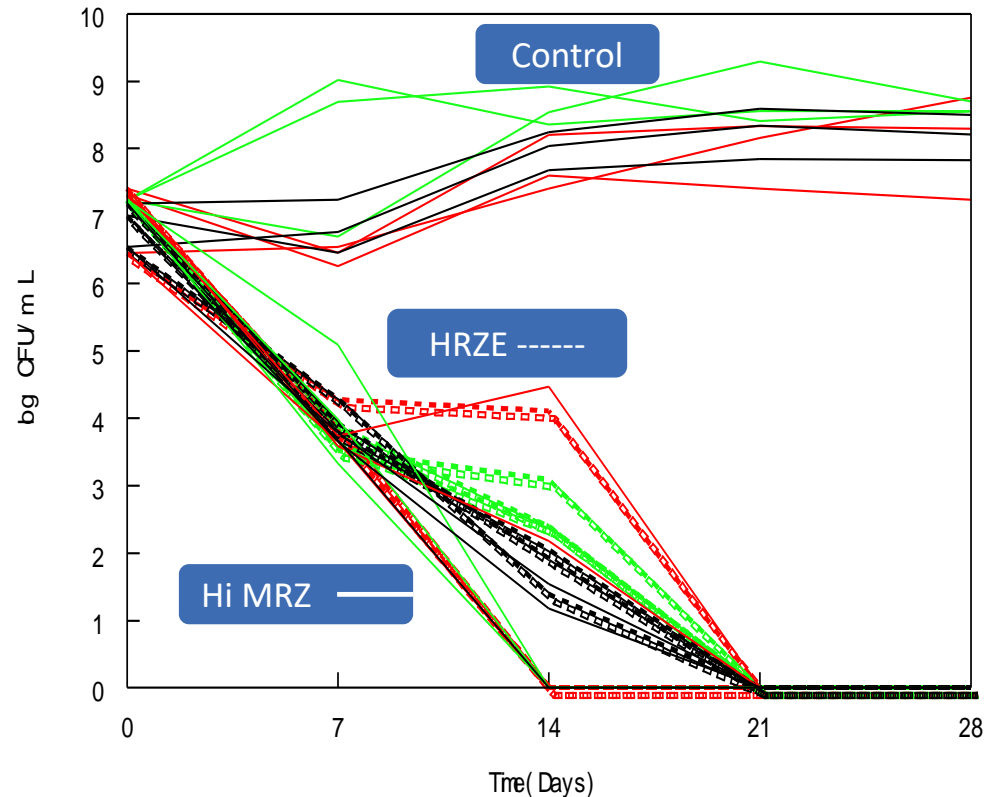
## System Reliably Achieves Targets for Cmax and AUC

Drug	PK parameter	Target	Observed Mean $\pm$ SD	%CV	MAPE (%)	% Accuracy (95% CI)	% Bias (95% CI)
E	Peak (mcg/mL)	6.30	6.29 $\pm$ 0.07	1.18	0.88	99.10 (98.90-99.30)	0.11 (-0.22 to 0.43)
E	AUC <sub>0-24</sub> (mcg*hr/mL)	23	22.50 $\pm$ 0.55	2.44	2.72	97.30 (96.80-97.80)	2.34 (1.68 to 2.99)
M	Peak (mcg/mL)	4.2	4.10 $\pm$ 0.09	2.16	2.66	97.30 (96.90-97.80)	2.30 (1.73 to 2.88)
M	AUC <sub>0-24</sub> (mcg*hr/mL)	45	42.30 $\pm$ 2.80	6.60	7.50	92.50 (91.40-93.60)	5.93 (4.24 to 7.62)
Hi M	Peak (mcg/mL)	8.4	8.13 $\pm$ 0.20	2.50	3.39	96.60 (96.0-97.20)	3.20 (2.54 to 3.86)
Hi M	AUC <sub>0-24</sub> (mcg*hr/mL)	90	84.50 $\pm$ 4.88	5.70	6.45	93.50 (92.20-94.90)	6.14 (4.67 to 7.62)



# Log Phase Time Course

(Control, HRZE, High Dose MRZ)

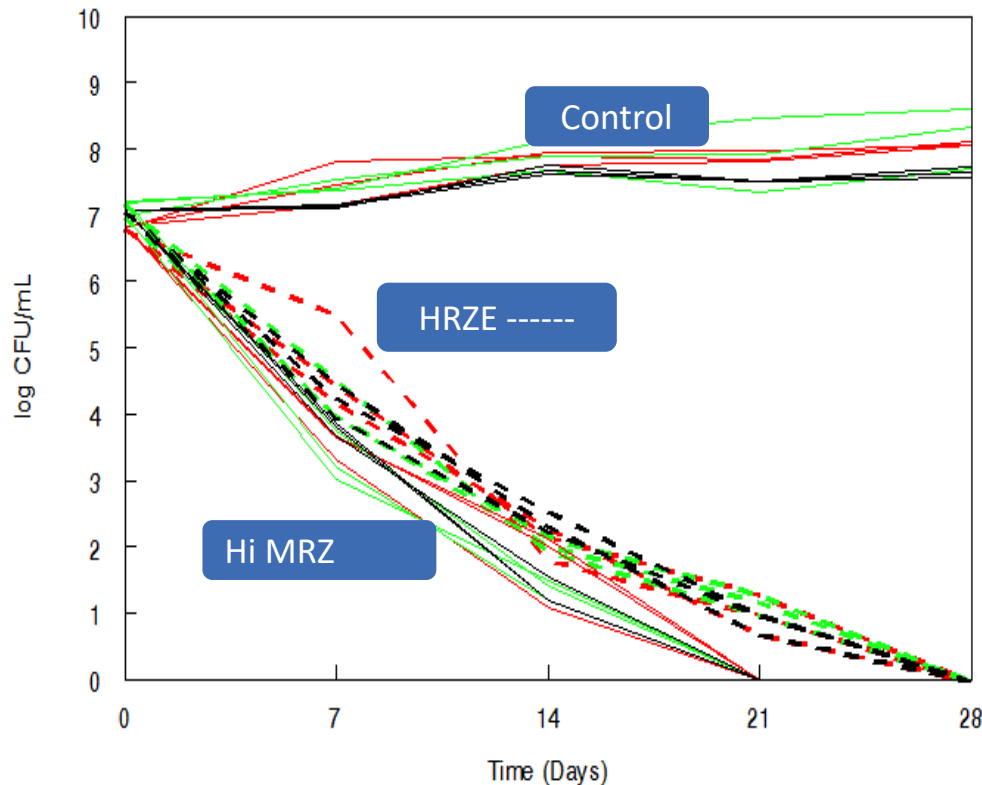


- All teams found equivalent results for the different groups
- The HRZE and High MRZ groups showed a marked difference against the untreated control
- However, the HRZE and High MRZ groups achieved similar treatment effects

\*Updated 20JAN2017

# Semi-Dormant Time Course

(Control, HRZE, High Dose MRZ)

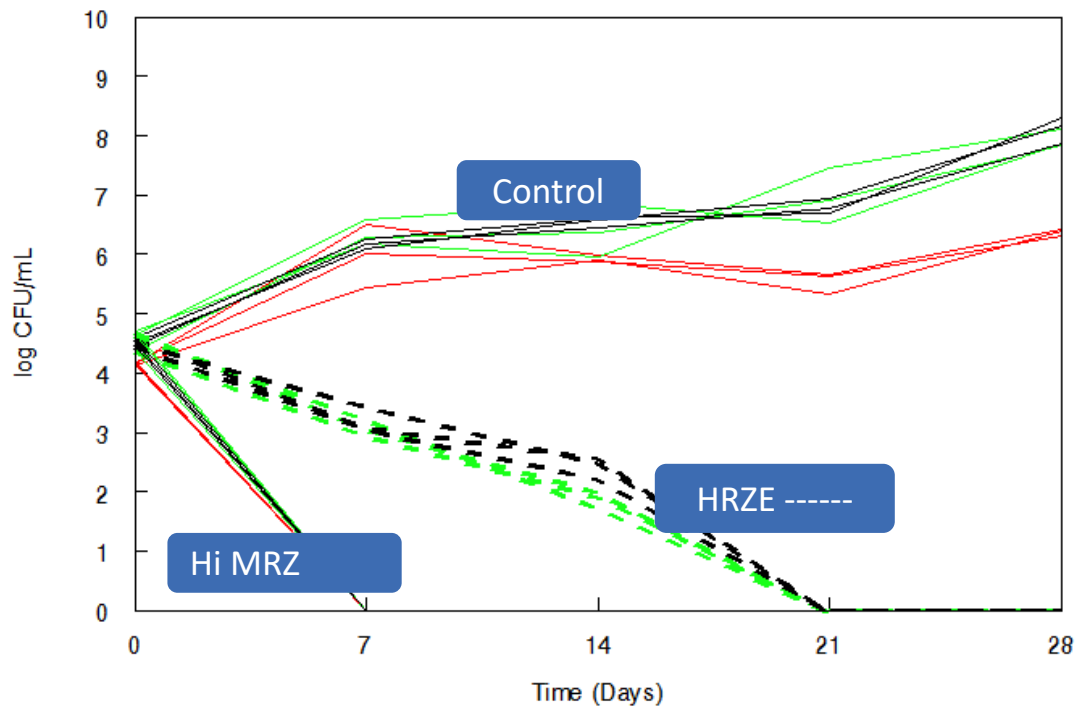


- As in the Log-Phase condition, the experimental teams found equivalent results for the different groups
- The HRZE and High MRZ groups showed a marked difference against the untreated control
- However, the HRZE and High MRZ groups achieved similar treatment effects
- Given the reduced bacterial activity in semi-dormant condition, the overall variability in results was reduced across groups

\*Updated 20JAN2017

# Intracellular Time Course

(Control, HRZE, High Dose MRZ)



- The results were consistent across teams
- The HRZE and High MRZ groups showed a marked difference against the untreated control
- However, unlike in the other two metabolic conditions, the intracellular experiments showed a marked difference between all three groups, favoring the High MRZ regimen

\*Updated 20JAN2017

Data Issue: HRZE

## When to Apply HFS-TB

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- ✓ To provide preliminary proof of concept for developing a specific drug or combination to treat tuberculosis
  - ✓ To select the pharmacodynamic target (e.g. T/MIC, AUC/MIC)
  - ✓ To provide data to support PK/PD analyses leading to initial dose selection for non-clinical and clinical studies, with the aim of limiting the number of regimens that are to be tested in vivo
  - ✓ To assist in confirming dose regimens for clinical trials taking into account the accumulated human PK data in healthy volunteers, patients & available information on exposure-response relationships
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## New HFS-TB Work

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- Prospective combination studies (1000 + HFS TB Units)
  - Ex: PaMZ
- New, emerging drugs of interest aligning with Pharma and TB Accelerator partners
- Expand capacity with partner lab

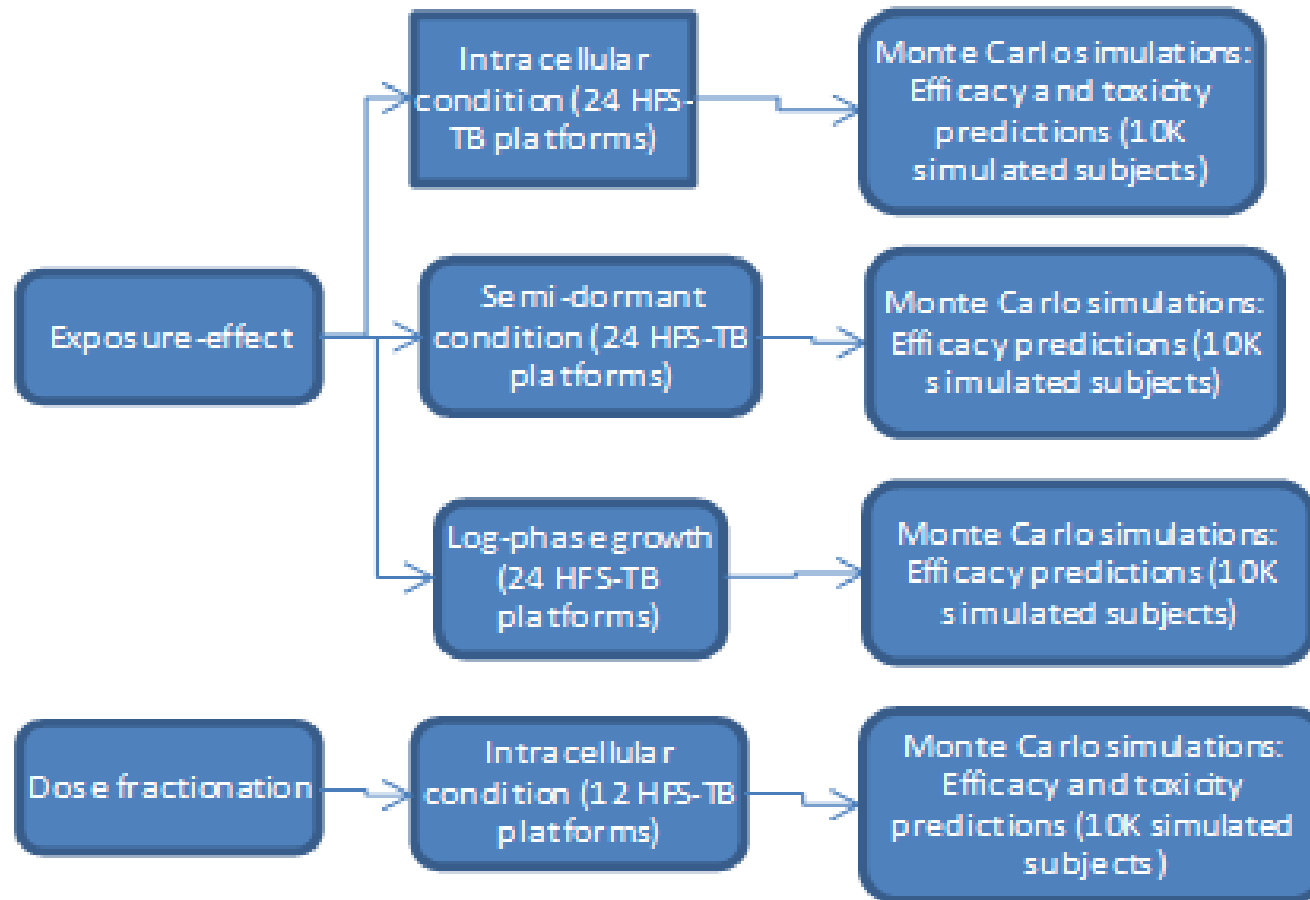
# Thank you

***A special thank you to the Baylor laboratory  
team and the CPTR Pre-Clinical and Clinical  
Sciences Workgroup***

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# HFS-TB New Work Experimental Schema

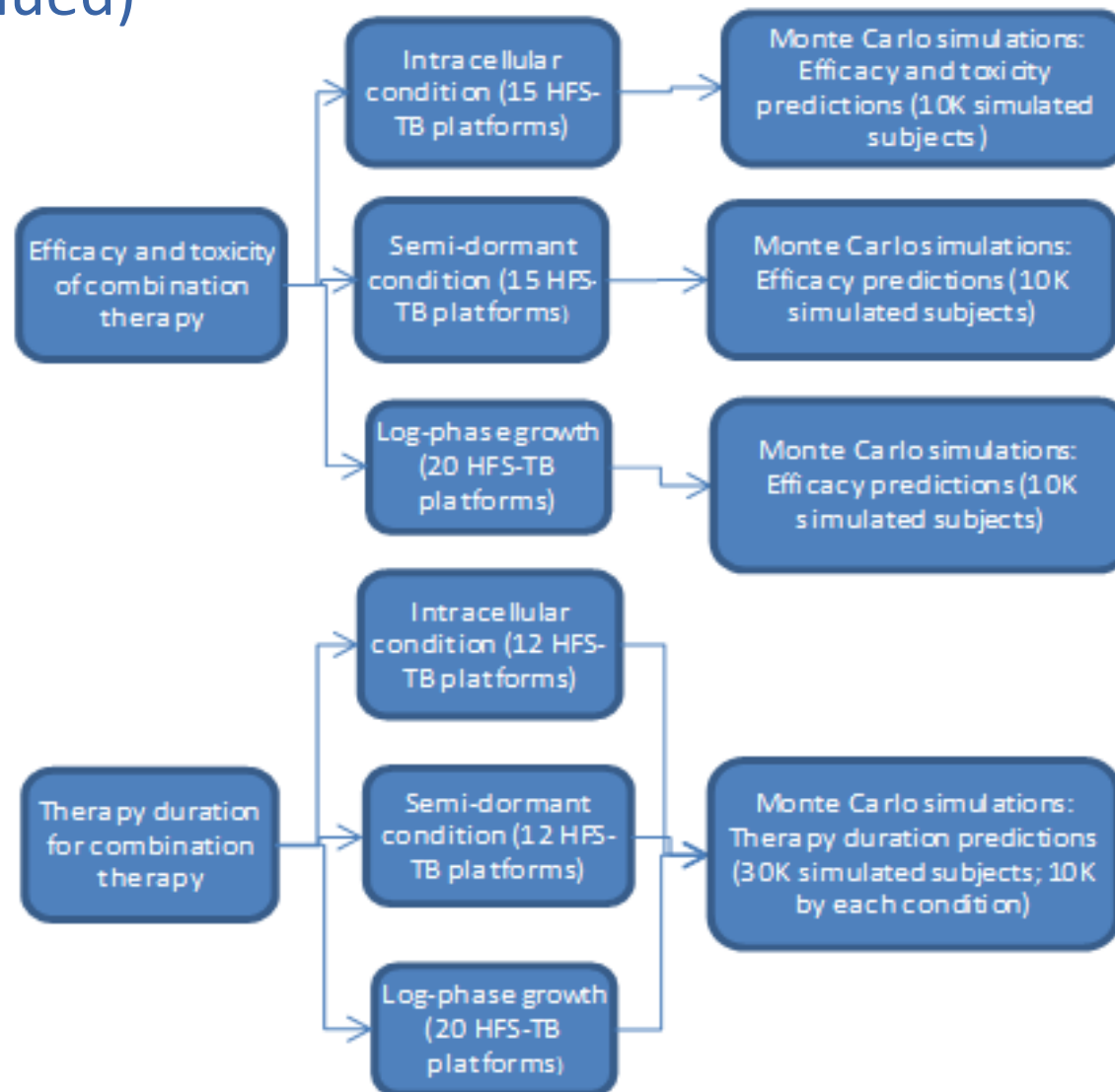


# HFS-TB Stage New Work Experimental Schema

## (continued)



Critical Path to  
TB Drug Regimens

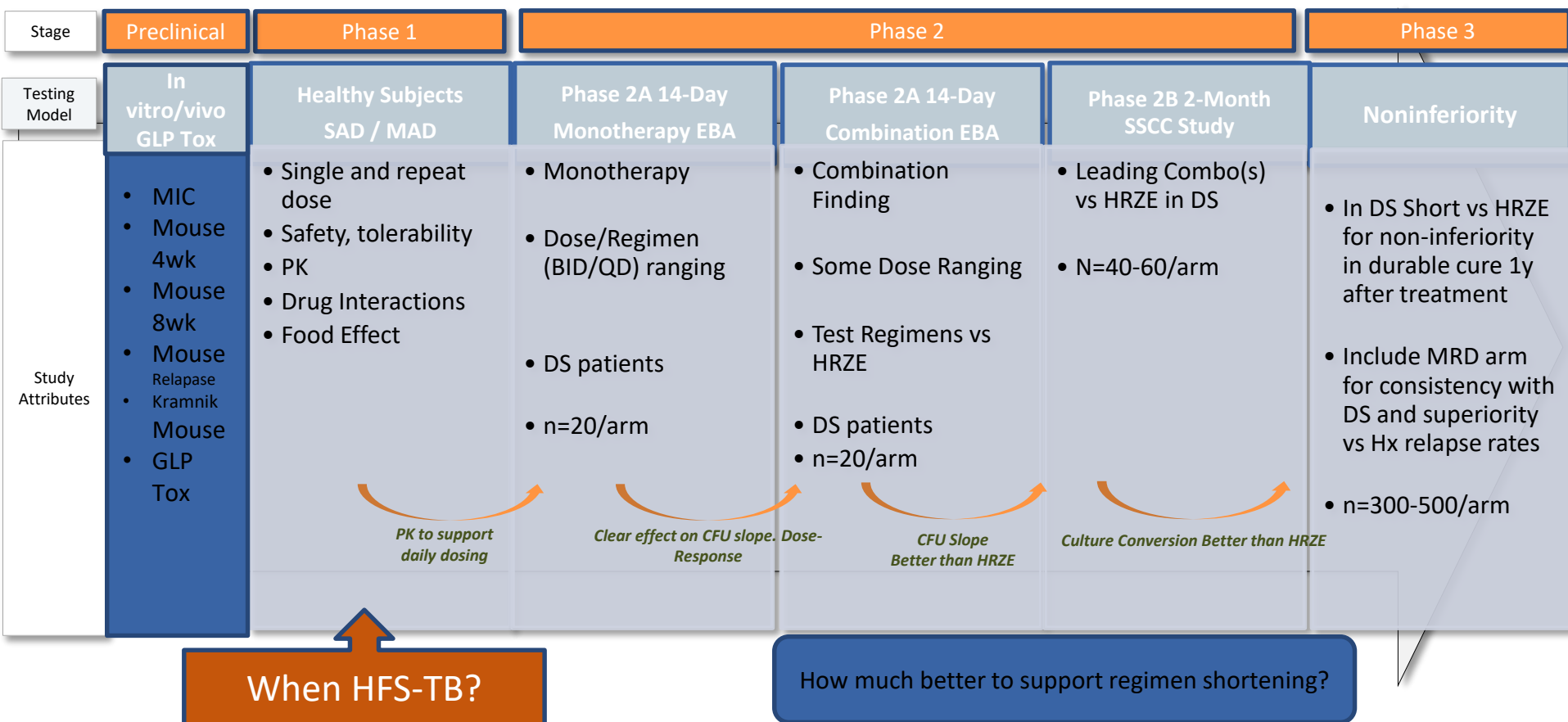


- Initial step to address the “translational gap” is to learn what data from what models analyzed in what way informs key trial design decisions
- Evidence-based validation of preclinical models is important:
  - To confidently place preclinical models on the critical development path
  - To increase the efficiency of regulatory interactions
  - To set a precedent for objective, data-driven process to apply to other models and tools (e.g., C3HeB/FeJ mouse, marmoset)
  - To identify/clarify knowledge and tool gaps to drive future research
- The successful HFS-TB qualification process has accomplished each of these goals
- Evaluation of sterilizing mouse model is the appropriate next step, with other models to follow



- **Analysis Objective** to determine predictive accuracy of HFS-TB outputs for clinical trial results
- **Literature Search** to identify relevant HFS-TB and clinical data from published literature
- **Systematic Review** to summarize HFS-TB-generated hypotheses and outcomes of clinical trials
- **Quality of Evidence Scoring** to provide basis for weighting in the predictive accuracy analysis
- **Statistical Analysis** comparing HFS-TB predictions with clinical findings to examine:
  - descriptive correlations where HFS-TB studies post-dated clinical studies
  - predictive accuracy where HFS-TB studies pre-dated clinical studies

# Unified Development Pathway

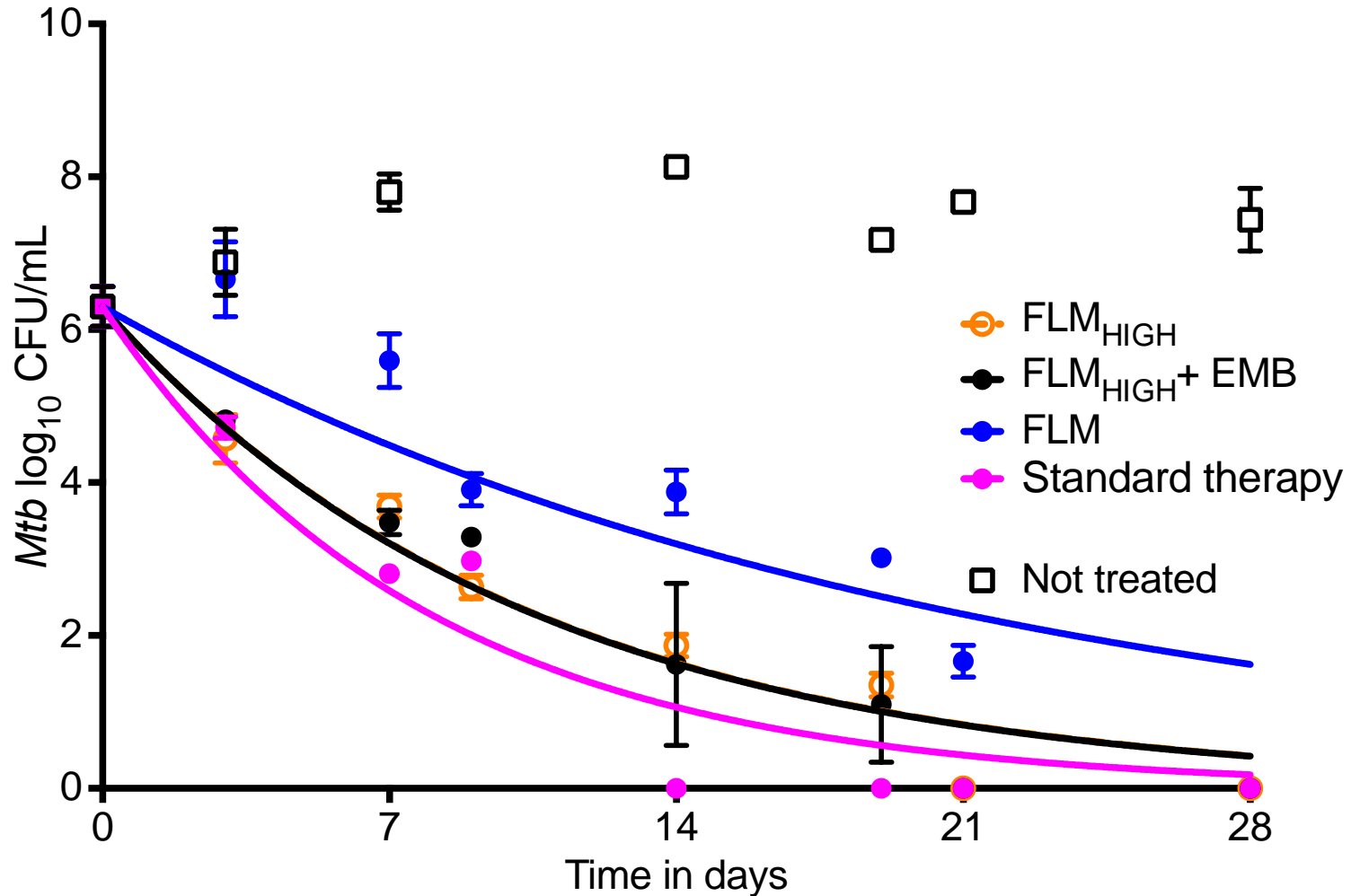


# HFS-TB Stage II – Project Plan

	Intracellular	SDB	Log-phase	# of experiments	# HFS	Time in weeks	Start Date	End Date
PaMZ	9	9	15	3	33	8	1/2/2017	2/24/2017
Del PK	8	5	5	3	18	1	2/27/2017	3/6/2017
Del	36	24	24	3	84	8	2/27/2017	4/21/2017
OPC-167832	36	24	24	3	84	8	4/24/2017	6/16/2017
Break						2	6/19/2017	6/30/2017
OPC-167832 + Del	36	24	24	3	84	8	7/3/2017	8/25/2017
OPC-167832 + Del + Oxa	36	24	24	3	84	8	8/28/2017	10/20/2017
Del & Ba	36	24	24	3	84	8	10/23/2017	12/8/2017
OPC-167832 + Del + Ba	36	24	24	3	84	8	12/11/2017	2/2/2018
Break						2	2/5/2018	2/16/2018
Sutezolid	36*	24	24	3	84	8	2/19/2018	4/13/2018
BaPZ	9	9	15	3	33	8	4/16/2018	6/8/2018
Oxa + Far + Del	18	18	30	3	66	8	6/11/2018	8/3/2018
Break						2	8/6/2018	8/17/2018
Oxa + Pa + Ba	18	18	30	3	66	8	8/20/2018	10/12/2018
AZD5847	36	24	24	3	84	8	10/15/2018	12/7/2018
Oxa + Far + Pa	18	18	30	3	66	8	12/10/2018	2/1/2019
Break						2	2/4/2019	2/15/2019
Oxa + Pa + Ba + Far	15	15	20	3	50	8	2/18/2019	4/12/2019
Oxa +M + Far + Z	15	15	20	3	50	8	4/15/2019	6/7/2019
Final Report Preparation						8	6/10/2019	8/1/2019
<b>TOTAL</b>	<b>362</b>	<b>299</b>	<b>357</b>	<b>48</b>	<b>1054</b>	<b>137</b>		



# New Regimen Design: “FLAME”



Deshpande et al. A faropenem, linezolid, and moxifloxacin regimen for both drug susceptible and multidrug-resistant tuberculosis in children. Clin Infect Dis. 2016;63:S95



- ❑ Objectives: Characterize reproducibility and signal to noise in HFS-TB system under different growth conditions (Jan 2014-Feb 2016)
- ❑ Design:
  - ❑ 6 treatment arms
    - Positive Control – HRZE
    - REMox 1 – MRZE
    - REMox 2 – HRZM
    - Hi Dose MRZ
    - H 3 Days+Hi Dose MRZ
    - Control
  - ❑ Three conditions: Log-Phase, Semi-dormant, Intracellular
  - ❑ Three separate teams (Each team included a Team Leader and 4 supporting lab techs)
  - ❑ Each team runs each experiment in triplicate
  - ❑ Total of 162 HFS-TB experiments (6 regimens x 3 conditions x 3 teams x 3 replicates)

# Results: HFS-TB REMox Reproducibility

- ❑ Typical inter-day assay variability (CV%) is 5-10% for a PK assays
- ❑ In HFS experiments CV% in drug concentration across time were typically  $\leq 5\%$  (Passed Go / No Go)
- ❑ At end of dosing intervals for INH and RIF CV% up to 25% were observed, however SDs remained consistent with other time points (e.g., this is a function of low mean conc where CV% is SD/mean)
- ❑ Variability in PK concentrations attributed to TEAM was very low across drugs ( $<0.1\%$  of total variance).
- ❑ Low variability expected due to administration via programmed syringe pump

