

Critical Path to TB Drug Regimens Role in Facilitating TB Drug Development

Debra Hanna, Executive Director, Critical Path to TB Drug Regimens 19 July 2017



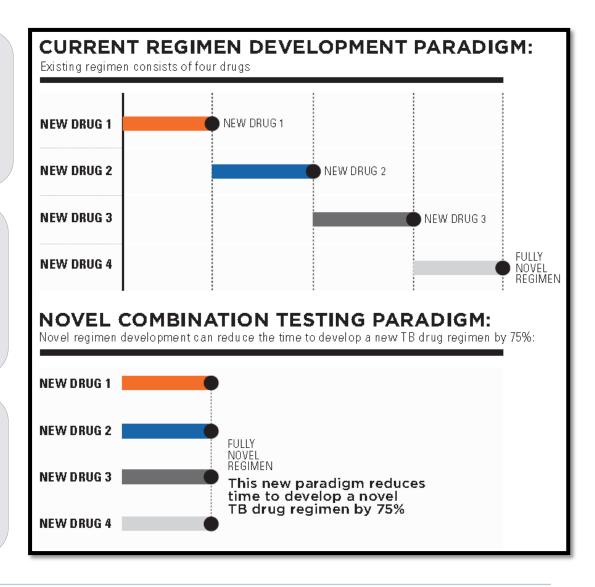




Novel regimen development requires emphasis on combination study approaches

Define, based on evidence, best drug development tools to de-risk compounds and improve understanding of efficacy

Define, based on evidence, novel biomarkers to inform improved trial design and adaptivity

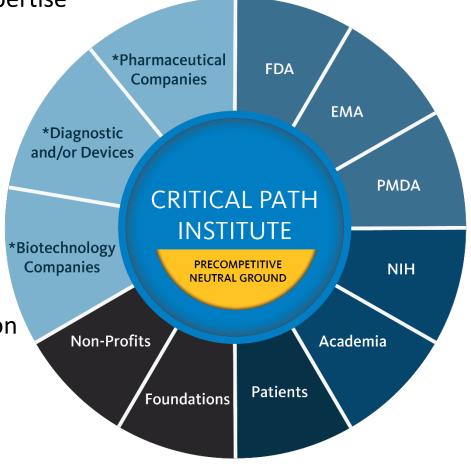




- Mission: The Critical Path to TB Drug Regimens (CPTR) is a crosssector initiative that aims to speed the development of safer and shorter duration anti-tuberculosis (TB) drugs. Focus on:
 - drug development tools and methodologies to support go/no-go decisions during each stage of research and development
 - curation of supportive data through establishment of collaboration network to support new methods and tool validation (and ensure public access wherever possible)
 - developing pathways for new TB treatment regimens that include drugs that are not yet individually approved
 - providing regulatory excellence in the development, validation, and advancement of these drug development tools and methodologies
- CPTR Partners and Members: Consortium of 8 pharma / 18 diagnostic companies, 26 academic institutions, 20 NGOs, and 5 governmental bodies.

PUBLIC PRIVATE PARTNERSHIP MODEL

- Act as a trusted, neutral third party
- Convene scientific consortia of industry, academia, and government for pre-competitive sharing of data/expertise
 - ✓ The best science
 - The broadest experience
 - Active consensus building
 - Shared risk and costs
- Enable iterative EMA/FDA/PMDA participation in developing new methods to assess the safety and efficacy of medical products
- Pursue official regulatory recognition through "qualification" of Novel Methodologies and Drug Development Tools



TICAL PATH

CPTR LEVERAGES REGULATORY QUALIFICATION STRATEGY



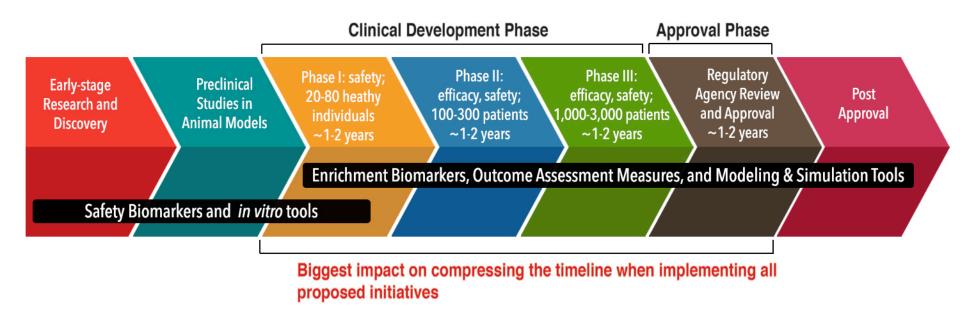


<u>Start at the end approach</u>: Up-front conversations around the context of use (COU) since the COU drives the level of evidence needed

SHARED LEARNING CAN SHORTEN THE TIMELINE



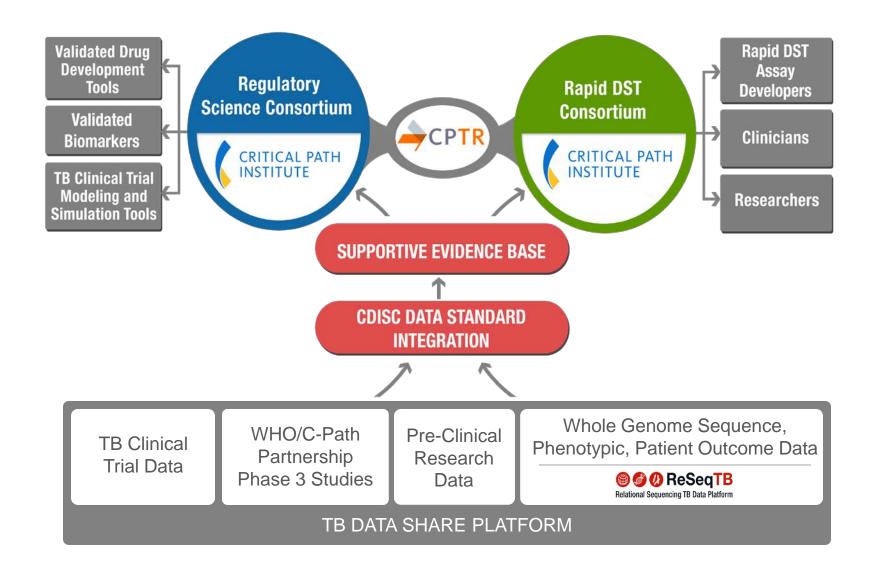
- Data Standardization and Sharing
- Biomarker Development and Qualification
- Clinical Outcome Assessment Measures
- Modeling and Simulation Tools



Adapted from *"A virtual space odyssey "*, Cath O'Driscoll (2004) <u>http://www.nature.com/horizon/chemicalspace/background/odyssey.html</u>

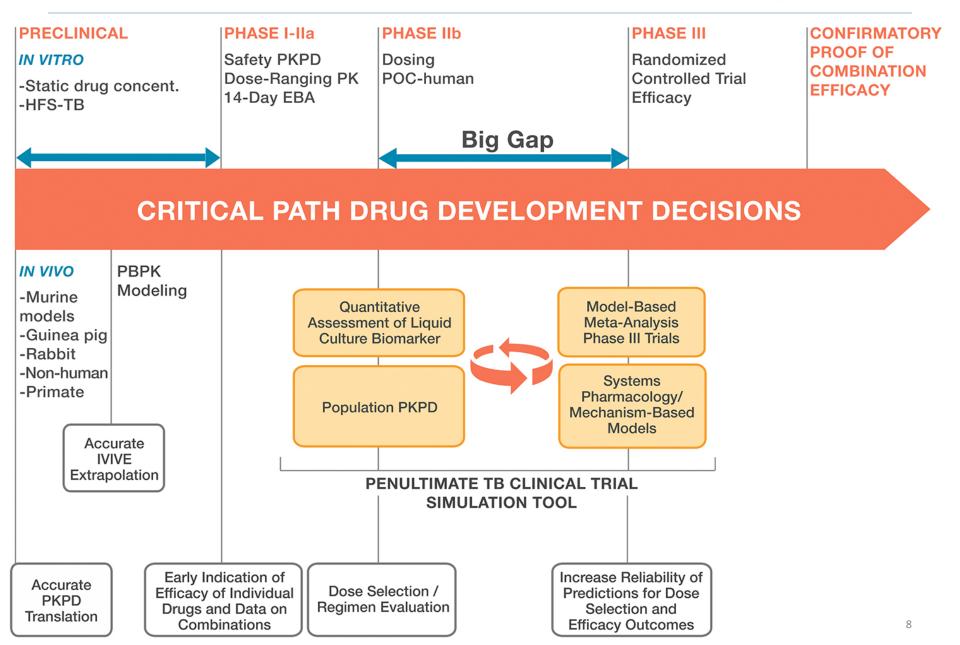
DATA COLLABORATION IS CRITICAL





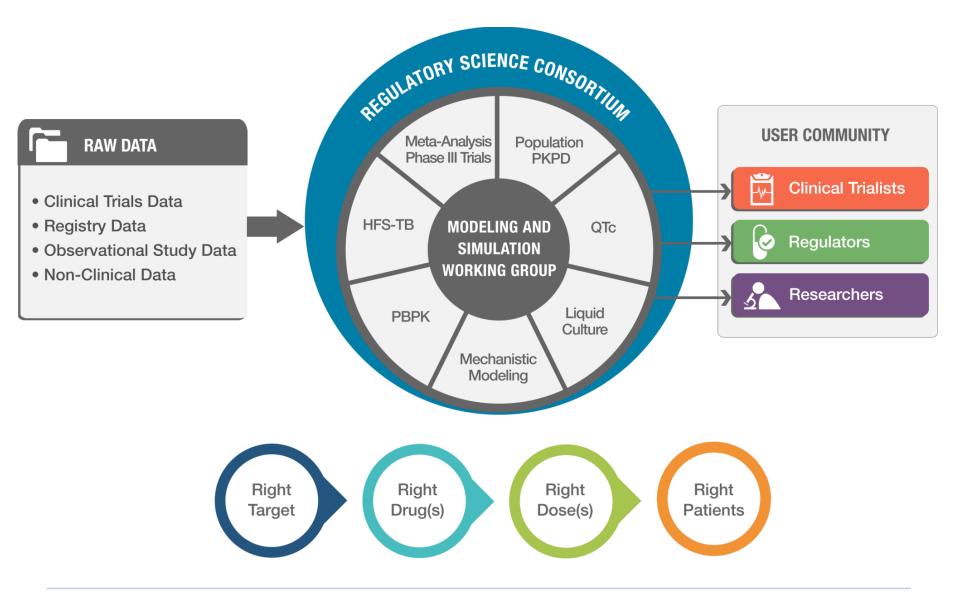
GAPS IN THE TB DRUG DEVELOPMENT PROCESS

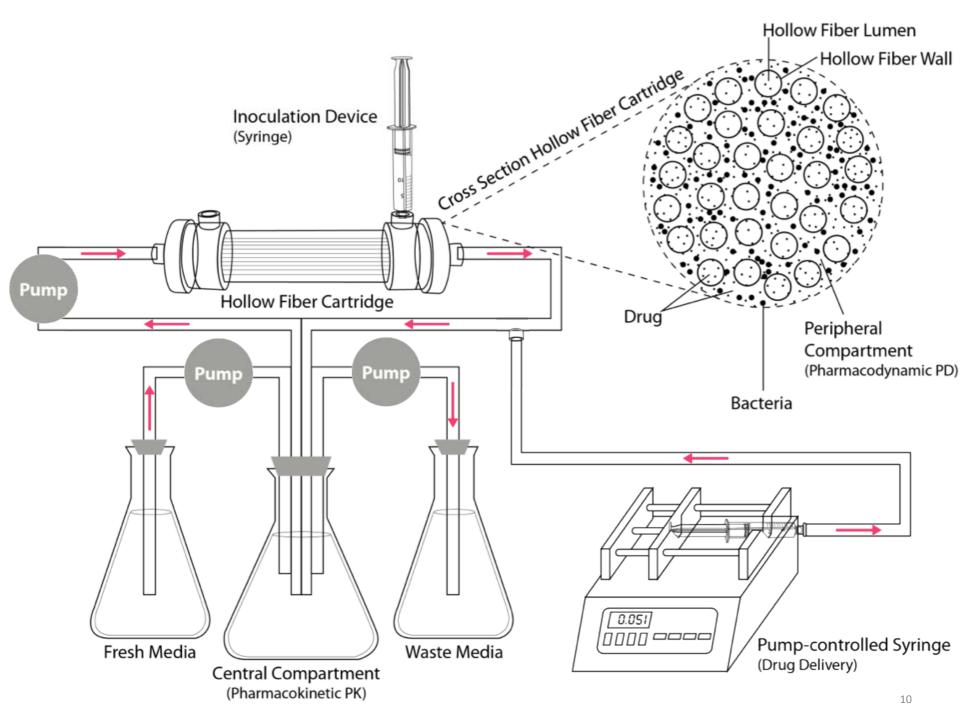




CPTR MODELING AND SIMULATION PROGRAMS

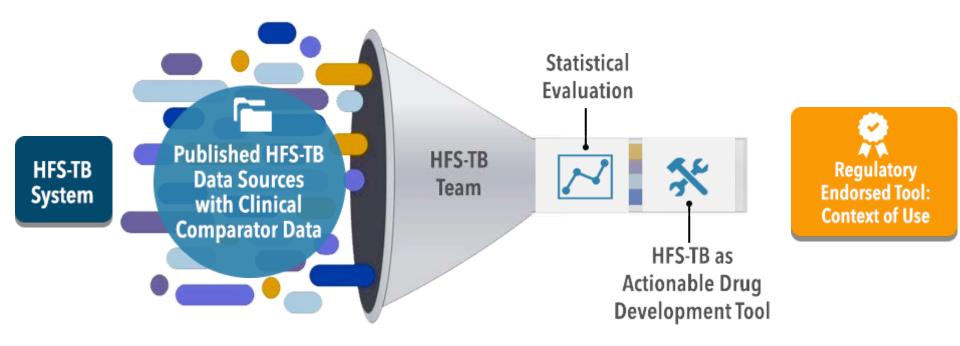






EVIDENCE BASED EVALUATION OF HOLLOW FIBER SYSTEM MODEL FOR TB





REGULATORY INTERACTIONS ON HFS-TB QUALIFICATION



	FEBRUARY 20, 2013	FEBRUARY 27, 2013	OCTOBER 16, 2013	NOVEMBER 15, 2013	FEBRUARY 4, 2014	
FDA	LOI submission	LOI discussion	VXDS document submission	VXDS meeting	Submission of comments to FDA draft guidance	
EUROPEAN MEDICINES AGENCY						
EUROPEAN MEDICINES AGENCY	Briefing document submission (for qualification opinion)	SAWP meeting	Draft qualification opinion	Public comment period	Final qualification opinion	

- 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:

 \checkmark To provide preliminary proof of concept for developing a specific drug or combination to treat TB

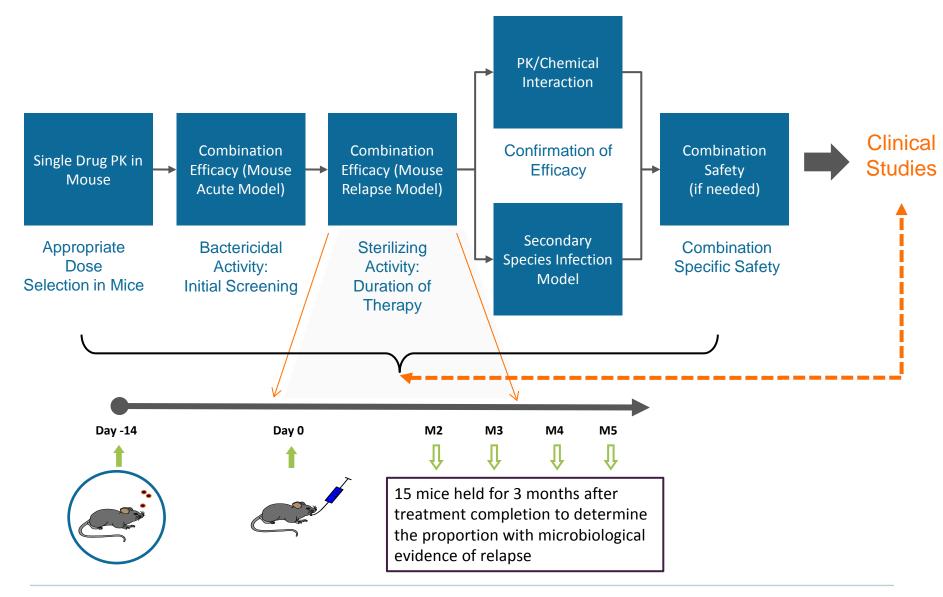
 \checkmark To select the pharmacodynamic target (e.g. T_{>MIC}, AUC/MIC)

✓ 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

MOUSE MODEL OF STERILIZING ACTIVITY





EVIDENCE-BASED EVALUATION OF STERILIZING MOUSE MODEL

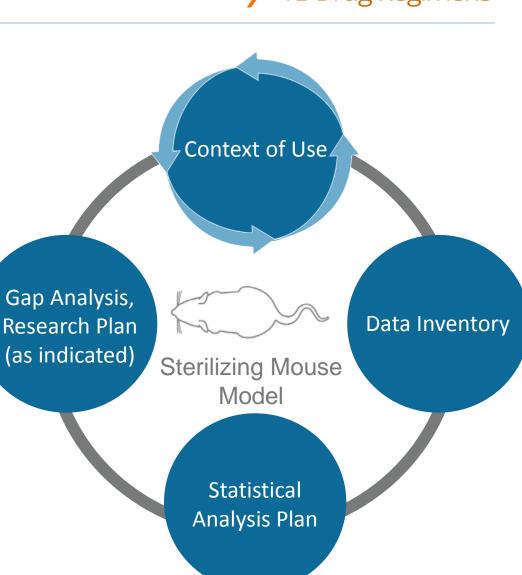
General Aim:

Quantify the predictive accuracy of mouse TB efficacy models to estimate the treatment-shortening potential of a test regimen, by evaluating differences in the treatment duration necessary to prevent relapse compared to control (standard TB regimen).

Intended Application:

The data from experiments in mice infected with *M. tuberculosis*, using relapse as the main endpoint will be used to:

- Calculate treatment effect sizes, to then rank-order regimens
- Estimate clinical treatment duration







High unmet need for real-time assessment of efficacy in TB drug development trials

- Field requires a tool that:
 - Assesses Early Bactericidal Activity (EBA) and Sputum Culture Conversion (SCC), endpoints recommended by FDA and EMA, in real-time, allowing for quick decision making
 - Reduces cost associated with delayed results in development of drugs for TB, a therapeutic area with limited treatment options and few commercial incentives
 - Can be easily utilized in any laboratories that are suitable for clinical trials
 - Is not affected by contamination or drug carry-over effect
- EBA and SCC are useful but challenging to conduct
 - Time delays and labor intensive
 - Issues with contamination and drug carry over effects



- LAM: Lipoarabinomannan; a major cell wall component
- A new immunoassay was developed (LAM-ELISA) that measures sputum LAM
 - Specific for LAM from MTB and a few slow growing mycobacterium strains
 - No cross-reactivity with oral bacteria
 - Strong correlation between sputum LAM and cfu counts/TTD
- Not affected by <u>contamination</u> or <u>drug carry-over</u>
- LAM-ELISA: 20 min LAM extraction; 5 hours ELISA
- Quicker tests being developed (results in <1 hour)

LAM BIOMARKER EFFORT



- An expert team convened to assess lipoarabinomannan (LAM) as a pharmacodynamic biomarker for quantitative measurement of bacterial load in sputum.
- This is one of the first pharmacodynamic biomarkers C-Path has advanced to a proposed Context of Use discussion with FDA.
- A Letter of Intent was submitted to FDA on June 9, 2017 to pursue regulatory qualification.





Use Statement

- LAM is a pharmacodynamic biomarker for quantitative measurement of bacterial load in sputum. A decrease of LAM in sputum likely reflects the reduction of bacterial load in the lung.
- This pharmacodynamic biomarker should be considered with other microbiological measurements, such as culture, as a realtime evaluation of treatment response in clinical trials of patients with pulmonary tuberculosis and positive smears and cultures, such as:
 - 14-day early bactericidal activity (EBA) trials,
 - Clinical trials of pulmonary tuberculosis up to 56 days, or
 - Clinical trials to provide evidence for early decision making in adaptive trial designs.

THE ENVISIONED IMPACT: POTENTIALLY SHORTENS DEVELOPMENT TIME BY 2-3 YEARS



Traditional	<u>EBA</u> Regimen	1		<u>Phase 2</u> (2-mo SCC)			<u>Phase 3</u> <u>Pivotal endpoint</u>		
	Regimen Regimen Regimen	3	12-18 months for regulatory approvals in many TB	Regimen Regimen Regimen	3	12-18 months for regulatory approvals in many TB	Regi	men 3	
	Regimen	5	endemic countries			endemic countries			total
	12-18 months			18-24 months			36-48 months	5	7-10 years
With qualified biomarker			hase 2/Pha al. BMC Med. (201 Real-time	6) 14:51, Bratto	n et al. B		hodol (2013) 13:		
Regimen 1> STOP									
Regimen 2						Evaluate fo	or		
Regimen 3			P			>	pivotal endpoint	5-7	
	Regimen		/ 0.0	→ STOP					years

CPTR INITIATIVE MEMBERS AND PARTNERS





Academic Partners					
Baylor Institute for Immunology Research	O'Neill Institute at Georgetown Law Center	University College of London			
Case Western Reserve University TB Research Unit	Radboud University	University of Arkansas for Medical Sciences			
Colorado State University	RESIST-TB [Boston University]	University of Cape Town			
Duke University	Rutgers [University Of Medicine & Dentistry]	University of Liverpool			
Forschungszentrum Borstel	St. George's, University of London	University of St. Andrews			
Harvard University	Stanford University	University of Virginia			
Johns Hopkins University	Stellenbosch University	University of Texas Health Science Center at San Antonio			
London School of Hygiene and Tropical Medicine	University of Florida	University of Toronto			
Munich University	University of California, San Diego	Uppsala University, Dept. of Pharmaceutical Biosciences			
NYU		Vanderbilt University School of Medicine			



Thank you