



Drug discovery for Tuberculosis

Vincent Delorme, PhD
Tuberculosis Research Laboratory

2018, October 17th



Institut Pasteur Korea

We are Pasteurians fighting disease in Korea for all mankind

Contents

Introduction

1. Tuberculosis and disease progression
2. Drug discovery in TB field

Phenotypic screening assays for Tuberculosis research

1. Chemical modulators
2. Genetic modulators (Bacteria)
3. Genetic modulators (Host)



Introduction

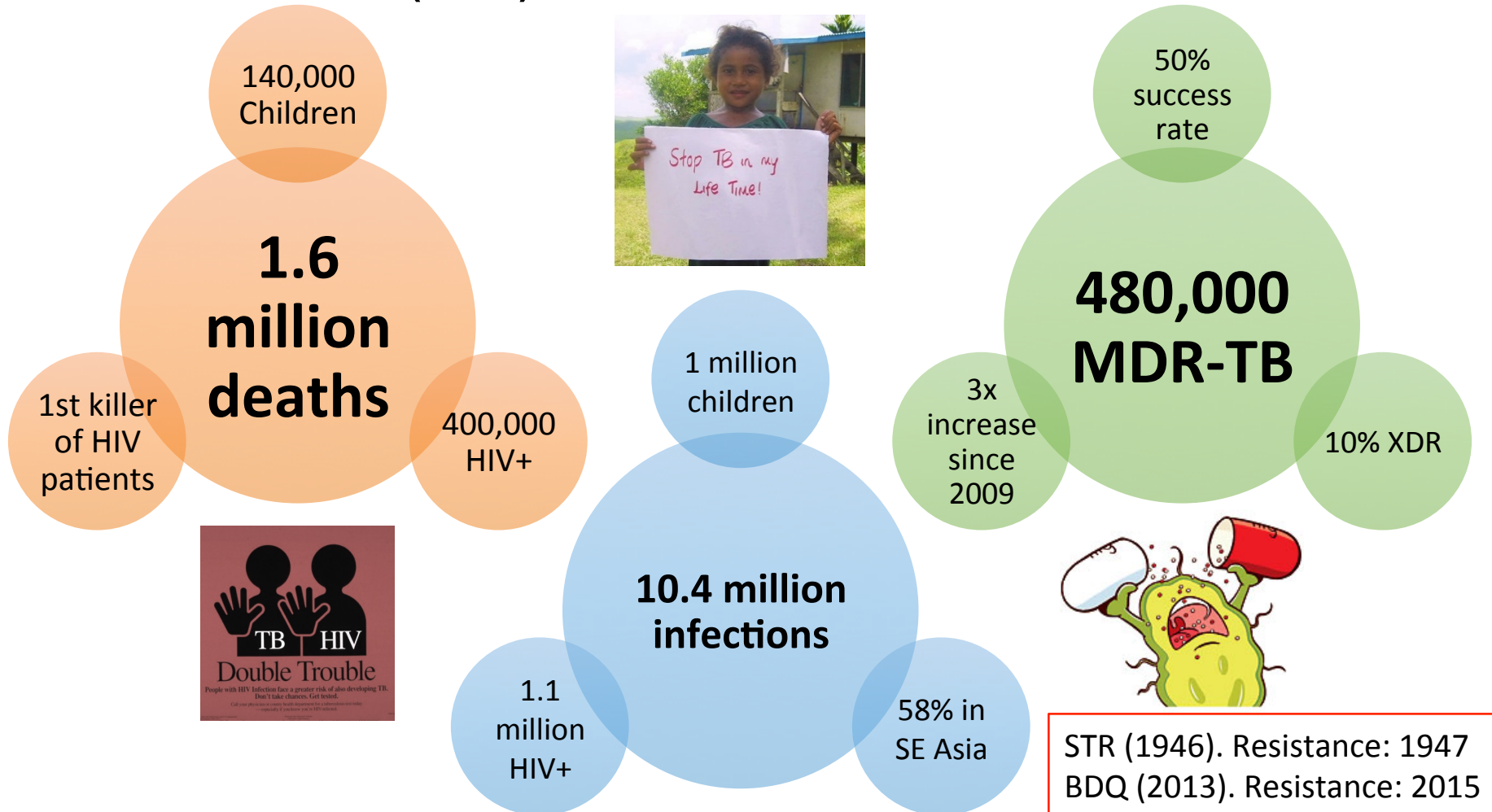


Part 1.

TUBERCULOSIS AND DISEASE PROGRESSION

Facts on Tuberculosis

Facts about TB (2017)



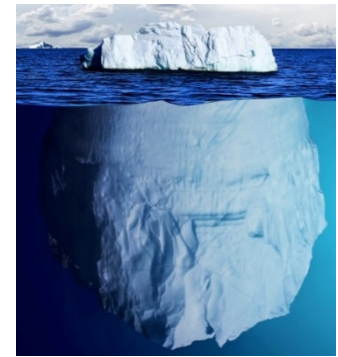
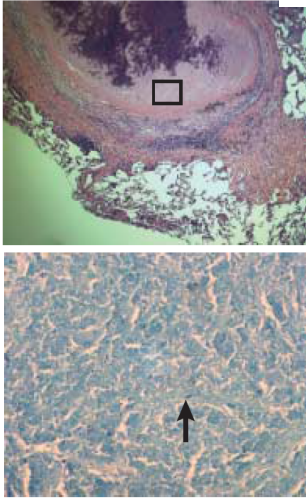
Disease progression

Primo
infection

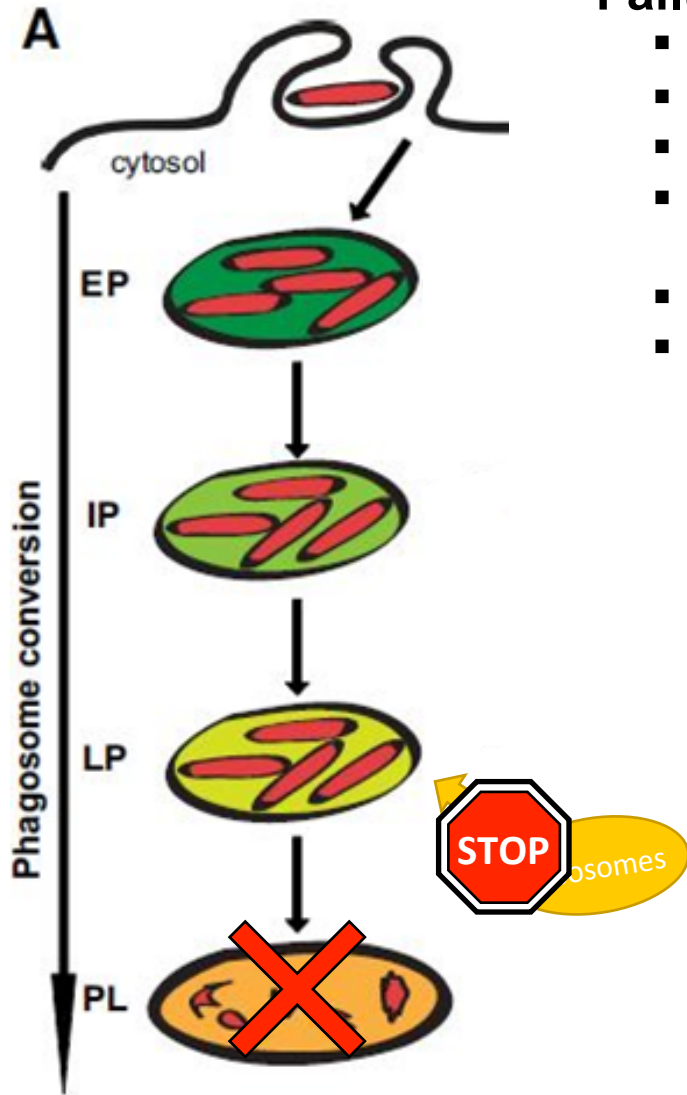
**1/3 of the
global
population!**

**1.6
million
deaths**

Boshoff *et al.* 2005 Nat. Rev. Microbio. 3: 70-80

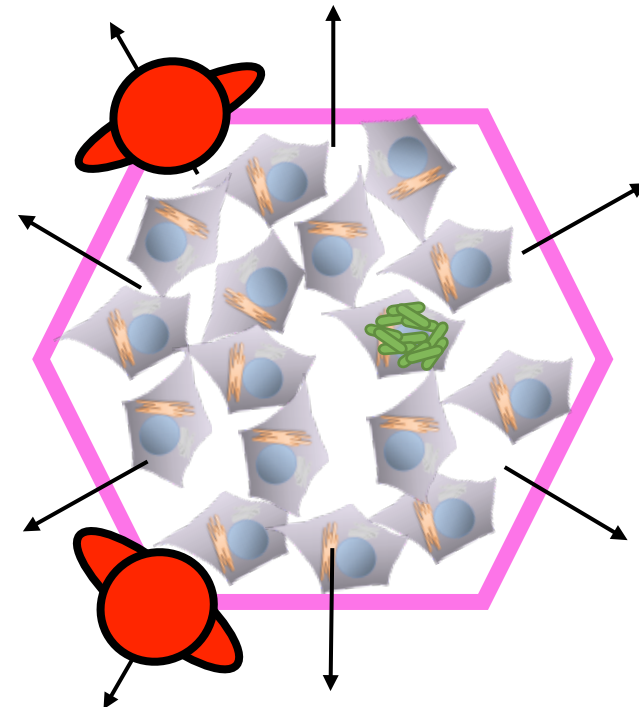


Phagosomal escape



Failure of bacterial killing by macrophages

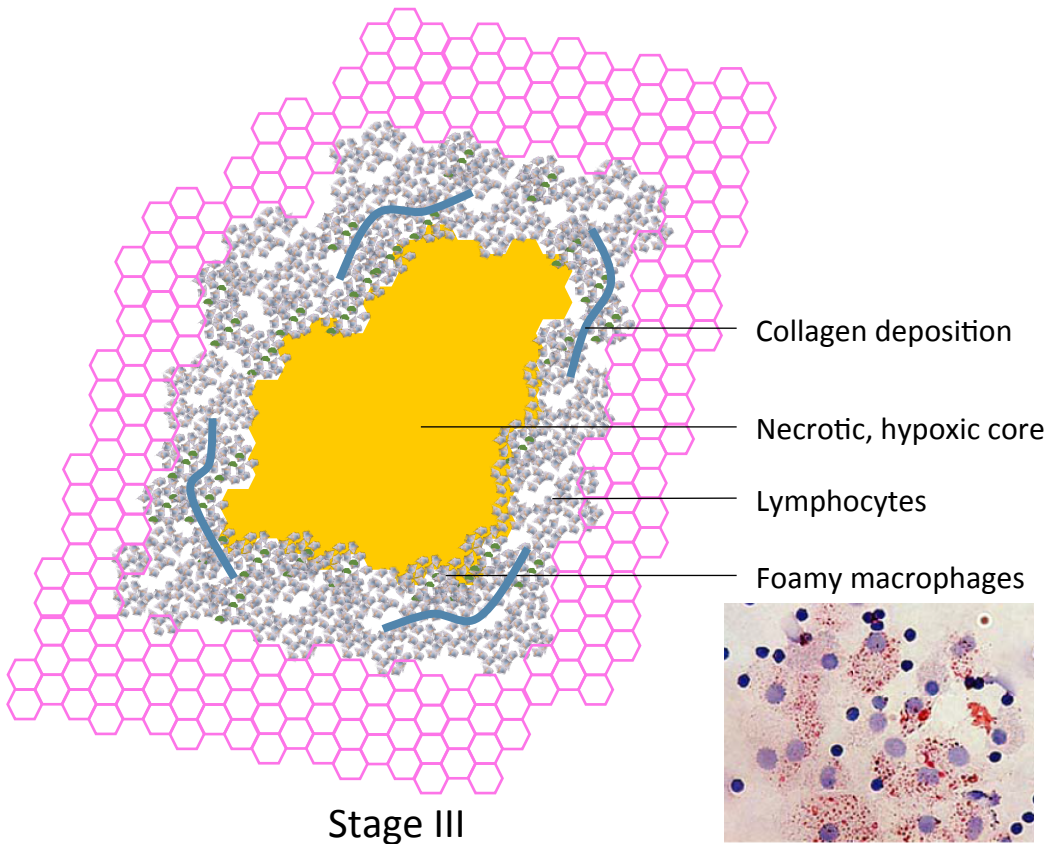
- Bacterial replication
- Cell death
- Inflammation
- Large influx of cells
 - Mostly macrophages and neutrophils
- High Local pressure on epithelium and capillary
- Collapsing of capillary and alveolus



Granuloma

Granuloma with a necrotic core

Necrotic core surrounded by foamy macrophages

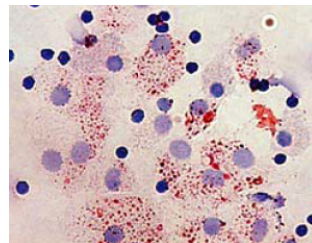


Stage III

31 days in guinea pig model

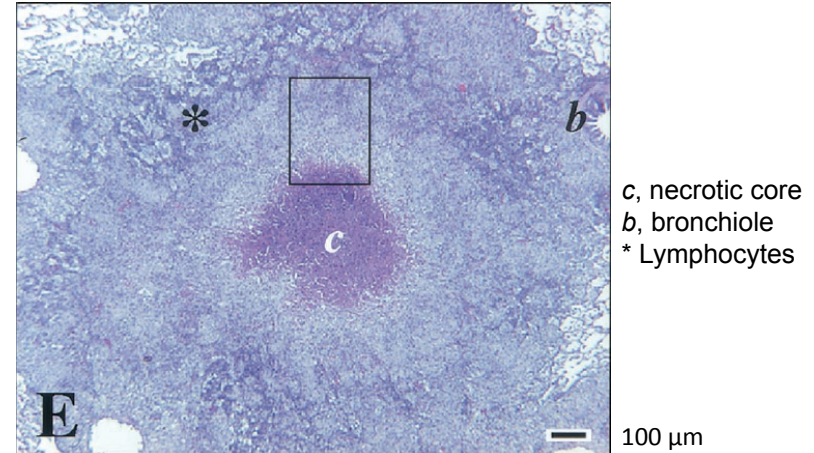
Large amount of lymphocytes in the edge

Bacterial propagation stopped



Peyron *et al.* **2008** PLoS Pathog
4(11): e1000204

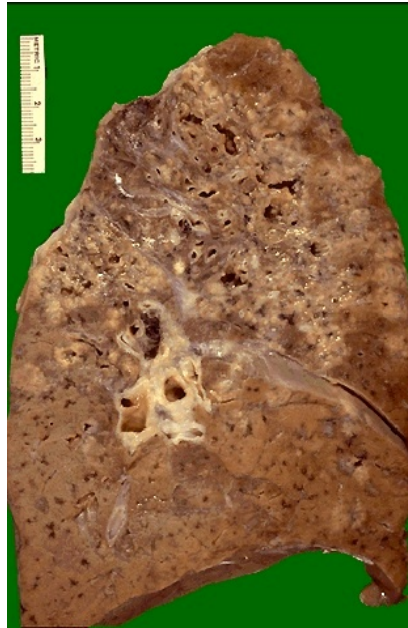
Turner *et al.* **2003** Infect Immun (71) 864-871



c, necrotic core
b, bronchiole
* Lymphocytes

100 µm

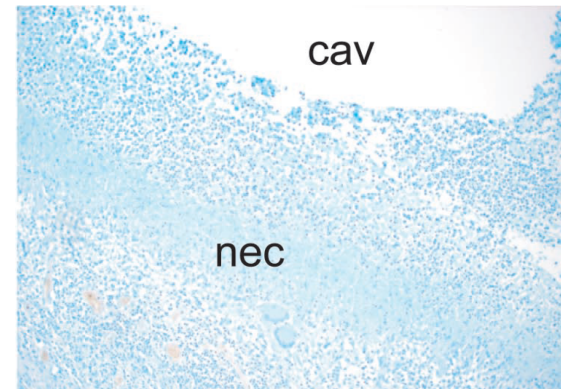
10% of TB cases: cavitation and propagation



Case courtesy of Dr Frank Gaillard, Radiopaedia.org



Boshoff *et al.* **2005** Nat Rev Microbio (3) 70-80

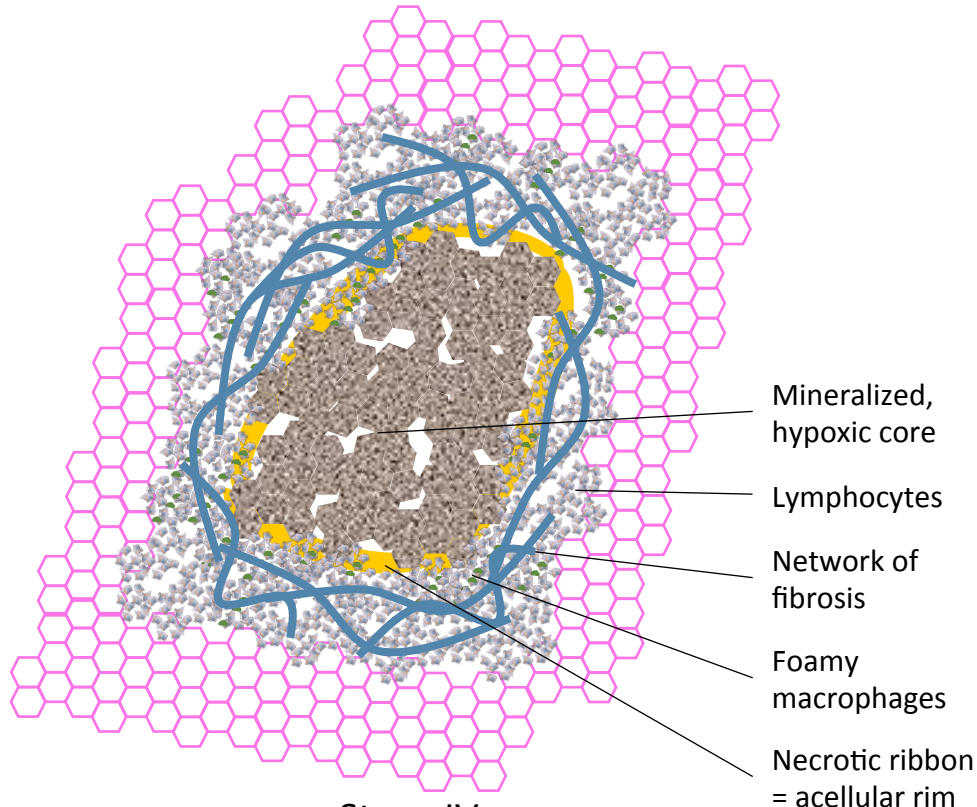


cav, cavity
nec, necrotic area

Kaplan *et al.* **2003** Infec Immun (71) 7099-7108

90% of cases: Granuloma calcification

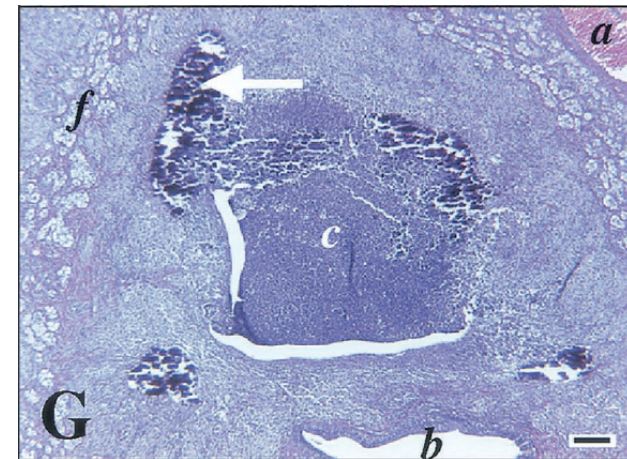
Mature granuloma with a mineralized core
Surrounded by foamy macrophages and lymphocytes



Stage IV

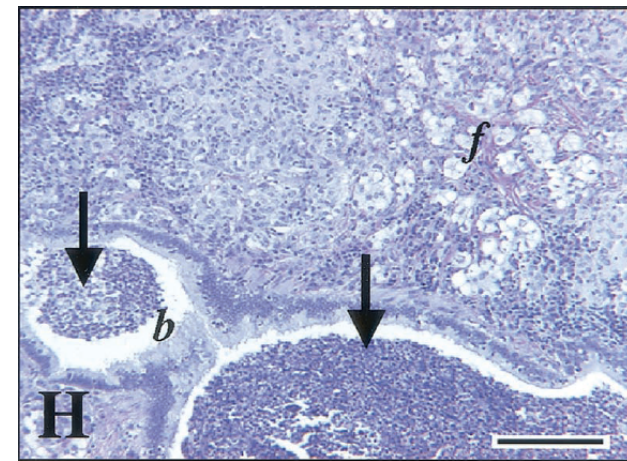
93 days in guinea pig model
Fibrosis and mineralization,
healing process by
calcification

Turner *et al.* 2003 Infect Immun (71) 864-871



c, mineralized core
f, fibrosis
a, artery
b, bronchiole

100 μ m



b, bronchiole
f, fibrosis

100 μ m

Arrows: Necrotic debris

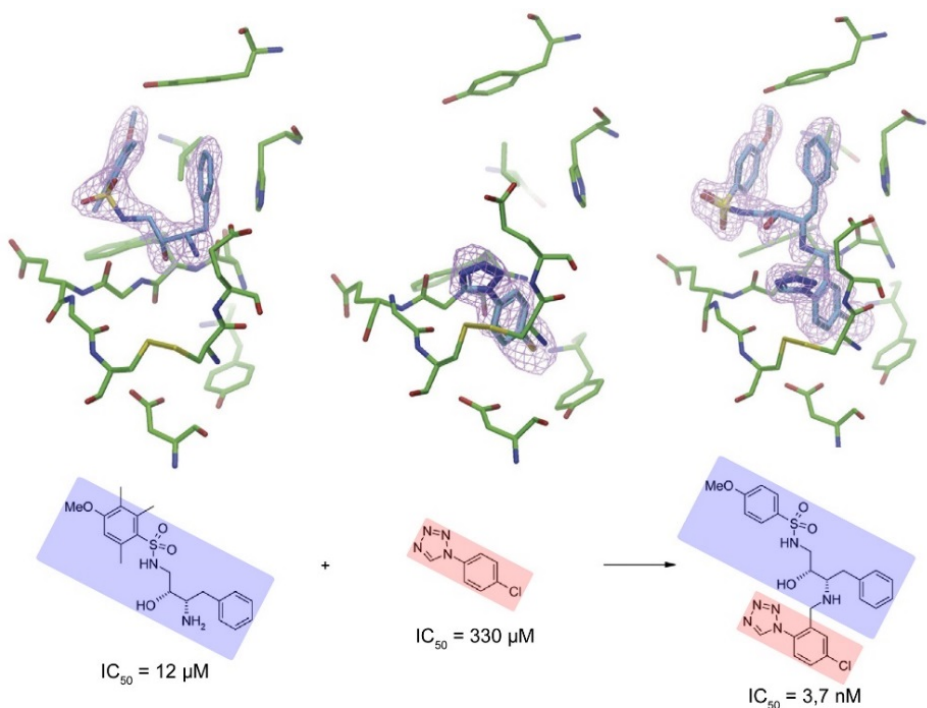
Part 1.

DRUG DISCOVERY IN TB FIELD

Screening strategies

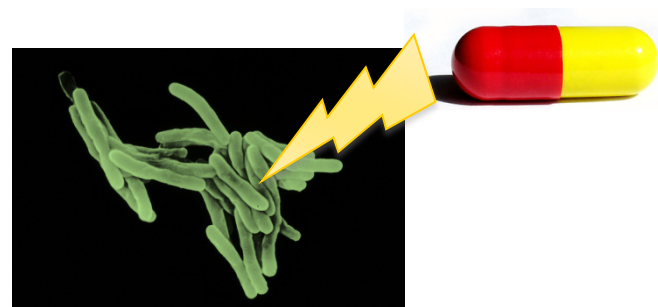
Target-based screening

- Enzymatic assays
- Fragment / *in silico* approaches
- Need a protein/crystal/model!

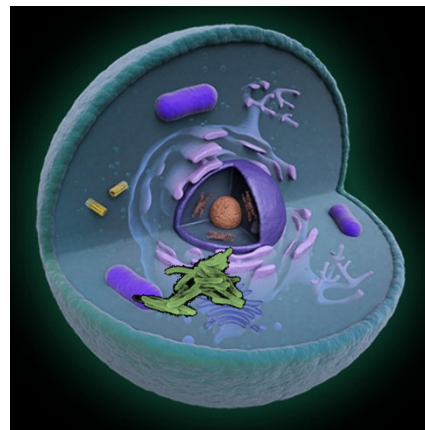


Target-free / Phenotypic screening

- Bacterial replication / survival assays

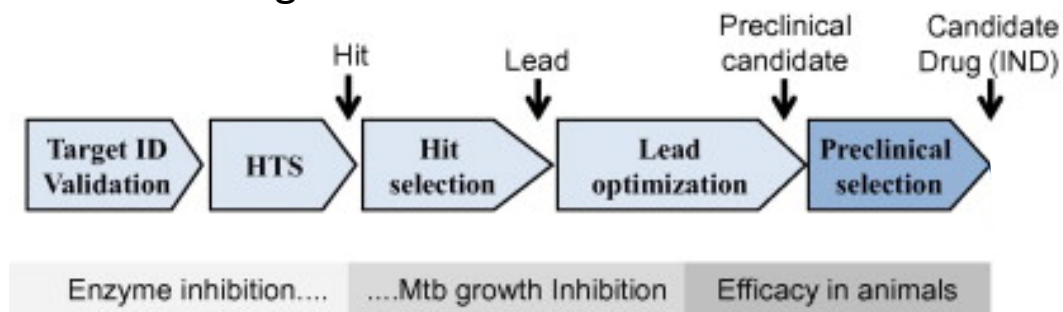


- Infected cells



Drug discovery pipeline

Target-based screening



Target



Hit selection

Primary screening

- Single dose
- Select compounds with activity > Threshold



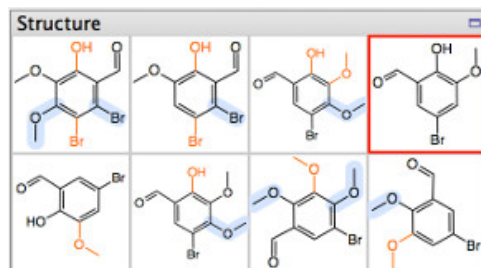
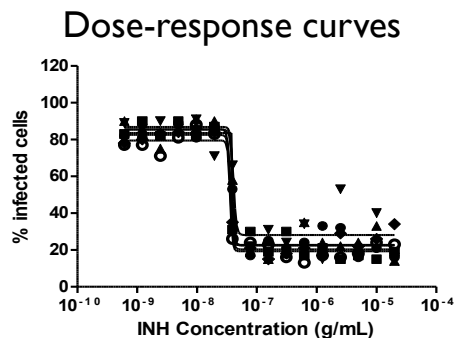
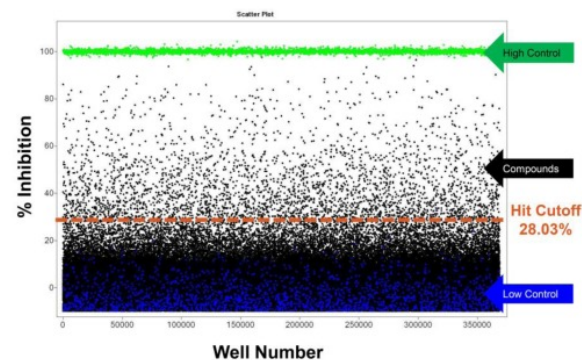
Secondary screening

- Cherry-picking & validation by dose-response curve (DRC)
- Repurchase & Resynthesize, validate again by DRC

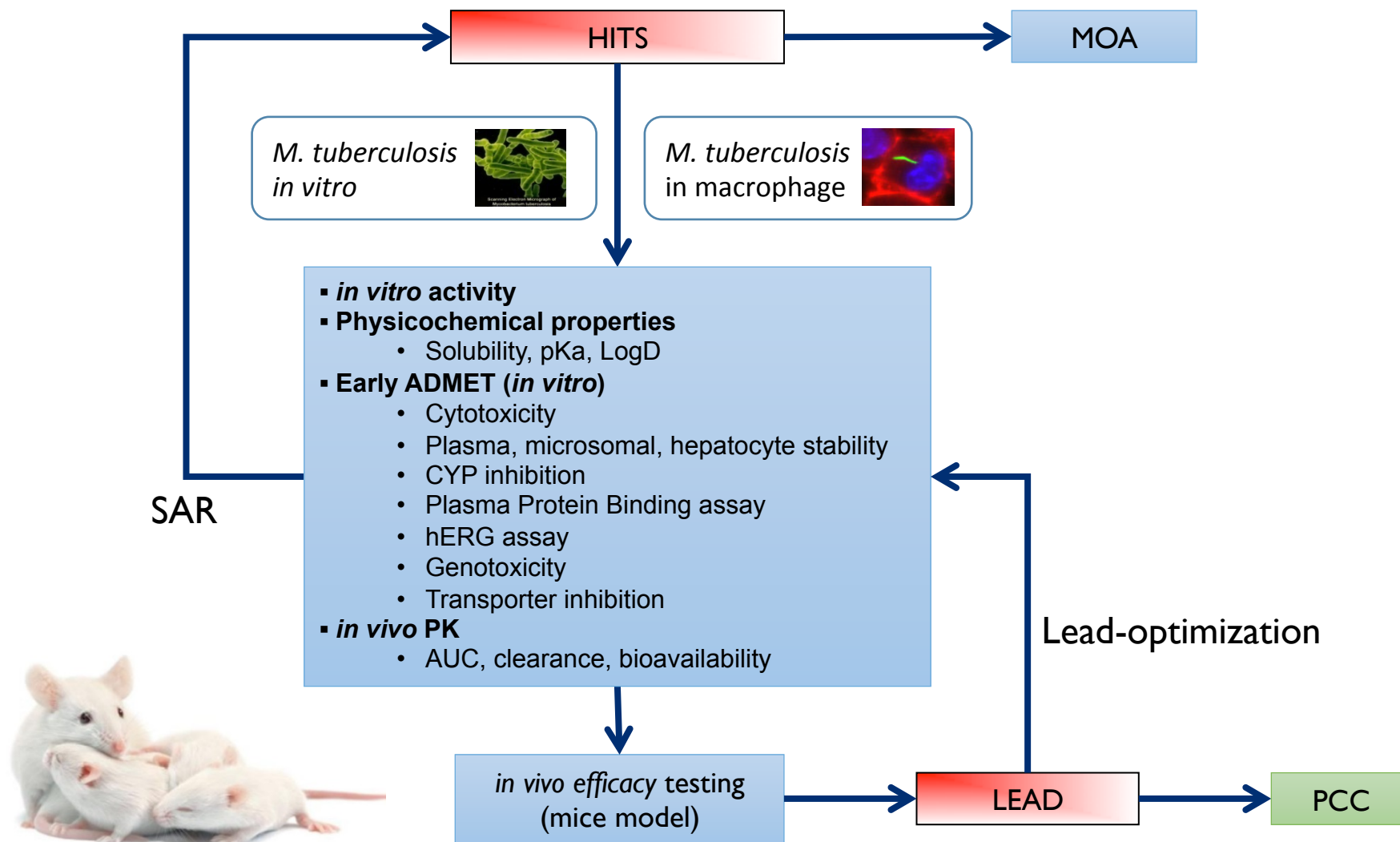


Hits

- Structure and activity confirmed
- Clustering & prioritization



Hit-to-lead



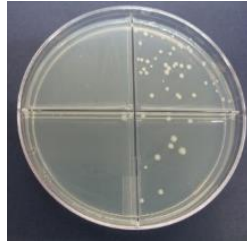
Phenotypic screening assays for Tuberculosis research



Phenotypic assays – Quantification?

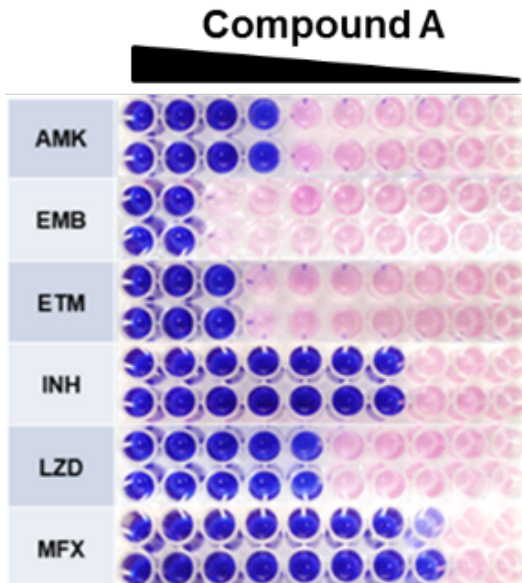
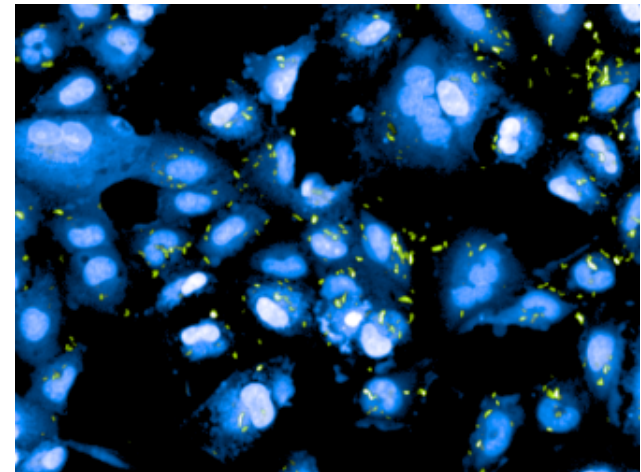
Wild-Type bacteria

- CFU counting
 - Agar
 - Bactec MGIT
- OD reading
- Resazurin



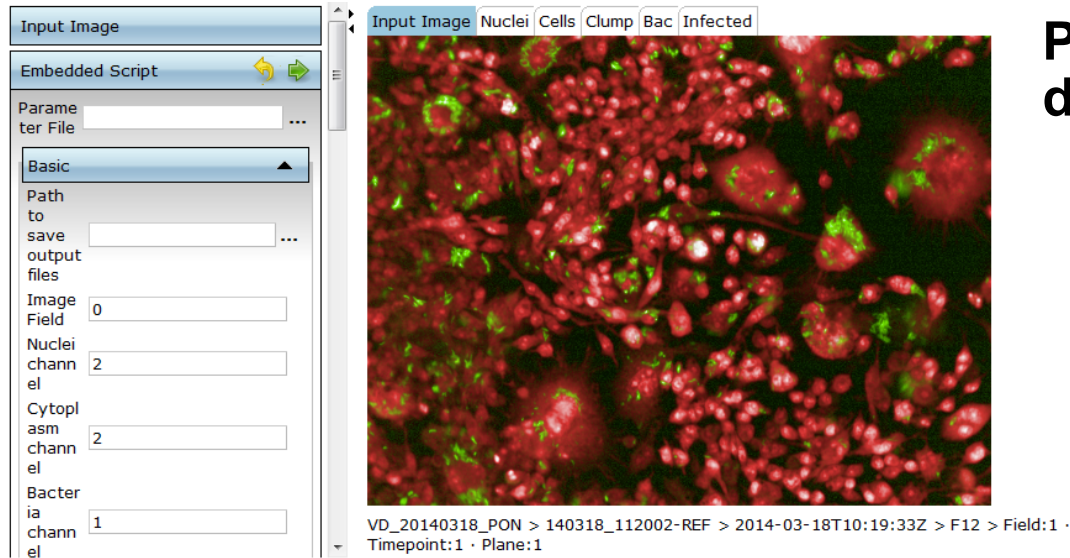
Reporter strains

- Luminescence, fluorescence (GFP, RFP...)
- Constitutive vs stimuli-driven expression



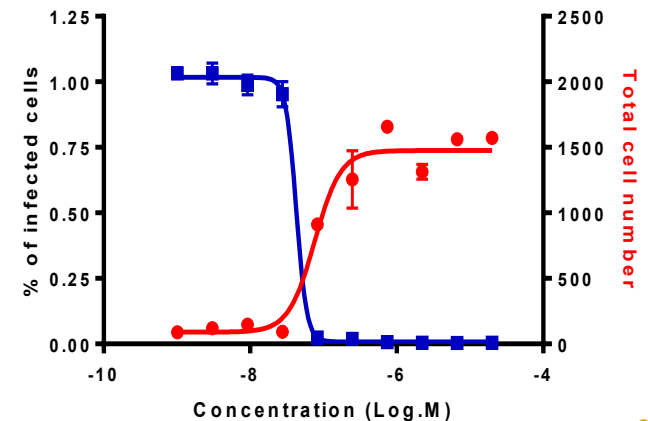
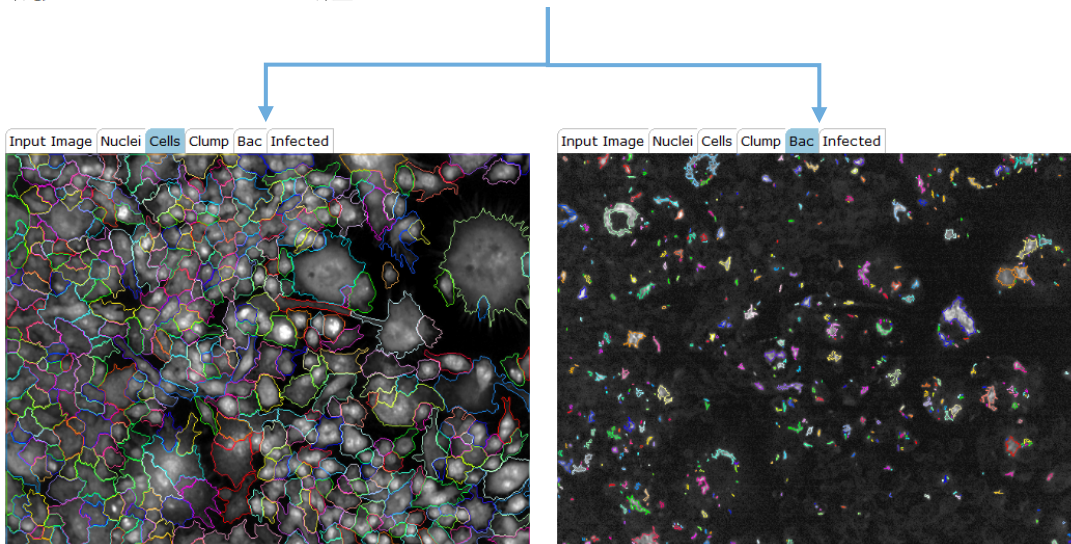
Dyes or antibodies

Multiparametric image analysis

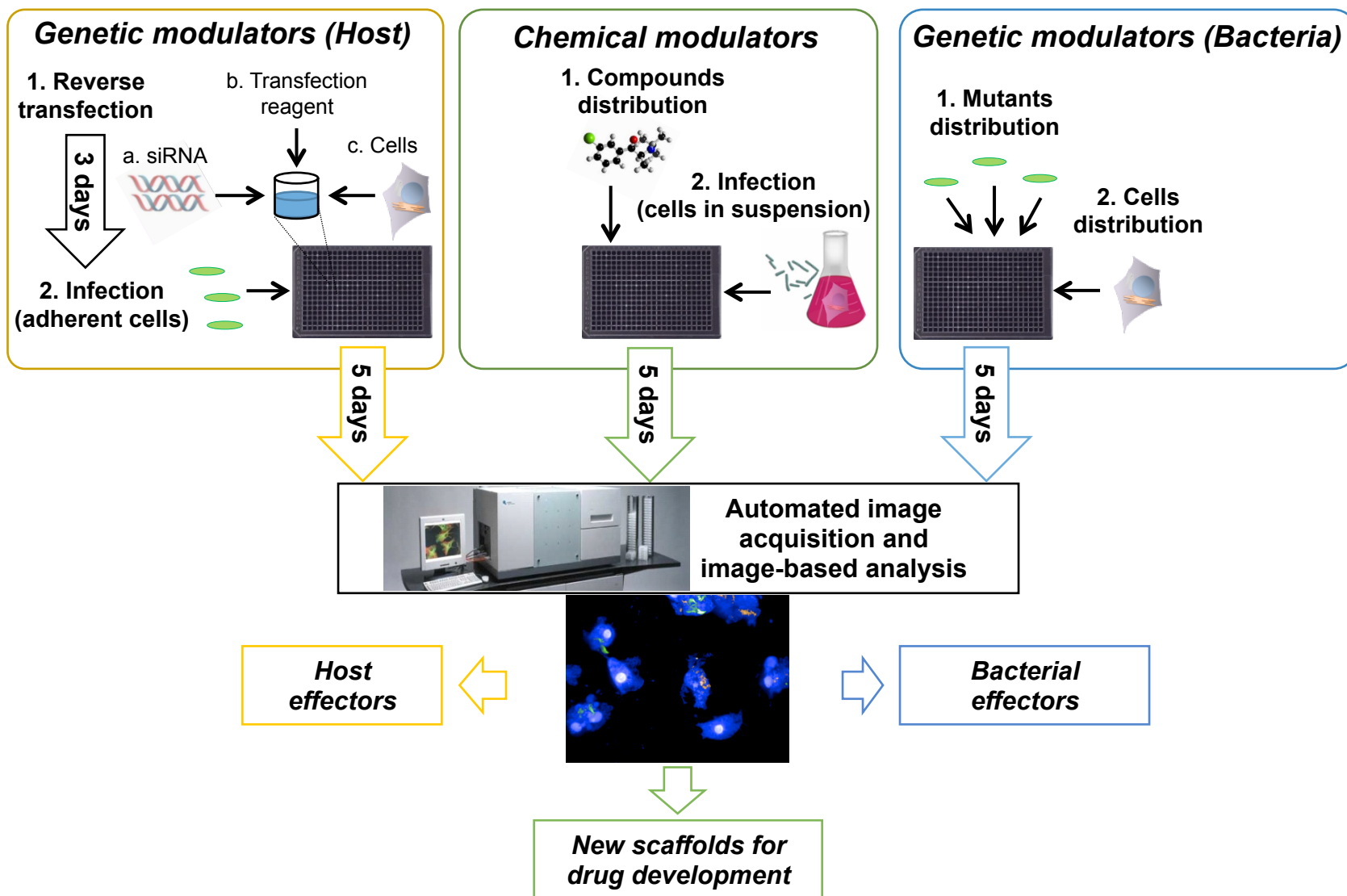


Pictures analyzed by dedicated scripts

- Definition of two Populations: cells & bacteria
- Several parameters used to quantify compound efficiency:
 - Number of cells
 - Total area of bacteria (px)
 - Average area of bacteria per infected cells (px)
 - Ratio of infected cells



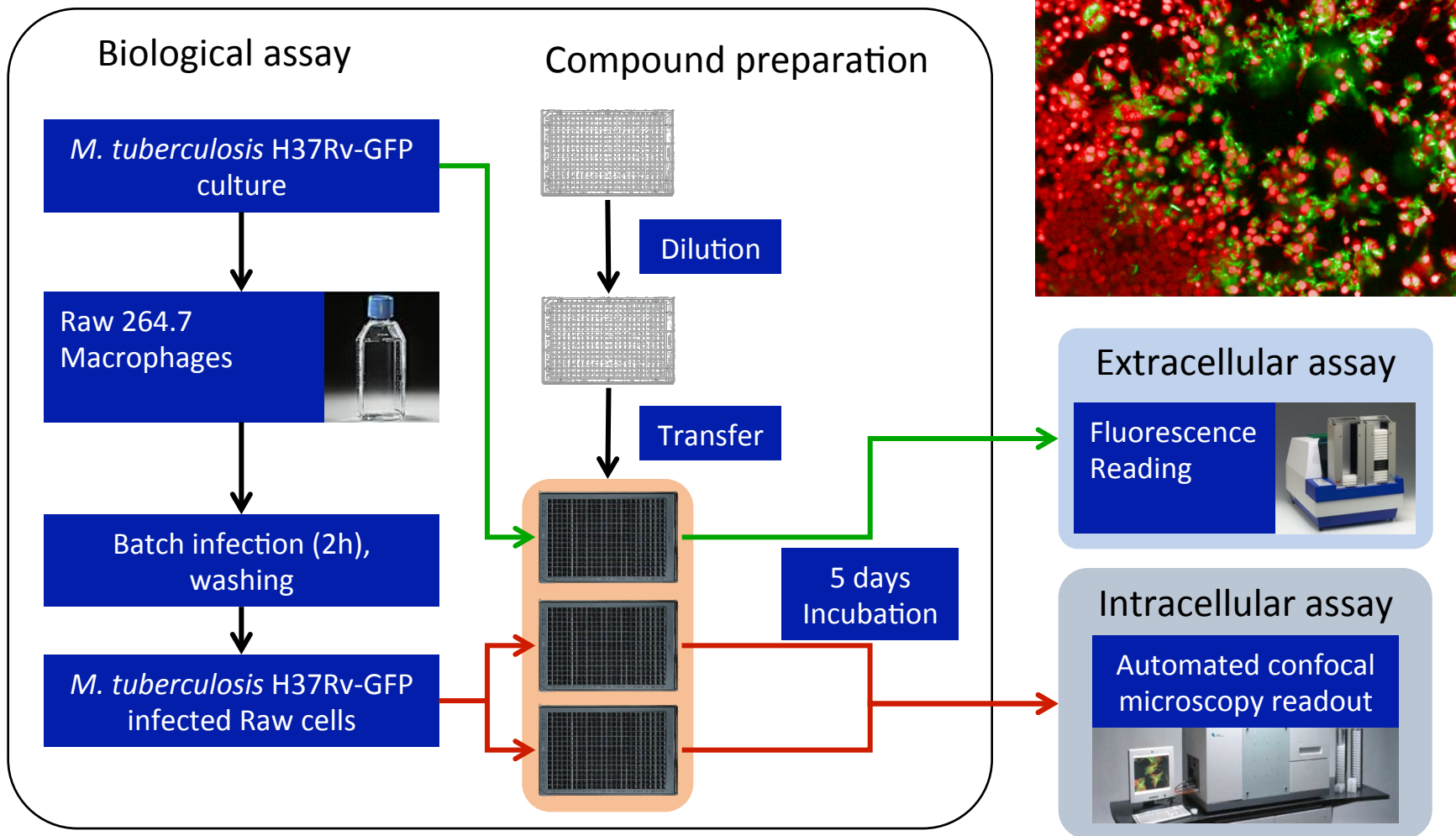
Phenotypic screening assays - overview



Part 1.

CHEMICAL MODULATORS

Chemical modulators - Screening process



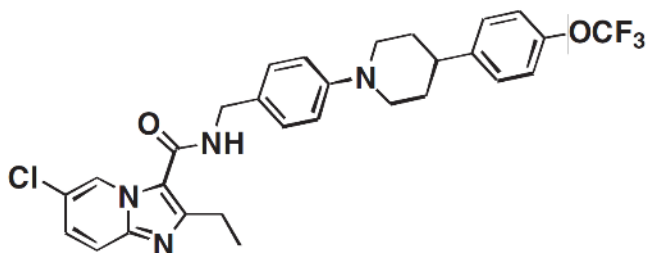
Christophe *et al.* **2010** Future Med. Chem. 2: 1283-1293

Chemical modulators: Q203

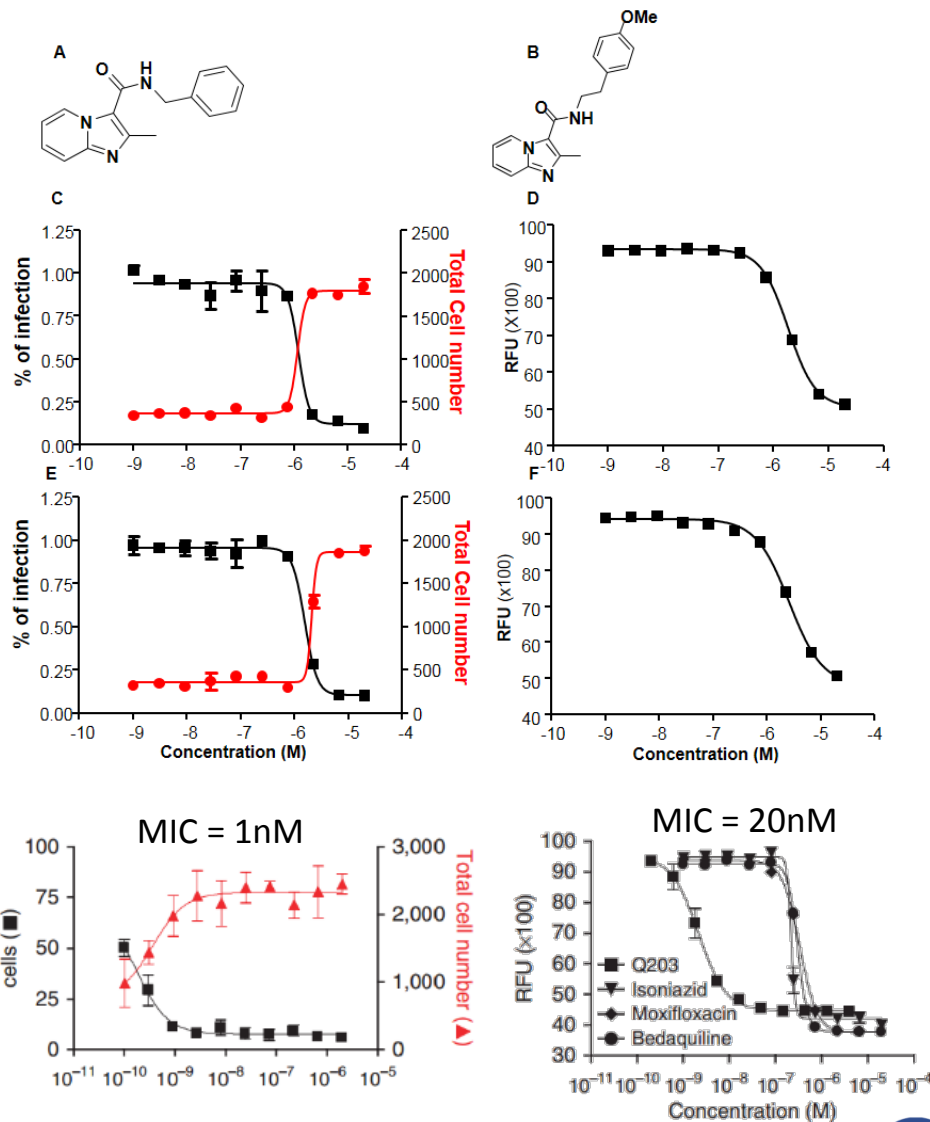
Q203

- Imidazo[1,2-a]pyridine amide (IPA) Family
- From a screening of 120,000 compounds
- Identified by infected macrophage assay (QIM)
- Hit compounds were active in the μM range

Optimizations gave preclinical candidate



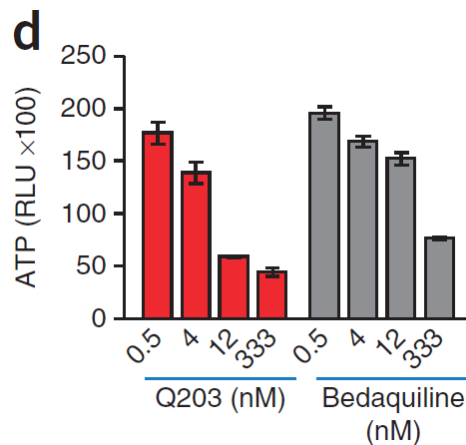
Pethe *et al.* 2013 Nat. Med



Q203 - Overview

Triggers ATP depletion

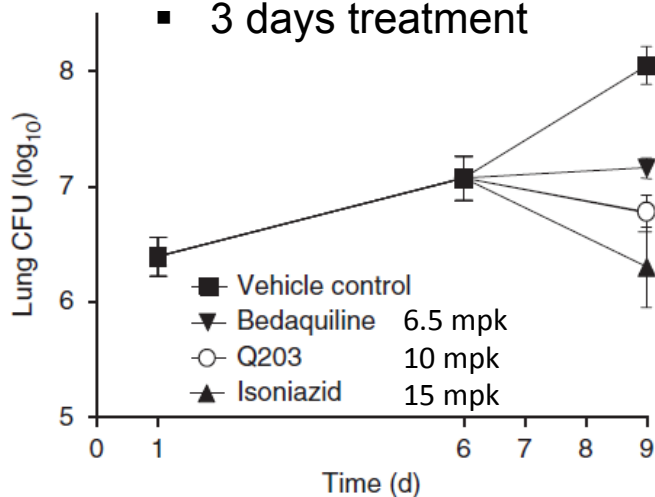
- More potent than BDQ
- Active against MDR-XDR



Type	#	Family	INH	Rif	Strept	Oflox	MIC ₉₀ (nM)
	4	Beijing	R	R	R	S	0.43
	5	Beijing	R	R	S	R	<0.43
	6	Beijing	R	R	S	R	<0.43
	7	Beijing	R	R	S	R	<0.43
	8	-	R	R	S	R	<0.43
MDR	9	Beijing	R	R	R	S	<0.43
	10	Beijing	R	R	S	S	<0.43
	11	Beijing	R	R	S	R	<0.43
	12	Beijing	R	R	S	R	0.98
	13	Beijing	R	R	S	S	<0.43
	14	Beijing	R	R	S	S	0.88
	15	Beijing	R	R	S	R	3.51
	16	Beijing	R	R	R	S	<0.43
	17	Beijing	R	R	R	R	0.43
	18	Beijing	R	R	R	R	<0.43
	19	-	R	R	R	R	<0.43
	20	-	R	R	R	R	<0.43
	21	Beijing	R	R	R	R	<0.43
	22	Beijing	R	R	R	R	<0.43
	23	Beijing	R	R	R	R	<0.43
	24	Beijing	R	R	R	R	<0.43
	25	Beijing	R	R	R	R	<0.43
	26	Beijing	R	R	R	R	<0.43
	27	Beijing	R	R	R	R	7.02
	28	Beijing	R	R	R	R	<0.43
		Beijing	R	R	R	R	7.02
		Beijing	R	R	R	R	<0.43
		Beijing	R	R	R	R	28

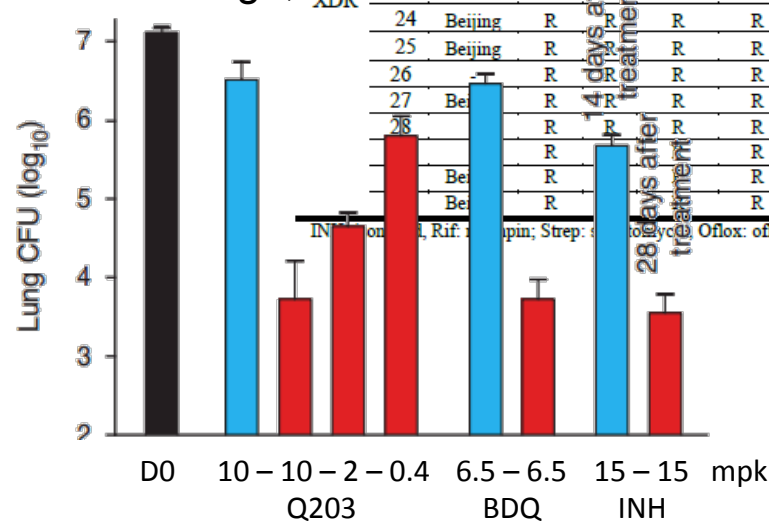
In vivo efficacy in mice

- Acute model (GSK)
- 3 days treatment



Chronic model

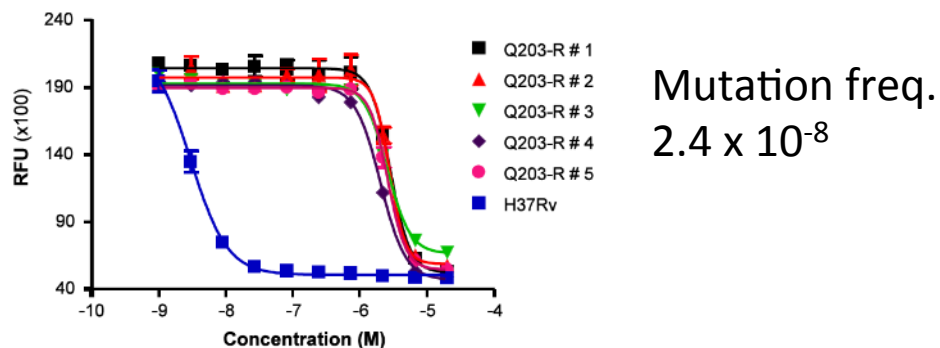
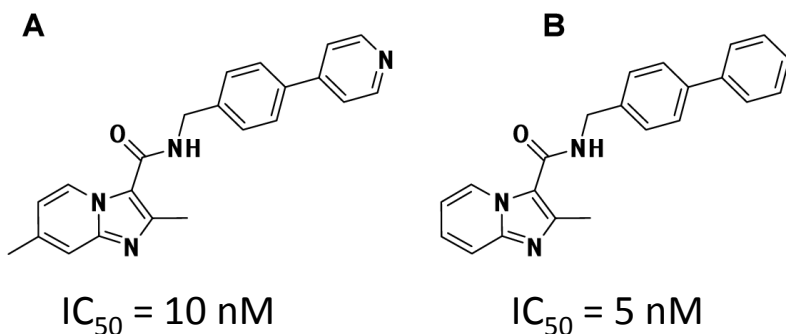
- Gavage, 5x / weeks



Q203 & QcrB

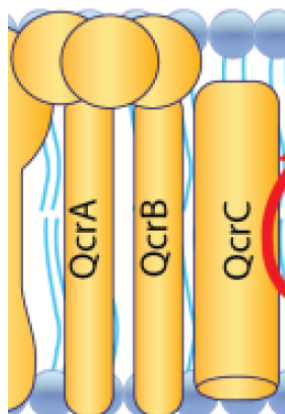
Target identification

- Generation of spontaneous resistant mutants

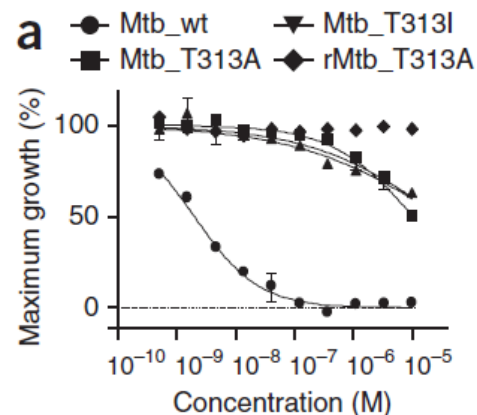


- Whole genome sequencing: *qcrB*

H37Rv	301	SAGSQPDFYMMWTEGLARI	319
CDC1551	301	SAGSQPDFYMMWTEGLARI	319
W4	301	SAGSQPDFYMMWTEGLARI	319
XDR #27	301	SAGSQPDFYMMWTEGLARI	319
XDR #29	301	SAGSQPDFYMMWTEGLARI	319
XDR #31	301	SAGSQPDFYMMWTEGLARI	319
Q203-R #1	301	SAGSQPDFYMMWTEGLARI	319
Q203-R #2	301	SAGSQPDFYMMWTEGLARI	319
Q203-R #3	301	SAGSQPDFYMMWTEGLARI	319
Q203-R #4	301	SAGSQPDFYMMWTEGLARI	319
Q203-R #5	301	SAGSQPDFYMMWTEGLARI	319



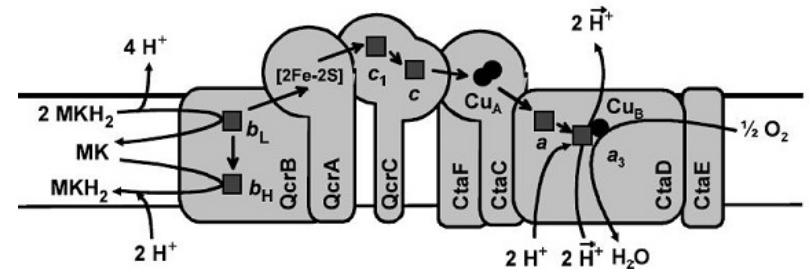
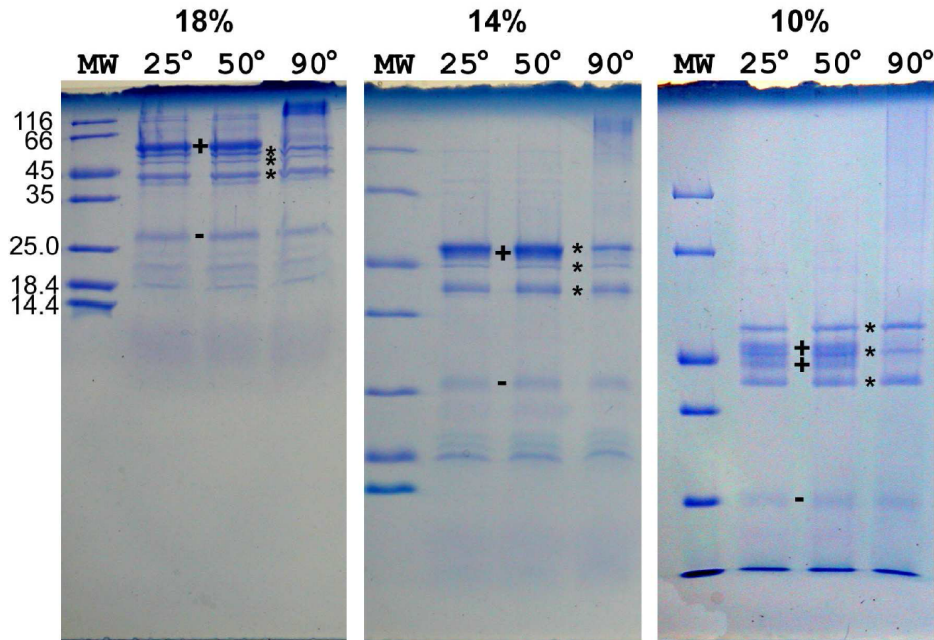
Respiratory complex III
(Cytochrome *bc* oxidase)



Q203 & QcrB

Target purification

- Collaboration with Dr. E. Berry (Upstate Univ)
- Hybrid supercomplex
 - M. smegmatis* cytochrome aa_3 + *M. tuberculosis* QcrCAB (bcc)



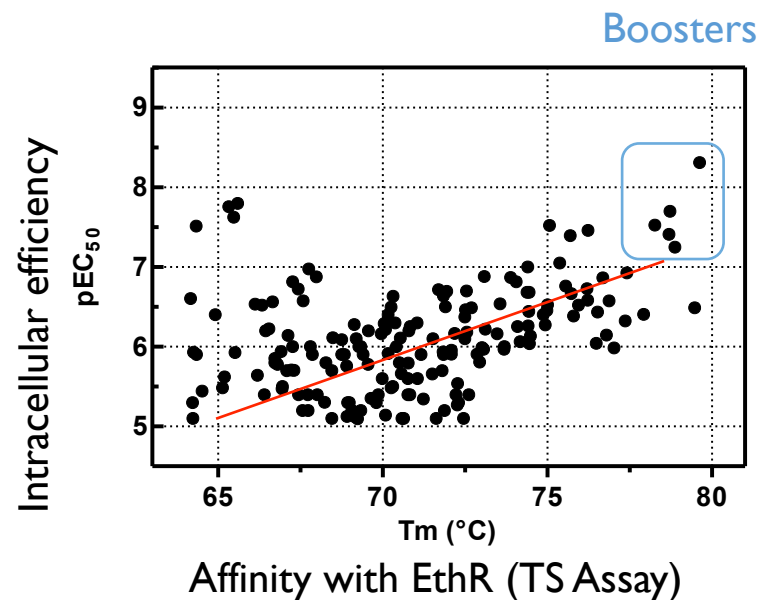
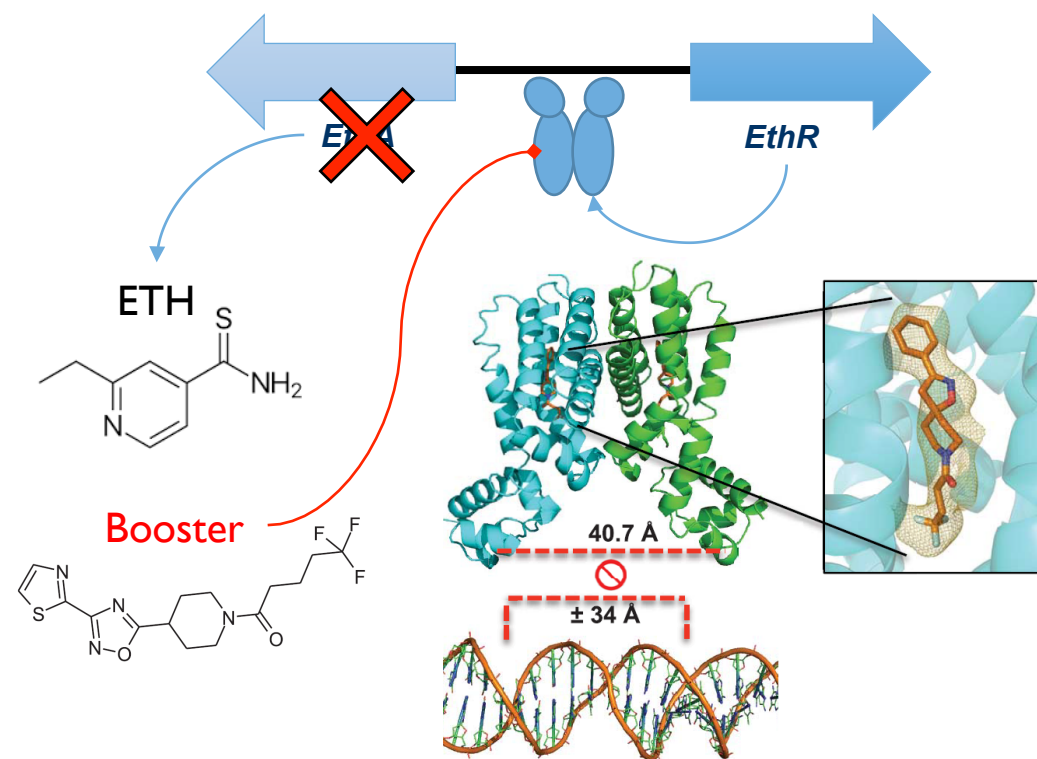
Kim et al. 2015 J Biol Chem

+, QcrB, Cox1; -, QcrC; *, QcrA, Cox2, Rieske

Chemical modulators - optimization

Adding value by combining screening strategies

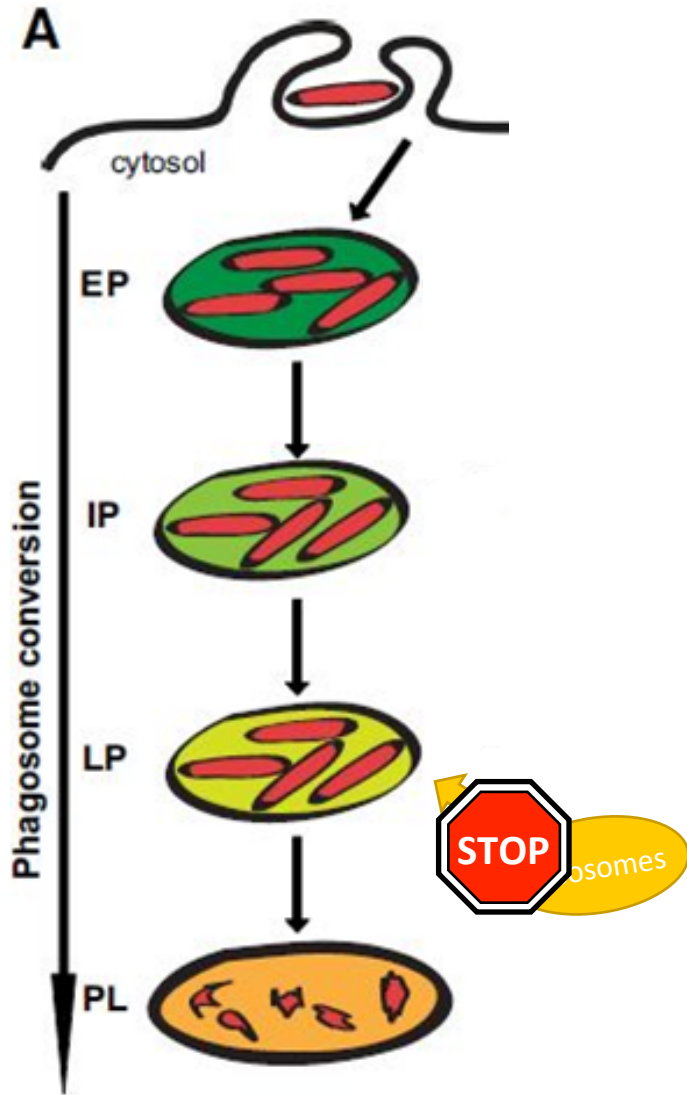
- Finding compounds active in multiple models



Part 3.

GENETIC MODULATORS (BACTERIA SIDE)

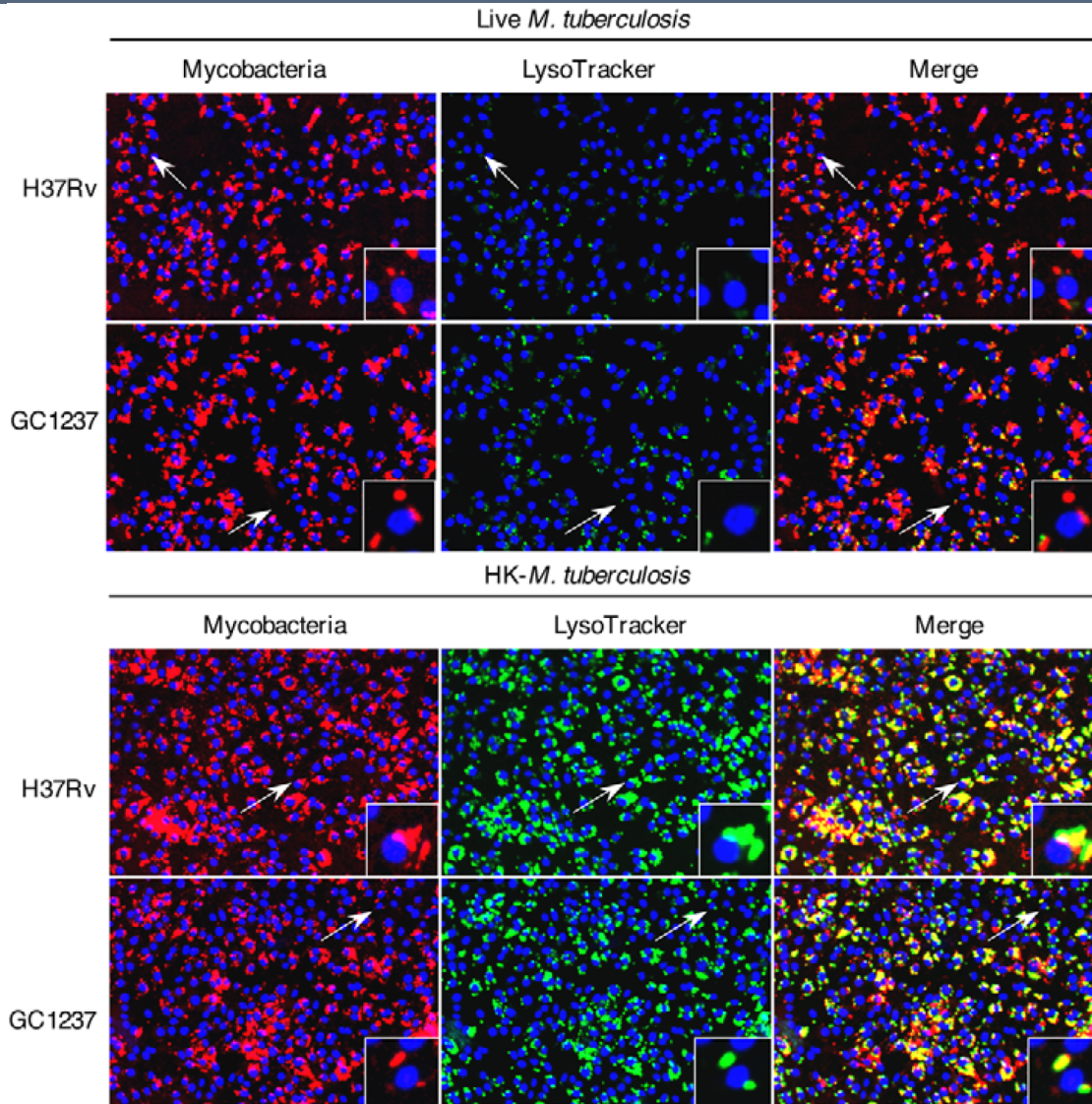
Genetic modulators – Bacteria side



Failure of lysosomal fusion with *M. tuberculosis* containing phagosomes

- Which bacterial effectors are implied?
- Use of a library of ~11,000 transposon mutants build in *M. tuberculosis* Beijing GC1237 strain
- Follow lysosomal fusion using the pH-sensitive LysoTracker probe
- Read-out: number of bacteria in acidified compartments

Genetic modulators - Screening principle



Validation using H37Rv and Beijing GC1237 fluorescent (RFP) strains

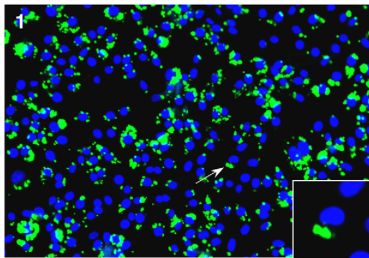
- Live and heat killed (HK) bacteria
- Increase in the number of acidified compartment with the HK strains

Brodin *et al.* 2010 Plos Pathogen 6(9): e1001100

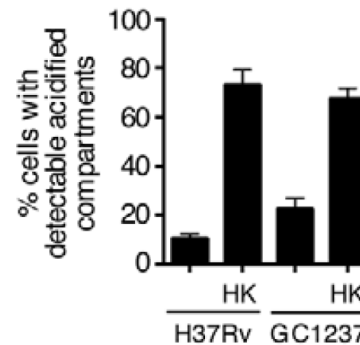
Genetic modulators - Quantification

Image analysis

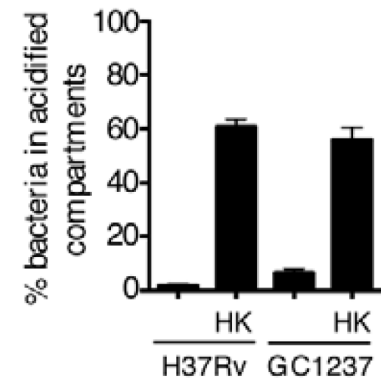
- Detection of cell nuclei
- Detection of acidified compartment
 - Detection of bacteria is optional,
 - Quantification gave similar results



Lysotracker



Bacteria + Lysotracker



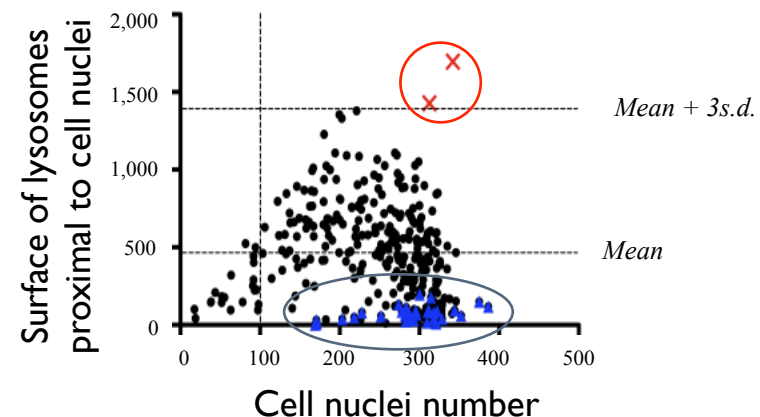
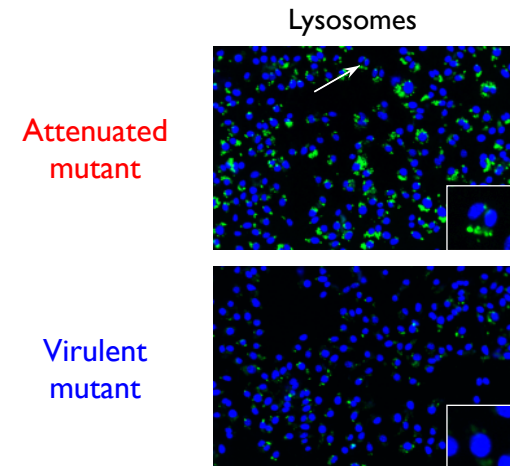
Genetic modulators - Process and screening

Process:

- Mutant distribution
- Cell distribution
- 2h incubation
- Staining, fixation, reading

Results:

- Two extreme phenotypes:



Mutant library

Transfer

Primary
macrophages
(BMDM)

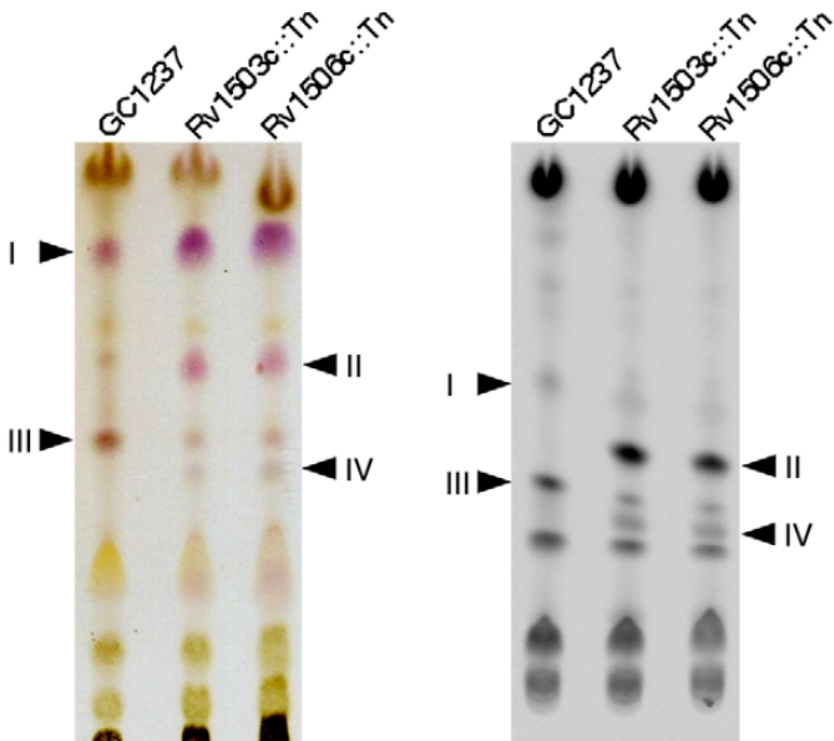
2 hours
infection

Lysotracker assay

Automated confocal
microscopy readout

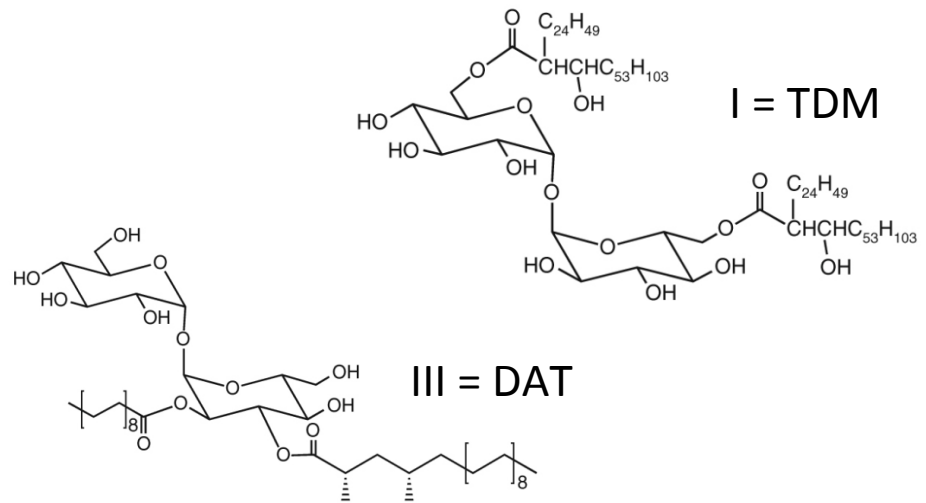
Genetic modulators – Results

Set of 10 mutants with defect in escaping lysosomal fusion



Defect in DAT production

Mutant id	Gene (nt ^a)	Putative function
P69D07	<i>pst53</i> (482)	Phosphate transport
P117C08	Rv1503c (186)	TDP-4-oxo-6-deoxy-D-glucose transaminase (glycosyl aminotransferase)
P2E07	Rv1506c (211)	Methyltransferase
P65B12	<i>lppM</i> (272)	Lipoprotein of unknown function
P1E07	Rv2295 (-35)	Unknown
P58C04	<i>fadD28</i> (1252)	DIM biosynthesis
P32E07	<i>moaC1</i> (507)	Molybdopterin biosynthesis
P36D07	<i>moaD1</i> (49)	Molybdopterin biosynthesis
P55C04	<i>ppe54</i> (1126)	Unknown
P39E07	Rv3880c (269)	Unknown



Part 3.

GENETIC MODULATORS (HOST SIDE)

Genetic modulators – host side

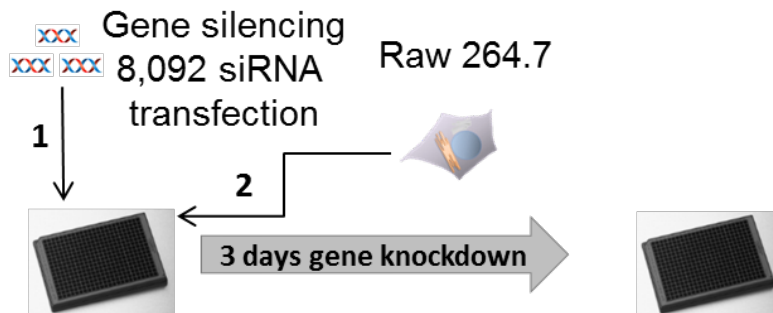
Objectives:

- To identify essential pathways manipulated by the bacteria

Method:

- Bacterial replication assay in siRNA transfected cells

Reverse transfection

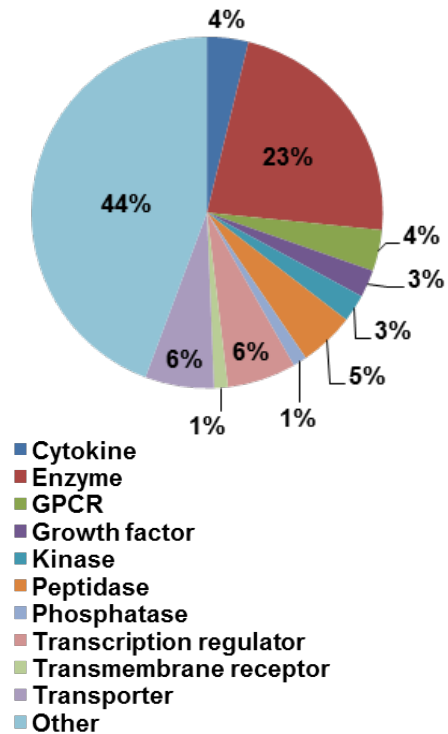
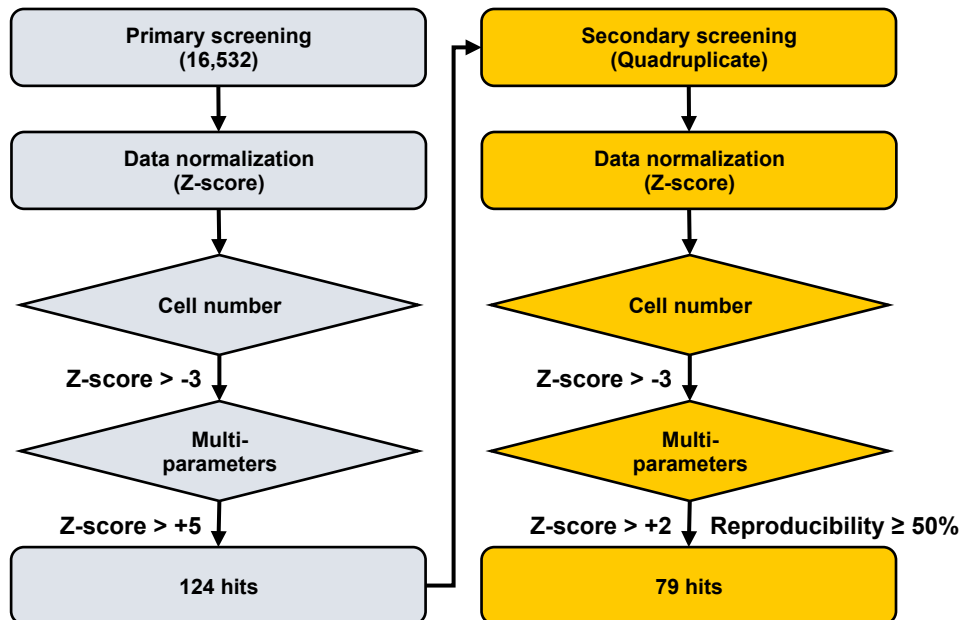


Genetic modulators – Host side

Strategy can be applied to different cell types

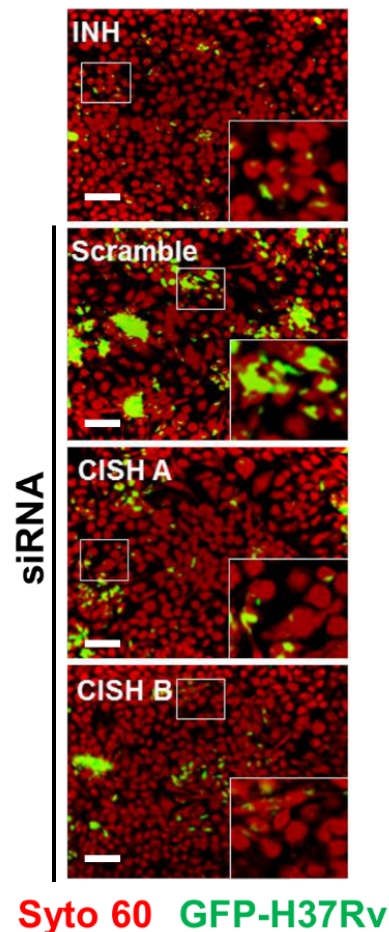
- Macrophages
- Epithelial cells

Validation process:

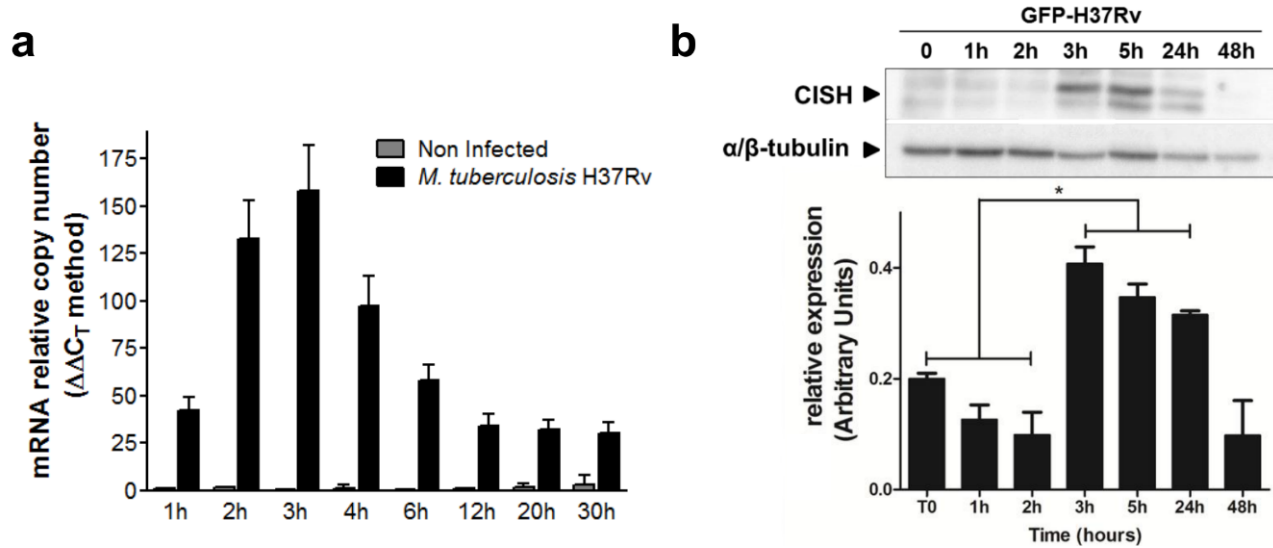


Identification of CISH

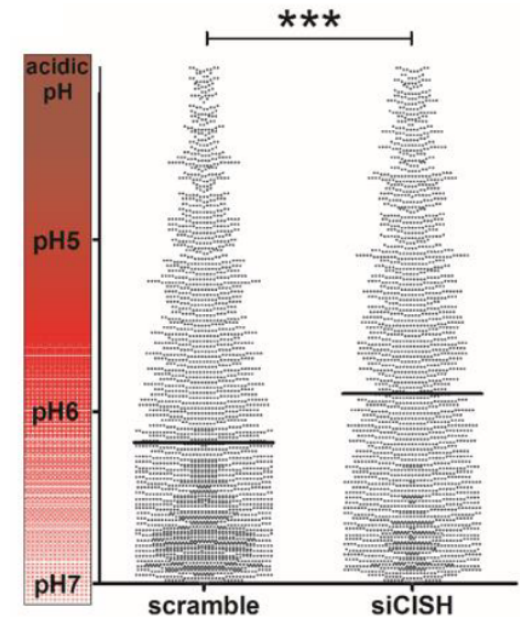
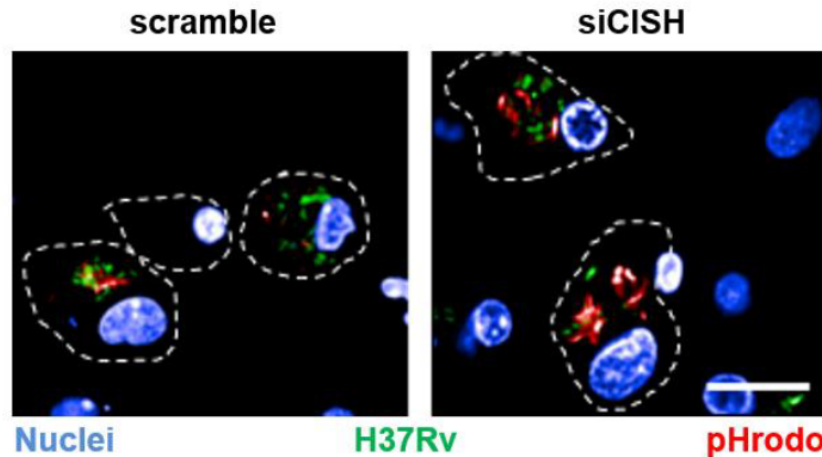
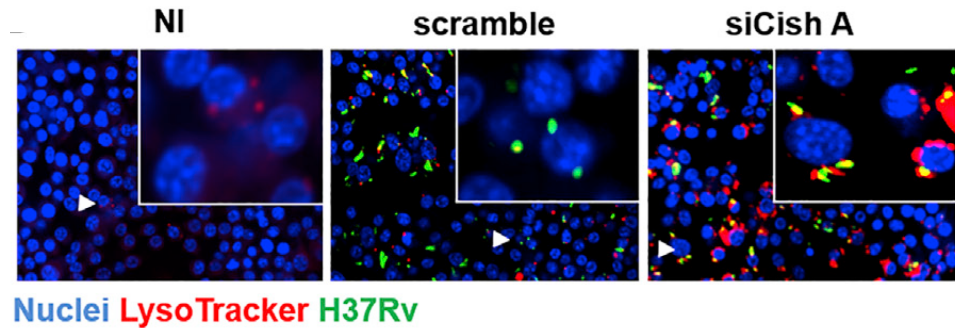
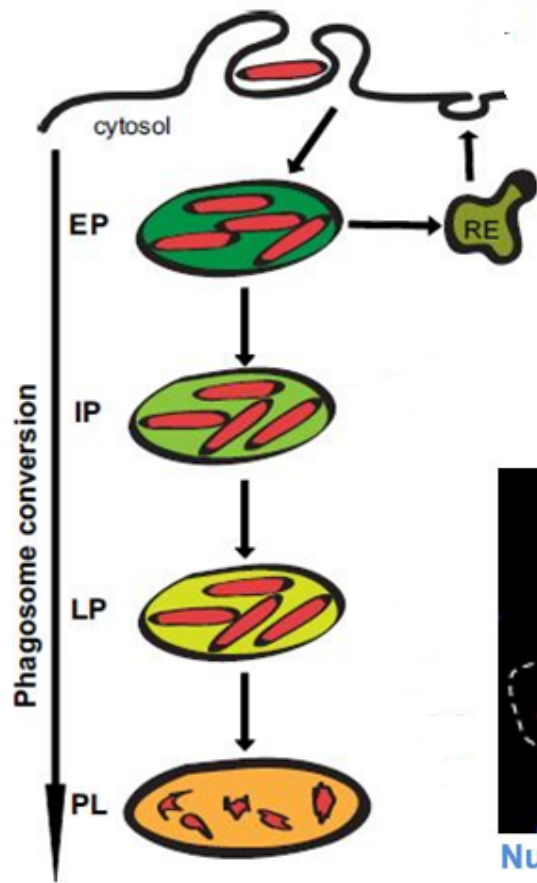
CISH is required for efficient *Mtb* replication in macrophages



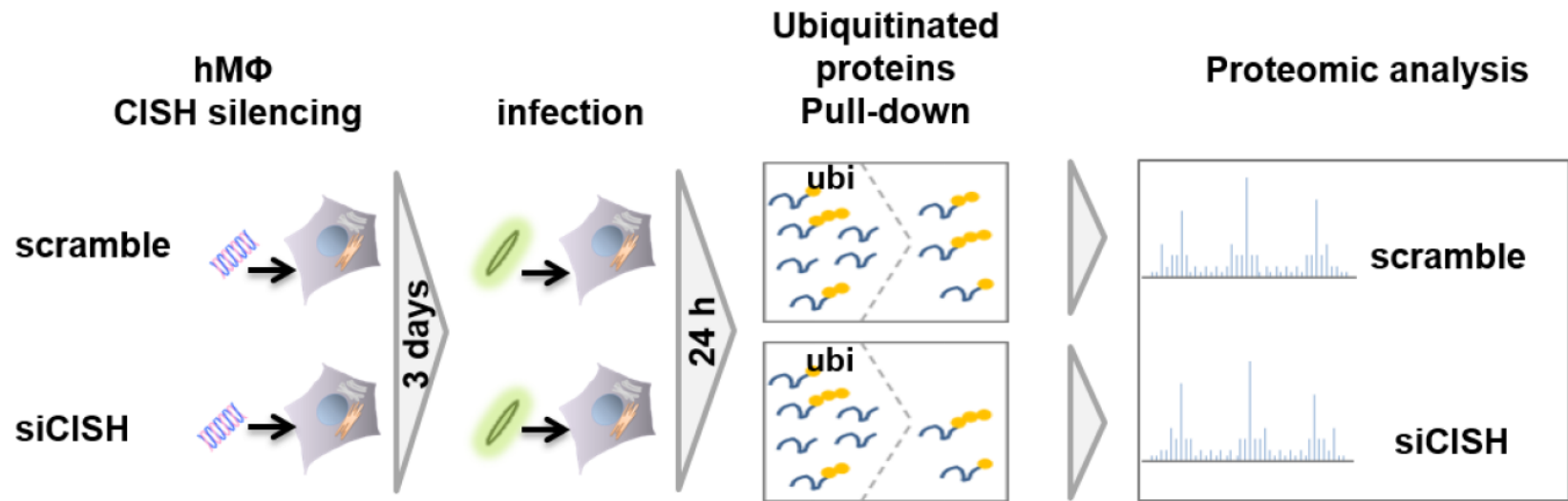
CISH expression pattern



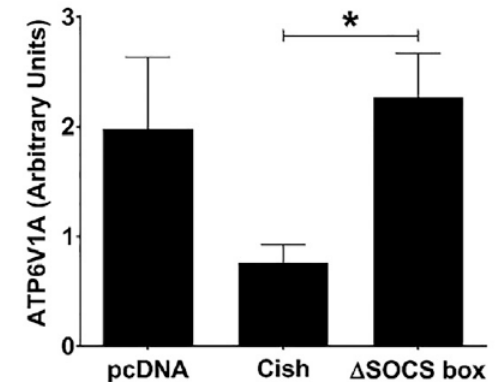
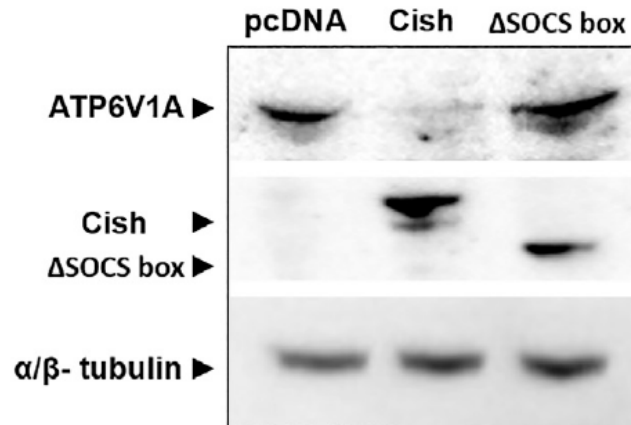
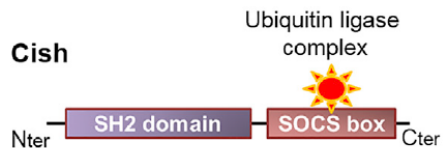
CISH interferes with phagosome acidification



CISH targets V-ATPase



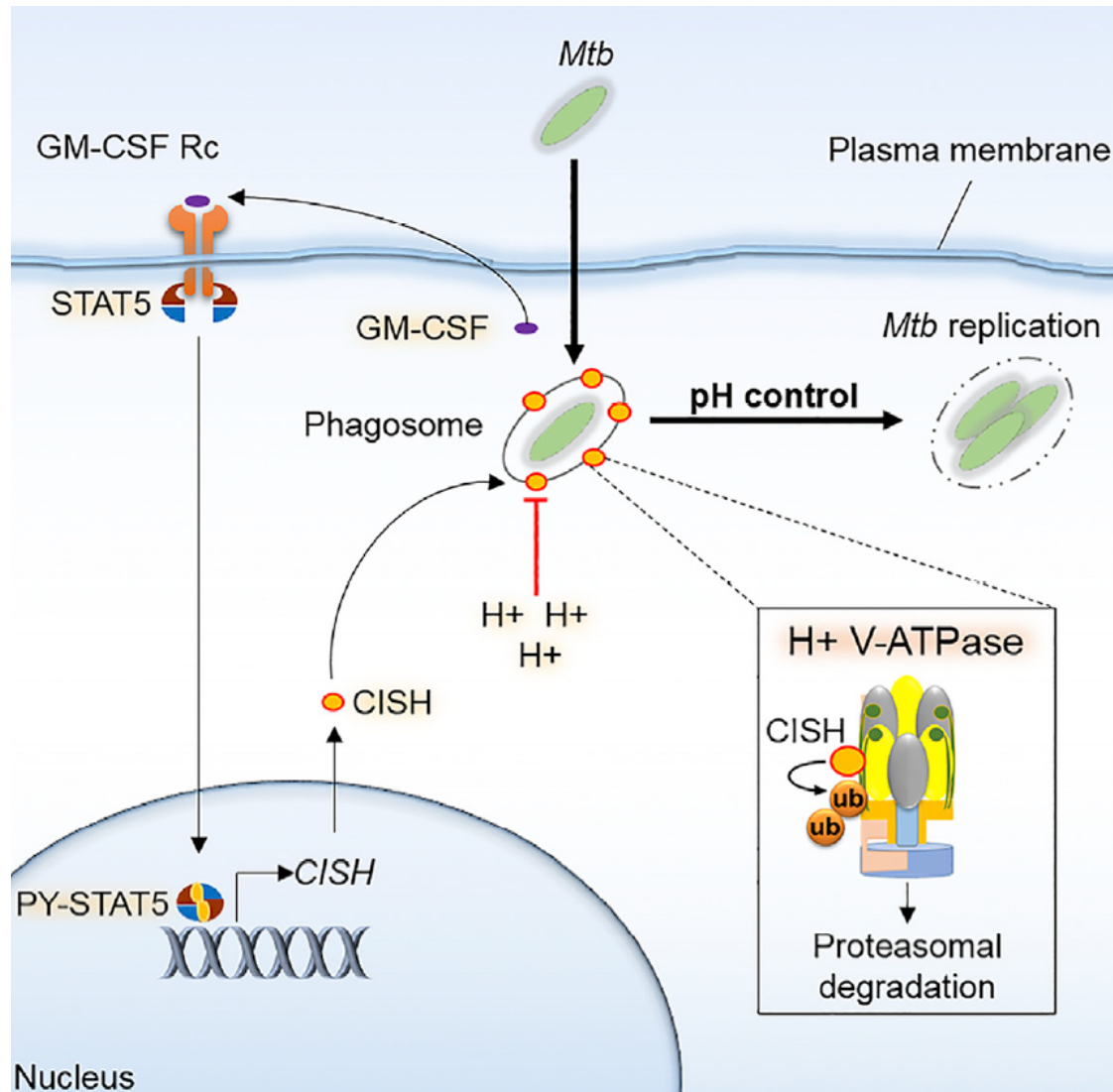
Expression vector : pcDNA 3.1



Queval *et al.* 2017 Cell Reports



CISH – A summary



Thank you!

Q & A

Vincent Delorme
Team Head
Tuberculosis Research laboratory
+82 31-8018-8190
vincent.delorme@ip-korea.org

