INTERNATIONAL RESEARCHERS

Current Alarming Status, Strategies and Drug Development Pipeline for Tuberculosis

S S Pekamwar, Aakansha Singh, S R Lokhande, M S Attar

Volume No 1 Issue No.4 December 2012

www.iresearcher.org

ISSN 227-7471

THE INTERNATIONAL RESEARCH JOURNAL "INTERNATIONAL RESEACHERS"

www.iresearcher.org

© 2012 (individual papers), the author(s)

© 2012 (selection and editorial matter)

This publication is subject to that author (s) is (are) responsible for Plagiarism, the accuracy of citations, quotations, diagrams, tables and maps.

All rights reserved. Apart from fair dealing for the purposes of study, research, criticism or review as permitted under the applicable copyright legislation, no part of this work may be reproduced by any process without written permission from the publisher. For permissions and other inquiries, please contact

editor@iresearcher.org

INTERNATIONAL RESEARCHERS is peer-reviewed, supported by rigorous processes of criterion-referenced article ranking and qualitative commentary, ensuring that only intellectual work of the greatest substance and highest significance is published.

Current Alarming Status, Strategies and Drug Development Pipeline for Tuberculosis

¹SS Pekamwar, ² Aakansha Singh, ³SR Lokhande, ⁴MS Attar

School of Pharmacy, SRTM University, Nanded-431606 (MS), India.

(INDIA)

² singhakansha20@yahoo.in

ABSTRACT

From the past few decades and today tuberculosis ranks at the top amongst the world's deadliest disease's causing morbidity. Due to lack of mandatory primary health care and cost effective treatment, slow emergence of resistant strains of mycobacterium tuberculosis combat against the disease is still a matter of fact globally.Latent infection TB patients are at a risk of reactivation and this is one of the major barriers in controlling tuberculosis. Except pyrazinamide and to some extent INH none of the drugs currently available act on latent M. tuberculosis. Therefore, there is an urgent need to develop novel drugs that can act against both actively growing and dormant bacteria. The ameliorations in the strategies that paved the way in the search of new drugs involves improvement in the existing drugs, systematic screening, exploitation of the biological information and planned research and rational approaches. New promising drug candidates involve the recent discovery of diarylquinoline, new fluoroquinolones, rifamycin, oxazolidinones and nitroimidazopyran. Desirable targets should be involved in vital aspects of bacterial growth, metabolism and viability, whose inactivation will lead to bacterial death. India accounts for nearly one third of global TB problem. Tuberculosis Kills 14 times more people than all tropical diseases and 4.5 million TB cases with 1.8 million new cases being reported each year. Approximately 50 % of the India's population is reported to be tuberculin test positive. Annually about 0.4 million death and one million new cases of TB are reported and one person dies every minute by TB. This article intends to focus the current trends and new pipeline drug development including the new drug targets and the novel promising compounds under clinical trials.

Keywords: Mycobacterium tuberculosis, chemotherapy, anti-tuberculosis, drug candidates, tuberculin test

INTRODUCTION

Tuberculosis, the preeminent health concern of this century, effectuated by Mycobacterium Tuberculosis the etiological agent of TB, has currently affected two billion people with more than 1.6 million deaths and 9.2 million new cases being reported each year 1. With one-third of the world's population harbouring the tuberculosis bacillus, 2 million deaths due to TB occurring each year, 3 million people becoming infected with both HIV and Mycobacterium tuberculosis, and the augmenting incidence of both multidrug-resistant TB (MDR-TB) and extensively drug-resistant TB (XDR-TB), the need for more effective chemotherapy for the treatment of TB has never been greater 2,3,4. The biggest conundrum of intensive sprawl of HIV infection and the ascendant problem of multi-drug resistant TB (MDR-TB) plays a consequential role for the need of new drugs to more effectively treat the disease in all patients 5,6,7. Current survey demonstrates that in 2008, there were estimated to be 9.4 million new M. tuberculosis infections and 1.8 million deaths as a result of TB, the most by any single infectious agent ^{8,9}.

PATHOGENESIS

Mycobacterium tuberculosis is a large ,complex, slow growing, gram-positive, rod-shaped, obligate ,aerobic bacteria belonging to the order actinomycitales family mycobactericae whose cell is highly rich in lipids, evolved from soil bacterium more than 10,000 years ago is an enormous contagious and respiratory transmitted disease that sprawls from person to person when a contagious person oozes TB bacilli into the air by coughing or sneezing which are then inhaled into the lungs by another individual, mainly 1–2 mm in size. ^{10, 11}.

```
{}^{\rm Page}33
```

Once inside the lungs, the dynamic interplay between the host and pathogen can have any of the four outcomes:

- (a) The initial host response may be completely effective and kill the bacilli ¹¹;
- (b) The organisms can grow and multiply immediately after infection, resulting in primary TB,
- (c) The bacilli may become dormant and never cause disease at all.

(d) The latent bacilli can eventually become active and progress to disease condition. The bacilli do not remain in the airways in the lung; they enter the lung parenchyma and replicate in the tissue macrophages and monocyte-derived macrophages.

MYCOBACTERIUM TUBERCULOSIS (MTB) BACETERIUM

It is a small aerobic non-motile bacillus with high lipid content and a thick cell wall. It has a very slow generation time as compared to other bacteria like Escherichia coli and Plasmodium falciparum. As a result of the fact that MTB has a cell wall but lacks a phospholipids 'outer, it is classified as a Gram-positive bacterium ¹³. However, accomplishment of Gram stain, MTB either stains very weakly Gram-positive, cell clump together due to hydrophobicity or becomes impermeable to stain. They give acid fast stain hence called as acid fast bacilli.

SYMPTOMS ASSOCIATED WITH TUBERCULOSIS INFECTION^{14, 15}

- Fever
- Night-time sweating
- Unintentional loss of weight
- Persistent cough
- Constant tiredness
- Loss of appetite

DIAGNOSIS

Diagnosis of active TB disease remains difficult as clinical symptoms of tuberculosis are common to other diseases. Tests that are used for diagnosis are:

- Tuberculin Skin Test (TST) or Monteux test
- Chest Radiograph (X-ray)
- Sputum Smear Microscopy (SSM) culture
- Polymerase Chain Reaction (PCR)
- Blood Test (e.g. T-SPOT.TB)
- Expert MTB/RIF Test

MANAGEMENT OF TUBERCULOSIS¹⁶

Vaccination

Active immunization is one of the essential components to control tuberculosis, although the vaccination is ineffective nowadays. Until now one billion people have been vaccinated with BCG. In general, four classes of vaccine candidates are being focused and these are; rationally attenuated strains of *Mycobacterium tuberculosis*, Bacilli Calmette Guerin (BCG) vaccine, protein subunit vaccines and nucleic acid vaccines.

Chemotherapy

Chemotherapy of TB started in the 1940s. In 1943, anti-TB research resulted in discovery of the active anti-TB agent Streptomycin.17 Since then, a number of agents have been discovered, like para-aminosalicylic acid (PAS), isoniazid

(INH), pyrazinamide (PZA), cycloserine, ethionamide (ETA), rifampicin (RMP), and ethambutol (EMB). Drugs used for treatment of tuberculosis can be divided into 3 classes:

- 1) First line anti-tubercular agents
- 2) Second line anti-tubercular agents
- 3) Third line anti-tubercular agents

FIRST LINE ANTI-TB AGENTS¹⁰

Important first-line antitubercular drugs are:

- Streptomycin,
- Isoniazid,
- Rifampicin,
- Ethambutol,
- Pyrazinamide,





Isoniazid

Pyrazinamide



Streptomycin



Rifampicin

Fig 1: First line antitubercular drugs

SECOND LINE ANTI-TB AGENTS¹⁰

As per WHO second line anti-tb agents fall into six classes of second line drugs that are used in the treatment of tuberculosis.

A drug may be classified as a second-line because of either of two possible reasons:

- it may have lower efficacy compared to the first-line drugs or it may have toxic side-effects
- It may be inaccessible in most developing countries.
- These comprise of different classes namely

•Amino Glycosides: (Amikacin, Kanamycin),

- Polypeptides: (Capreomycin, Viomycin),
- •Fluoroquinolones (Fqs): (Ciprofloxacin, Moxifloxacin),
- •Thioamides: (Ethionamides, Prothioamide),
- •Cycloserine And *P*-Amino Salicylic Acid





Para amino salicylic acid



Cyclosrine

Ethionamide

Fig 2: Second line antitubercular drugs

THIRD LINE ANTI-TB DRUGS¹⁰

These are not in WHO list:

- Rifabutin
- Macrolides: e.g., clarithromycin (CLR)
- Linezolid (LZD)
- Thioacetazone (T)
- Thioridazine
- Arginine
- Vitamin D
- R207910



Fig 3: Third line antitubercular drugs

VEXATION IN THE WAY OF DEVELOPMENT OF NEW TREATMENT

The biggest conundrums in the advancement of new treatments are:

- a. The crave for novel and ingenious drug regimens,
- b. Novel trial designs,
- c. Studies in paediatric populations,
- d. Amplify clinical trial capacity,
- e. Clear regulatory guidelines,
- f. Biomarkers for prediction of the long-term outcome^{.18}.



CHEMOTHERAPY PROBLEMS AND REQUIREMENTS ¹⁹

Current requirements

- a) Shortening the duration of treatment
- b) New quality drugs or its combination with new agents
- c) Find out new drugs for persistent infection
- d) Economical methods of production of quality drugs
- e) Public private partnership programme.

Limitations of old drugs

- a) Drug resistance (MDR & XDR TB)
- b) TB & HIV synergy
- c) Latent tuberculosis and persistent infection
- d) Problems with DOTS
- e) Long treatment schedule
- f) Paediatric tuberculosis
- g) Tuberculosis in diabetic cases

TREATMENT REGIMENS

Current short-course TB therapy used to treat drug-susceptible *Mycobacterium tuberculosis* consists of 2 months treatment with four drugs so-called first-line drugs including rifampin (RIF), isoniazid (INH), pyrazinamide (PZA), and ethambutol (EMB), followed by 4 months treatment with RIF and INH. Infection by MDR-TB strains requires treatment with second-line drugs such as kanamycin, amikacin, capreomycin, *p*-aminosalicylic acid (PAS), fluoroquinolones (e.g., levofloxacin), ethionamide and cycloserine where treatments often extend for as long as 2 years. All of these drugs are aged and unappealing by today's standards. Many patients fail to complete therapy because of the complicated drug regimen.









Isoniazid

Ethionamide

Pyrazinamide

R = CH3, Rifampicin





Fig 4: Currently used first line antitubercular and second line antitubercular drugs







Fig 5: Molecules under preclinical and clinical trials

TARGETS UNDER STUDY²⁴

Cell wall biosynthesis

a) Peptidoglycan biosynthesis b) Arabinogalactan biosynthesis c) Mycolic acid biosynthesis Amino acid biosynthesis a) Shikimic acid pathway b) Arginine biosynthesis c) Branched amino acid biosynthesis **Cofactor biosynthesis** a) Folic acid biosynthesis b) Pantothenic acid biosynthesis c) Riboflavin biosynthesis d) Reductive sulphur assimilation **Mycothiol biosynthesis Terpenoid biosynthesis DNA synthesis ATP biosynthesis** Menaquinoneoxidoreductase (Type II NADH) inhibition **Tubulin polymerase inhibition**

EXTENSIVELY DRUG RESISTANT TUBERCULOSIS (XDR-TB): A GLOBAL THREAT

The term XDR Tuberculosis, firstly coined by the US Centre for Disease Control and Prevention. It is defined as infection caused by Mycobacterium tuberculosis that is resistant to isoniazid, rifampin any fluoroquinoline and any injectable drug (amino glycosides or polypeptides)^{20, 21, 22.} At the outset XDR-TB was defined as malady caused by *M. tuberculosis* not alone resistant to isoniazid and rifampin but rather to at least 3 of the 6 classes of second-line agents approved for the treatment of TB (amino glycosides, polypeptides, fluoroquinolones, thioamides, cycloserine, and para-aminosalicyclic acid). The World Health Organization Global Task Force on XDR-TB modified the definition in October 2006 following the report of the outbreak in Kwazulu Natal. Since drug susceptibility testing is reliable only for amikacin, kanamycin, capreomycin, and fluoroquinolones among the second-line agents, the definition of extensively drug-resistant TB is now resistance to isoniazid and rifampin in addition to resistance to any fluoroquinolone and any of the second-line injectable drugs: amikacin, kanamycin, and capreomycin²².

Medications Currently Available for the Treatment of Extensively Drug-resistant TB ²³

CATEGORY	AGENT	MECHANISM OF ACTION	TOXICITY AND COMMENTS
Ethambutol		Inhibits arabinosyl	Optic neuritis
		transferase,an important	
		enzyme in the synthesis of	
		Arabinogalactan, a	
		mycobacterium cell wall	
		constituent.	
Pyrazinamide		Inhibits fatty acid	Hyperuricemia, arthralgia, hepatitis
		synthetase I	
Amino	Streptomycin	Misreading of the genetic	Commonest toxicities: nephrotoxicity,
glycosides	Kanamycin	code and Inhibition of	ototoxicity
	Amikacin	translocation by binding to	Amikacin is the preferred drug.Cross-
		the 30S ribosomal bacterial	resistance is common.
		subunit.	
Polypeptides	Capreomycin	Unknown	Nephrotoxicity, ototoxicity Cross-
			resistance with amino glycosides is
			common.
Quinolones	Ofloxacin	Inhibits DNA gyrase	Confusion, seizures, tendinopathy, QT
	Moxifloxacin		prolongation.
	Gatifloxacin		Level of cross resistance with other
			quinolones is uncertain.
Thioamides	Ethionamide	Inhibits synthesis of	Gastrointestinal toxicity, peripheral
	Prothionamide	mycolic acid	neuropathy. Requires use of
			concomitant vitamin B6.
	Para-amino	Competitive antagonism	Gastrointestinal toxicity, haemolytic
	salicylic	with Para amino benzoic	anaemia, especially with G6PD
	Acid	Acid	deficiency.
	Cycloserine	For incorporation into the	Peripheral neuropathy, psychosis,
		bacterial cell wall competes	seizures. Use of concomitant vitamin
		with D-alanine	B6 is required.

(Abbreviation: G6PD_glucose-6-phosphate dehydrogenase)

Agents in Clinical Evaluation ²⁴

DRUG	CHEMICAL CLASS	CELLULAR TARGET	
Gatifloxacin	Fluoroquinolone	DNA gyrase, DNA replication and transcription	
Moxifloxacin	Fluoroquinolone	DNA gyrase, DNA replication and transcription	
Levofloxacin	Fluoroquinoline	uinoline DNA gyrase, DNA replication and transcription	
Linezolid	Oxazolidinones	protein synthesis	
Metronidazole	Nitroimidazole	cytochrome P450a	
TMC207	Diarylquinoline	ATP synthesis	
PA-824 Nitroimadazooxazine mycolic aci protein syn		mycolic acid biosynthesis and protein synthesis	
OPC-67683	Nitroimidazo-Oxazole	cell wall biosynthesis	
LL-3858	Pyrrole	Unknown	
SQ109	2109 Ethylenediamine fatty acid biosynthesis		
SCV-07 Dipeptide none, immunomo		none, immunomodulator	

POTENTIAL DRUG	ACTIVE OR PRO- DRUG	DESCRIPTION	CELLULAR PROCESS INHIBITED	STAGE OF DEVELOPMENT
PA-824	Pro-drug	Nitroimidazo- oxazine	Mycolic acid synthesis	Phase II
OPC-67683	Pro-drug	Nitroimidazo- oxazole	Mycolic acid synthesis	Phase II
R207910	Active	Diarylquinoline	ATP synthesis	Phase II
SQ109	Active	Ethylenediamine derivative	Lipid/cell wall synthesis	Phase I
Compound 5	Pro-drug	Quinoxaline-oxide derivative	Unidentified	Preclinical
Compound 7	Unknown	Quinoline-isoxazole derivative	Unidentified	Preclinical

Tuberculosis drug candidates in development ²⁵

CONCLUSION

Many M. tuberculosis biochemical transformations have been identified as potential targets at the bench level for antituberculosis treatment today. The key issue is to control both drug-susceptible and drug-resistant tuberculosis, which could be successfully achieved through the sound implementation of the *DOTS* strategy and the careful introduction of second-line drugs. To come to grips with TB and with resurgent infectious diseases as a whole, more effective policies are urgently needed. In particular, it is essential to develop active agents that are potentially bactericidal against persistent and dormant organism of *MTB*, in order to shorten the duration of directly observed, short-course of TB patients and to eliminate the reservoir of *MTB*, especially in developing countries. The most urgent goal of chemotherapy for TB is to develop highly active but low-cost drugs that can be used not only in industrialized but also developing countries.

REFERENCES

- Tripathi R.P, Pandey Jyoti, Kukshal Vandana, Arya Ajay, Mishra Mridul, Dubey Divya, Deepti Chopra, Dwivedi R, Chaturvedi Vinita, R. Ravishankar. 2011. Synthesis, in silico screening and bioevaluation of dispiro-cycloalkanones as antitubercular and mycobacterial NAD+-dependent DNA ligase inhibitors. *Med. Chem. Commun.*2: 378–384.
- 2. Spigelman MK. 2007. New Tuberculosis Therapeutics: A Growing Pipeline. *The Journal of Infectious Diseases***196:** S28–S34.
- 3. Jiang J, Chai Ying, Cui Hua. 2011. The electrogenerated chemiluminescence detection of IS6110 of Mycobacterium tuberculosis based on a luminol functionalized gold nano probe. *RSC Adv.*1: 247–254.
- 4. Siawaya J F D. 2007. Correlates for disease progression and prognosis during concurrent HIV/TB infection. *International Journal of Infectious Diseases***11**: 289–299.
- 5. Kaufmann S.H.E. 2007. Tuberculosis and AIDS a devilish liaison. Drug Discovery Today12: 891-892.
- Bogatcheva E , Hanrahan Colleen, Chen Ping, Gearhart Jacqueline, Sacksteder Katherine, Einck Leo, Nacy Carol, Protopopova Marina. 2010. Discovery of dipiperidines as new antitubercular agents. *Bioorg. Med. Chem. Lett.*, 20: 201–205.

- Santos DS, Vasconcelos IB, Meyer E, Sales F, Moreira A.M, Basso I.S. 2008. The Mode of Inhibition of Mycobacterium tuberculosis Wild-Type and Isoniazid-Resistant 2-Trans-Enoyl-ACP(CoA) Reductase Enzymes by An Inorganic Complex. Anti-Infective Agents in Medicinal Chemistry 7:50-62.
- 8. Ahmad S ,Mokkaddas E. 2010. Recent advances in the diagnosis and treatment of mu ltidrug-resistant tuberculosis. *Respiratory Medicine***12:** 1777-90.
- AzizMA, Wright Abigail, Laszlo Adalbert, Muynck A.D, Portaels F, Deun A.V, Wells Charles. 2006. Epidemiology of antituberculosis drug resistance (the Global Project on Anti-tuberculosis Drug Resistance Surveillance): an updated analysis. *The Lancet* 368: 2142-54.
- 10. Tripathi RP, Tewari Neetu, Dwivedi Namrata, Tiwari V.K. 2005. Fighting Tuberculosis: An Old Disease with New Challenges. *Medicinal Research Reviews*, **25**: 93-131.
- 11. Raman K, Bhat A.G, Chandra N. 2010. A systems perspective of host-pathogen interactions: predicting disease outcome in tuberculosis. *Mol. BioSyst.* 6: 516–530.
- 12. Parida SK. 2010. The quest for biomarkers in tuberculosis. Drug Discovery Today 15: 148-157.
- 13. http://www.who.int/mediacentre/factsheets/fs104/en/
- 14. http://www.tuberculosis.net/symptoms-of-tuberculosis.html.
- 15. http://www.nlm.nih.gov/medlineplus/ency/article/000077.html.
- 16. Waksman S. A. 1969. Successes and failures in the search for antibiotics. Ad. Appl. Microbiol.11: 1-16.
- 17. Zhang Y,Martens K.P, Denkin Steven. 2006. New drug candidates and therapeutic targets for tuberculosis therapy. *DDT***11:** .
- 18. Ashforth EJ, Chengzhang Fu, Xiangyang Liu, Huanqin Dai, Fuhang Song, HuiGuo and Lixin Zhang. 2010. Bioprospecting for antituberculosis leads from microbial metabolites. *Nat. Prod. Rep.* **27:** 1709–1719.
- 19. Frieden TR ,Driver C.R. 2003. Tuberculosis control: past 10 years and future progress. *Tuberculosis*83: 82 -85.
- 20. McCarthy M. 2002. Tuberculosis experts review 10 years of progress. The Lancet Infectious Diseases2: 387.
- 21. Belisle J. T, Vissa V.D Brennan P.J Besra G.S. 1997. Role of the major antigen of Mycobacterium tuberculosis in cell wall biogenesis. *Science* **276**: 1420-1422.
- 22. Frieden TR, Sterling TR, Munsiff SS, Watt CJ, Dye C. 2003. Tuberculosis. *Lancet* **363**: 887–899.
- 23. JaninYL. 2007. Antituberculosis drugs: ten years of research. *Bioorg Med Chem* **15**: 2479–2513.
- 24. Rivers EC, Mancera RL. 2008. New anti-tuberculosis drugs with novel mechanisms of action. *Curr Med Chem***15**: 1956–1967.
- McLean KJ, Marshall K R, Richmond A, Hunter IS, Fowler K, Kieser T, Gurcha SS, Besra GS, Munro AW. 2002. Azole antifungals are potent inhibitors of cytochrome P450 mono-oxygenases and bacterial growth in mycobacteria and streptomycetes. *Microbiology* **148**: 2937–2949.