

Mini Review





Pyrazinamide drug resistance in *Mycobacterium tuberculosis*: a mini review

Abstract

Pyrazinamide (PZA) is a nicotinamide analog that is used as a frontline drug to treat tuberculosis (TB). It has special place in modern TB therapy, as it appears to kill a population of semi dormant tubercle bacilli persisting in the body. It is important drug because of its sterilizing activity against semi dormant tubercle bacilli. Although PZA has a very high in vivo activity, it's in vitro activity is not apparent unless an acidic environment is available, which makes PZA drug susceptibility testing (DST) more difficult by conventional methods. Being a nicotinamide analogue, PZA needs to be converted into its active bactericidal form pyrazinoic acid (POA) by the mycobacterial enzyme pyrazinamidase (PZase). PZase is encoded by pncA, and mutations in this gene have been demonstrated as the major mechanism of PZA resistance. PZA-resistant Mycobacterium tuberculosis strains are usually correlated well with defective PZase activity. The rapid detection of PZA resistance is of utmost importance for an effective treatment and also to avoid further spread of MDR strains. Analysis of the pncA gene provides rapid and useful information regarding PZA susceptibility in M. tuberculosis and may therefore contribute to early optimization of treatment. The diversity of methods currently used in clinical laboratories for the detection of PZA resistance in M. tuberculosis isolates causes inconsistent results of PZA DST and such inconsistent results of PZA DST have been reported by a number of laboratories by various methods, including the qualitative BACTEC test. Therefore, it is essential to elucidate the genetic basis of clinical resistance and to correlate phenotypic and molecular resistance data.

Keywords: pyrazinamide resistance, *mycobacterium tuberculosis*, qualitative bactec test, rifampicin, isoniazid

Abbreviations: TB, tuberculosis; PZA, pyrazinamide; DST, drug susceptibility testing; POA, pyrazinoic acid; MDR, multi drug-resistant

Introduction

The addition of PZA with isoniazid and rifampicin, forms the cornerstone of modern TB therapy, shortening the TB therapy from previously 9-12months to current 6months. It kills a semi-dormant tubercle bacilli population in acidic pH environments that are not killed by any other anti- TB drugs.1 Though PZA has a remarkable role in shortening the treatment duration, it remains a difficult paradox because of its incompletely understood mode of action as well as mechanism of resistance.² Basically PZA is bacteriostatic, but can become bactericidal on actively replicating tubercle bacteria.³ It is a nicotinamide analogue, a prodrug which gets converted into the active bactericidal form pyrazinoic acid (POA) by the mycobacterial enzyme pyrazinamidase (PZase).² However, an exact mechanism of its action is unknown. It has been suggested that the accumulation of POA in acidic conditions (from lactic acid produced by inflammation of cells) leads to acidification of the cytoplasm and results in cell death.⁴ PZase is encoded by pncA, and mutations in this gene have been demonstrated as the major mechanism of PZA resistance. Several mutations, including missense, insertion, deletion and nonsense mutations have been reported and located in both the putative promoter and coding regions of pncA.5,6 PZA-resistant Mycobacterium tuberculosis strains are usually correlated well with defective PZase activity, but some

Volume 3 Issue 2 - 2016

Vaishali R Wabale, Ameeta A Joshi

Department of Microbiology, Grant Government Medical College, India

Correspondence: Vaishali R Wabale, Assistant Professor, Department of Microbiology, Grant Government Medical College & Sir J.J. Group of Hospitals, Byculla, Mumbai, Maharashtra, India, Tel 8828050378, Email vrwabale@gmail.com

Received: October 23, 2016 | Published: November 25, 2016

PZA resistant strains have been reported to contain wild-type and to maintain PZase activity suggesting that there might be other unknown resistance mechanisms could be responsible for the resistance phenotype.^{6–8}

Anti-tuberculosis drugs are known as two-edged sword. Although, these drugs are capable to destroy pathogenic M. tuberculosis, on the other hand, they also go for selection of drug resistant tubercle bacteria against which those drugs are ineffective. Tuberculosis (TB) global surveillance has shown that such drug resistant TB is widespread and is an alarming threat to TB control programmes in many countries.9 Although consistent effort in monitoring and TB treatment is going on still the disease remains as a major public health issue.¹⁰ In year 1982, the World Health Organization (WHO) estimated worldwide every year occurrence around 4-5million new highly infectious, smear-positive cases of pulmonary infection with M. tuberculosis. PZA was synthesized way before 1940, though its antituberculous activity was not recognized until the early 1950s.11 WHO estimated around 8.8million new cases of TB globally in year 2003 within an expected 1% increase every year. However, appropriate therapies and rapid diagnosis become the first priorities in controlling the growing TB epidemics.¹⁰ Even, after decades of decline, rates of TB cases are still increasing worldwide. Today TB is the most frequent cause of death from a single infectious disease in persons aged 15-49years, causing a total of 2-3million deaths annually. One- third of the human population is thought to be infected by the M. tuberculosis, and about 200million additional people are at risk of developing disease

J Bacteriol Mycol Open Access. 2016;3(2):203-205.



© 2016 Wabale et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and build upon your work non-commercially.

in the next 20years, as emerging coinfection of TB with human immunodeficiency virus (HIV) and resistance of M. tuberculosis to the presently used TB regimen.¹² In both low- and high- income countries Multidrug-resistant (MDR) strains of M. tuberculosis have been emerging worldwide.13,14 Since long PZA has been known to be active against pulmonary TB in humans and is used as a first-line antituberculosis drug in routine TB therapy. It's mechanism of action is least understood. Because of its outstanding sterilizing activity against semi dormant tubercle bacilli, it plays an important role in the treatment of both drug-susceptible and drug-resistant TB. The MDR-TB inclusive of PZA drug resistance definitely worsen prognoses, and the rising prevalence of such MDR-TB and HIV- MDR -TB coinfection is alarming.4,15,16 The need for rapid methods of the diagnosis and determination of drug susceptibility testing (DST) is very important. While the procedures for DST of the most of the first-line and second-line antituberculosis drugs have been well standardized in both liquid and solid media, the main problem comes with the PZA DST, is the requirement of acidic pH for PZA activity.¹⁷ It also becomes difficult because of the acidic pH of culture medium restricts the growth of organism. In addition, the use of large inoculum sizes results in the release of NH3 that leads to increased pH and inactivated PZA.¹⁵ The application of various rapid phenotypic methods such as radiometric BACTEC 460, flourimetric Mycobacteria Growth Indicator Tube (MGIT) 960 (Becton Dickinson), ESP Culture System II (Trek Diagnostic Systems, West-lake, OH), and the colorimetric BacT/ALERT 3D system (bioMerieux Inc., Durham, NC), previously designated MB/BACT (Organon Teknika, Boxtels, The Netherlands) has been reported to be very useful for rapid and reliable susceptibility testing of *M. tuberculosis*isolates.¹⁷ Presently many laboratories performing PZA liquid DST by the non-radiometric, fully automated, continuous-monitoring MGIT 960 system (Becton Dickinson) that gives reliable result with the turnaround time of 15-22days.^{18,19} Number of studies revealed a good correlation between loss of PZase activity and resistance to PZA.4,6,20,21 However, the rapid detection of PZA resistance is of utmost importance for an effective treatment and also to avoid further spread of MDR strains. Molecular assays those detect the genetic variants that mediate resistance constitute a rapid alternative to conventional DST and may even be performed directly on clinical specimens without culture. Therefore, it is essential to elucidate the genetic basis of clinical resistance and to correlate phenotypic and molecular resistance data. It is considered that the analysis of the pncA gene provides rapid and useful information regarding PZA susceptibility in M. tuberculosis and may therefore contribute to early optimization of treatment. The most PZA-resistant M. tuberculosis strains have mutations in the pncA gene has implications for developing a rapid test for detecting PZAresistant *M. tuberculosis* strains.²¹ The diversity of methods currently used in clinical laboratories for the detection of PZA resistance in M. tuberculosis isolates causes inconsistent results of PZA DST.22 Inconsistent results of PZA DST have been reported by a number of laboratories by various methods, including the qualitative BACTEC test.23

Almost all PZA-resistant *M. tuberculosis* strains have mutations in the *pncA* gene points the necessity for developing a rapid test for detecting such PZA-resistant *M. tuberculosis* strains globally.

Acknowledgements

None.

Conflict of interest

The author declares no conflict of interest.

References

- Zhang Y, Wade MM, Scorpio A, et al. Mode of action of pyrazinamide: disruption of *Mycobacterium tuberculosis* membrane transport and energetics by pyrazinoic acid. *J Antimicrob Chemother*. 2003;52(5):790– 795.
- Singh P, Mishra AK, Malonia SK, et al. The paradox of pyrazinamide: An update on the molecular mechanisms of pyrazinamide resistance in Mycobacteria. *J Communc Dis.* 2006;38(3):288–298.
- 3. http://en.wikipedia.org/wiki/Pyrazinamide.
- Yeager RL, Munroe WG, Dessau FI. Pyrazinamide (aldinamide) in the treatment of pulmonary tuberculosis. *Am Rev Tuberc*. 1952;65(5):523– 546.
- Ando H, Mitarai S, Kondo Y, et al. Pyrazinamide resistance in multidrug– resistant *Mycobacterium tuberculosis* isolates in Japan. *Clin Microbiol Infect*. 2010;16(8):1164–1168.
- Scorpio A, Zhang Y. Mutations in *pncA*, a gene encoding pyrazinamidase/ nicotinamidase, cause resistance to the antituberculous drug pyrazinamide in tubercle *bacillus*. *Nat Med.* 1996;2(6):662–667.
- Singh R, Manjunatha U, Boshoff HI, et al. PA-824 kills non-replicating *Mycobacterium tuberculosis* by intracellular NO release. *Science*. 2008;322(5906):1392–1395.
- Mestdagh M, Fonteyne PA, Realini L, et al. Relationship between pyrazinamide resistance, loss of pyrazinamidase activity, and mutations in the *pncA* locus in multidrug–resistant clinical isolates of *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother*. 1999;43(9):2317–2319.
- Johnson R, Streicher EM, Louw GE, et al. Drug Resistance in Mycobacterium tuberculosis. Curr Issues Mol Biol. 2006;8(2):97–111.
- Yam WC. The HONG KONG Medical Diary. *Medical Bulletin*. 2006;2:6–7.
- Max Salfinger, Alfred JC, Barth RL. Pyrazinamide and pyrazinoic acid activity against tubercle bacilli in cultured human macrophages and in the BACTEC system. *The Journal of Infectious Diseases*. 1990;162(1):201– 207.
- Van Soolingen D. Molecular epidemiology of tuberculosis and other mycobacterial infections: main methodologies and achievements. *Journal of Internal Medicine*. 2001;249(1):1–26.
- McCune RM, Tompsett R, McDermott W. The fate of *Mycobacterium tuberculosis*in mouse tissues as determined by the microbial enumeration technique II. The conversion of tuberculous infection to the latent state by administration of pyrazinamide and a companion drug. *J Exp Med.* 1956;104:763–802.
- Mitchison DA. The action of anti– tuberculosis drugs in short course chemotherapy. *Tubercle*. 1985;66:219–225.
- Zhang Y, Wade MM, Scorpio A, et al. Mode of action of pyrazinamide: disruption of *Mycobacterium tuberculosis* membrane transport and energetics by pyrazinoic acid. *J Antimicrob Chemother*. 2003;52(5):790– 795.
- Miller MA, Thibert L, Desjardins F, et al. Testing of susceptibility of *Mycobacterium tuberculosis* to pyrazinamide: comparison of BACTEC method with pyrazinamidase assay. J Clin Microbiol. 1995;33(9):2468– 2470.
- Singh P, Wesley C, Jadaun GPS, et al. Comparative evaluation of Lowenstein–Jensen proportion method, BacT/ALERT 3D system, and enzymatic pyrazinamidase assay for pyrazinamide susceptibility testing of *Mycobacterium tuberculosis*. J Clin Microbiol. 2007;45(1):76–80.
- Scarparo C, Ricordi P, Ruggiero G, et al. Evaluation of the fully automated BACTEC MGIT 960 system for testing susceptibility of *Mycobacterium tuberculosis* to pyrazinamide, streptomycin, isoniazid, rifampin, and ethambutol and comparison with the radiometric BACTEC 460 TB method. *J Clin Microbiol.* 2004;42(3):1109–1114.

Citation: Wabale VR, Joshi AA. Pyrazinamide drug resistance in *Mycobacterium tuberculosis*: a mini review. J Bacteriol Mycol Open Access. 2016;3(2):203–205. DOI: 10.15406/jbmoa.2016.03.00054

- Pfyffer GE, Palicova F, Rüsch Gerdes S. Testing of susceptibility of Mycobacterium tuberculosis to pyrazinamide with the non-radiometric BACTEC MGIT 960 System. J Clin Microbiol. 2002;40(5):1670–1674.
- Davies AP, Billington OJ, McHugh TD, et al. Comparison of phenotypic and genotypic methods for pyrazinamide susceptibility testing with *Mycobacterium tuberculosis. J Clin Microbiol.* 2000;38(10):3686–3688.
- Wabale VR, Joshi AA, Muthaiah M, et al. *pncA* gene sequence analysis for pyrazinamide resistance in *Mycobacterium tuberculosis* from highincidence setting. *Int J Pharm Bio Sci.* 2016;7(4):B648–B654.
- 22. Scorpio A, Lindholm Levy P, Heifets L, et al. Characterization of *pncA* mutations in pyrazinamide–resistant *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother*. 1997;41(3):540–543.
- 23. Jureen P, Werngren J, Toro JC, et al. Pyrazinamide resistance and *pncA* gene mutations in *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother*. 2008;52(5):1852–1854.