Minireview

Metabolism and hepatotoxicity of pyrazinamide, an anti-tuberculosis drug

Zahir Hussain, Junjie Zhu, and Xiaochao Ma

Center for Pharmacogenetics, Department of Pharmaceutical Sciences, School of Pharmacy,
University of Pittsburgh, Pittsburgh, Pennsylvania, USA (Z.H., J.Z., X.M.)

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Corresponding author: Dr. Xiaochao Ma, Department of Pharmaceutical Sciences, School of

Pharmacy, University of Pittsburgh; 309 Salk Pavilion, 335 Sutherland Drive, Pittsburgh, PA

15261. Phone: 412-648-9448. Email: mxiaocha@pitt.edu

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Abbreviation list

5-OH-PA, 5-hydroxy-pyrazinoic acid; 5-OH-PZA, 5-hydroxy-pyrazinamide; AO, aldehyde oxidase; BNPP, bis-*p*-nitrophenyl phosphate; FXR, farnesoid X receptor; IFN, interferon; INH, isoniazid; PA, pyrazinoic acid; PXR, pregnane X receptor; PZA, pyrazinamide; RIF, rifampicin; ROS, reactive oxygen species; TB, tuberculosis; XDH, xanthine dehydrogenase; XO, xanthine oxidase

Abstract

Pyrazinamide (PZA) is an important component of a standard combination therapy against tuberculosis. However, PZA is hepatotoxic and the underlying mechanisms are poorly understood. Biotransformation of PZA in the liver was primarily suggested behind its hepatoxicity. This review summarizes the knowledge of the key enzymes involved in PZA metabolism and discusses their contributions to PZA hepatotoxicity.

Significance statement

This review outlines the current understanding of PZA metabolism and hepatotoxicity. This work also highlights the gaps in this field, which can be used to guide the future studies on PZA-induced liver injury.

1. Introduction

Tuberculosis (TB), mostly caused by *Mycobacterium tuberculosis*, is a leading infectious disease killing around 4000 people worldwide daily (WHO, 2020). Pyrazinamide (PZA), first synthesized in 1936 as a structural analog of nicotinamide, made its way into clinical use against TB since 1952. PZA has still been in use till today as an essential element of a standard anti-TB combination therapy together with isoniazid (INH), rifampicin (RIF), ethambutol or streptomycin (Stout, 2004; WHO, 2020). In the recommended 6-month anti-TB regimen by WHO, PZA is used in the intensive phase of 2 months (**Table 1**). Same as INH and RIF, PZA is on the WHO's list of essential medicines for TB treatment.

Despite the important contribution of PZA to anti-TB success for past seven decades, it can cause liver injury and even liver failure (LiverTox, 2020). In comparison with other potential hepatotoxic anti-TB drugs such as INH and RIF, PZA was initially reported safe (Girling, 1982). However, according to recent studies, PZA turns out to be more hepatotoxic than previously considered and could be more hepatotoxic than either INH or RIF (Schaberg et al., 1996; Yee et al., 2003; Chang et al., 2007; Tostmann et al., 2008). Unfortunately, very little is known about the mechanisms of PZA hepatotoxicity. One reason behind this is that PZA is often used together with other anti-TB drugs which are also hepatotoxic making it difficult to distinguish its individual contribution to liver damage (Table 1). Evidence from animal studies and clinical trials suggests that metabolites derived from PZA biotransformation are related to its hepatoxicity (Kudo et al., 2008; Shih et al., 2013; Rawat et al., 2018). In this review, we summarize the major enzymes involved in PZA metabolism and discuss the current understanding of PZA hepatotoxicity.

2. Metabolism and disposition of PZA

The oral dosage of PZA for adults is 20–30 mg/kg/day (NCBI, 2021). It is well absorbed from the gastrointestinal tract and widely distributed in the body including the liver. PZA is mainly metabolized in the liver by amidase, which converts PZA to pyrazinoic acid (PA) (Aoki et al., 1957). PA can be further oxidized by xanthine oxidase (XO) to form 5-hydroxy-pyrazinoic acid (5-OH-PA). Alternatively, PZA can first be oxidized to 5-hydroxy-pyrazinamide (5-OH-PZA) by XO followed by amidase-mediated hydrolysis to form 5-OH-PA (**Figure 1**). Furthermore, PA can be conjugated with glycine to form trace amounts of pyrazinuric acid (Lacroix et al., 1989). PZA and its metabolites are predominantly excreted by the kidney. Within 36 hours, ~70% of an oral dose of PZA is excreted in urine, and the relative abundances are as follows: PA (36%), 5-OH-PZA (15.4%), 5-OH-PA (13.8%), and PZA (3.8%) (Lacroix et al., 1988).

In subjects with normal renal and hepatic functions, the plasma half-life of PZA is 9.6 h (Lacroix et al., 1989). The half-life of PZA is significantly prolonged in patients with pre-existing liver or kidney diseases. In subjects with insufficient hepatic functions, a marked reduction of PZA clearance was observed and the half-life of PZA was increased to 15.07 h. In addition, the clearance rate of PA, a major and active metabolite of PZA, was also significantly decreased in patients with hepatic insufficiency (Lacroix et al., 1990). In chronic uremic patients, bioavailability of PZA was slightly increased, but bioavailability of PA was markedly

increased (Stamatakis et al., 1988). These data suggest that a reduction of PZA dosage is needed in patients with hepatic or renal insufficiency.

2.1. Amidase and its role in PZA metabolism

Amidase, classified as EC 3.5.1.4, has several aliases such as acylamide amidohydrolase (systematic name), acylamidase, fatty acylamidase, acylase (though misleading) as well as some ambiguous names including amidohydrolase, deaminase, and *N*-acetylaminohydrolase (IUBMB Enzyme Nomenclature). Enzymes of this group not only hydrolyze amide compounds but can also hydrolyze carboxyl esters in several cases and share a similar catalytic mechanism with esterases (Wang, 1994; Wang et al., 2016b). Amidases exhibit broad substrate specificity and have diverse biological roles in mammals including control of pain and neuromodulation by fatty acid amide hydrolase as well as regulation of inflammation by *N*-acylethanolamine-hydrolyzing acid amidase (Cravatt et al., 1996; Tsuboi et al., 2005).

A hepatic amidase has been proposed to metabolize PZA to produce PA and ammonia (Aoki et al., 1957). Later, the activity of PZA amidase was found in various tissues of mice, rats, guinea pigs, and rabbits (Toida, 1973). The activity of PZA amidase was the highest in the liver of all tested tissues, and the rabbit liver showed an outstandingly high activity of PZA amidase among these tested species. In addition, the PZA amidase was mostly localized in the microsomal fraction of the liver (Toida, 1973). Despite significant importance of hepatic microsomal amidase in PZA metabolism, the genetic identity of this enzyme is still unknown. Nicotinamidase/pyrazinamidase (EC 3.5.1.19, PNC1, I6XD65) is responsible for bacterial PZA

metabolism (French et al., 2010). NCBI blast search with the amino acid sequence of mycobacterial pyrazinamidase showed 40% homology to yeast nicotinamidase and 27% homology to *A. Thaliana* nicotinamidase, while no homology was found to any major mammalian proteins. Thus, discovering the mammalian amidase responsible for PZA metabolism remains a great challenge for the researchers in this field.

2.2. XO and its role in PZA metabolism

XO (EC 1.17.3.2) is expressed in almost all tissues with high levels in the intestine and the liver (Harrison, 2004). XO oxidizes hypoxanthine/xanthine to uric acid (Wang et al., 2016a). In addition, XO hydroxylates heterocyclic compounds including PZA and PA to produce 5-OH-PZA and 5-OH-PA, respectively (**Figure 1**). In mammals, XO predominantly exists as xanthine dehydrogenase (XDH, EC 1.17.1.4) which essentially acts on same substrates of XO but can use either NAD⁺ or O₂ as an electron acceptor while XO can only use O₂ as an electron acceptor resulting in reactive oxygen species (ROS) formation. The conversion from XDH to XO proceeds either reversibly by the oxidation of its certain cysteine thiols to form cystine disulfide bonds or irreversibly by specific proteolysis. Despite these differences between the two forms of this enzyme, the term XO was often used as a general name for the both throughout the literatures (Harrison, 2004).

Differences in XO activity among individuals and various ethnicities as evaluated by caffeine metabolic rate suggest interesting polymorphic behavior of this enzyme. A low activity of liver XO was observed in 20% Caucasian as well as in 11% Japanese (Kudo et al., 2008). In

addition, males showed higher XO activities than females. Classical xanthinuria type 1 is a rare autosomal recessive disorder, caused by Arg149Cys substitution resulting in loss of activity of XO leading to chronic renal failure (Kudo et al., 2008). On the contrary, two different point mutations, Ile703Val or His1221Arg, were reported to increase XO activity by around two-fold in a study conducted in Japanese population (Kudo et al., 2008). However, limited information is available for the associations of XO polymorphisms with PZA metabolism. Furthermore, enhanced XO expression and activity were found in mouse liver and other tissues treated with interferon (IFN) and IFN-inducers (Ghezzi et al., 1984). Human XO gene was also reported to contain IFN-gamma response elements (Xu et al., 1996). Therefore, further studies are needed to determine the potential impact of XO polymorphisms and/or induction on PZA metabolism.

It is here worthwhile to mention that rabbits do not express XO and thereby do not respond to allopurinol, a XO inhibitor, which explained the unique pharmacokinetics and high exposure of PA in rabbits when compared to mice and other mammals (Via et al., 2014). This fact is consistent with previous studies in human volunteers who showed increased levels of PA and decreased levels of 5-OH-PZA as well as 5-OH-PA when co-treated with PZA and allopurinol (Lacroix et al., 1988; Naftalin et al., 2017). Allopurinol is commonly used to treat hyperuricemia or gout but is not suggested to treat PZA-induced hyperuricemia, a common side effect of PZA, which occurs due to inhibition of renal excretion of uric acid by PA and thus counterbalancing the effect of allopurinol (Lacroix et al., 1988).

The ability of xanthinuria patients caused by XO deficiency to oxidize PZA raised the possibility of the existence of an alternative enzyme other than classical XO in PZA metabolism. Moriwaki *et al* showed that aldehyde oxidase (AO, EC 1.2.3.1) could potentially convert PZA to 5-OH-PZA but not PA to 5-OH-PA (**Figure 1**). In addition, the K_m value of XO for PZA was about 10 times higher than that of AO (Moriwaki et al., 1993). These results indicate that AO is catalytically distinct from XO. However, the relative contribution of AO to PZA metabolism in mammals was not investigated so far.

3. Mechanisms of PZA-induced hepatotoxicity

The contribution of PZA to liver damage during anti-TB therapy is not fully clear because PZA is used only in combination with other anti-TB drugs that are hepatotoxic, such as INH and RIF. However, mounting evidence supports that PZA is hepatotoxic. For example, the use of combination therapy with RIF and PZA for latent TB was abandoned because of the frequency of severe liver injury (LiverTox, 2020). The pattern of PZA hepatotoxicity is typically acute hepatitis with hepatocellular necrosis, inflammation, and variable degrees of cholestasis, which negatively impacts the outcomes of anti-TB therapy (LiverTox, 2020). Unfortunately, no approach is currently available to predict and prevent PZA hepatotoxicity because its mechanisms are poorly understood, especially when compared to that of INH and RIF.

PZA hepatotoxicity is dose-dependent especially at daily doses above 40 mg/kg, and the extent of PZA hepatotoxicity is correlated with its hepatic metabolism, suggesting a direct toxic effect, but not a hypersensitive or immune-mediated effect (Tostmann et al., 2008; Shih et al.,

2013). Recent *in vitro* and *in vivo* reports suggest that amidase-mediated production of PA from PZA was responsible for PZA hepatotoxicity (**Figure 1**). Experiments with Wistar rats treated with PZA or PA showed hepatotoxicity as observed in elevated serum levels of alanine aminotransferase, aspartate transaminase, and galactose single point (Shih et al., 2013). Amidase inhibitor, bis-*p*-nitrophenyl phosphate (BNPP), decreased PZA-induced, but not PA-induced, hepatotoxicity, suggesting amidase as the initiator of PZA hepatotoxicity (Shih et al., 2013). Consistently, PA levels in the urine were highly correlated with PZA hepatotoxicity in TB patients (Shih et al., 2013). However, because the genetic identity of PZA amidase remains unknown, it is difficult to investigate the role of PZA amidase in PZA hepatotoxicity using genetic approaches.

PA can be further metabolized by XO to produce 5-OH-PA that was suggested to be more hepatotoxic than PA (**Figure 1**) (Shih et al., 2013; Rawat et al., 2018). Shih *et al* treated HepG2 cells with PZA and its metabolites and found 5-OH-PA being the most toxic metabolite of PZA (Shih et al., 2013). A recent study in rats showed that 5-OH-PA caused liver damage accompanied with aberrant metabolic shifts (Rawat et al., 2018). In addition, the patients with severe hepatotoxicity showed much higher ratio of 5-OH-PA to PZA in the urine than other patients with mild or no hepatotoxicity (Shih et al., 2013; Rawat et al., 2018). These data indicate that XO may play an important role in PZA hepatotoxicity by producing 5-OH-PA. However, inhibition of XO by allopurinol increased PZA toxicity in HepG2 cells, suggesting that the hydroxy metabolites of PZA and/or PA, products of XO, were not responsible for PZA hepatotoxicity (Tostmann et al., 2010). With these controversial data, further studies are needed to determine the role of XO in PZA hepatotoxicity.

Physiological significances of XO-catalyzed reactions are to increase hydrophilicity of purine catabolic end products by forming uric acid to be excreted through urine. XO-null mice developed renal interstitial fibrosis through aberrant lipid and purine accumulation in renal tubules resulting in premature death (Ohtsubo et al., 2009). In a recent study, hepatocyte-specific ablation of XO in mice was found to correct obesity-induced systemic hyperuricemia despite other metabolic abnormalities being unchanged (Harmon et al., 2019). In addition, ROS produced as a by-product of XO, especially in immune cells, are used to kill microbes as a part of innate immunity of the host. However, due to be responsible for ROS formation, XO is also implicated to several pathophysiological processes such as oxidative stress, hypertension, and ischemia reperfusion injury (Wang et al., 2016a). Therefore, it is worthwhile to investigate the contributions of XO-mediated production of ROS as well as 5-OH-PA to PZA hepatotoxicity. In addition, further studies are also suggested to explore whether PZA and/or PA disrupt XO-dependent metabolism of endobiotics in the liver and result in liver dysfunction.

Genetic studies have been conducted to explore the impact of PZA on the liver. In rats treated with PZA for 28 days, it was found that PZA upregulates cytochrome P450 2b1, epoxide hydrolase 1, and heme oxygenase, and down-regulates two peroxisome proliferator activated receptor (PPAR)-dependent genes including carnitine palmitoyltransferase 1b and fatty acid binding protein 7 (Zhang et al., 2013). In a follow-up study using the same model, PPARα expression was shown to be inversely correlated with PZA-induced liver injury (Zhang et al., 2016). However, in a very recent study conducted in TB patients with or without liver injury taking standard anti-TB drug regimen, no association was found between various single

nucleotide polymorphisms in PPARα gene and liver injury (Zhang et al., 2020). Interestingly, two polymorphic variants in pregnane X receptor (PXR), a nuclear transcription factor known to regulate the expression of various drug metabolizing enzymes, were associated with the decreased risk of anti-TB drug-induced hepatotoxicity, suggesting that drug-metabolizing enzymes regulated by PXR are involved in the hepatotoxicity of the standard anti-TB drug regimen (Wang et al., 2019).

Since some case reports of PZA-induced hepatotoxicity showed evidence of cholestasis, a recent study investigated its mechanisms in rats (Guo et al., 2016). When rats were orally treated with PZA (2g/kg/day) for 1-week, total bile acids increased 10-fold in the serum while ALT and AST increased 2-fold. The farnesoid X receptor (FXR), a bile acid-responsive transcription factor, which plays a key role in the regulation of bile acid synthesis, excretion and transport, was found to be downregulated. Interestingly, treatment with the FXR agonist obeticholic acid attenuated PZA hepatotoxicity, suggesting that PZA-induced cholestatic liver injury was related to FXR suppression (Guo et al., 2016).

4. Conclusion

Metabolism of PZA, a first-line anti-TB drug, is worth interesting in clinical context because increased PZA metabolites such as PA and 5-OH-PA were found to be highly correlated to the extent of hepatotoxicity. Out of two major enzymes metabolizing PZA, amidase catalyzed the production of PA, but the molecular identity of amidase is unknown. In addition, XO is suggested to play an important role in PZA hepatotoxicity by producing 5-OH-PA. However, the

preclinical data of 5-OH-PA- and/or PA-mediated hepatotoxicity are not conclusive. Thus, more research is needed to elucidate the metabolic pathways of PZA and determine their contributions to PZA hepatotoxicity.

Authorship Contributions

Writing the manuscript: Z.H., J.Z, X.M.

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Footnotes

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Figure legends

Figure 1. Metabolic map of PZA and proposed mechanisms for hepatotoxicity. Amidase is the primarily enzyme in PZA metabolism that produces PA. PA can be further metabolized by XO to produce 5-OH-PA. 5-OH-PA can also be produced from PZA through XO and AO-mediated oxidation followed by amidase-mediated hydrolysis. PA and 5-OH-PA are proposed as the causes of PZA hepatotoxicity.

Table 1. The preferred regimen for TB treatment in adults and the risk of drug-induced liver injury (DILI) from each anti-TB drug.

	mg/kg/day	Treatment period	DILI score*
Rifampin (RIF)	8–12	6 months	A
Isoniazid (INH)	4–6	6 months	A
Pyrazinamide (PZA)	20–30	first 2 months	A
Ethambutol (EMB)	15–25	first 2 months	С

^{*}DILI scores were adapted from LiverTox (<u>LiverTox - NCBI Bookshelf (nih.gov)</u>). A, well established cause of clinically apparent liver injury; C, probable cause of clinically apparent liver injury.

