

Effect of Oxidative Stress on AhpC Activity and Virulence in *katG* Ser315 Thr *Mycobacterium tuberculosis* Mutant.

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Abstract

Mycobacterium tuberculosis strains resistance to INH is mainly caused by the alteration in several genes encoding the molecular targets. Mutation of *katG* at codon 315 especially Ser315Thr are responsible for INH resistance in a large proportion of TB cases. The aim of this study is to evaluate the influence of stress oxidative on AhpC activity of *katG* Ser315Thr of *M.tuberculosis*, and to find out the relation of AhpC and the virulence of this mutant. The study design was laboratoric experimental, subjects of study were *M.tuberculosis* INH resistance strains, and the treatment were serial dose of H₂O₂. Eighty five *M.tuberculosis* INH resistant clinical strain were screened for mutation of *katG*Ser315Thr by PCR/RFLP and characterized on the basis of phenotypic properties (catalase activity and AhpC activity). AhpC activity of *katG* Ser315Thr *M.tuberculosis* strains in response to oxidative stress condition was evaluated by culturing the strains on liquid culture medium containing 1mM H₂O₂. To ascertain role of AhpC in the virulence of *katG*Ser315Thr mutant strains, the mutants were infected into human macrophages culture, and several indicator of virulence were observed (i.e: replication competence, and apoptosis induction on human macrophages). The results showed that *katG* Ser 315Thr were identified in 23 (27,05%) of 85 INH resistance strains, all mutant strains had decrease of catalase activity. AhpC activity of *katG* Ser315Thr of *M.tuberculosis* increased significantly with increase of hydrogen peroxide dose. In addition, it has been shown that increased AhpC activity related to replication ability of mutant, and reduction of apoptosis macrophages induction significantly. We conclude that the production of AhpC of *katG* Ser315Thr *M.tuberculosis* induced by oxidative stress. There was a role of AhpC in virulence of the *M.tuberculosis katG* Ser315Thr strains by replication capability and macrophages apoptosis.

Keywords : *katG* Ser315Thr *Mycobacterium tuberculosis*- oxidative stress - AhpC - virulence

Introduction

Tuberculosis (TB) is a chronic infectious disease in human caused by *Mycobacterium tuberculosis* (*M.tuberculosis*) and a leading cause of death in community. The disease is still a major health problem in the world, even though ten years ago, in 1993, WHO

declared TB as a global emergency (Zhang *et al.*, 2005). The emergence of HIV and low socioeconomic status has greatly influenced the development of the disease. In most developing countries, TB is still continuing problem, because it is an endemic disease which is worsened by HIV pandemic and social changes due to industrialization (Rattan *et al.*, 1999). Another obstacle in overcoming TB problem is the emergence of *M.tuberculosis* mutant that is resistant to anti TB drugs. The emergence of resistant strain which is caused

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by inadequate treatment induce changes in bacterial properties which in turn become resistant to anti TB drugs. The changes include multi drug resistant (MDR), which is resistant to isoniazid and rifampicin, and extreme drug resistant (XDR) which is MDR plus resistant to one of fluoroquinolones, and one or more of injectable drugs, such as kanamycin, amikacin, capreomycin (WHO, 2009). Tuberculosis treatment began in 1944, following the discovery of streptomycin and para-aminosalicylic acid (PAS), and isoniazid, and other drugs (rifampicin and pirazinamid). Soon after the use of first anti TB drug, strains of resistant bacteria appeared. Previously, there was a hope that TB could be eradicated at the end of 20th century, but in fact the disease re-emerge in 1980 with increase of incidence and appearance of multi drug resistant of *M.tuberculosis* strain (da Silva and Ainsa, 2007).

The search for anti TB drugs has been undergoing slow moving phase. Commercially available anti TB drugs have been used for 40 years. Drug resistant *M.tuberculosis* strain has emerged since many years, however, until recently new and potent anti TB drug have not yet discovered. Slow development of anti TB drug is presumably caused by uncertainty in virulence and survival mechanism of *M.tuberculosis*. Besides, knowledge of anti TB drugs resistance mechanism is still limited. The rate of resistance is in imbalance with discovery of new drugs. The search of new and effective drugs is crucial, therefore it is necessary to study bacterial components which become target of anti TB drugs.

Several mycobacterial enzymes necessary for intracellular survivor include *catalase-oxidase*, *alkyl hydroperoxide reductase*, *enoyl-ACP (acyl carrier protein) reductase*, β -ketoacyl-ACP synthetase. Synthesis of these enzymes is coded by *katG*, *ahpC*, *inhA* and *kasA*, respectively, in which they could be target of anti TB drug (Musser, 1995; Pym *et al.*, 2002; Rouse *et al.*, 1995; Costa *et al.*, 2009).

Isoniazid (INH) is one of effective anti TB drugs in which the rate of resistance to

this drug is increasing. Isoniazid is pro-drug which is highly dependent on KatG to become reactive radical that attack target in *M.tuberculosis* cell (WHO/IUATLD, 2004; Zhang *et al.*, 2005).

The mechanism of resistance of *M.tuberculosis* to INH is very complex, involving mutation on several genes, such as *katG*, *inhA*, and *kasA* (Rattan *et al.*, 1999; Ozturk *et al.*, 2005; Zhang *et al.*, 2005). Previous studies has shown that every region has a tendency different mutation frequency in *M.tuberculosis*. It is suggested that this condition is caused by differences in TB control program (Bulatovic *et al.*, 2002; Pym *et al.*, 2002; Sajduda *et al.*, 2004). Several reports has shown that mutation of *katG* is the main resistance mechanism of INH, and mutation on codon 315 (Ser315Thr) is the most frequent. This mutation has been found in more than 50-90% clinical strain, and caused increase in MIC 200 times compared to that of H37Rv *M.tuberculosis*. This mutation is strongly associated with MDR mutant and ability to be transmitted easily. Mutation of *katG* may result in decrease activity of peroxidase-catalase and deletion of this gene causes loss of activity of the enzyme (Moukrousov *et al.*, 2002; Xue-qiong *et al.*, 2006; Hu *et al.*, 2010).

KatG product has two important roles; altering INH (prodrug) into isonicotinic acid (active drug) and protect bacteria from intracellular oxidative stress (Master *et al.*, 2001; Musser, 1995; Bulatovic *et al.*, 2002). Based on concept that peroxidase-catalase can protect bacteria from phagocytosis by macrophages, decrease or deletion of this enzyme will affect bacterial virulence. *katG* mutant of *M.tuberculosis* should not be able to survive inside macrophages, the disease will not develop/progress and may transmission to other hosts does not occur. In fact this concept has not been the case.

Previous studies has shown that mutation on *katG* does not affect bacterial virulence. It is believed that *M.tuberculosis* has developed another mechanism to overcome

its drawback. It is suggested that alkyl hydroperoxide reductase (AhpC) takes place of KatG for maintaining bacterial virulence. There has been a controversial evidence on the role of *katG* and *ahpC* in bacterial virulence of *M.tuberculosis*. In *M.tuberculosis*, *katG* deletion have an impact on *ahpC* mutation that results in over production of this enzyme. In normal *M.tuberculosis* this enzyme is produced minimally (Pym *et al.*, 2002; Hillas *et al.*, 2000; Zhang *et al.*, 2005). Other studies has shown that *katG* mutant did not influence on AhpC activity and there was no mutation on *ahpC* (Heym *et al.*, 1997; Guimaraes *et al.*, 2005). Influence of AhpC on *katG* Ser 315 Thr mutant, factors effecting AhpC production and its relation with virulence of *M.tuberculosis* is not understood with certainty.

Recently, peroxidase-catalase and alkyl hydroperoxide reductase in INH-resistant mutant have been studied at genetic and molecular levels. Previous studies on *katG* showed that the role of this enzyme is not simple, involving role of AhpC (Pym *et al.*, 2002). Both enzymes in concert may function in resistance against intracellular oxidative stress. If these enzymes prove to be the main virulence factor and intracellular resistance of *M.tuberculosis*, they may be the target of anti tuberculosis drugs development in the future.

This study is aimed to understand the role of AhpC, in *katG* mutant of INH-resistant *M.tuberculosis* specifically the role of AhpC in virulence of isoniazid resistant *katG* Ser315Thr *M.tuberculosis* under oxidative stress condition.

Materials and Methods

Population and samples

In this study 85 INH-resistant *M.tuberculosis* strain collected from several laboratory were used (laboratorium Rumah Sakit Persahabatan, Jakarta, Balai Pengembangan Laboratorium Kesehatan (BPLK) Bandung, Balai Besar Laboratorium Kesehatan (BBLK) Surabaya, Balai Besar Kesehatan Paru Masyarakat (BBKPM) Surakarta, Balai Kesehatan Paru Masyarakat

(BPKM) Jogjakarta, and Laboratorium Mikrobiologi Fakultas Kedokteran UGM). For control purpose, *M.tuberculosis* H37Rv strain was included in this study.

Isolation of *M.tuberculosis*

Sputum samples from TB patient were homogenized and deconcentrated with modified method of Petrof. Sputum was added with equal volume of 4% NaOH containing phenol red and flash mixed for 15-30 sec. Samples were added with phosphate buffer and spinned at 3000 g for mins. Supernatant was discarded, pellet was added with NaOH until red colour appeared. Samples were examined microscopically after Ziehl Neelsen staining and cultured on Lowenstein Jensen (LJ) medium for 3-6 weeks at 37^o C.

Susceptibility tes

Susceptibility test was done with modified indirect proportional using LJ and Middlebrook 7H10 media. Each sample was cultured on a series of LJ medium containing INH (0.2 mg/L dan 1.0 mg/L). Sample was also cultured on LJ without drug for control purpose. Inoculated media were incubated at 37^o C for 3 weeks, and the presence of bacterial growth was examined every week. Confluent growth on control was considered as 100 % growth. Sensitive or resistant determination was based on comparison between growth on medium with and without drug. An resistant isolate was determined by number of colonies on drug containing medium, if was 1% or more of number of colonies on control medium.

DNA isolation

DNA isolation was performed for INH resistant *M. tuberculosis* strain. Isolation was carried out with boiling method. Samples were scrapped from LJ and placed in tube containing 500 µL of sterile aquadest and heated at 95^o C for 5 min. Samples were then spinned at 12.000 x g for 5 min, supernatants were saved on -20^o C and used as DNA template.

Mutation analysis of *katG* Ser315Thr

Primers for *katG* 315, *katG*IF (5'-AGCTCGTATGGCACCGGAAC) and *katG* 4r (5'-AACGGGTCCGGGATGGTG), were used for *katG* fragment amplification (position 904 to 1103 in H37Rv) (Mokrousov *et al.*, 2002).

RFLP analysis codon 315

Amplified fragments were digested with *Msp*I. Fifteen μ L PCR product was added with 10 μ L of buffer and 2.7 μ L of the enzyme. The mixtures were placed in waterbath at 37°C for 8-16 h. Digested fragments and marker V (Roche) were subjected to 4% agarose gel electrophoresis and results were visualized with UV light. Mutated codon 315 will be digested by *Msp*I, resulting in 7 fragments ; 14, 6, 21, 6, 11, 10, and 132 bps. Digestion of non mutant *katG* Ser315Thr resulted in 6 fragments; 14 , 6, 6,11,10, and 153 bps (adapted from Mokrousov *et al.*, 2002).

Catalase activity assay

Two weeks old of *M.tuberculosis* in LJ medium were added with mixture solution of 1:2 tween and hydrogen peroxide, incubated in room temperature for 5 min. The height of developed bubbles was measured.

Exposure of *M.tuberculosis* to hydrogen peroxide

M.tuberculosis was grown in 7H9 Middlebrook liquid medium to density was adjusted to 10^8 cfu/mL (Mc Farland I standard), then aliquoted into 5 mL portions in tubes and treated with 0.02, 0.2, 0.1, 1 and 2 mM H_2O_2 (Sigma). All tubes were incubated at 37°C for 4x 24 h. (modified from Pagan-Ramos *et al.*, 2006).

AhpC activity assay

Five hundred microliters of isolate suspension was added with 500 μ L of DTT solution, incubated at room temperature for 120 min. The rate of DTT oxidation was measured spectrophotometrically at 310 nm

at room temperature (modified Chauhan, 2002). The experiments were repeated three times.

Virulence study in human macrophage cultures

To find out the effect of increased AhpC activity (after oxidative stress) on virulence of *katG* Ser315 Thr mutant, several indicator of virulence were examined, i.e: replication ability of *M.tuberculosis* after were phagocyted by human macrophage and macrophage apoptosis induction. Dose of H_2O_2 that gave the highest of AhpC activity was used in the experiment.

Macrophage preparation: macrophages were obtained from 4 blood donors. After macrophages were isolated from donors, these cells were mixed in RPMI containing fetal bovine serum. Isolated macrophages was counted with hemocytometer to get cell density of 5×10^4 /mL. Cells suspension was the divided into 24-well plates with cover slip inside. Each well was filled with 200 μ L of macrophages suspension and incubated at 37°C with 5% CO_2 for 4 X 24 h.

Inoculation of *M.tuberculosis* into macrophage cultures: bacterial suspension was diluted into density of 10^6 cfu/mL, placed in a tube containing glass beads. Resuspension was done using 27-G syringe until homogenous suspension was obtained. A hundred μ L of suspension was added into each wells of microplates that contained of 5×10^4 /mL macrophage cells density, so that 1: 100 multiplicity of infection (MOI) was expected. Microplates were incubated at 37°C with 5% CO_2 for 2 h. The presence of phagocytosis was confirmed with microscopic examination after auramin-acridine orange staining.

Replication ability of *M.tuberculosis*: after 4 X 24 h incubation, macrophages were lysed using lysis buffer. A hundred μ L of lysate was inoculated on LJ medium and incubated at 37°C. Growth of *M.tuberculosis* was examined after 4 weeks, it was confirmed with Ziehl Neelsen staining. Replication

ability was determined by comparing the number of colonies growth of strain tested with the number of colonies before infected.

Apoptosis assay: assay was done on day 4 after infection, cover slips were taken out of the well and stained with hematoxylin-eosin, and microscopically examined. To avoid bias, examination was performed by more than one observer and in inter observer blinded manner. Cells undergoing apoptosis will be seen as a round or oval mass, eosinophilic cytoplasm. Chromatin will be condensed and aggregated with clear

border, nuclei fragmented and wrapped with cell membrane. In acute apoptosis, edge part of nuclei absorb stain strongly and there is vacuole in the middle. Cells with late phase apoptosis absorb stain weakly compared to that of cells with no apoptosis.

Data analysis

Student T-test was used for AhpC activity between oxidative stress-treated and non treated groups. Linier regression correlation test was used in analysing association between increased activity with virulence factors.

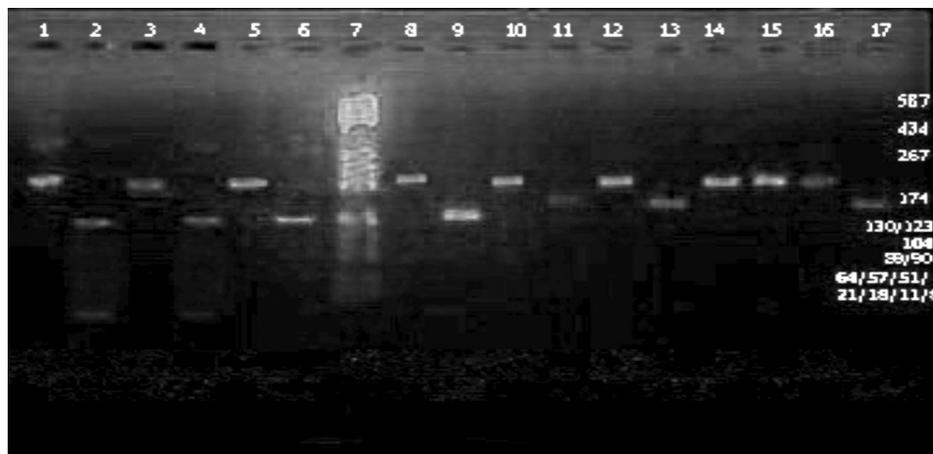


Figure1. A-lane: PCR/RFLP product of *katG* kodon 315. B- lane: PCR product after digested by *MspI*.

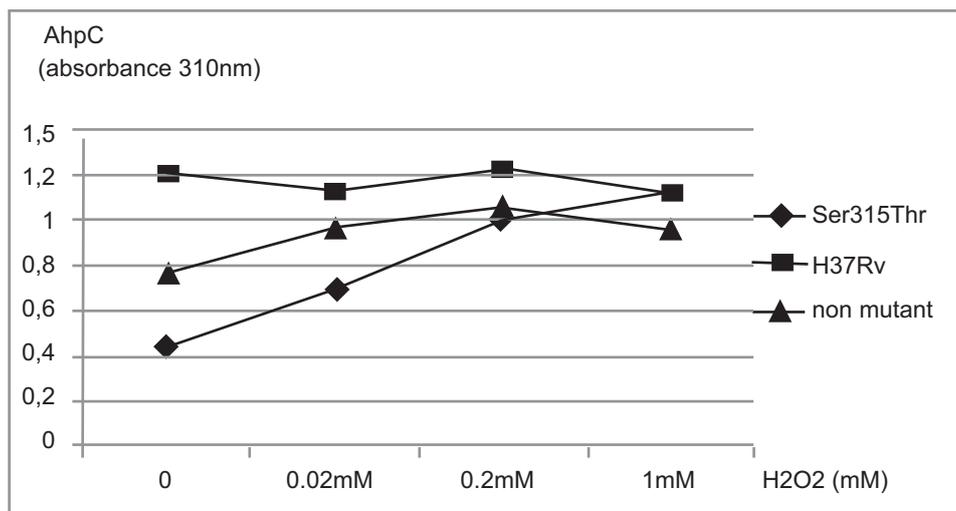


Figure 2. Mean of AhpC activity after induction with H₂O₂.

Results

Frequency of *katG Ser315Thr* mutant *M.tuberculosis*

Mutation on *katG Ser315Thr* was investigated from 85 PCR products using RFLP method. Twenty three out of 85 strain (27.05%) were indicated as *katG Ser315Thr* mutants. As predicted that PCR/RFLP product was 199 bps. Mutated DNA was digested into 7 fragments with the longest was 132 bps. In non mutant, product was digested into 6 fragments and the longest was 153 bps.

Catalase activity in *katG Ser315Thr* mutant strain

To test the catalase activity level, the semiquantitative assay was used. The H37Rv strain was included in this test to find out the basal level of AhpC. Data of Table 1 demonstrated that all of *katG Ser315Thr* mutant strains had height of bubbles less than 4 mm, that indicated the catalase activity strongly decreased (6-7% remained relative to the non mutant and H37Rv strain). In contrast, non mutant strains showed high catalase activity. The deviation standard of non mutant was 14,993, these value indicated the range variation of the catalase activities of

these groups were very width, possibly due to variation of the non mutant strains.

Table 1. Mean catalase activity in strain examined

Strain	Number	Mean bubbles heights (mm)	Significancy
Mutant	23	3,38 ± 2,599	p = 0,00
Non mutant	62	46,06 ± 14,993	t = 20, 963

Note : the bubble height of H37Rv strain is 55 mm

Effect of oxidative stress on increase AhpC activity

In order to understand the role of AhpC as compensator of low catalase activity in *katG Ser315Thr* mutant, inducing expression of AhpC was examined by exposure to various concentration of H₂O₂. Then the dithiothreitol (DTT) oxidation as indicator of AhpC activity was measured by t-butyl hydroperoxide decreased that observed by spectrophotometrically at absorbance 310 nm.

Inspection of AhpC level in response to H₂O₂ induction in *katG Ser315 Thr* of *M.tuberculosis* mutant strain showed increase of the H₂O₂ level have an effect on of AhpC activity, with peak induction at 1 mM H₂O₂. However, in non mutant of *M.tuberculosis*

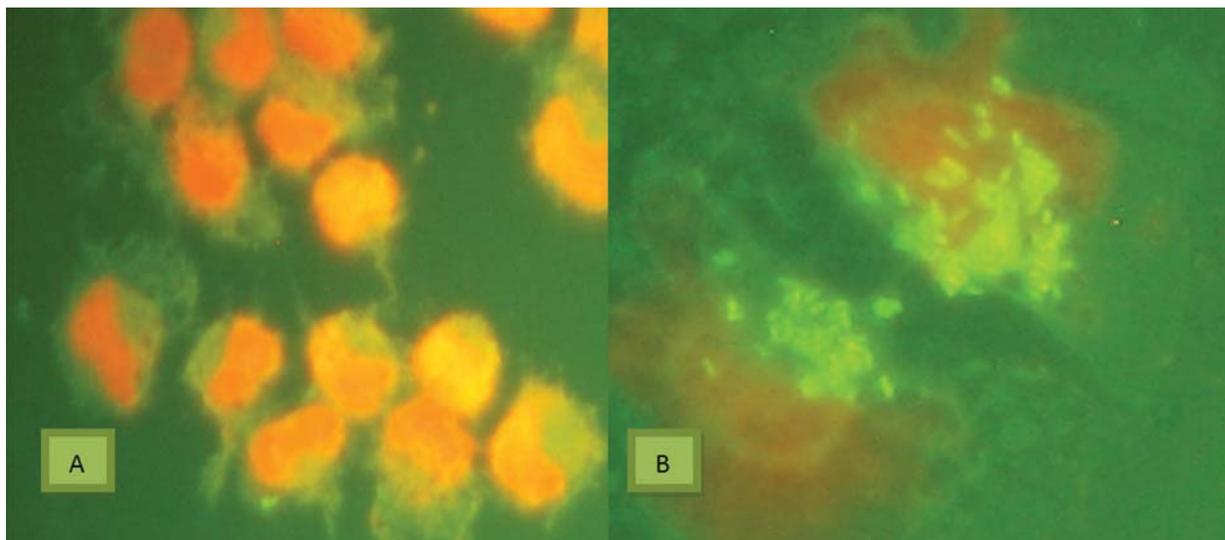


Figure 3. Macrophages phagocytosis after infected by *M.tuberculosis* under florescens microscope by auramine-acridine orange 10xobjective. A. 4-days macrophages culture without infection. B. yellow-green appearance of mutant *M.tuberculosis* no 43 were phagocytied by 4-days macrophages culture by MOI 1:50, after 2 hr incubation.

strains showed that all of the H₂O₂ levels have no effect on AhpC activity, and AhpC level in H37Rv strain remained similar and were not induced by H₂O₂ (Figure 2).

Association between increase of AhpC activity and virulence of katG Ser 315Thr M.tuberculosis mutant

To reach the aim of the study in understanding association between the increased of AhpC activity and virulence of mutant we observed the several indicator, ie: ability to replicate and reducing of apoptosis macrophages induction after the mutant strain infected to peripheral blood monocytes cultures.

Replication ability of mutant.

To observe replication ability in vitro of mutant, strain were inoculated into 4 days old macrophages culture. Strain were divided into two groups; those pretreatment with oxidative stress inoculated into culture of macrophages and those without oxidative stress treatment (normal AhpC activity).

Replication ability of *M.tuberculosis* after phagocytized by macrophages indicate intracelullar survival capability. In this study shown in Table 2, *katG Ser315Thr* mutant replication before obtained stress was average

66.88 cfu/mL, but after AhpC developed (under oxidative stress), its replication ability increase twice was 144.52 cfu/mL.

Table 2. Increased and replication ability in *katG Ser315Thr M.tuberculosis* in groups with and without oxidative stress treatment

Groups	Mean of replication increased (cfu/mL)	Significancy
Without stress oxidatives	66,88± 58,824	p=0,001
Under stress oxidatives	141,52 ±118, 857	t =3,233

These results demonstrate that the high level of AhpC affected to increase of mutant replication ability. It arisen a guess that by KatG catalase weakness compensation by AhpC development, intra cell oxidative stress effect could decrease so that bacterial replication could run optimal (see Figure 4).

Macrophages apoptosis after infection with katG Ser315Thr mutant of M.tuberculosis

Apoptosis or programmed cell death in alveolar macrophages can be induced by *M.tuberculosis* infection. In this study, apoptosis was observed by microscopic examination after Hematoxillin-eosin staining. Proportion of cell undergoing apoptosis per 100 macrophages cells was calculated.

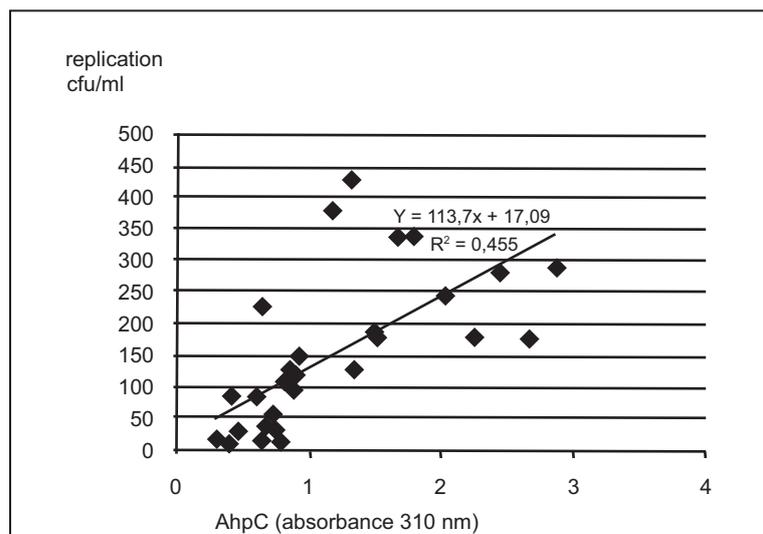


Figure 4. Correlation between increased AhpC and replication ability

The results presented in this study indicate a link between increased level of AhpC and decrease of macrophage apoptosis

induction, see Table 3. The correlation between increased and macrophages apoptosis after

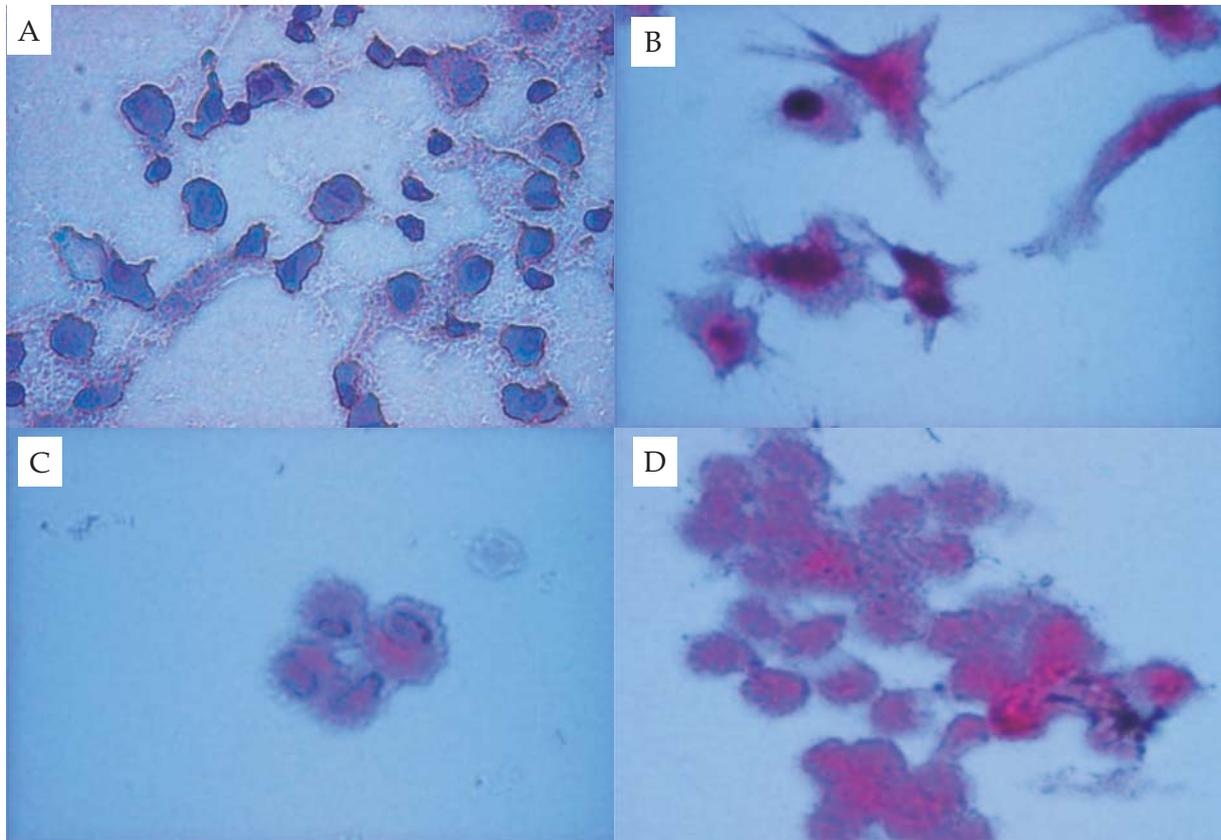


Figure 5. Appearance of macrophages by Haematoxilin-Eosine, under microscope observation (400 x objective). A. 7 days macrophages culture without infection. B. macrophages culture after infected by *M.tuberculosis* C. Early apoptosis macrophages after infected by mutant, D. late apoptosis macrophages after infected by isolate of mutant.

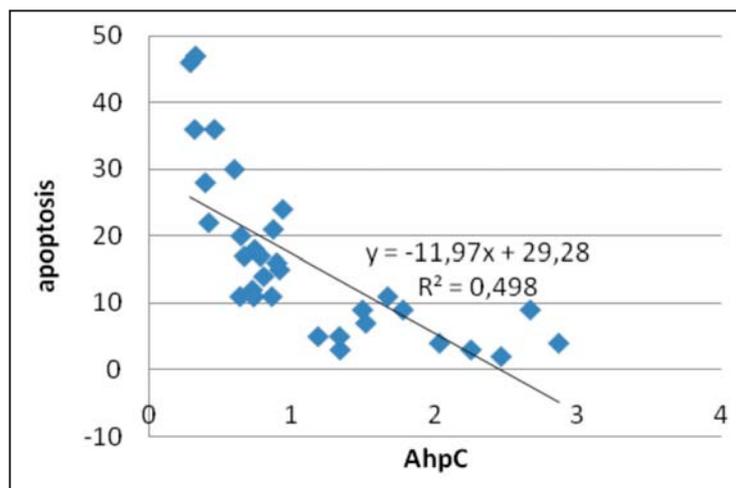


Figure 6. Correlation between increased AhpC and macrophages apoptosis induced by *katG* Ser315Thr *M.tuberculosis* mutant

Table 3. Mean of apoptosis macrophages after infected with *katG* Ser315Thr *M.tuberculosis* mutant without oxidative stress under and oxidative stress

Macrophages groups	Mean of apoptosis/ 100 macrophage cells	significancy
Without oxidative stress	57,7976 ±11,04650	p = 0,00 t =-3,706
Under oxidative stress	33,0167±11,30805	

infected by *katG* Ser 315Thr mutant of *M.tuberculosis* showed by Figure 5.

Discussion

In order to understand the role of AhpC as compensator of low catalase activity in *katG* Ser315Thr mutant, AhpC activity was examined with oxidative stress treatment by inducing with H₂O₂ at different concentrations. A preliminary study was conducted by using dose of 1, 2, 3, 5, and 10mM. Result showed that significant increase only occur at dose of 1 and 2 mM. At 3 mM AhpC decreased in accordance with decreased bacterial growth. At high peroxide levels (i.e:10 mM), some loss activity was detected, possibly due to increased degradation and/or inability to increase production due to excessive damage. Based on this study dose of 0.2, 0.2, and 1 mM were used in AhpC activity assay.

The highest AhpC activity of the mutant was reached after treatment with 1 mM of H₂O₂. In control isolate H37Rv there was no significant difference in AhpC activity after treatment with different doses. This control isolate seemed to have a relatively high AhpC activity without H₂O₂ exposures, and the treatment did not affect AhpC activity.

Results in this study showed that there was selectivity in AhpC regulation in *M.tuberculosis*. When the bacteria was under oxidative stress and KatG was decreased, compensation by AhpC occurred. It was known that during evolution process the bacteria had lost *oxyR* as *ahpC* regulator, and up to now mechanism of *ahpC* regulation

in *M.tuberculosis* is not yet understood. AhpC is probably induced by H₂O₂ in the environment.

In this study virulence of the isolate was determined from ability to replicate intracellular, and apoptosis of macrophage after infected in culture of peripheral blood monocytes. Oxidative stress was able to increase AhpC activity in *katG* Ser315Thr mutants with decreased KatG .

Ability of *M.tuberculosis* to manipulate macrophage is the key of its survival. This is associated with bacterial ability to avoid death inside macrophage and to induce delayed hypersensitivity. To be able to remain alive in the host, *M.tuberculosis* have to develop strategies; replication inside macrophage, resistance to system immune response or enable the host to control bacterial growth without destroying the bacteria. In certain condition, *M.tuberculosis* have to live in inactive state without losing ability to be reactive (Glickman and Jacobs, 2001).

Data showed that mutant strain was able to replicate inside macrophage. Results also showed that significant increase of AhpC was influence the intracellular replication ability. It means that *katG*Ser315Thr mutant can resist intracellular environment and phagocytosis by macrophage although there is a decrease of KatG, because of AhpC compensation.

Regarding the macrophages apoptosis induction by *katG* Ser315Thr mutant *M.tuberculosis* as one of virulence factor, this results indicated there was relationship of increased AhpC activity with macrophages apoptosis induction.

Based on results in this study it can be concluded that *katG*Ser315Thr mutant *M.tuberculosis* is a dynamic mutant that keep balance between resistance to INH and inactivated peroxide-catalase. When peroxide-catalase is weak, this enzyme can still protect bacteria from organic peroxide produced by the host. If there is strong oxidative stress beyond peroxide-catalase strength, then compensation by AhpC occurs. Beside compensating low KatG activity,

AhpC enhanced the replicating ability and also reduce induction of apoptosis of macrophages, so that survival of the bacteria can be maintained.

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