Bovine serum albumin promotes reactivation of viable but non-culturable *Mycobacterium tuberculosis* via activation of protein kinase-dependent cell division processes

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Abstract

Objective: *Mycobacterium tuberculosis* (Mtb) H37Ra strain has been reported to rapidly enter the viable but non-culturable (VBNC) state following treatment with an NADH oxidase inhibitor (diphenyleneiodonium [DPI]) and to be reactivated by fetal bovine serum (FBS). However, the mechanism underlying FBS-induced reactivation is unclear. We tried to reveal the mechanism of FBS-induced reactivation using *M. tuberculosis* H37Rv.

Methods: First, we evaluated the effect of DPI on culturability, viability and changes of acid-fastness toward H37Rv. Secondly, we measured the reactivation-promoting effects of human serum albumin, egg-white albumin and antioxidative agents in DPI-induced VBNC cells. We also inhibited adenylyl cyclase and protein kinase which is the downstream of adenylyl cyclase to evaluate the influence to reactivation capacity of bovine serum albumin (BSA).

Results: DPI treatment induced VBNC state in H37Rv, resulting in a high proportion of viable cells but a low proportion of culturable cells, loss of acid-fastness and lipid-accumulation. Not only FBS but also BSA alone could reactivate H37Rv. Contrary to our expectation, only human serum albumin had a similar restorative effect to BSA. The inhibition of adenylyl cyclase by SQ22536 did not have a significant effect on reactivation; however, the inhibition of mycobacterial protein kinase by H89 and staurosporine strongly suppressed the BSA-induced reactivation.

Conclusion: DPI-induced VBNC Mtb cells may be reactivated via the activation of protein kinase-dependent cell division processes through interaction with BSA.

Introduction

Mycobacterium tuberculosis (Mtb) is known to as one of the intracellular parasitic bacteria that can survive inside host macrophages. It develops a latent phenotype, dormancy, due to various stresses such as host immune response, oxygen depletion, nutritional starvation, *etc.*

The dormant Mtb cells can persist inside the host and cause latent tuberculosis infection (LTBI),

which seeks reactivation. Lowering the risk of Mtb reactivation in LTBI patients is considered

to be one of the important strategies to reduce the incidence of TB (1). Dormancy phenomenon

in non-sporulating bacteria have described as two major states: the viable but non-culturable

(VBNC) state and bacterial persistence(2). VBNC cells are the nonculturable population that

can be induced by exposure to stress. In addition, they are unable to grown on nutrient media

even after removal of stress and require much longer time or specific stimulus for reactivation

(2, 3). On the other hand, persistent cells are stochastically arising cells within growing cultures

that is tolerant to stress including high-dose antibiotics, and they are also growth-arrested.

However, they are typically able to grow on nutrient media following after the removal of stress

with short period (2).

Many pathogenic bacteria, including Mtb, can enter into the VBNC state, and it is considered

to be their natural state in the environment including human body (4). Various *in vitro* models,

such as models of hypoxia, nutritional starvation (including potassium-deficiency), a lipid-rich

environment, and other multiple stresses are reported to induce a VBNC state in Mtb cells (5–

9). A deep understanding of the mechanisms of reactivation is necessary for the prevention of

active tuberculosis.

One of the major hypotheses regarding the mechanism underlying the formation of VBNC

bacteria involves oxidative damage generated by harsh external conditions (10). Thus, some

studies have reported that the antioxidation during culture is one of the key mechanisms

through which VBNC bacteria are reactivated (11, 12). Some mycobacterial culture media

contain bovine serum albumin (BSA) and BSA on mycobacterial culture has been considered

to be growth-supporting agent that is involved in detoxification by the absorption of free fatty

acid and other growth-suppressing agents that are spontaneously generated during culturing

(13–15). For this reason, the mycobacterial culture media that are most frequently used at the

present time are BSA-containing systems, such as Middlebrook 7H series or Dubos medium.

However, few studies have focused on the effect of such media on VBNC cells.

Mukamolova *et al*. revealed that among patients with active TB, there is certain

subpopulation for whom sputum is not culturable in conventional medium, but which becomes

to be culturable in the presence of resuscitation promoting factor (Rpf), which is a

peptidoglycan-hydrolyzing protein that is highly conserved in actinobacteria (16). Rpf

reactivates non-culturable Mtb cells both *in vitro* and *in vivo* (17), suggesting the presence of

VBNC Mtb cells in clinical specimens.

Previously, NADH oxidase inhibitor diphenyleneiodonium chloride (DPI) was reported to

simply and rapidly induce a VBNC state in the Mtb H37Ra strain, and incubation with fetal

bovine serum, which is a common supplement for mammalian cell cultures, could facilitate the

reactivation of VBNC cells, while incubation with OADC (oleate-albumin-dextrose-catalase)

supplementation could not (18). This report gave us an important clue for constructing a simple

and rapid assay system to induce and reactivate VBNC Mtb cells. However, when we tried to

apply this system to H37Rv, we happened to find that DPI-induced VBNC cells that could be

reactivated by not only fetal bovine serum but also by the addition of BSA—this was observed

for both H37Rv and H37Ra (internal data). Then, we hypothesized that BSA would act as a

reactivation-promoting agent toward DPI-induced VBNC cells.

One possibility of the reactivation-promoting mechanism of BSA is its antioxidative

property, which has been well-studied (19). Another possibility is that the cellular processes

involving cyclic AMP, an important second messenger of the cell. Shleeva *et al*. showed that

the presence of high amounts of cAMP provided by adenylyl cyclase is also essential for

reactivation of *M. smegmatis* and Mtb from the VBNC state (20, 21). Notably, overexpression

of Rv2212 gene, which encodes adenylyl cyclase in Mtb (22), significantly affects both entry

into and reactivation from the VBNC state (21).

Understanding the detailed mechanism of reactivation from the VBNC state is crucial to

reduce the risk of reactivation of Mtb cells in LTBI patients. In this study, we examined the

effects of antioxidative agents and the inhibition of adenylyl cyclase to reveal the roles of these

factors in the reactivation of VBNC Mtb cells.

Materials and Methods

Bacterial strains, growth media and culture conditions

This work was carried out using *Mycobacterium tuberculosis* H37Rv (ATCC 27294) and

H37Ra (ATCC 25177). Both cells were cultured for 3–5 days in a 60 mL glycol-modified

polyethylene terephthalate (PETG) bottle containing 20 mL Dubos broth (2.5 g/L Na_2HPO_4,

2.0 g/L L-asparagine, 1.0 g/L KH_2PO_4, 0.5 g/L pancreatic digest of casein, 0.2 g/L Tween 80,

0.5 mg/mL $CaCl_2 \cdot 2H_2O$, 0.1 mg/mL $CuSO_4$, 0.1 mg/mL $ZnSO_4 \cdot 7H_2O$, 0.05 g/L ferric

ammonium citrate, 0.01g $MgSO_4 \cdot 7H_2O$) with 5% (v/v) glycerol, and 10% (v/v) ADC

supplementation (5.0 g bovine serum albumin fraction V (BSA), 2.0 g dextrose, 0.003 g

catalase in 100 mL distilled water) at 37˚C in an orbital shaker (100 rpm) until the OD_{600}

reached up to 0.35. All procedures were performed at a BSL-3 facility. For further analysis, we

used commercially available BSA Cohn fraction V (Merck, Darmstadt, Germany), unless

otherwise stated. In addition, we used molecular biology grade Tween 80 (Merck product

number P5188-100ML) without any predilution.

Counting the number of culturable cells

The cultured bacterial suspension was serially diluted 10-fold in phosphate buffered saline (pH

6.8) with 0.1%(w/v) Tween 80 (PBS-T) and 25 µL of each diluent was inoculated onto a

Middlebrook 7H10 agar plate supplemented with oleic-albumin-dextrose-catalase (OADC)

(Becton Dickinson, Sparks, MD) in duplication. Colonies were counted after at least 3 weeks'

incubation at 37˚C under 5% CO_2. The limit of detection was determined to be 20 CFU/mL

because of the diluent factor.

Effect of BSA on the growth of Mtb cells in Dubos medium

Ten milliliters of log phase culture of Mtb H37Rv that reached an OD_{600} of up to 0.35 in Dubos

medium, as described previously, was sedimented at 3,000 $\times g$ for 10 minutes at 4˚C and the

supernatant was discarded. The sediment was washed twice with fresh Dubos broth and

resuspended with 10 mL of Dubos broth. The suspension was diluted to 1:10 with Dubos broth

with or without the addition of BSA solution (5.0 g BSA fraction V in 100 mL distilled water)

at the final concentration of 0.1% (w/v) BSA. The diluted suspension was dispensed into 24-

well plates (Sarstedt, Nümbrecht, Germany) in 1.5 mL increments and sealed with gas-

permeable film (Breathe-Easy, Diversified Biotech Inc, Dedham, MA, United States). Cultures

were incubated for 5, 10, 15 and 20 days at 37˚C under 5% CO_2. At each time-point, the number

of culturable cells was measured as described above.

Induction of the VBNC cells

Induction of the VBNC cells were performed according to the previous report (18) with some

modification. In short, the log phase culture of Mtb H37Rv or H37Ra that reached an OD_{600} of

up to 0.35 in Dubos medium, as previously described, was evenly divided into 30 mL PETG

bottles and 5 mg/mL diphenyleneiodonium chloride (DPI) in dimethyl sulfoxide (DMSO)

solution was directly added at the final concentration of 4 μg/mL with immediate agitation.

The same volume of DMSO was added to the untreated control sample. Cultures were further

incubated for 24 h at 37˚C without shaking. After incubation, their culturability and viability

were measured as described above.

Cell viability assay

Mtb cell viability was measured using esterase activity and membrane integrity of the cells

using dual staining with 10 mg/mL carboxyfluorescein diacetate (CFDA, Dojindo, Kumamoto,

Japan) and ethidium bromide (EtBr, Dojindo, Kumamoto, Japan). Briefly, 500 μL of DPI-

treated or untreated Mtb culture was sedimented by centrifugation at 15,000 $\times g$ for 5 minutes

at 4˚C and washed twice with the same volume of PBS-T. The washed bacterial sediment was

resuspended in 200 µL of PBS-T with CFDA and EtBr incubated at room temperature (RT) for

30 minutes in shade. The final concentration of CFDA and EtBr were 60 µg/mL and 2 µg/mL,

respectively. The stained Mtb cells were washed twice with PBS-T and fixed with 3.7%

formaldehyde solution at RT for at least 2 h. After fixation, 10 µL of bacterial suspension was

smeared on three APS-coated 8-well slides (Matsunami glass, Osaka, Japan) and air-dried.

Then the slides were mounted with DPX new non aqueous mounting medium (Merck,

Darmstadt, Germany) and a 150 µm-thick cover glass (Matsunami glass, Osaka, Japan) prior

to the microscopic observation. Specimens were examined with a BX53 fluorescence

microscope (Olympus, Tokyo, Japan) at a magnification of 100× for green (esterase-active with

intact membrane; live) or red (esterase-negative with damaged membrane; dead) fluorescing

cells. The excitation source for CFDA and EtBr was generated from a mercury-arc lamp (blue

beam, 470–495 nm; green beam, 530–550nm) using a corresponding band-pass filter. The

green-fluorescence emission from the CFDA-positive cells was collected through a 510–550

nm band-pass filter. The red-fluorescence emission from EtBr-positive cells was collected

through a 575 nm long-pass filter. At least 10 random fields were observed for each sample and

the images (TIFF format) were analyzed using the Fiji/ImageJ software program (23). The

live/dead ratio was calculated by direct counting of green or red fluorescing cells.

Auramine-O/Nile Red dual staining

Loss of acid-fastness and accumulation of neutral lipids are distinctive features of VBNC

mycobacteria (24). To detect the phenotype, fluorescent acid-fast staining with auramine-O and

neutral lipid staining with Nile Red (9-dimethylamino-5H-benzo-α-phenoxadine-5-one) were

performed using a previously described method (9) with some modification. Briefly, 10 µL of

DPI-treated or untreated culture were smeared onto an APS-coated slide (Matsunami glass,

Osaka, Japan) and air-dried. Then, the slide was heat-fixed and cooled down to room

temperature before staining. The smear was flooded with fluorochrome staining solution (10

μg/mL auramine-O in 5% [w/v] phenol solution) and incubated at room temperature for 20 min

in shade. Excessive dye was removed using 3% hydrochloric acid ethanol for 15 minutes. Next,

the smear was covered with 10 μg/mL Nile Red in ethanol and incubated at room temperature

for 15 minutes. Finally, the smear was counterstained with 0.1% (w/v) potassium permanganate

solution with 1 minute. Stained slides were air-dried and mounted as described previously.

Fluorescence microscopy was performed at 100× magnification for green (acid-fastness

positive, lipid accumulation negative; active) or red (acid-fastness negative, lipid accumulation

positive; VBNC) fluorescing cells. The excitation source for auramine-O and Nile Red was

generated from a mercury-arc lamp (blue beam, 470–495 nm; green beam, 530–550 nm) using

a corresponding band-pass filter. The green-fluorescence emission from auramine-O-positive

cells was collected through a 510–550 nm band-pass filter. The red-fluorescence emission from

Nile Red-positive cells was collected through 575 nm long-pass filter. At least 5 random fields

were observed for each sample and images (TIFF format) were analyzed by the Fiji/ImageJ

software program (23) and the proportion of acid-fast-positive cells and lipid-accumulation-

positive cells was calculated by direct counting of green or red fluorescing cells.

Ziehl-Neelsen staining

Ziehl-Neelsen staining was also performed to confirm the loss of acid-fastness resulting in the

VBNC state by DPI treatment, according to the standard method.(25) The slide was observed

by light microscope under 100x magnification with an oil immersion lens (BX53, Olympus,

Tokyo, Japan).

Confocal super-resolution microscopy

Detailed images of accumulated lipids were also obtained by a confocal laser-scanning

microscope (LSM900 with Airyscan 2, Carl Zeiss Microscopy, Jena, Germany). Briefly, culture

slides of DPI-treated cells dual stained with auramine-O and Nile Red, as described above,

were observed at 63x magnification with oil immersion lens. The excitation source of

auramine-O and Nile Red was 488 nm (blue beam) and 563 nm (green beam), from a solid-

state laser unit. The green-fluorescence emission from auramine-O and red fluorescence from

Nile Red were sequentially collected through a variable dichroic beamsplitter and an Airyscan

2 detector. Super-resolution images were processed with the ZEN Blue software program (Carl

Zeiss, Jena, Germany).

VBNC reactivation assay

The VBNC reactivation assay was performed as described elsewhere (18) with some

modification. DPI-treated culture was sedimented by centrifugation at 3,000 xg for 10 minutes

at 4°C and washed twice with fresh Dubos broth, then resuspended in ten-milliliters of Dubos

broth. The bacterial suspension was diluted to 1:10 with fresh Dubos medium with or without

0.1% (w/v) BSA. For both BSA-containing and BSA-free series, OADC supplementation

(Becton Dickinson, Sparks, MD), fetal bovine serum (FBS; Moregate BioTech, Bulimba,

Queensland, Australia), or sodium pyruvate (PA; Merck) was added. The final concentration

of each reagent was as follows: 10% (v/v) for OADC supplementation, 2% (v/v) for FBS, and

3 mM for PA. Then, the diluted suspension was dispensed into 24-well plates in 1.5 mL

increments and sealed with gas-permeable film. Cultures were incubated for 5, 10, 15 and 20

days at 37°C under 5% CO_2. At each time-point, the number of culturable cells was measured

as described previously.

Evaluation of the reactivation-promoting effect of human serum albumin, egg-white

albumin (ovalbumin) and antioxidative agents

To determine whether the reactivation-promoting effect of BSA is its antioxidative property,

we measured the reactivation-promoting activity of other albumins and antioxidative agents

toward DPI-treated Mtb cells. Washed DPI-treated cells were resuspended with fresh Dubos

broth without BSA and the suspension was diluted to 1:10 with Dubos broth containing BSA,

human serum albumin (HSA), ovalbumin (OVA), N-acetyl-*L*-cysteine (NAC) or D-mannitol

(MAN). The final concentration of each reagent was 0.1% (w/v) for BSA, HSA and OVA, 500

μM for NAC, and 50 mM for MAN, respectively. Then, the diluted suspension was dispensed

into 24-well plates in 1.5 mL increments and sealed with gas-permeable film. The plate was

incubated for 20 days at 37°C under 5% CO_2. At the end of incubation, the number of colonies

grown was measured as described previously.

Evaluation of the reactivation-promoting effect of fatty acid and globulin-free BSA

To determine the effect of BSA Cohn fraction V contaminants, we measured the reactivation-

promoting activity of fatty acid and globulin-free BSA toward DPI-treated VBNC Mtb cells.

Washed DPI-treated cells were resuspended with fresh Dubos broth without BSA and the

suspension was diluted to 1:10 with Dubos broth containing BSA Cohn fraction V or fatty acid

and globulin-free BSA. Fatty acid and globulin-free BSA were acquired from Merck. The final

concentration of both BSAs was 0.1% (w/v). The number of culturable cells was measured as

described previously.

Effects of adenylyl cyclase or protein kinase on BSA-induced reactivation

Washed DPI-treated VBNC Mtb cells were resuspended with fresh Dubos broth and the

suspension was diluted to 1:10 with Dubos broth with 0.1% (w/v) BSA containing adenylyl

cyclase inhibitor SQ22536 (Merck, Darmstadt, Germany) or protein kinase inhibitor H89

(Abcam, Cambridge, United Kingdom) or staurosporine (Merck). The final concentration of

each inhibitor was as follows: 0.1 to 10 mM for SQ22536, 1 to 30 μM for H89 and 1 to10 μM

for staurosporine. Then, the number of culturable cells were measured as described previously.

Molecular docking simulation

Molecular docking simulation was performed by using AutoDock Vina software, an open-

source molecular docking program designed by the Scripps Research Institute (26, 27). 3D

structures of receptor proteins, PknA and Rv2212, were obtained from Protein Data Bank (28)

and AlphaFold Database (29, 30), respectively. PDB ID for PknA was 4OW8 (31) and

AlphaFold Database identifier for Rv2212 was AF-P9WMU7-F1. We removed water

molecules from proteins, added polar hydrogen and charge, and adjusting X-Y-Z coordinates

and grid size for optimizing molecular docking. 3D structures of ligand molecules, H89,

SQ22536 and ATP were obtained from PubChem Database. PubChem CID of each ligand were

449241, 5270 and 5957, respectively. Graphical processing was done by Chimera software (32).

Statistical analysis

The statistical analysis was performed with R (33) using the EZR on R package (34). *P* values

of < 0.05 were considered statistically significant.

Results

I.   BSA is not essential for the growth of MTB in Dubos medium with purified non-

**   prediluted Tween 80**

Figure 1 shows the different effects of BSA-containing and BSA-free Dubos medium on the

growth of Mtb H37Rv. As shown in Fig. 1(A), there was no significant difference in the rate

of growth between the two conditions. Fig. 1(B) also shows the sufficient growth of cells. The

cells were more aggregative without BSA, however, the aggregation was easily dispersed by

pipetting and there was no significant difference in the OD_{600} value. The albumin component

was inessential for Mtb growth.

II.   DPI treatment induced a VBNC state in Mtb H37Rv

Figure 2 shows the DPI treatment results for the induction of VBNC state in H37Rv and

H37Ra (18). As shown in Fig. 2(A), the numbers of culturable cells of untreated control

population and DPI-treated population were significantly different: $4.43 \pm 1.43 \times 10^8$ CFU/mL

for the untreated control population and $4.64 \pm 6.10 \times 10^3$ CFU/mL for DPI-treated population.

Fig. 2(B) shows that the CFDA-positive cells in the untreated control and DPI-treated

populations were $81.8 \pm 11.4\%$ and $65.9 \pm 11.8\%$, respectively. These results suggested that

the majority of DPI-treated H37Rv cells were alive ($p = 0.169$), while their culturability was

significantly reduced ($p = 0.033$). Representative images of CFDA/EtBr dual-stained cells of

the untreated control and DPI-treated population also showed that there was little difference in

the proportions of CFDA-positive and EtBr-positive cells. (Fig. 2[C])

Auramine-O/Nile Red dual staining showed the reduction of acid-fastness and the

accumulation of the neutral lipid. As shown in Fig. 2(D), the proportion of acid-fast-

negative/lipid accumulation-positive cells was $84.0 \pm 14.0\%$ in the DPI-treated population, and

$12.1 \pm 11.9\%$ in the untreated control population ($p < 0.01$). Representative images of

auramine-O/Nile Red dual-stained cells of the untreated control and DPI-treated populations

also showed that there were distinctive differences in the proportions of acid-fast-positive cells

and lipid-accumulation-positive cells (Fig. 2[E]). The loss of acid-fastness was also confirmed

by Ziehl-Neelsen staining; the majority of DPI-treated population were found to have lost their

acid-fastness (Suppl. Fig. 1). Interestingly, Airyscan microscopy revealed some DPI-treated

cells were both acid-fast stain-positive and neutral lipid stain-positive (Fig. 2[F]). These results

indicated that DPI treatment induced the VBNC state in H37Rv as well as in H37Ra.

III.   The effects of BSA, FBS, OADC and sodium pyruvate on the reactivation of DPI-

treated VBNC Mtb cells

As shown in Fig. 3(A) and (B), our results partially support the previous study (18) with

some differences; although the culturability of DPI-treated H37Rv cells was restored by

incubation with FBS-supplemented medium, OADC-supplemented medium could also induce

culturability. The addition of albumin alone into the medium could also induce reactivation.

The addition of FBS in BSA-free Dubos medium slightly reduced the regrowth rate; however,

cells were successfully reactivated at the end of incubation with or without FBS. These

phenomena were also observed in H37Ra (Suppl. Fig 2). Thus, we used the H37Rv strain for

the further analyses in this study.

Incubation with sodium pyruvate, which was reported to have a reactivation-promoting

effect on VBNC cells elsewhere, led to transient regrowth by day 15 and a slight reduction at

day 20. (Suppl. Fig. 3[A] and [B]).

We also confirmed that the presence of small population of intact Mtb cells (10^3 CFU/mL)

could grow normally in BSA-free Dubos medium, suggesting that the reactivation might not

be due to the presence of small number of culturable cells after DPI-treatment (data not shown).

IV.   The antioxidative property or the fatty acid from BSA did not promote

reactivation.

As shown in Fig. 4(A) and (B), the reactivation-promoting effect of BSA was specific to

bovine and human serum albumin. Ovalbumin did not show a reactivation-promoting effect

but maintained the number of culturable cells in this system. NAC (antioxidative agent) and

D-mannitol (free radical scavenger) did not show any reactivation capacity.

We also checked whether the purity of albumin affects the promotion of reactivation using

fatty acid and globulin-free BSA and confirmed that there was no significant difference in

reactivation-promoting effects (Suppl. Fig 4). These results suggest that commercially

available albumin including fatty acids do not affect the reactivation of DPI-treated Mtb.

V.   The reactivation-promoting effect of albumin is canceled by treatment with

eukaryotic protein kinase inhibitor.

As shown in Fig. 5(A) and (B), SQ22536, adenylyl cyclase inhibitor, only suppressed

reactivation at a very high concentration (10 mM) in this study. However, H89, a eukaryotic

protein kinase inhibitor, suppressed reactivation in a dose-responsive manner. In detail,

incubation with 10 µM or 30 µM H89 resulted in a CFU/mL value for $3.42 \times 10^6 \pm 7.45 \times 10^5$

CFU/mL and $3.63 \pm 1.70 \times 10^2$ CFU/mL, while incubation without H89 resulted in 3.00×10^8

$\pm 5.07 \times 10^7$ CFU/mL. We also confirmed that staurosporine, which is known as mycobacterial

protein kinase PknB inhibitor (35), suppressed reactivation at 10 µM and resulted in a CFU/mL

value for $2.09 \times 10^3 \pm 7.62 \times 10^2$ CFU/mL, while incubation without staurosporine resulted in

$3.58 \times 10^8 \pm 5.78 \times 10^7$ CFU/mL (Fig 5[A] and [B]). All inhibitors used in this study did not

cause a reduction of the growth of intact MTB cells (Suppl. Fig. 5A and 5B).

We also performed molecular docking simulation of these inhibitors toward their

considerable targets on Mtb. As shown in Suppl. Table 1, both SQ22536 and H89 have

sufficient affinity toward Rv2212 (adenylyl cyclase) and PknA (protein kinase), respectively.

Discussion

In this study, we confirmed that DPI could induce a VBNC state in H37Rv as well as H37Ra.

The mechanism underlying the effect of DPI is considered to involve the inhibitory effect of

NADH oxidase, which results in the inhibition of the electron transport system of Mtb. This

may induce a hypoxic response in the liquid culture, resulting in significant decrease of

culturability (Fig.2[A]). However, viability was highly retained (65.9%), suggesting that

majority of the cells transformed to VBNC (Fig. 2[B]). Additionally, VBNC transformation

was confirmed by auramine-O and Nile Red staining. The loss of acid-fastness and the

accumulation of neutral lipids are distinctive features of VBNC mycobacterial cells.

Our data also demonstrated that the majority of DPI-treated Mtb H37Rv cells were stained

with Nile Red alone (Fig. 2[D]). Airyscan super-resolution microscopy showed the presence of

cells that were stained with auramine-O and which contained an intracellular lipid body stained

with Nile Red (Fig. 2[F]). This could also reveal the distribution of the lipid body in the cell as

some foci of relatively strong signals of Nile Red, suggesting that the transformation from

growing state to VBNC, with drastic alteration of the lipid metabolism.

Secondly, we found that DPI-induced VBNC could facilitate reactivation not only by

incubation with FBS but also with OADC supplementation and BSA alone, suggesting albumin

might act as a reactivation-promoting agent in both H37Rv and H37Ra (Fig .3 and Suppl. Fig.

2). These findings were contrary to the findings of the previous study, which showed that only

FBS could facilitate reactivation (18). We considered the reason for the difference may be due

to contaminant in Tween 80, presence of excessive amount of oleic acid. We also confirmed

that the reactivation was not facilitated by the use of prediluted Tween 80 for the preparation

of Dubos medium (internal data) for both H37Rv and H37Ra. Thus, our results suggested that

the key component of reactivation-promoting agent in FBS might be serum albumin, and

further analyses were performed using H37Rv.

We also tested the reactivation-promoting effect of BSA in Wayne's hypoxic culture, which

is widely used for inducing a VBNC of Mtb, and found that there was no significant difference

with or without BSA (data not shown).

Pyruvate, which is known to act as reactivation promoting agent for both gram-negative

bacteria, such as *Escherichia, Legionella, Vibrio* and gram-positive bacteria such as

Staphylococcus (36–39), did not show any reactivation-promoting effect in our experiment

(Suppl. Fig.3). Vilhena *et al.* reported that their proteomic analysis of VBNC *E. coli* cells

during reactivation revealed that enzymes involved in pyruvate metabolism, pyruvate formate-

lyase-activating protein (PflA), phosphoenolpyruvate carboxykinase (PckA) and lactate

dehydrogenase (LldD), were significantly upregulated (40), while Wagley *et al.* also reported

that VPA1499, an LldD homologue of Vibrio parahaemolyticus, was significantly upregulated

in VBNC *V. parahaemolyticus* and that depletion of LldD led to quicker entry to the VBNC

state (41). Interestingly, they showed that the addition of pyruvate, the product of lactate

oxidation, to LldD-depleted mutant restored the wild-type characteristics of cells in the VBNC

state. Their studies might give some crew of the lack of reactivation-promoting effect toward

mycobacterial VBNC. Since Mtb does not have a catalytic center of pyruvate formate-lyase

(PflB), pyruvate might not be able to act as reactivation-promoting agent, although pyruvate

enhances the growth of Mtb and *M. bovis* both intracellular and extracellular growth by

promoting TCA cycle and other energy-producing pathways. (42, 43).

Our results showed that the reactivation promoting effect of albumin might be structure-

specific because ovalbumin (OVA) did not facilitate reactivation, but bovine (BSA) and human

serum albumin (HSA) did. Focusing on their amino acid sequences, BSA and HSA share 76%

sequence homology (44); however, BSA and OVA share less sequence homology (16%). One

of the considerable roles of albumin may be as an antioxidative agent due to its cysteine

residues with free thiols (19, 45). BSA and HSA have only one cysteine residue with free thiol,

while OVA has four cysteines with free thiols (46). OVA therefore seems to be more reductive

than BSA and HSA. Our results were paradoxical in relation to this hypothesis; however, Davis

and Dubos showed the differential growth-promoting effect of albumin toward Mtb. Their

study supplied some interesting clues for our present study; ovalbumin does not show a growth-

promoting effect, while bovine and human serum albumin showed the effect (15). Moreover,

the antioxidative agents used in this study, NAC and D-mannitol, showed any restorative

effects (Fig. 4). We also examined catalase, a component of OADC supplementation, and its

reactivation promoting effect toward VBNC *Corynebacterium, Lactobacillus, Salmonella,*

Ralstonia and *Vibrio* (47–52); no restorative effects were observed (data not shown). These

results suggest that the effect of albumin may be due to complex functions rather than its

antioxidative property.

Serum albumin is known to act as a good carrier of many kinds of biologically active

substances, including free fatty acid. It is also known as a detoxifier of growth-arresting

substances in culture medium. BSA is added to mycobacterial culture medium as both a carrier

and detoxifier of growth-supporting and growth-arresting substances, respectively. Shleeva *et*

al. showed that several types of unsaturated free fatty acid, such as oleic, linoleic, and

arachidonic acids, can promote the reactivation of VBNC *M. smegmatis* in modified Sauton's

medium and oleic acid also works for Mtb (20). In this study, we should note that impurities

of albumin did not affect reactivation. Although the underlying mechanism is still unclear, both

fatty acid and globulin-free BSA and BSA Cohn fraction V showed similar reactivation-

promoting effect toward DPI-treated Mtb (Suppl. Fig. 4).

The reactivation inhibition assay by SQ22536, H89 and staurosporine gave us an important

clue for understanding the effect of BSA. In the present study, we could suppress the

reactivation of DPI-induced VBNC cells by inhibiting adenylyl cyclase only with a very high

concentration of SQ22536. Although the underlying mechanism is unclear, adenylyl cyclase

may not play a key role in reactivation from a DPI-induced VBNC state, while the inhibition

of protein kinase by H89 and staurosporine clearly suppressed reactivation. These results

suggest that protein kinase might mainly be involved with the reactivation of VBNC Mtb cells

triggered by BSA and the contribution of adenylyl cyclase may be smaller than that of protein

kinase.

Mtb has the eukaryotic-type Ser/Thr protein kinases PknAB and PknD-L. For example,

PknA regulates major cell processes, regulates several types of proteins involved with

mycobacterial cell division and morphogenesis via phosphorylation of such proteins in

coordination with PknB (Manuse *et al*., 2016; Carette *et al*., 2018); phosphorylation of GlmU

(Rv1018c) and MurD (Rv2155c), Wag31 (Rv2145c), HupB (Rv2986c) and ParB (Rv3917c),

influence the production and cross-linking of peptidoglycan precursors (55–58), the spatial

localization of peptidoglycan synthesis (59), DNA condensation and partitioning (60, 61),

respectively. It should be noted that the phosphorylation of FtsZ and FipA by PknA controls

cell division under oxidative stress (62, 63). In addition, mycobacterial protein kinase has

cyclic AMP-dependent protein kinase domains which is very similar to eukaryotic version,

which was shown in PknB (64) and staurosporine, which act as PknB inhibitor, could also

inhibit reactivation of DPI-induced VBNC Mtb (Fig. 5).

Our study suggested that the inhibition of mycobacterial protein kinase by H89 and

staurosporine seems to critically affect several important cellular processes, followed by

reactivation (Fig. 6). To confirm the essentiality of mycobacterial protein kinase and other

involved proteins toward reactivation from this DPI-induced VBNC state by albumin, further

studies using experiments such as knock-down of genes using the CRISPR interference

(CRISPRi) technique (65), because such proteins—with the exception of FipA (Rv0019c)—

are known to be essential for *in vitro* growth (66–69); thus, manipulating their coding genes is

not permissible.

Conclusion

Our data showed that the presence of BSA in Dubos medium is not only for depletion of

growth-supressing substances but also could trigger the reactivation of the DPI-induced VBNC

state in both H37Rv and H37Ra strains. The reactivation-promoting effect of other kinds of

albumins and antioxidative agents, including NAC and D-mannitol was tested; however, only

human serum albumin showed a reactivation-promoting effect. The results of the inhibition

assay of reactivation using SQ22536, H89 and staurosporine suggests the possibility that the

inhibition of mycobacterial protein kinase suppressed reactivation to a much greater extent than

adenylyl cyclase, indicating the presence of a reactivation-promoting pathway which is protein

kinase-dependent.

To the best of our knowledge, few studies have focused on the reactivation-promoting effect

of albumin toward VBNC Mtb. Taken together, our findings indicated that the reactivation of

MTB from a DPI-induced VBNC state could be obtained by interaction with bovine serum

albumin and the mechanism of albumin-induced reactivation may involve the activation of

mycobacterial protein kinase, which regulates several important processes of cellular

component construction and cell division. Although it remains unclear how albumin interacts

with VBNC cells, our study may supply some important clues to understanding the physiology

of the mycobacterial VBNC state.

Author Contributions

YMori, YMura and SM conceived and designed the experiments. YMori performed

experiments. YMura, KC, AA, YI, YS, MH, KK, HY, AT, and SM assisted in the experiments.

YMori acquired data and YMori and YMura analyzed data. YMori performed molecular

docking simulation and statistical analysis. YMori and SM wrote the manuscript. All authors

contributed to the article and approved the submitted version.

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Conflict of Interest Statement

The authors declare that the research was conducted in the absence of any commercial or

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References

1.   World Health Organization. 2020. TB disease burden, p. Chapter 4, 23. *In* Global

459   Tuberculosis Report 2020.

460   2.   Ayrapetyan M, Williams T, Oliver JD. 2018. Relationship between the Viable but

461   Nonculturable State and Antibiotic Persister Cells. J Bacteriol 200.

462   3.   Li L, Mendis N, Trigui H, Oliver JD, Faucher SP. 2014. The importance of the viable

463   but non-culturable state in human bacterial pathogens. Front Microbiol 5:258.

464   4.   Oliver JD. 2010. Recent findings on the viable but nonculturable state in pathogenic

465   bacteria. FEMS Microbiol Rev 34:415–425.

466   5.   Wayne LG, Hayes LG. 1996. An in vitro model for sequential study of shiftdown of

467   *Mycobacterium tuberculosis* through two stages of nonreplicating persistence. Infect

468   Immun 64:2062–2069.

469   6.   Betts JC, Lukey PT, Robb LC, McAdam RA, Duncan K. 2002. Evaluation of a

470   nutrient starvation model of *Mycobacterium tuberculosis* persistence by gene and

471   protein expression profiling. Mol Microbiol 43:717–31.

472   7.   Salina EG, Waddell SJ, Hoffmann N, Rosenkrands I, Butcher PD, Kaprelyants AS.

473   2014. Potassium availability triggers *Mycobacterium tuberculosis* transition to, and

474   resuscitation from, non-culturable (dormant) states. Open Biol 4.

475   8.   Aguilar-Ayala DA, Cnockaert M, Vandamme P, Palomino JC, Martin A, Gonzalez-Y-

476   Merchand J. 2018. Antimicrobial activity against *Mycobacterium tuberculosis* under in

477   vitro lipid-rich dormancy conditions. J Med Microbiol 67:282–285.

478   9.   Deb C, Lee CM, Dubey VS, Daniel J, Abomoelak B, Sirakova TD, Pawar S, Rogers L,

479   Kolattukudy PE. 2009. A novel in vitro multiple-stress dormancy model for

480   *Mycobacterium tuberculosis* generates a lipid-loaded, drug-tolerant, dormant pathogen.

481   PLoS One 4:e6077.

482   10.   Desnues B, Cuny C, Grégori G, Dukan S, Aguilaniu H, Nyström T. 2003. Differential

483   oxidative damage and expression of stress defence regulons in culturable and non-

484   culturable *Escherichia coli* cells. EMBO Rep 4:400–404.

485   11.   Imazaki I, Nakaho K. 2009. Temperature-upshift-mediated revival from the sodium-

486   pyruvate-recoverable viable but nonculturable state induced by low temperature in

487   *Ralstonia solanacearum*: linear regression analysis. J Gen Plant Pathol 2009 753

488   75:213–226.

489   12.   Mizunoe Y, Wai SN, Ishikawa T, Takade A, Yoshida S. 2000. Resuscitation of viable

490   but nonculturable cells of *Vibrio parahaemolyticus* induced at low temperature under

491   starvation. FEMS Microbiol Lett 186:115–120.

492   13.   Lynn M, Wilson AR, Solotorovsky M. 1979. Role of bovine serum albumin in the

493   nutrition of *Mycobacterium tuberculosis*. Appl Environ Microbiol 38:806–810.

494   14.   Dubos RJ. 1947. THE EFFECT OF LIPIDS AND SERUM ALBUMIN ON

495   BACTERIAL GROWTH. J Exp Med 85:9–22.

496   15.   Davis BD, Dubos RJ. 1947. THE BINDING OF FATTY ACIDS BY SERUM

497   ALBUMIN, A PROTECTIVE GROWTH FACTOR IN BACTERIOLOGICAL

498   MEDIA. J Exp Med 86:215–228.

499   16.   Mukamolova G V, Murzin AG, Salina EG, Demina GR, Kell DB, Kaprelyants AS,

500   Young M. 2006. Muralytic activity of *Micrococcus luteus* Rpf and its relationship to

501   physiological activity in promoting bacterial growth and resuscitation. Mol Microbiol

502   59:84–98.

503   17.   Mukamolova G V., Turapov O, Malkin J, Woltmann G, Barer MR. 2010.

504   Resuscitation-promoting factors reveal an occult population of tubercle bacilli in

505   sputum. Am J Respir Crit Care Med 181:174–180.

506   18.   Yeware A, Gample S, Agrawal S, Sarkar D. 2019. Using diphenyleneiodonium to

507   induce a viable but non-culturable phenotype in *Mycobacterium tuberculosis* and its

metabolomics analysis. PLoS One 14:e0220628.

19.   Roche M, Rondeau P, Singh NR, Tarnus E, Bourdon E. 2008. The antioxidant

properties of serum albumin. FEBS Lett 582:1783–1787.

20.   Shleeva M, Goncharenko A, Kudykina Y, Young D, Young M, Kaprelyants A. 2013.

Cyclic amp-dependent resuscitation of dormant mycobacteria by exogenous free fatty

acids. PLoS One 8:e82914.

21.   Shleeva MO, Kondratieva TK, Demina GR, Rubakova EI, Goncharenko A V., Apt

AS, Kaprelyants AS. 2017. OVerexpression Of adenylyl cyclase encoded by the

Mycobacterium tuberculosis Rv2212 gene confers improved fitness, accelerated

recovery from dormancy and enhanced virulence in mice. Front Cell Infect Microbiol

7:370.

22.   Knapp GS, Lyubetskaya A, Peterson MW, Gomes ALC, Ma Z, Galagan JE,

McDonough KA. 2015. Role of intragenic binding of cAMP responsive protein (CRP)

in regulation of the succinate dehydrogenase genes Rv0249c-Rv0247c in TB complex

mycobacteria. Nucleic Acids Res 43:5377–93.

23.   Schindelin J, Arganda-Carreras I, Frise E, Kaynig V, Longair M, Pietzsch T, Preibisch

S, Rueden C, Saalfeld S, Schmid B, Tinevez J-Y, White DJ, Hartenstein V, Eliceiri K,

Tomancak P, Cardona A. 2012. Fiji: an open-source platform for biological-image

analysis. Nat Methods 2012 97 9:676–682.

24.   Garton NJ, Christensen H, Minnikin DE, Adegbola RA, Barer MR. 2002. Intracellular

lipophilic inclusions of mycobacteria in vitro and in sputum. Microbiology 148:2951–

2958.

25.   Weyer K. 1998. Laboratory services in tuberculosis control.Part II. WHO Tech Bull

98:258.

26.   Trott O, Olson AJ. 2010. AutoDock Vina: improving the speed and accuracy of

docking with a new scoring function, efficient optimization, and multithreading. J

Comput Chem 31:455–61.

27.   Eberhardt J, Santos-Martins D, Tillack AF, Forli S. 2021. AutoDock Vina 1.2.0: New

Docking Methods, Expanded Force Field, and Python Bindings. J Chem Inf Model

61:3891–3898.

28.   Berman HM, Westbrook J, Feng Z, Gilliland G, Bhat TN, Weissig H, Shindyalov IN,

Bourne PE. 2000. The Protein Data Bank. Nucleic Acids Res 28:235–242.

29.   Jumper J, Evans R, Pritzel A, Green T, Figurnov M, Ronneberger O, Tunyasuvunakool

K, Bates R, Žídek A, Potapenko A, Bridgland A, Meyer C, Kohl SAA, Ballard AJ,

Cowie A, Romera-Paredes B, Nikolov S, Jain R, Adler J, Back T, Petersen S, Reiman

D, Clancy E, Zielinski M, Steinegger M, Pacholska M, Berghammer T, Bodenstein S,

Silver D, Vinyals O, Senior AW, Kavukcuoglu K, Kohli P, Hassabis D. 2021. Highly

accurate protein structure prediction with AlphaFold. Nat 2021 5967873 596:583–589.

30.   Varadi M, Anyango S, Deshpande M, Nair S, Natassia C, Yordanova G, Yuan D,

Stroe O, Wood G, Laydon A, Zídek A, Green T, Tunyasuvunakool K, Petersen S,

Jumper J, Clancy E, Green R, Vora A, Lutfi M, Figurnov M, Cowie A, Hobbs N,

Kohli P, Kleywegt G, Birney E, Hassabis D, Velankar S. 2022. AlphaFold Protein

Structure Database: massively expanding the structural coverage of protein-sequence

space with high-accuracy models. Nucleic Acids Res 50:D439–D444.

31.   Ravala SK, Singh S, Yadav GS, Kumar S, Karthikeyan S, Chakraborti PK. 2015.

Evidence that phosphorylation of threonine in the GT motif triggers activation of

PknA, a eukaryotic-type serine/threonine kinase from *Mycobacterium tuberculosis*.

FEBS J 282:1419–31.

32.   Pettersen EF, Goddard TD, Huang CC, Couch GS, Greenblatt DM, Meng EC, Ferrin

TE. 2004. UCSF Chimera - A visualization system for exploratory research and

analysis. J Comput Chem 25:1605–1612.

33.   R Core Team. 2021. R: A Language and Environment for Statistical Computing. 4.1.0.

R Foundation for Statistical Computing, Vienna, Austria.

34. Kanda Y. 2013. Investigation of the freely available easy-to-use software "EZR" for

medical statistics. Bone Marrow Transplant 48:452–8.

35. Ortega C, Liao R, Anderson LN, Rustad T, Ollodart AR, Wright AT, Sherman DR,

Grundner C. 2014. *Mycobacterium tuberculosis* Ser/Thr Protein Kinase B Mediates an

Oxygen-Dependent Replication Switch. PLOS Biol 12:e1001746.

36. Asakura H, Igimi S, Kawamoto K, Yamamoto S, Makino S-I. 2005. Role of in vivo

passage on the environmental adaptation of enterohemorrhagic *Escherichia coli*

O157:H7: cross-induction of the viable but nonculturable state by osmotic and

oxidative stresses. FEMS Microbiol Lett 253:243–9.

37. Ducret A, Chabalier M, Dukan S. 2014. Characterization and resuscitation of "non-

culturable" cells of *Legionella pneumophila*. BMC Microbiol 14:3.

38. Asakura H, Ishiwa A, Arakawa E, Makino S, Okada Y, Yamamoto S, Igimi S. 2007.

Gene expression profile of *Vibrio cholerae* in the cold stress-induced viable but non-

culturable state. Environ Microbiol 9:869–79.

39. Pasquaroli S, Zandri G, Vignaroli C, Vuotto C, Donelli G, Biavasco F. Antibiotic

pressure can induce the viable but non-culturable state in *Staphylococcus aureus*

growing in biofilms.

40. Vilhena C, Kaganovitch E, Grünberger A, Motz M, Forné I, Kohlheyer D, Jung K.

2019. Importance of Pyruvate Sensing and Transport for the Resuscitation of Viable

but Nonculturable *Escherichia coli* K-12. J Bacteriol 201.

41. Wagley S, Morcrette H, Kovacs-Simon A, Yang ZR, Power A, Tennant RK, Love J,

Murray N, Titball RW, Butler CS. 2021. Bacterial dormancy: A subpopulation of

viable but non-culturable cells demonstrates better fitness for revival. PLoS Pathog

17:e1009194.

42. Osada-Oka M, Goda N, Saiga H, Yamamoto M, Takeda K, Ozeki Y, Yamaguchi T,

Soga T, Tateishi Y, Miura K, Okuzaki D, Kobayashi K, Matsumoto S. 2019.

Metabolic adaptation to glycolysis is a basic defense mechanism of macrophages for

Mycobacterium tuberculosis infection. Int Immunol 31:781–793.

43. Schaefer WB. 1952. GROWTH REQUIREMENTS OF DYSGONIC AND EUGONIC

STRAINS OF MYCOBACTERIUM TUBERCULOSIS VAR. BOVIS. J Exp Med

96:207–219.

44. Huang BX, Kim HY, Dass C. 2004. Probing three-dimensional structure of bovine

serum albumin by chemical cross-linking and mass spectrometry. J Am Soc Mass

Spectrom 15:1237–1247.

45. Rombouts I, Lagrain B, Scherf KA, Lambrecht MA, Koehler P, Delcour JA. 2015.

Formation and reshuffling of disulfide bonds in bovine serum albumin demonstrated

using tandem mass spectrometry with collision-induced and electron-transfer

dissociation. Sci Rep 5:12210.

46. Broersen K, Van Teeffelen AMM, Vries A, Voragen AGJ, Hamer RJ, De Jongh HHJ.

2006. Do sulfhydryl groups affect aggregation and gelation properties of ovalbumin? J

Agric Food Chem 54:5166–74.

47. Hamabata T, Senoh M, Iwaki M, Nishiyama A, Yamamoto A, Shibayama K. 2021.

Induction and Resuscitation of Viable but Nonculturable *Corynebacterium*

diphtheriae. Microorganisms 9.

48. Liu J, Li L, Peters BM, Li B, Chen L, Deng Y, Xu Z, Shirtliff ME. 2017. The viable

but nonculturable state induction and genomic analyses of *Lactobacillus casei* BM-

LC14617, a beer-spoilage bacterium. Microbiologyopen 6.

49. Morishige Y, Koike A, Tamura-Ueyama A, Amano F. 2017. Induction of viable but

nonculturable *Salmonella* in exponentially grown cells by exposure to a low-humidity

environment and their resuscitation by catalase. J Food Prot 80:288–294.

50. Kong HG, Bae JY, Lee HJ, Joo HJ, Jung EJ, Chung E, Lee S-W. 2014. Induction of

612   the viable but nonculturable state of *Ralstonia solanacearum* by low temperature in the

613   soil microcosm and its resuscitation by catalase. PLoS One 9:e109792.

614 51. Kong I-S, Bates TC, Hülsmann A, Hassan H, Smith BE, Oliver JD. 2004. Role of

615   catalase and oxyR in the viable but nonculturable state of *Vibrio vulnificus*. FEMS

616   Microbiol Ecol 50:133–142.

617 52. Senoh M, Hamabata T, Takeda Y. 2015. A factor converting viable but nonculturable

618   *Vibrio cholerae* to a culturable state in eukaryotic cells is a human catalase.

619   Microbiologyopen 4:589–96.

620 53. Manuse S, Fleurie A, Zucchini L, Lesterlin C, Grangeasse C. 2016. Role of

621   eukaryotic-like serine/threonine kinases in bacterial cell division and morphogenesis.

622   FEMS Microbiol Rev 40:41–56.

623 54. Carette X, Platig J, Young DC, Helmel M, Young AT, Wang Z, Potluri L-P, Moody

624   CS, Zeng J, Prisic S, Paulson JN, Muntel J, Madduri AVR, Velarde J, Mayfield JA,

625   Locher C, Wang T, Quackenbush J, Rhee KY, Moody DB, Steen H, Husson RN. 2018.

626   Multisystem Analysis of *Mycobacterium tuberculosis* Reveals Kinase-Dependent

627   Remodeling of the Pathogen-Environment Interface. MBio 9.

628 55. Thakur M, Chakraborti PK. 2008. Ability of PknA, a mycobacterial eukaryotic-type

629   serine/threonine kinase, to transphosphorylate MurD, a ligase involved in the process

630   of peptidoglycan biosynthesis. Biochem J 415:27–33.

631 56. Parikh A, Verma SK, Khan S, Prakash B, Nandicoori VK. 2009. PknB-mediated

632   phosphorylation of a novel substrate, N-acetylglucosamine-1-phosphate

633   uridyltransferase, modulates its acetyltransferase activity. J Mol Biol 386:451–64.

634 57. Jagtap PKA, Soni V, Vithani N, Jhingan GD, Bais VS, Nandicoori VK, Prakash B.

635   2012. Substrate-bound crystal structures reveal features unique to *Mycobacterium*

636   *tuberculosis* N-acetyl-glucosamine 1-phosphate uridyltransferase and a catalytic

637   mechanism for acetyl transfer. J Biol Chem 287:39524–37.

58. Gee CL, Papavinasasundaram KG, Blair SR, Baer CE, Falick AM, King DS, Griffin
JE, Venghatakrishnan H, Zukauskas A, Wei J-R, Dhiman RK, Crick DC, Rubin EJ,
Sassetti CM, Alber T. 2012. A phosphorylated pseudokinase complex controls cell
wall synthesis in mycobacteria. Sci Signal 5:ra7.

59. Kang C-M, Nyayapathy S, Lee J-Y, Suh J-W, Husson RN. 2008. Wag31, a homologue
of the cell division protein DivIVA, regulates growth, morphology and polar cell wall
synthesis in mycobacteria. Microbiology 154:725–735.

60. Gupta M, Sajid A, Sharma K, Ghosh S, Arora G, Singh R, Nagaraja V, Tandon V,
Singh Y. 2014. HupB, a nucleoid-associated protein of *Mycobacterium tuberculosis*, is
modified by serine/threonine protein kinases *in vivo*. J Bacteriol 196:2646–57.

61. Baronian G, Ginda K, Berry L, Cohen-Gonsaud M, Zakrzewska-Czerwińska J,
Jakimowicz D, Molle V. 2015. Phosphorylation of *Mycobacterium tuberculosis* ParB
participates in regulating the ParABS chromosome segregation system. PLoS One
10:e0119907.

62. Thakur M, Chakraborti PK. 2006. GTPase activity of mycobacterial FtsZ is impaired
due to its transphosphorylation by the eukaryotic-type Ser/Thr kinase, PknA. J Biol
Chem 281:40107–13.

63. Sureka K, Hossain T, Mukherjee P, Chatterjee P, Datta P, Kundu M, Basu J. 2010.
Novel role of phosphorylation-dependent interaction between FtsZ and FipA in
mycobacterial cell division. PLoS One 5:e8590.

64. Wright DP, Ulijasz AT. 2014. Regulation of transcription by eukaryotic-like serine-
threonine kinases and phosphatases in gram-positive bacterial pathogens. Virulence
5:863–885.

65. Rock JM, Hopkins FF, Chavez A, Diallo M, Chase MR, Gerrick ER, Pritchard JR,
Church GM, Rubin EJ, Sassetti CM, Schnappinger D, Fortune SM. 2017.
Programmable transcriptional repression in mycobacteria using an orthogonal CRISPR

interference platform. Nat Microbiol 2:16274.

66.   Minato Y, Gohl DM, Thiede JM, Chacón JM, Harcombe WR, Maruyama F, Baughn

666   AD. 2019. Genomewide Assessment of *Mycobacterium tuberculosis* Conditionally

667   Essential Metabolic Pathways. mSystems 4.

668   67.   DeJesus MA, Gerrick ER, Xu W, Park SW, Long JE, Boutte CC, Rubin EJ,

669   Schnappinger D, Ehrt S, Fortune SM, Sassetti CM, Ioerger TR. 2017. Comprehensive

670   Essentiality Analysis of the *Mycobacterium tuberculosis* Genome via Saturating

671   Transposon Mutagenesis. MBio 8.

672   68.   Sassetti CM, Boyd DH, Rubin EJ. 2003. Genes required for mycobacterial growth

673   defined by high density mutagenesis. Mol Microbiol 48:77–84.

674   69.   Griffin JE, Gawronski JD, Dejesus MA, Ioerger TR, Akerley BJ, Sassetti CM. 2011.

675   High-resolution phenotypic profiling defines genes essential for mycobacterial growth

676   and cholesterol catabolism. PLoS Pathog 7:e1002251.

Figure legends

Figure 1. Growth curves of Mtb H37Rv cells in Dubos medium in the presence or absence of

BSA

(A) Number of culturable cells in Dubos medium in the presence (closed circle) or

absence (open circle) of 0.1% BSA at the indicated days of incubation. The CFU/mL

values were determined by plating cells onto 7H10 plates in duplicate. Data represent

the mean ± SD from three independent experiments.

(B) Representative images of Mtb growth captured at the end of incubation.

Figure 2. Viability analyses of DPI-treated Mtb

(A) Number of culturable cells after DPI treatment. The CFU/mL values were determined

by plating cells onto 7H10 plates in duplicate. Asterisk indicates a statistically

significant difference (*$p < 0.05$, Welch's *t*-test). Data represent the mean ± SD from

three independent experiments.

(B) Percentages of esterase active cells after DPI treatment. Around 850 cells were

directly counted under 100× objective lens. Data represent the mean ± SD from three

independent experiments.

(C) Merged fluorescence micrographs of Mtb H37Rv cells stained with

carboxyfluorescein diacetate (CFDA) and ethidium bromide (EtBr). Cells stained with

CFDA (green) are esterase-positive; those stained with EtBr (red) are membrane-

damaged. Images are representative images from three individual experiments. Scale

bar: 2μm

(D) Percentage of acid-fast-positive cells (open column) or acid-fast-negative but lipid-

accumulation-positive cells (gray column). Approximately 1,100 cells were directly

counted under 100× objective lens. Data represent the mean ± SD from four

independent experiments. Asterisk indicates significant difference (***$p < 0.005$,

Welch's *t*-test).

(E) Merged fluorescence micrographs of Mtb H37Rv cells stained with auramine-O and

Nile Red. Cells stained with auramine-O (green) are acid-fast-positive; those stained

with Nile Red (red) are acid-fast-negative but lipid accumulation-positive. Images are

representative images from three individual experiments. Scale bar: 2µm

(F) Magnified views of Airyscan confocal super-resolution microscopy observation of

three different Mtb cells obtained from DPI-treated culture. Only acid-fast-positive

cells (left), both acid-fast-positive and lipid accumulation-positive cells (center,

counted as acid-fast-positive) and acid-fast-negative cells with only lipid

accumulation-positive cells (right). Dual-positive cells probably represent the

transition to a non-culturable state. Indicated foci with arrowheads represent a neutral

lipid body. Scale bar: 1 µm

Figure 3. Reactivation of DPI-treated Mtb H37Rv cells by FBS and OADC supplementation

(A) Number of culturable cells at the indicated days of incubation, without

supplementation or those cells supplemented with 2% FBS or 10% OADC in the

presence (closed circle) or absence (open circle) of 0.1% BSA. The CFU/mL values

were determined by plating cells onto 7H10 plates in duplicate. Data represent the

mean ± SD from three independent experiments. U.D. means under the limit of

detection (20 CFU/mL).

(B) Representative images of regrowth under the indicated conditions captured at the end

of incubation. None, FBS, and OADC represent no supplementation, 2% FBS and

10% OADC supplementation, respectively.

Figure 4. Effect of human serum albumin, ovalbumin and antioxidants on the reactivation of

DPI-treated Mtb.

(A) Number of culturable cells at the end of incubation, supplemented with albumins or

antioxidative agents. The CFU/mL values were determined by plating cells onto 7H10

plates in duplicate. Data represent the mean ± SD from three independent

experiments. Asterisk indicates a statistically significant difference between Day 0

and Day 20 by Student's *t*-test. (*$p < 0.05$)

(B) Representative images of regrowth captured at the end of incubation.

Abbreviations:

BSA, bovine serum albumin; HSA, human serum albumin; OVA, ovalbumin; NAC, N-

acetyl-L-cysteine; MAN, D-mannitol.

The final concentrations of each reagent were as follows:

BSA, HSA and OVA: 0.1% (w/v), NAC: 500 µM, MAN: 50 mM.

Figure 5. The effect of SQ22536 or H89 or staurosporine on the reactivation of DPI-treated

Mtb

(A) Number of culturable cells at the end of incubation, supplemented with 0.1% BSA

and SQ22536 or H89 or staurosporine. Inhibitor was added at the indicated

concentration.

(B) Representative images of regrowth captured at the end of incubation.

The CFU/mL values were determined by plating cells onto 7H10 plates in duplicate. Data

represent the mean ± SD from three independent experiments.

Figure 6. Schematic representation of the mechanism of BSA-induced reactivation of Mtb.

Although the detailed mechanism of interaction between BSA and PknA/PknB is not yet

clear, BSA may activate PknA/PknB and positively regulate target proteins that are involved

in the localized synthesis of peptidoglycan, cell division control, and DNA condensation and

partitioning through their phosphorylation, resulting in the reactivation of VBNC Mtb cells.

(A)

(B)

BSA (-)   BSA (+)

Fig 1.

(A)

(B)

(C)

Control   DPI

(D)

(E)

Control   DPI

(F)

Fig 2.

Fig 3.

Fig 4.

(A)

(B)

Fig 5.

Fig. 6