Human and bovine serum albumin, but not mouse serum and egg-white albumin, promote reactivation of viable but non-culturable *Mycobacterium tuberculosis* via the activation of protein kinase-dependent cell division processes

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Abstract

We investigated the mechanisms underlying the fetal bovine serum (FBS)-induced reactivation

of the viable but non-culturable (VBNC) state of *Mycobacterium tuberculosis* (Mtb) using the

H37Rv strain. We found that bovine serum albumin (BSA), a major component of FBS, could

reactivate the VBNC state of Mtb cells without other reactivation-promoting agents. Human

serum albumin also had a restorative effect similar to BSA, but mouse serum albumin and egg-

white albumin (ovalbumin) did not, suggesting that a common protein structure between bovine

and human serum albumin is essential for reactivation. In addition, antioxidative agents, such

as N-acetyl-*L*-cysteine, showed no restorative effect, suggesting that the restoration of

culturability might not be due to the antioxidative property of BSA. BSA-mediated reactivation

is inhibited by H89 and staurosporine. These inhibitors are known to inhibit multiple protein

kinases, including serine/threonine protein kinases, in Mtb, suggesting the involvement of

mycobacterial protein kinases in cell division during reactivation. These findings provide new

insights into the mechanisms of Mtb reactivation, and may contribute to the development of

novel strategies for the treatment of latent tuberculosis.

Importance

Reactivation of dormant, including viable but non-culturable (VBNC) *Mycobacterium*

tuberculosis (Mtb) cells, represents a critical step in the progression from latent infection to

active tuberculosis. However, the molecular mechanisms that trigger this transition are not yet

fully understood. We herein report that bovine serum albumin (BSA), a major component of

fetal bovine serum, can restore the culturability of VBNC Mtb cells. Human serum albumin

showed a similar effect, whereas mouse and egg-white albumins did not, suggesting that the

specific structural features shared between bovine and human albumins are essential for

reactivation. The lack of reactivation by the antioxidant N-acetyl-*L*-cysteine indicates that the

mechanism is not related to redox balance. Inhibition of BSA-induced reactivation by H89 and

staurosporine implicates mycobacterial serine/threonine protein kinases in this process. These

findings highlight the previously unrecognized role of albumin-kinase signaling in Mtb

reactivation and suggest new molecular targets for preventing tuberculosis relapse.

Introduction

Mycobacterium tuberculosis (Mtb) develops a latent phenotype, dormancy, due to various

stresses, such as host immune response, oxygen depletion, and nutritional starvation, *etc*. The

Dormant Mtb cells can persist inside the host and may establish latent tuberculosis infection

(LTBI), which requires activation. Lowering the risk of Mtb activation in patients with LTBI

is an important strategy to reduce the incidence of TB [1]. The viable but nonculturable (VBNC)

state is a variant of dormancy in non-sporulating bacteria [2]. VBNC cells are a non-culturable

population that can be induced by exposure to certain stressors. They cannot grow on nutrient

media even after the removal of stress and require a much longer time or specific stimulus for

growth [2, 3]. It is increasingly being recognized that the VBNC state may play an important

role in the survival of Mtb in human hosts.

Many pathogenic bacteria, including Mtb, can enter the VBNC state and are considered to be

in a protective state in the host environment [4]. Various *in vitro* models, such as models of

hypoxia, nutritional starvation, a lipid-rich environment, and other multiple stresses, have been

reported to induce a VBNC state in Mtb cells [5–9]. A deeper understanding of the activation

mechanisms of these cells is necessary for the prevention of active tuberculosis.

One of the major hypotheses underlying the formation of VBNC bacteria involves the

oxidative damage generated by harsh external conditions [10]. Some studies have reported that

antioxidation during culture is one of the key mechanisms by which VBNC bacteria are

reactivated [11, 12]. Some mycobacterial culture media containing bovine serum albumin

(BSA) and BSA are considered to be involved in the detoxification of growth-suppressing

agents that are spontaneously generated during culture [13–15]. However, few studies have

focused on the effect of such media on the reactivation of VBNC cells or as an antioxidative

agent in mycobacterial cultures.

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

to simply and rapidly induce a VBNC state in Mtb H37Ra, and incubation with fetal bovine

serum (FBS) facilitated the reactivation of VBNC cells, whereas incubation with oleic-

albumin-dextrose-catalase (OADC) supplementation did not (16). This report provides

important clues for constructing a simple and rapid assay system to reactivate VBNC Mtb cells.

However, we found that the whole body of FBS might not be essential for the reactivation of

DPI-induced H37Rv VBNC cells, but the presence of BSA, a major component of FBS, seemed

to be sufficient to reactivate the VBNC state, as observed for both H37Rv and H37Ra. Because

FBS contains a high concentration of BSA, we hypothesized that BSA would act as a

reactivation-promoting agent for DPI-induced VBNC cells.

One possible reactivation mechanism of BSA is its antioxidative properties, which have been

well studied (17). Another possibility is cellular processes involving cAMP, an important

second messenger in the cell. Shleeva *et al*. showed that the presence of high amounts of cAMP

provided by adenylyl cyclase is essential for the reactivation of *M. smegmatis* and Mtb from

the VBNC state (18, 19). Notably, overexpression of R*v2212* gene, which encodes adenylyl

cyclase in Mtb (20), significantly affects both entry into and reactivation from the VBNC

state (19).

Understanding the detailed reactivation mechanism of VBNC is crucial for reducing the risk

of Mtb disease development. In this study, we reported that the reactivation-promoting

mechanisms of BSA against DPI-induced VBNC Mtb might not be due to its antioxidative

property or the activation of adenylyl cyclase.

Results

I.   DPI treatment induced a VBNC state in Mtb H37Rv

Yeware *et al.* previously reported that DPI treatment, which inhibits the electron transport

system, resulting in the inhibition of aerobic respiration, could effectively induce the VBNC

state in Mtb H37Ra cells (16). In our study, we confirmed that H37Ra and H37Rv cells could

effectively enter the VBNC state with DPI treatment, as evaluated by culturability, loss of acid-

fastness, and accumulation of neutral lipids. Figure 1 shows the DPI treatment results for the

induction of the VBNC state in H37Rv cells. As shown in Fig. 1(A), the numbers of culturable

cells in the 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 the DPI-treated population ($p = 0.033$). Fig. 1(B) shows that the proportion of

CFDA-positive cells in the untreated control and DPI-treated populations was not significantly

different ($81.8 \pm 11.4\%$ vs $65.9 \pm 11.8\%$, $p = 0.169$). Representative images of CFDA/EtBr

dual-stained cells from the untreated control and DPI-treated populations are shown in Fig.

1(C). These results indicate that the majority of DPI-treated H37Rv cells were alive, while their

culturability deteriorated significantly.

Auramine-O/Nile Red dual staining showed a reduction in acid fastness and accumulation

of neutral lipids. As shown in Fig. 1(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 in the untreated control and DPI-treated populations also showed distinctive

differences in the proportions of acid-fast-positive cells and lipid-accumulation-positive cells

(Fig. 1[E]). The loss of acid-fastness was also confirmed by Ziehl-Neelsen staining; the

majority of DPI-treated populations were found to have lost their acid-fastness (Suppl. Fig. 1).

Interestingly, Airyscan microscopy revealed that some DPI-treated cells were positive for acid-

fast and neutral lipid staining, suggesting the presence of cells in the transitional state from

culturable to VBNC (Fig. 1[F]). These results indicate that DPI treatment induced the VBNC

state in H37Rv and H37Ra.

II.   BSA is essential on the reactivation of DPI-induced VBNC Mtb cells.

To evaluate whether DPI-treated H37Rv cells could regain culturability by supplementing

the medium with FBS, we performed a reactivation assay on DPI-treated H37Rv cells

supplemented with FBS or OADC. As shown in Fig. 2(A) and (B), our results partially support

the previous study (16) with some differences; not only were the DPI-treated H37Rv cells

reactivated by incubation with FBS-supplemented medium, but also the OADC-supplemented

medium. Addition of albumin alone to the medium can also induce reactivation. The addition

of FBS to the BSA-free Dubos medium slightly reduced the regrowth rate; however, cells were

successfully reactivated at the end of incubation with or without FBS. This phenomenon was

also observed for H37Ra (Suppl. Fig. 2). Thus, we used H37Rv for further analyses in this

study.

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

effect on VBNC cells elsewhere (21–23), 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 growth rates under the two conditions did not differ significantly

($p = 0.428$ on day 5, Suppl. Fig. 4[A]). Supplementary Figure 4(B) also shows sufficient cell

growth. The cells were more aggregative without BSA; however, aggregation was easily

dispersed by pipetting, and there was no significant difference in the OD_{600} value. Albumin is

not essential for Mtb growth.

III.   Reactivation-promoting effect by BSA is not attributed to its antioxidative property.

It is well known that antioxidants are effective as reactivation-promoting agents for bacteria

in the VBNC state (3). We hypothesized that the reactivation-promoting effect of BSA might

be due to its antioxidative properties and assumed that human serum albumin (HSA), which

has a similar structure, and ovalbumin (OVA), which has a different structure but a stronger

antioxidative effect, as well as NAC (antioxidant) and D-mannitol (free radical scavenger),

would also reactivate VBNC Mtb. As shown in Fig. 3(A) and (B), the reactivation-promoting

effect of albumin was specific to BSA and HSA. Contrary to our expectation, OVA did not

show a reactivation-promoting effect but maintained the number of culturable cells in this

system. Contrary to our expectations, neither NAC nor D-mannitol showed reactivation-

promoting activity.

Interestingly, we found that mouse serum albumin (MSA) did not show a reactivation

promoting effect (Suppl. Fig. 5). Although only a subset of commercially available species was

analyzed, these results suggest that the reactivation promoting effect of serum albumin on DPI-

treated Mtb may be specific to certain mammalian species.

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

fatty acid- and globulin-free BSA and confirmed that there was no significant difference ($p =$

0.986) in the reactivation-promoting effects (Suppl. Fig. 6). These results suggest that

commercially available albumin, including fatty acids, does not affect reactivation of DPI-

treated Mtb.

IV.   The reactivation-promoting effect of albumin was canceled by treatment with

eukaryotic protein kinase inhibitor.

As Shleeva *et al.* reported that cAMP production from adenylyl cyclase is essential for the

reactivation of VBNC Mtb [18], we hypothesized that the inhibition of adenylyl cyclase and its

downstream enzyme, protein kinase, could modulate the reactivation-promoting effect of BSA.

As shown in Fig. 4(A) and (B), SQ22536, an adenylyl cyclase inhibitor, only suppressed

reactivation at a very high concentration (10 mM). In contrast, H89, a eukaryotic protein kinase

inhibitor, suppressed reactivation in a dose-dependent 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

± 1.70 × 10² CFU/mL, while incubation without H89 resulted in 3.00 × 10⁸ ± 5.07 × 10⁷

CFU/mL. We also confirmed that staurosporine, a known mycobacterial protein kinase PknB

inhibitor (24), suppressed reactivation at a concentration of 10 μM and resulted in a CFU/mL

value of 2.09 × 10³ ± 7.62 × 10² CFU/mL, whereas incubation without staurosporine resulted

in 3.58 × 10⁸ ± 5.78 × 10⁷ CFU/mL (Fig. 4[A] and [B]). None of the inhibitors used in this

study reduced the growth of intact Mtb cells (Suppl. Fig. 7A and 7B).

Since both SQ22536 and H89 are targeted to eukaryotic enzymes, we also performed a

molecular docking simulation of these inhibitors against their equivalent targets on Mtb to

validate the inhibition assay. As shown in Supplemantary Table 1, SQ22536 and H89 have

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

Discussion

In this study, we found that BSA, a supplement to mycobacterial culture, could act as a

reactivation-promoting agent of the VBNC state of Mtb. We also showed that this effect of

serum albumin might be species-specific: bovine and human serum albumin had the effect,

while mouse serum albumin did not. The predictive mechanism of BSA-induced reactivation

was also proposed to involve the activation of mycobacterial protein kinase, not its

antioxidative property or the activation of adenylyl cyclase.

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

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

induce an unbalanced NADH/NAD ratio and hypoxic response in liquid culture, resulting in a

significant decrease in culturability (Fig. 1[A]) (16). However, viability was highly retained

(65.9%), suggesting that the majority of the cells were transformed into VBNC (Fig. 1[B]).

VBNC transformation was confirmed by auramine-O and Nile Red staining. The loss of acid-

fastness and 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. 1[D]). Airyscan super-resolution microscopy showed the presence of

cells stained with auramine-O, which contained an intracellular lipid body stained with Nile

Red (Fig. 1[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 a transformation from a growing state to

VBNC, with drastic alterations in lipid metabolism.

We also found that DPI-induced VBNC could facilitate reactivation not only by incubation

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

act as a reactivation-promoting agent in both H37Rv and H37Ra (Fig. 2 and Suppl. Fig. 2).

These findings were contrary to those of a previous study, which showed that only FBS can

facilitate reactivation (16). We considered that the reason for this difference may be due to a

contaminant in Tween 80: the presence of an excessive amount of oleic acid. We also

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

preparation of Dubos medium (unpublished data) for both H37Rv and H37Ra, probably due to

extended hydrolyzation of Tween 80 (25). Thus, our results suggest that the key reactivation-

promoting agent in FBS is serum albumin, and further analyses were performed using H37Rv.

It should be noted that not all cells completely transitioned to the VBNC state in this study.

In general, *in vitro* cultured cells are thought to be a non-uniform population due to the

difference in microenvironment; oxygen concentration and nutritional conditions might vary

at the local level, which could make the variable subpopulations in metabolic state, as Yeware

et al. also indicated (16). In our study, a small subpopulation (10^3 CFU/mL) was found that

could not proliferate on Dubos broth without BSA but could proliferate on agar medium (Fig.

2). The results of acid-fast staining also suggested the presence of cells that appeared to be in

the process of transitioning from a culturable state to a VBNC state (Fig. 1F). These results

suggest that, although the VBNC state could be induced by DPI treatment, the level of

induction might vary from cell to cell. Therefore, the results of an *in vitro* VBNC model should

be carefully interpreted and the final conclusions must be verified in *in vivo* model with

albumin administration.

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

specific because OVA did not facilitate reactivation, whereas BSA and HSA did. Regarding

their amino acid sequences, BSA and HSA share 76% sequence homology (26); however, BSA

and OVA share less sequence homology (16%). One of the considerable roles of albumin is as

an antioxidative agent because of its cysteine residues with free thiols (17, 27). BSA and HSA

have only one cysteine residue with a free thiol, whereas OVA has four cysteines with free

thiols (28). Although OVA therefore seems to be more reductive than BSA and HSA, their

reactivation-promoting effect is paradoxical, indicating that antioxidative properties might not

be essential for this effect. This hypothesis was considered to support the reduced effect of

NAC and D-mannitol. Moreover, this paradoxical result confirmed that certain common

structures of BSA and HSA may be essential. Additionally, we would like to note that MSA

did not show the reactivation promoting effect, suggesting that this effect might have species-

dependency (Suppl. Fig. 5). At this point, we have found that there are certain differences

between bovine and mouse serum albumins (internal data) and analyzed their contribution to

this property for the subsequent study, including measurement of the reactivation-promoting

effect of MSA on VBNC *M. microti*, a pathogen of small field rodents (29). These results

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

properties.

Serum albumin is known to act as a good carrier for many biologically active substances,

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

culture medium. BSA is added to the 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 acids, including oleic, linoleic, and

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

medium, and oleic acid also works in Mtb (18). It should be noted that albumin impurities did

not affect reactivation in this study. Although the underlying mechanism is still unclear, both

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

promoting effect toward DPI-treated Mtb (Suppl. Fig. 6).

The reactivation inhibition assay using SQ22536, H89, and staurosporine provided an

important clue for understanding the effect of BSA. In the present study, we suppressed the

reactivation of DPI-induced VBNC cells by inhibiting adenylyl cyclase at 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, whereas the inhibition

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

suggest that protein kinase might be involved in the reactivation of VBNC Mtb cells triggered

by BSA, and that the contribution of adenylyl cyclase may be smaller than that of protein kinase.

Mtb encodes eukaryotic Ser/Thr protein kinases including PknA, PknB, and PknD-L.

Among them, 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 (30, 31); phosphorylation of GlmU (Rv*1018c*) and MurD (Rv*2155c*),

Wag31 (Rv*2145c*), HupB (Rv*2986c*) and ParB (Rv*3917c*), influence the production and cross-

linking of peptidoglycan precursors (32–35), the spatial localization of peptidoglycan

synthesis(36), DNA condensation and partitioning (37, 38), respectively. Notably, the

phosphorylation of FtsZ and FipA by PknA controls cell division under oxidative stress(39, 40).

In addition, mycobacterial protein kinase has cyclic AMP-dependent protein kinase domains,

which are very similar to the eukaryotic version, which was shown in PknB (41), and

staurosporine, which acts as a PknB inhibitor, could also inhibit the reactivation of DPI-induced

VBNC Mtb (Fig. 4).

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

staurosporine critically affects several important cellular processes that underlie reactivation

(Suppl. Fig. 8). To confirm the role of mycobacterial protein kinases and related proteins in the

reactivation of DPI-induced VBNC cells by albumin, further studies involving gene

knockdown using CRISPR interference (CRISPRi) (42) are necessary. This is particularly

relevant because most of these proteins, except FipA (Rv*0019c*), are essential for *in vitro*

growth; (43–46) therefore, genetic manipulation using conventional methods is not permissible.

Conclusion

Our data showed that the presence of BSA in Dubos medium not only depleted growth-

suppressing substances, but also triggered the reactivation of the DPI-induced VBNC state in

both H37Rv and H37Ra strains. The reactivation-promoting effects of other kinds of albumins

and antioxidative agents, including NAC and D-mannitol, were tested; however, only bovine

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

assay of reactivation using SQ22536, H89, and staurosporine suggest that the inhibition of

mycobacterial protein kinase suppressed reactivation to a much greater extent than adenylyl

cyclase, indicating the presence of a reactivation-promoting protein kinase-dependent pathway.

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

of albumin on 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 provides some important clues for understanding the physiology

of the mycobacterial VBNC state.

Author Contributions

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

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

experiments. YMori acquired the data, and YMori and YMura analyzed the data. YMori

performed a molecular docking simulation and statistical analysis. YMori, YMura, and SM

wrote the manuscript. All authors have contributed to the manuscript and approved the

submitted version.

Data availability

All data contained in this study can be obtained from the corresponding author upon reasonable

request.

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Conflict of Interest Statement

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

financial relationships that could be construed as potential conflicts of interest.

Materials and Methods

Bacterial strains, growth media and culture conditions

This study 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, 0.01 g $MgSO_4 \cdot 7H_2O$) 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 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 three

weeks of incubation at 37 °C under 5% CO_2. The limit of detection was determined to be 20

CFU/mL owing to the diluent factor.

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

Ten milliliters of log phase culture of Mtb H37Rv (OD_{600} = 0.35) in Dubos medium, as described previously, was sedimented at 3,000 $\times g$ for 10 min at 4 °C, and the supernatant was discarded. The sediment was washed twice with fresh Dubos broth and resuspended in 10 mL Dubos broth. The suspension was diluted 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 a 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 was performed according to a previous report (16) with some modifications. In short, the log phase culture of Mtb H37Rv or H37Ra (OD_{600} = 0.35) in Dubos medium was evenly divided into 30 mL PETG bottles, and 5 mg/mL diphenyleneiodonium chloride (DPI) in dimethyl sulfoxide (DMSO) solution was directly added at a final concentration of 4 µg/mL with immediate agitation. An equal volume of DMSO was added to the untreated control samples. The cultures were incubated for 24 h at 37 °C without shaking. After incubation, their culturability and viability were measured as described above.

Cell viability assay

Mtb cells was measured by esterase activity and membrane integrity of the cells using dual staining with 10 mg/mL carboxyfluorescein diacetate (CFDA, DOJINDO LABORATORIES, Kumamoto, Japan) and ethidium bromide (EtBr, DOJINDO LABORATORIES). Briefly, 500 µL of DPI-treated or untreated Mtb culture was sedimented by centrifugation at 15,000 $\times g$ for 5 min at 4 °C and washed twice with an equal volume of PBS-T. The washed bacterial sediment

was resuspended in 200 µL of PBS-T with CFDA and EtBr and incubated at room temperature

(RT) for 30 min in the shade. The final concentrations of CFDA and EtBr were 60 µg/mL and

2 µg/mL, respectively. 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 the bacterial suspension

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

The slides were then mounted with DPX new non-aqueous mounting medium (Merck,

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

to microscopic observation. Specimens were examined using a BX53 fluorescence microscope

(Olympus, Tokyo, Japan) with a 100× objective lens for green (esterase-active with intact

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

At least 10 random fields were observed for each sample and the images (TIFF format) were

analyzed using the Fiji/ImageJ software program[47]. The live/dead ratio was calculated by

directly counting green or red fluorescent cells.

Auramine-O/Nile Red dual staining

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

mycobacteria[48]. 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 modifications. Ten microliters

of DPI-treated or untreated cultures were smeared onto an APS-coated slide (Matsunami Glass,

Osaka, Japan) and air-dried. The slides were then heat-fixed and cooled 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 the

dark. The excess dye was removed using 3% hydrochloric acid in ethanol for 15 min. The

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

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

solution for 1 min. The stained slides were air-dried and mounted, as previously described.

Fluorescence microscopy was performed with a 100× objective lens for green (acid-fastness

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

positive; VBNC) fluorescent cells. At least five random fields were observed for each sample,

and images (TIFF format) were analyzed using the Fiji/ImageJ software program.(47) The

proportion of acid-fast-positive cells and lipid-accumulation-positive cells was calculated by

direct counting of green or red fluorescent cells.

Acid-fast 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(49). The slides were observed

under a light microscope with a 100× objective lens (BX53, Olympus, Tokyo, Japan).

Confocal super-resolution microscopy

Detailed images of accumulated lipids were obtained using 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 with a 63× objective lens. The green fluorescence emission from auramine-O

and red fluorescence from Nile Red were sequentially collected using a variable dichroic beam

splitter and an Airyscan 2 detector. Super-resolution images were processed using ZEN Blue

software (Carl Zeiss, Jena, Germany).

VBNC reactivation assay

The VBNC reactivation assay was performed as described previously (16) with some

modifications. DPI-treated cultures were sedimented by centrifugation at 3,000 x*g* for 10 min

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

The bacterial suspension was diluted 1:10 with fresh Dubos medium, with or without 0.1%

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

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

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

of each reagent was 10% (v/v) for OADC supplementation, 2% (v/v) for FBS, and 3 mM for

PA. The diluted suspension was then dispensed into 24-well plates in 1.5 mL increments and

sealed with a 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 previously

described.

Evaluation of the reactivation-promoting effect of human and mouse 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 that have variable

antioxidative properties and antioxidative agents toward DPI-treated Mtb cells. Washed DPI-

treated cells were resuspended in fresh Dubos broth without BSA, and the suspension was

diluted 1:10 with Dubos broth containing BSA, human serum albumin (HSA), mouse serum

albumin (MSA), 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, MSA, and OVA; 500 μM

for NAC; and 50 mM for MAN. The subsequent reactivation assay was performed according

to the method described above.

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 in fresh Dubos broth without BSA, and the

suspension was diluted 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 the BSAs was 0.1% (w/v). The subsequent reactivation assay was

performed according to the method described above.

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

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

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

adenylyl cyclase inhibitor SQ22536 (Merck, Darmstadt, Germany), protein kinase inhibitor

H89 (Abcam, Cambridge, United Kingdom), or staurosporine (Merck). The final

concentrations of each inhibitor were as follows: 0.1 to 10 mM SQ22536, 1–30 μM H89 and 1

to10 μM for staurosporine. The number of culturable cells was then measured as described

previously.

Molecular docking simulation

Molecular docking simulation was performed using the AutoDock Vina software program

(version 1.1.2), an open-source molecular docking program designed by the Scripps Research

Institute(50, 51). The 3D structures of the receptor proteins PknA and Rv*2212* were obtained

from the Protein Data Bank (52) and AlphaFold Database(53, 54), respectively. The PDB ID for

PknA was 4O*W*8 (55) and the AlphaFold Database identifier for Rv*2212* was AF-P9WMU7-

F1. We removed water molecules from the proteins, added polar hydrogen and charge, and

adjusted the X-Y-Z coordinates and grid size to optimize molecular docking. Each simulation

was performed using default options. The 3D structures of the ligand molecules, H89, SQ22536

and ATP were obtained from the PubChem Database. The PubChem CIDs of the ligands were

449241, 5270, and 5957, respectively. Graphical processing was performed using the UCSF

Chimera software program version 1.16(56).

Statistical analysis

The statistical significance of differences between culturable cell counts of Mtb cells in

untreated and DPI-treated cells was determined by Student's *t*-test. The statistical significance

of differences between staining of Mtb cells in untreated and DPI-treated cells was determined

using Welch's *t*-test. The statistical significance of differences between culturable cell counts

of DPI-treated Mtb cells at days 0 and 20 on reactivation was determined by the Student's *t*-

test. For multiple group comparisons, one-way analysis of variance (ANOVA) was performed,

followed by Tukey's honest significant difference (HSD) test for multiple comparisons. Only

comparisons with the control group (day 0) were extracted. Statistical analysis was performed

with R (57) using EZR in the R package(58). $P < 0.05$.

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Figure legends

Figure 1. Viability analyses of DPI-treated Mtb H37Rv

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

by plating the cells onto 7H10 plates in duplicate. Asterisks indicate statistically

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

three independent experiments.

(B) Percentage of esterase-active cells after DPI treatment. Approximately 850 cells were

counted directly using a 100× objective lens. Data represent mean ± SD from three

independent experiments.

(C) Fluorescence micrographs of Mtb H37Rv cells stained with carboxyfluorescein

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

were esterase-positive, and those stained with EtBr (red) were membrane-damaged.

Representative images from three individual experiments are shown. Scale bar: 2 μm

(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 counted

directly using a 100× objective lens. Data represent mean ± SD from four independent

experiments. Asterisks indicate significant differences (***$p < 0.005$, 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) were acid-fast-positive; those stained

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

Representative images from three individual experiments are shown. Scale bar: 2μm

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

three different Mtb cells obtained from DPI-treated cultures. 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 a transition

to a non-culturable state. The indicated foci with arrowheads represent the neutral

lipid bodies. Scale bar: 1 μm

Figure 2. Reactivation of DPI-treated Mtb H37Rv cells by FBS and OADC supplementation

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

supplementation, or 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 the cells onto 7H10 plates in duplicate. Data represent mean ±

SD from three independent experiments. U.D. indicates the limit of detection (20

CFU/mL).

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

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

supplementation, and 10% OADC supplementation, respectively.

Figure 3. Effect of human serum albumin, ovalbumin and antioxidants on the reactivation of

DPI-treated Mtb.

(A) The culturable cells at the end of incubation were supplemented with albumins or

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

7H10 plates in duplicate. Data represent mean ± SD from three independent

experiments. Asterisks indicate statistically significant differences between days 0 and

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; and MAN, 50 mM.

Figure 4. 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,

SQ22536, H89, or staurosporine. Inhibitors were added at the indicated

concentrations.

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

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

represent mean ± SD from three independent experiments.

Fig 1.

(A)

None | FBS | OADC

Culturable cells (CFU/mL)

U.D.

Regrowth (days)

(B)

None | FBS | OADC

BSA (-)

BSA (+)

Fig 2.

(A)

(B)

Fig 3.

(A)

(B)

SQ
22536
(mM)

H89
(µM)

Stauro-
sporine
(µM)

Fig 4.