

# Improving of pelB-Secreted MPT64 protein released by *Escherichia coli* BL21 (DE3) using Triton X-100 and Tween-80

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## ABSTRACT

pelB has been known as a successful signal peptide to translocate the protein target extracellularly in the *Escherichia coli* system. However, in our previous study, the yield of MPT64 protein extracellular recovery was still low and plenty of this protein was remain trapped in cytoplasm and periplasm. Recently, nonionic surfactants were efficiently reported to secrete recombinant protein extracellularly. Nonetheless, it must be clarified whether the surfactant supplementation can improve the yield of MPT64 extracellular protein significantly without giving impact on the structure of isolated MPT64 protein and can minimized the cell lysis effect. MPT64 protein secretion was carried out by comparing the effects of surfactants Tween 80 and Triton × 100 at various concentrations. Triton × 100 was able to increase the extracellular MPT64 protein gain up to 3 times higher than Tween 80 and it was in line with the greater level ratio of cell leakage of Triton × 100 compared to that of Tween 80. Similarly, the viable cell of the cultures decreased dramatically. However, both surfactants did not interfere the structure of MPT64 protein. In conclusion, Triton × 100 can be chosen as the supporting surfactant to assist the act of peptide signal in improving the resulting of MPT64 extracellular protein.

**Key words:** Extracellular, MPT64, pelB, Triton × 100, Tween 80

## INTRODUCTION

The strategy to evade the host's immune system is an important virulence factor of *Mycobacterium tuberculosis* in maintaining the multiplication of this bacterium in infected host cells. Several *M. tuberculosis* secreted proteins facilitate this bacterium by regulating antiapoptotic and

proapoptotic molecules to inhibit the apoptosis of infected macrophages.<sup>[1,2]</sup> MPT64 is one of the proteins secreted from *M. tuberculosis* with antiapoptosis mechanism on RAW264.7 macrophages by upregulating the Bcl-2, miRNA21, and nuclear factor-kappa B (NF-κB).<sup>[2]</sup> The most interesting of this protein is its specification to distinguish the *M. tuberculosis* complex from other mycobacteria bacterial species. This protein has great potential for *M. tuberculosis* identification using immune chromatographic methods through the antigen-antibody binding process.<sup>[3,4]</sup> Hence, this protein must be present in adequate concentration to generate antibodies against MPT64 to be constructed in immunochromatographic diagnostic kit. However, MPT64

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is difficult to be conventionally isolated from the original source due to the abundant of other secreted tuberculosis proteins which difficult to purify, and it is grouped as high-risk air infect pathogen.

In our previous study, this protein was produced by expressing the synthetic gene coded for the MPT64 protein in *Escherichia coli* BL21 (DE3) and secreted extracellularly by constructing the gene fused with pelB signal peptide.<sup>[5,6]</sup> We found that the pelB success to translocate the MPT64 extracellularly, but the yield was still low and plenty of this protein was remain trapped in cytoplasm and periplasm. Thus, to enhance the secretory potential, the growth medium of *E. coli* BL21 (DE3) was augmented with surfactants, which allows the production of highly efficient extracellular target proteins in *E. coli* but does not impair the activity of the target protein.<sup>[7,8]</sup> Triton × 100 and Tween 80 were reported effectively secreted protein targets in high levels compared to other surfactants such as CaCl<sub>2</sub>, sodium dodecyl sulfate (SDS), and glycine in *E. coli*.<sup>[8]</sup> Therefore, in this study, the optimized conditions related to the secretion process of MPT64 protein were conducted and the extracellular MPT64 structure was evaluated to validate the structural integrity of the protein required to maintain antigen epitopes for antibody production.

## MATERIALS AND METHODS

### Materials

The materials used are *E. coli* BL21 (DE3) transformant, Luria Bertani (LB) (Himedia), L-rhamnose (Sigma-Aldrich), Tween 80 (Sigma-Aldrich), Triton × 100 (Sigma-Aldrich), and SDS PAGE components.

### Optimization of recombinant MPT64 protein secretion

Optimization of MPT64 protein secretion was carried out by comparing the effects of surfactants Tween 80 and Triton × 100 at various concentrations as follows: 0%, 0.1%, 0.2%, and 0.5%. A total of 6 mL of bacterial starter were inoculated into four Erlenmeyer flasks containing 294 mL of liquid LB, then 300 L of kanamycin (100 g/mL) were added to each of the Erlenmeyer flasks and incubated at 37°C, 180 rpm, optical density OD<sub>600</sub> (0.6–0.8). After that, the bacterial culture was measured for its OD at a wavelength of 600 nm (t<sub>0</sub>) and 10 mL of the bacterial culture was taken and centrifuged at 6000 g at 4°C for 20 min. The cell culture was then induced with the addition of Rhamnose to a concentration of 4 mM, then incubated again at 37°C, 180 rpm for 24 h. After 24 h, the OD<sub>600</sub> of bacterial culture was remeasured (t<sub>24</sub>) and 10 mL of cell culture was taken for centrifugation at a speed of 6000 g at 4°C for 20 min. Then, the surfactant was added to each Erlenmeyer flask with each concentration of 0, 0.1, 0.2, and 0.5%, and incubated at 37°C, 100 rpm for 6 h. After that, the bacterial culture OD<sub>600</sub> was measured again (t<sub>30</sub>) and 10 mL of cell culture was taken and centrifuged at 6000 g

at 4°C for 20 min. Before the isolation of the media fraction, phenylmethylsulfonyl fluoride was added to each culture until the final concentration was 0.1 mM. A total of 10 mL of the culture taken at t<sub>0</sub>, t<sub>24</sub>, and t<sub>30</sub> were centrifuged to separate the supernatant from the pellets. Proteins in the media fraction were characterized using the SDS-PAGE method and quantified using ImageJ software.

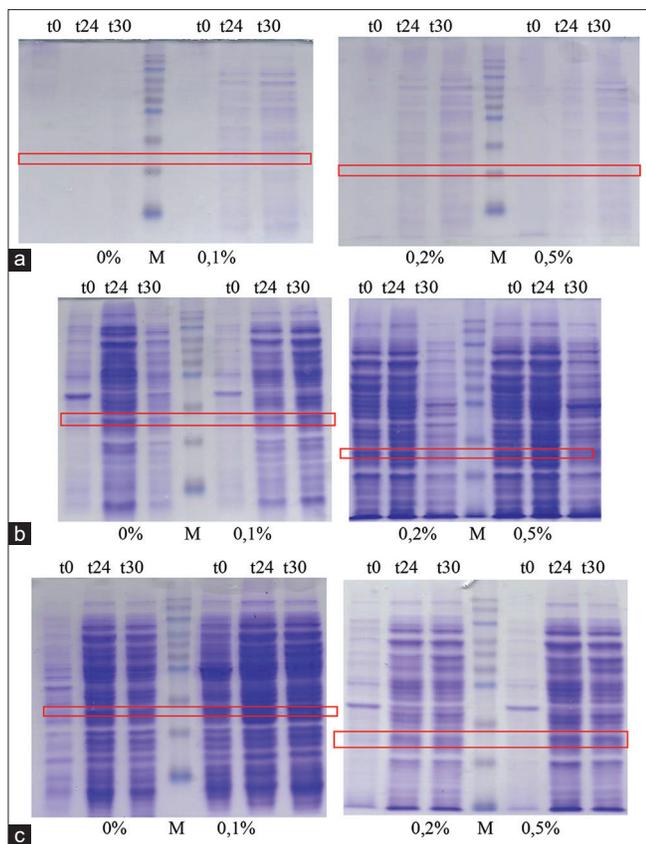
### Effects of surfactants

Ten milliliters of overproduced cultures at t<sub>0</sub>, t<sub>24</sub>, and t<sub>30</sub> were centrifuged at 10,000 g, 4°C, for 5 min. Then, its absorbance at OD<sub>260</sub> was checked to measure the DNA concentration in the transformant cultures. The higher the absorbance, the higher the concentration of DNA in the sample, which indicates the number of cells undergoing lysis. As confirmation of cell viability, each sample at t<sub>0</sub>, t<sub>24</sub>, and t<sub>30</sub> was inoculated as much as 10 L on the surface of solid LB medium containing kanamycin. The culture on solid media was then incubated at 37°C for 24 h and the number of colonies observed was counted.

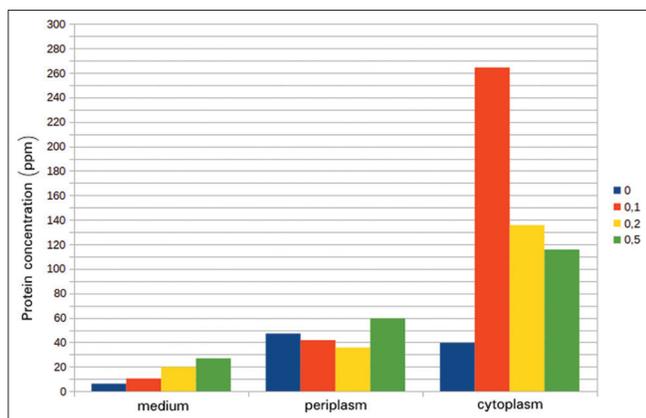
## RESULTS AND DISCUSSION

In our previous study, the optimal conditions for MPT64 gene expression have been conducted, but the acquisition of extracellular MPT64 protein as a protein target was still low.<sup>[9]</sup> In line with this, it has been widely reported that the challenge to express extracellular proteins in *E. coli* is always distributed in the periplasm.<sup>[10]</sup> To overcome this drawback, the acquisition of extracellular MPT64 protein can be increased by improving the leakage of specific host cell membranes by constructing the cell host into a leaky phenotype. Such phenotypes can be induced through several ways, as follows: mutation or deletion of membrane components, addition of permeability enhancers (glycine, calcium, Triton X-100, etc.), or coexpression of proteins with lytic activity.<sup>[11-15]</sup>

Triton × 100 and Tween 80 are the most widely used nonionic surfactants to increase the permeability of living cell membranes.<sup>[16-20]</sup> Therefore, this evidence based bring this study to investigate the effect of both surfactants in improving the yield of MPT64 extracellular recovery. As seen in Figures 1 and 2, the thickness of the protein band in each cell fraction described the different protein concentrations at various surfactant concentrations. The protein concentrations were demonstrated in Tables 1 and 2 and the different effects of each surfactant in various concentrations on the MPT64 extracellular protein recovery can be described clearly in Figures 3 and 4. The supplementation of Triton × 100 (0.5% v/v) into the growth medium could increase the extracellular MPT64 protein gain up to 3 times higher than using Tween 80 at the same concentration. The different effects of both surfactants on membrane permeability were predicted by the fact that nonionic surfactants do not dissociate when dissolved in

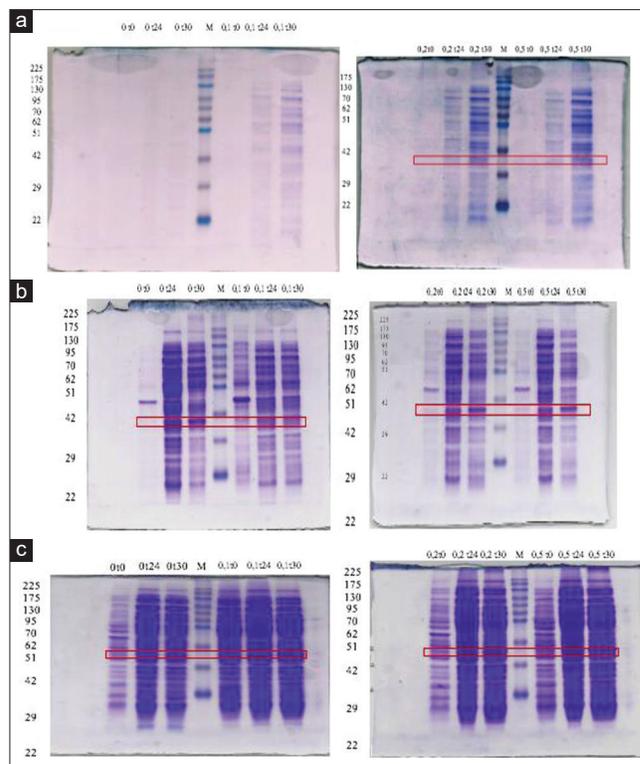


**Figure 1:** Effect of Tween 80 on MPT64 protein gain in different fractions (a) medium (b) periplasm (c) cytoplasm

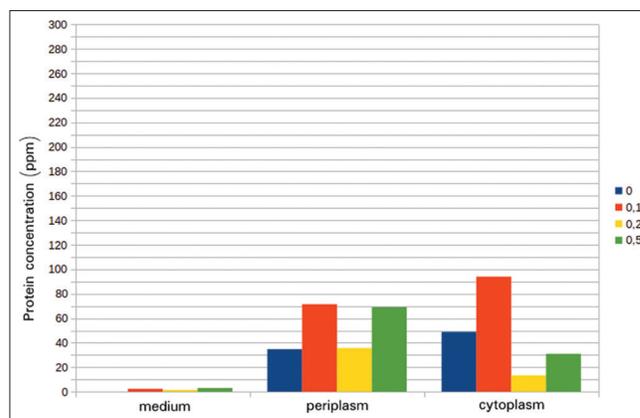


**Figure 3:** Distribution of MPT64 production after Triton x 100 treatment in different concentrations

water and have the widest range of properties depending on the hydrophilic-lipophilic equilibrium ratio (HLB).<sup>[21]</sup> If the HLB value of the surfactant is higher than 10, then the nonionic surfactant tends to be more hydrophilic. As a result, the surfactant layer lowers the surface tension of the water more than the surface tension of the oil. Surfactants with low HLB values are more soluble in oil (lipophilic), while surfactants with higher HLB values are more soluble in water (hydrophilic). This supports the evidence that Triton x 100 (HLB = 13.5) with a lower HLB value than



**Figure 2:** Effect of Triton x 100 on MPT64 protein gain in different fractions (a) medium (b) periplasm (c) cytoplasm



**Figure 4:** Distribution of MPT64 production after Tween 80 treatment in different concentrations

Tween 80 (HLB = 15.0) is able to interact more strongly with bacterial cell membranes composed of phospholipids.<sup>[22]</sup> In addition, we must confirm the effect of the surfactants to the MPT64 structure. From Figure 5, we can see that the antibody of the kit diagnostics can capture the MPT64 successfully. It is indicated that both surfactants did not interfere the MPT64 structure. Thus, the overproduction of MPT64 protein can be combined with Triton x 100 surfactant at the late stage of incubation.

However, the concentration used must be considered because several studies have reported the occurrence of cell

death after prolonged exposure to these surfactants.<sup>[23-25]</sup> From this study, both surfactants were shown to be able to release higher extracellular MPT64 protein compared to untreated cells, presented in Table 3. The level ratio of cell leakage between Triton × 100 (0.5%) and Tween 80 (0.5%) against the control cell was 86.07% and 66.35%, respectively. From Table 4, at the 30<sup>th</sup> h after the addition of surfactant,

the culture added with Triton × 100 experienced a large decrease number of viable cells for about 8%–20% at 24 h. Therefore, we hypothesized that both surfactants could interact with the cell membrane, but the high level of MPT64 protein is still trapped in the cytoplasm and periplasm. We predicted that this was due to the ineffectiveness of the pelB signal peptide used in translocating MPT64 protein



Figure 5: The identification of secreted MPT64 protein influenced by (A) Triton-×100; (B) Tween 80 (a) cytoplasm (b) periplasm, (c) medium

Table 1: Effect of Tween 80 on MPT64 protein levels

| Surfactant concentration (%v/v) | Time (h) | Quantification of protein content at various concentrations of Tween 80 |                     |           |                     |           |                     |
|---------------------------------|----------|---|---------------------|-----------|---------------------|-----------|---------------------|
|                                 |          | Medium  |                     | Periplasm |                     | Cytoplasm |                     |
|                                 |          | Band area   | Concentration (ppm) | Band area | Concentration (ppm) | Band area | Concentration (ppm) |
| 0                               | t0       | 0   | 0.00±0.000          | 22.282    | 22.48±0.002         | 23.717    | 24.16±0.001         |
|                                 | t24      | 0   | 0.00±0.000          | 78.863    | 88.61±0.001         | 50.809    | 55.82±0.002         |
|                                 | t30      | 0   | 0.00±0.000          | 32.805    | 34.78±0.002         | 44.78     | 48.78±0.001         |
| 0.1                             | t0       | 0   | 0.00±0.000          | 16.834    | 16.11±0.001         | 51.879    | 57.07±0.002         |
|                                 | t24      | 4.108   | 1.24±0.004          | 54.446    | 60.07±0.004         | 68.531    | 76.54±0.001         |
|                                 | t30      | 5.07  | 2.37±0.001          | 64.184    | 71.46±0.001         | 83.422    | 93.94±0.001         |
| 0.2                             | t0       | 0   | 0.00±0.000          | 54.633    | 60.29±0.002         | 53.396    | 58.85±0.001         |
|                                 | t24      | 3.997   | 1.11±0.001          | 52.447    | 57.74±0.001         | 62.834    | 69.88±0.001         |
|                                 | t30      | 4.254   | 1.41±0.002          | 33.571    | 35.68±0.001         | 14.481    | 13.36±0.001         |
| 0.5                             | t0       | 0   | 0.00±0.000          | 60.404    | 67.04±0.001         | 56.267    | 62.20±0.000         |
|                                 | t24      | 3.906   | 1.00±0.001          | 67.631    | 75.48±0.002         | 60.856    | 67.57±0.001         |
|                                 | t30      | 5.633   | 3.02 ± 0.004        | 62.166    | 69.10 ± 0.001       | 29.476    | 30.89 ± 0.001       |

Table 2: Effect of Triton X-100 on MPT64 protein levels

| Surfactant concentration (%v/v) | Time (h) | Quantification of protein content at various concentrations of Triton X-100 |                     |           |                     |           |                     |
|---------------------------------|----------|---|---------------------|-----------|---------------------|-----------|---------------------|
|                                 |          | Medium  |                     | Periplasm |                     | Cytoplasm |                     |
|                                 |          | Band area   | Concentration (ppm) | Band area | Concentration (ppm) | Band area | Concentration (ppm) |
| 0                               | t0       | 1.66  | -0.89±0.001         | 3.342     | 2.10±0.000          | 12.474    | 18.36±0.004         |
|                                 | t24      | 5.391   | 5.75±0.001          | 32.794    | 54.53±0.001         | 51.287    | 87.45±0.001         |
|                                 | t30      | 5.648   | 6.20±0.000          | 28.631    | 47.12±0.000         | 24.374    | 39.54±0.004         |
| 0.1                             | t0       | 1.478   | -1.21±0.001         | 12.503    | 18.41±0.001         | 16.521    | 25.56±0.004         |
|                                 | t24      | 5.675   | 6.25±0.000          | 29.644    | 48.92±0.002         | 81.168    | 140.65±0.001        |
|                                 | t30      | 8.052   | 10.48±0.002         | 25.595    | 41.71±0.001         | 150.719   | 264.47±0.001        |
| 0.2                             | t0       | 4.541   | 4.23±0.001          | 12.091    | 17.67±0.001         | 16.452    | 25.44±0.000         |
|                                 | t24      | 12.859  | 19.04±0.001         | 23.695    | 38.33±0.001         | 69.303    | 119.53±0.001        |
|                                 | t30      | 13.327  | 19.87±0.001         | 22.226    | 35.72±0.002         | 78.345    | 135.63±0.001        |
| 0.5                             | t0       | 5.191   | 5.39±0.001          | 10.984    | 15.70±0.000         | 33.447    | 55.69±0.001         |
|                                 | t24      | 8.925   | 12.04±0.001         | 34.819    | 58.14±0.001         | 68.55     | 118.19±0.001        |
|                                 | t30      | 17.257  | 26.87 ± 0.001       | 62.166    | 59.47 ± 0.001       | 67.15     | 115.69 ± 0.001      |

**Table 3: Release of intracellular components analysis**

| Sample | Surfactant  | DNA concentration ( $\mu\text{g/ml}$ ) |
|--------|-------------|--|
| 1      | 0%          | $3.55 \pm 0.001$                       |
| 2      | Triton 0.1% | $14.55 \pm 0.001$                      |
| 3      | Triton 0.2% | $15.90 \pm 0.000$                      |
| 4      | Triton 0.5% | $25.50 \pm 0.001$                      |
| 5      | Tween 0.1%  | $6.10 \pm 0.002$                       |
| 6      | Tween 0.2%  | $8.60 \pm 0.001$                       |
| 7      | Tween 0.5%  | $10.55 \pm 0.000$                      |

**Table 4: Effect of surfactants on the viable cell**

| Surfactant  | Incubation time (h)   |                       |                       |
|-------------|-----------------------|-----------------------|-----------------------|
|             | 0                     | 24 <sup>th</sup>      | 30 <sup>th</sup>      |
| 0%          | $2.05 \times 10^{11}$ | $4.59 \times 10^{12}$ | $3.84 \times 10^{11}$ |
| Triton 0.1% | $2.05 \times 10^{11}$ | $2.91 \times 10^{12}$ | $5.10 \times 10^{10}$ |
| Triton 0.2% | $3.84 \times 10^{11}$ | $4.14 \times 10^{12}$ | $4.46 \times 10^{10}$ |
| Triton 0.5% | $2.78 \times 10^{11}$ | $2.75 \times 10^{12}$ | $7.10 \times 10^{10}$ |
| Tween 0.1%  | $2.53 \times 10^{11}$ | $2.46 \times 10^{12}$ | $5.00 \times 10^{11}$ |
| Tween 0.2%  | $3.66 \times 10^{11}$ | $2.68 \times 10^{12}$ | $2.81 \times 10^{11}$ |
| Tween 0.5%  | $2.34 \times 10^{11}$ | $2.26 \times 10^{12}$ | $2.15 \times 10^{11}$ |

from the cytoplasm to the periplasm, so that the acquisition of MPT64 extracellular protein is still not optimal. In the future, it is possible to optimize the use of signal peptides to increase the acquisition of extracellular MPT64 protein with the support of surfactants to release the accumulated MPT64 protein from periplasm to medium.

## CONCLUSION

This study showed the efficiency of using Triton  $\times$  100 in supporting the action of the pelB signal peptide in translocating the extracellular MPT64 protein.

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Nil.

## Conflicts of interest

There are no conflicts of interest.

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