



## Neutrophils Are the Predominant Infected Phagocytic Cells in the Airways of Patients With Active Pulmonary TB

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**Background:** The exact role of neutrophils in the pathogenesis of TB is poorly understood. Recent evidence suggests that neutrophils are not simply scavenging phagocytes in *Mycobacterium tuberculosis* (*Mtb*) infection.

**Methods:** Three different types of clinical specimens from patients with active pulmonary TB who underwent lung surgery were examined: sputum, BAL fluid, and cavity contents. Differential cell separation and quantification were performed for intracellular and extracellular bacteria, and bacterial length was measured using microscopy.

**Results:** Neutrophils were more abundant than macrophages in sputum ( $86.6\% \pm 2.2\%$  vs  $8.4\% \pm 1.3\%$ ) and in BAL fluid ( $78.8\% \pm 5.8\%$  vs  $11.8\% \pm 4.1\%$ ). Inside the cavity, lymphocytes ( $41.3\% \pm 11.2\%$ ) were the most abundant cell type, followed by neutrophils ( $38.8\% \pm 9.4\%$ ) and macrophages ( $19.5\% \pm 7.5\%$ ). More intracellular bacilli were found in neutrophils than macrophages in sputum ( $67.6\% \pm 5.6\%$  vs  $25.2\% \pm 6.5\%$ ), in BAL fluid ( $65.1\% \pm 14.4\%$  vs  $28.3\% \pm 11.6\%$ ), and in cavities ( $61.8\% \pm 13.3\%$  vs  $23.9\% \pm 9.3\%$ ). The lengths of *Mtb* were shortest in cavities ( $1.9 \pm 0.1 \mu\text{m}$ ), followed by in sputum ( $2.9 \pm 0.1 \mu\text{m}$ ) and in BAL fluid ( $3.6 \pm 0.2 \mu\text{m}$ ).

**Conclusions:** Our results show that neutrophils are the predominant cell types infected with *Mtb* in patients with TB and that these intracellular bacteria appear to replicate rapidly. These results are consistent with a role for neutrophils in providing a permissive site for a final burst of active replication of the bacilli prior to transmission.

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**Abbreviations:** AFB = acid-fast bacilli; *Mtb* = *Mycobacterium tuberculosis*

Immunocompetent humans exposed to *Mycobacterium tuberculosis* (*Mtb*) mount an early innate and late adaptive immune response that ultimately destroys most *Mtb* bacilli and usually prevents the

development of clinical disease.<sup>1,2</sup> Whereas the late response is dependent on the acquisition of CD4<sup>+</sup> T-cell-mediated immunity and is characterized by granuloma formation involving epithelioid macrophages and multinucleated giant cells,<sup>3</sup> the early response is characterized by an influx of phagocytic cells. The early host response to *Mtb* infection involves primarily resident alveolar macrophages and recruited

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neutrophils and provides the first line of defense against mycobacterial infection. The role of alveolar macrophages in innate as well as adaptive immunity against *Mtb* infection is very well documented.<sup>4-6</sup>

Neutrophils are professional phagocytes that play important roles in many infections, and abundant neutrophils are observed in the BAL fluid of patients with pulmonary TB.<sup>7</sup> Some animal studies have also described acute and chronic neutrophil accumulation during mycobacterial infection,<sup>8,9</sup> leading to speculation on various roles for these cells in the development of TB.<sup>10,11</sup> Nevertheless, the exact role of neutrophils in the pathogenesis of *Mtb* infection in humans is still not completely defined. There are few reports describing neutrophil populations and their phagocytic activity in patients with active pulmonary TB.

The metabolism of *Mtb* is highly adaptable to the precise microenvironment in which the bacterium is residing.<sup>12</sup> Current efforts to develop new chemotherapies to shorten the course of TB treatment are dependent upon understanding the metabolism of *Mtb* in patients with uncontrolled disease.<sup>13</sup> Although there is no doubt of the importance of infected macrophages in successfully controlling initial infection with *Mtb*, the role of these cells in a patient with uncontrolled disease when initiating chemotherapy is far less clear. Nonetheless, new chemotherapeutic candidates with *in vitro* activity are often prioritized based on their activity in *Mtb*-infected macrophages.<sup>14-16</sup> The aim of this study was therefore to quantify the location of *Mtb* bacilli in various clinical specimens, including samples from sputum, BAL fluid, and cavities, reflecting the bacterial populations in patients with active TB. We examined whether the bacilli were intracellular or extracellular and which cell types were primarily infected. We also measured the lengths of bacilli in each specimen to investigate the metabolic status of *Mtb* bacilli in this environment.<sup>17</sup>

## MATERIALS AND METHODS

### Study Subjects

Human tissue collection from adult patients undergoing lung resection for the management of multidrug-resistant TB was approved by the National Masan Tuberculosis Hospital institutional review board and was granted an exemption by the US National Institutes of Health, Office of Human Subject Research. All subjects gave written informed consent for the collection and use of their specimens for research. Sputum was provided by 15 outpatients who visited National Masan Tuberculosis Hospital and received a diagnosis of active TB based on clinical findings and radiologic examination. Sputum was obtained for routine examination, and the remainder was used for this study. The mean age of the subjects was  $47.5 \pm 4$  years (range 29-47), and the average number of treatment episodes and average period of treatment history of the subjects were  $2.5 \pm 0.9$  episodes (range 0-9) and  $24.5 \pm 8.2$  months (range 0-87). All but one subject had cavitary

lesions. Observed comorbidities included two subjects with diabetes mellitus, two with hepatitis, and one with hypertension. Sputa from 11 of the 15 subjects were acid-fast bacilli (AFB) smear positive. BAL fluid was obtained from 10 hospitalized subjects with TB who were undergoing bronchoscopy and BAL for medical indications. The mean age was  $38.3 \pm 4.7$  years (range 17-60), and the average number of previous treatment episodes in these subjects was  $2.1 \pm 0.3$  (range 0-4). Seven subjects had cavitary lesions, and among them three subjects had taken no medication before BAL. The average duration of therapy for these subjects before BAL was  $13.9 \pm 3.7$  months (range 6-45). Observed complications included two cases of diabetes mellitus, one hepatitis case, and one case of COPD. Five of the 10 subjects who underwent BAL were AFB negative. Necrotic contents from nine cavities were collected from waste lung tissue that had been surgically resected for the debulking of multidrug-resistant disease in four patients. The mean age was  $36 \pm 3.9$  years (range 29-47), and the average number of previous treatments for these patients was  $1.5 \pm 0.6$  (range 0-3). The average duration of therapy prior to surgery for these patients was  $11.1 \pm 1.5$  months (range 9.5-15.5). No other comorbidities were observed in these patients.

### Preparation of Clinical Specimens

Sputum specimens were diluted 1:1 using RPMI 1640 medium (Gibco; Gaithersburg, MD), and collagenase type 4 (2 mg/mL) was added to digest the adhesive materials. After slight vortexing, the sputum was incubated in a shaking water bath at 37°C for 1 h and filtered through sterile gauze. The filtered sputum was overlaid carefully on Ficoll-Hypaque and centrifuged at 624g for 20 min at room temperature. After removing the supernatant, the buffy coat, ficoll, and pellet layers were recovered and transferred to three separate 50-mL conical tubes and washed with HBSS (SIGMA H2387; St. Louis, MO) at room temperature. The cell pellet of each tube was resuspended, and cytopsin slides were prepared from each tube. All the cytopsin slides in this study were stained with Ziehl-Neelsen and Hemacolor (Merck; Darmstadt, Germany). Few cells were observed among the sticky debris in the buffy coat and pellet layers, so only the cytopsin from the ficoll layer was used in this study.

BAL fluid was obtained by bronchoscopy. The bronchoscope was inserted near a region of affected lung and about 15 to 30 mL of sterile 0.9% saline was infused and recovered through the aspiration port. The collected BAL fluid was filtered through sterilized gauze and centrifuged at 699g for 10 min at room temperature. After removing the supernatants, the pellets were resuspended and washed with HBSS (SIGMA H2387). Cytopsin slides were prepared according to the manufacturer's instructions.

Cavity caseum was carefully removed using a 10-mL disposable syringe and spatula while dissecting the lung tissue. The caseum was transferred into a 50-mL conical tube, mixed with 10 mL of RPMI 1640 medium (Gibco), overlaid on Ficoll-Hypaque (GE Healthcare; Uppsala, Sweden), and centrifuged onto cytopsin slides, as above. Because only dead cell debris was found in the pellet layer, the slides from the buffy coat and ficoll layers were counted and then pooled and evaluated.

### Counting Infected Cells and Determination of Bacterial Length

Cells and *Mtb* bacilli were counted using light microscopy. First, differential cell counts were performed using a set of standardized morphometric appearance criteria (magnification  $\times 400$ ), and the percentage of each cell population was recorded after counting more than 200 cells. Second, the percentage of extracellular and intracellular *Mtb* bacilli were counted (magnification  $\times 1,000$ ) in more than 100 fields from the same slides. Finally, the number of cells containing bacilli was counted

(magnification  $\times 1,000$ ) in at least 100 fields, and the percentage of each phagocytic cell type was calculated. Differential cell counting, intracellular and extracellular localization, and determination of the number of cells with bacilli were all done at least twice by two independent, blinded readers.

The length of at least 200 *Mtb* bacilli within each sample was measured (magnification  $\times 1,000$ ) independently by two different persons using the AxioImage A1 and AxioVision Rel. 4.5 software (Zeiss; Jena, Germany). For statistical analysis, the precise lengths were used. For histographic representation, individual bacilli were regrouped based on a range of lengths (eg.  $\leq 1 \mu\text{m}$ ,  $\leq 1.5 \mu\text{m}$ ), calculated as a percentage.

#### Statistical Analysis

The statistical analysis was performed using GraphPad PRISM (Graph Pad Software, Inc.; La Jolla, CA), version 4.0. The data among groups were analyzed using one-way analysis of variance, and the differences between two groups were tested using the unpaired *t* test. Differences were considered statistically significant when  $P \leq .05$ .

## RESULTS

### Differential Cell Counts in Sputum, BAL Fluid, and Cavity Caseum

Differential cell counts were done using cytopspins obtained from sputum (Fig 1A), BAL fluid (Fig 1B), and cavity caseum (Fig 1C) from patients who were either AFB positive or AFB negative (for all samples from 15 subjects). In sputum and BAL, the most abundant cells observed were neutrophils, which comprised  $86.6\% \pm 2.2\%$  and  $78.8\% \pm 5.8\%$  of the total cells in each sample, respectively (Fig 2). Macrophages were much less numerous than neutrophils in both of these samples ( $8.4\% \pm 1.3\%$  in sputum and  $11.8\% \pm 4.1\%$  in BAL), and lymphocytes comprised only about 1% of the total cells found in both sputum and BAL. The remaining cells were epithelial cells and amounted to  $3.9\% \pm 1.9\%$  and  $7.8\% \pm 2.9\%$  in sputum and BAL, respectively (Fig 2). In the cavity caseum, however, 41% of the cells observed were lymphocytes, similar to the number of neutrophils ( $38.8\% \pm 9.4\%$ ), whereas macrophages comprised only about  $19.5\% \pm 7.5\%$  of the identifiable cells (Fig 2).

### Intracellular vs Extracellular Localization of *Mtb* Bacilli in Sputum, BAL, and Cavity Caseum

For these analyses, only the AFB-positive cytopspins (11 of the initial 15) from sputum, BAL fluid, and cavity caseum were counted. The number of bacilli was individually tallied based on whether they were intracellular or extracellular by manually scoring at least 100 fields of the AFB-stained cytopspin preparation. In both sputum and BAL, the bacilli were evenly distributed between extracellular and intracellular (Fig 3). Strikingly, in the cavity caseum, more than 80% of bacilli were extracellular, which was statisti-

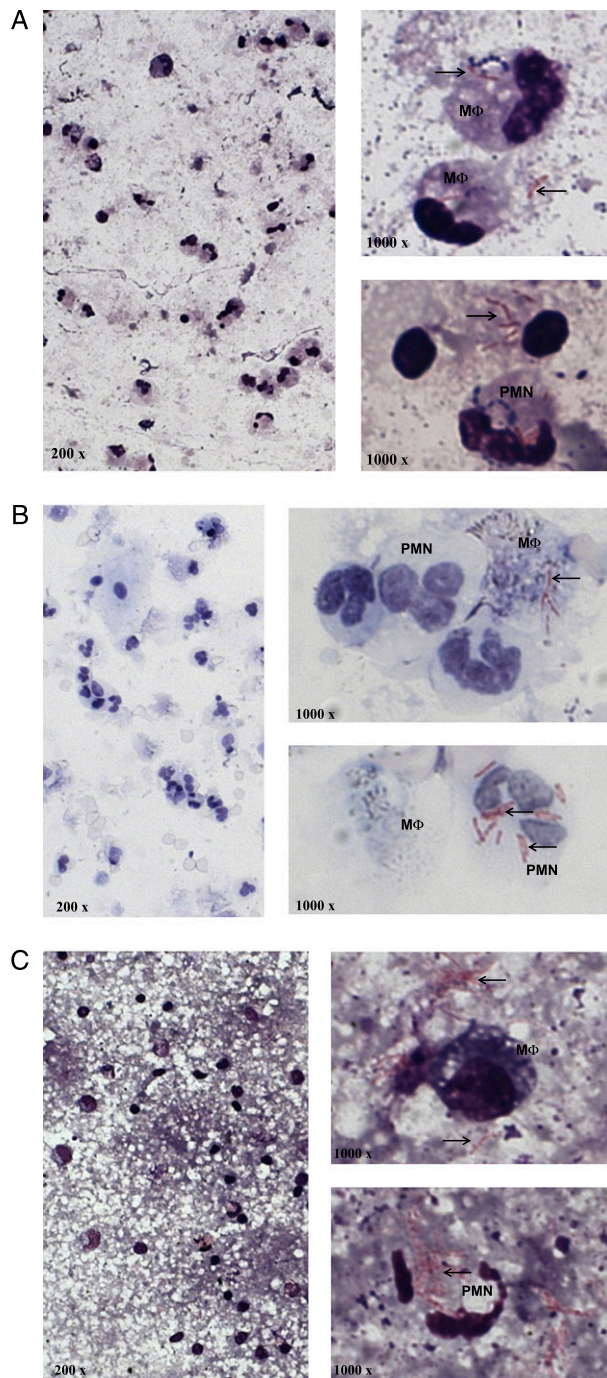


FIGURE 1. Localization of *Mtb* bacilli in sputum (A), in BAL (B), and in cavity caseum (C). Cytopspin slides of each sample were prepared and stained for acid-fast bacilli (Ziehl-Neelsen, original magnification  $\times 200$ ) and human cell morphology (Hemacolor, original magnification  $\times 1,000$ ). The bacilli (arrows) collocate with macrophages as well as neutrophils and the extracellular matrix. M $\Phi$  = macrophages; *Mtb* = *Mycobacterium tuberculosis*; PMN = neutrophils.

cally significant compared with both sputum and BAL ( $P < .001$ ) (Fig 3).

The type of cells engulfing bacilli was also quantified from each source. Neutrophils were more likely to

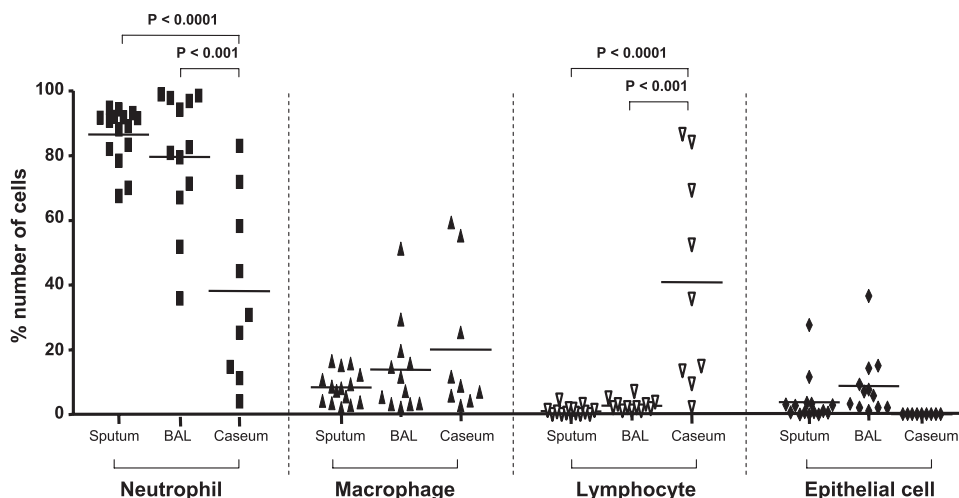


FIGURE 2. Differential cell counts in sputum, in BAL, and in cavity caseum. The percentage of each cell type was calculated when the counted total cell numbers were more than 200 and represented as neutrophil, macrophage, lymphocyte, and epithelial cells. Horizontal bars indicate mean value. The data among groups were analyzed using one-way analysis of variance, and the differences between two groups were tested using the unpaired *t* test;  $P \leq .05$  was considered significant.

have phagocytosed bacilli than macrophages in sputum ( $67.6\% \pm 5.6\%$  vs  $25.2\% \pm 6.5\%$ ), in BAL ( $65.1\% \pm 14.4\%$  vs  $28.3\% \pm 11.6\%$ ), and in cavity caseum ( $61.8\% \pm 13.3\%$  vs  $23.9\% \pm 9.3\%$ ) (Fig 4). Some bacilli were also found inside of epithelial cells in sputum and BAL ( $6.3\% \pm 3.9\%$  and  $6.6\% \pm 4.0\%$ , respectively) (Fig 4). No epithelial cells were found in the cavity caseum.

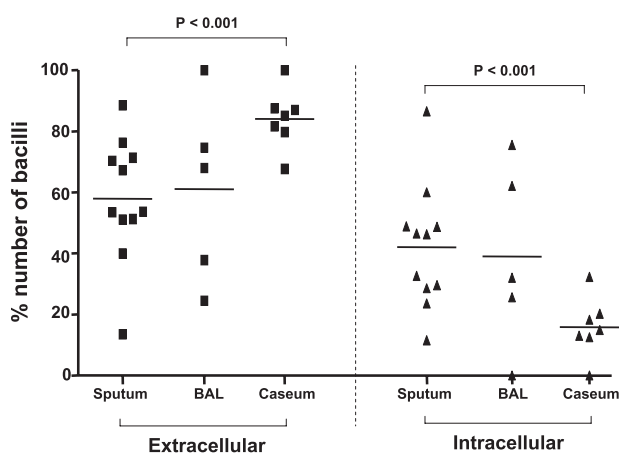


FIGURE 3. Localization of *Mtb* bacilli in sputum, in BAL, and in cavity caseum. Cytospin slides were prepared and stained by the Ziehl-Neelsen method followed by Hemacolor staining. The number of bacilli was counted both inside and outside of the cells in more than 100 fields (original magnification  $\times 1,000$ ) under a light microscope, and the total number ranged from 5 to 855. The percentage of extracellular and intracellular bacilli was calculated and represented in the extracellular (bacilli in extracellular space) and intracellular (bacilli in intracellular space) columns. Horizontal bars indicate mean value. The data among groups were analyzed using one-way analysis of variance, and the differences between two groups were tested using the unpaired *t* test;  $P \leq .05$  was significant. See Figure 1 legend for expansion of abbreviation.

### Bacterial Length Within Samples

Bacterial cell length was directly measured to the nearest tenth of a micron in each specimen. Figure 5 shows the frequency of *Mtb* bacilli in each 0.5- $\mu\text{m}$  increment. A broad distribution of *Mtb* lengths was observed, and the average value of *Mtb* cell lengths was  $2.9 \pm 0.1 \mu\text{m}$ ,  $3.6 \pm 0.2 \mu\text{m}$ , and  $1.9 \pm 0.1 \mu\text{m}$  for sputum, BAL, and cavity caseum, respectively. A statistically significant difference between the primarily extracellular cavity organisms and both BAL and sputum organisms (which contain primarily organisms within neutrophils) was observed ( $P < .001$ ).

### DISCUSSION

Neutrophils are the first defensive cells recruited to tissue following infection, where their role has been thought to involve eliminating invading pathogens via mechanisms such as the generation of reactive oxygen species<sup>18</sup> and the release of preformed oxidants and proteolytic enzymes from granules.<sup>19,20</sup> However, the proteolytic enzymes released by degranulation may also cause the destruction of neighboring cells and the dissolution of tissue.<sup>21,22</sup> This neutrophil-dependent tissue damage is known as the “neutrophil paradox,” in which the defending cells become an enemy.<sup>23</sup> Thus, a strict regulation of neutrophil influx and their turnover in infected tissues is essential. Neutrophils also have immunomodulatory function; when stimulated by *Mtb*, they release an array of cytokines and chemokines that attracts other inflammatory cells.<sup>24,25</sup> In addition, the interaction of neutrophils with *Mtb* triggers apoptosis of neutrophils,<sup>26,27</sup>

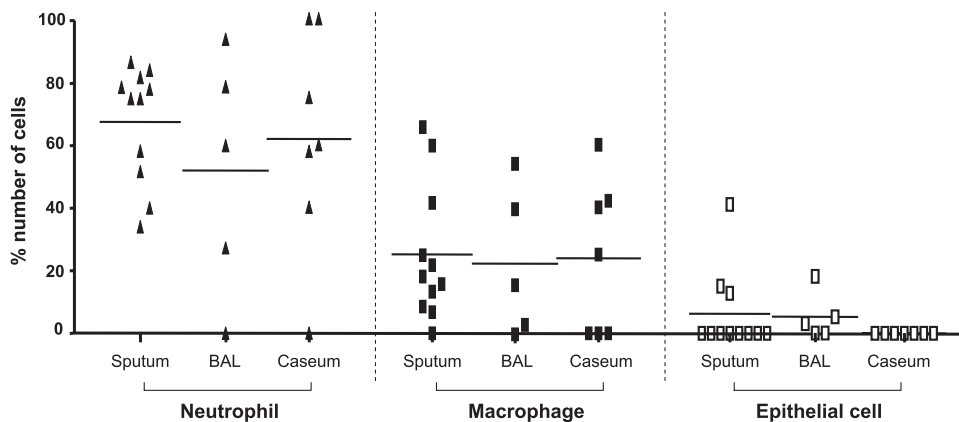


FIGURE 4. Differential counts of the cells ingesting *Mtb* bacilli in sputum, in BAL, and in cavity caseum. Cytospin slides were prepared and stained by the Ziehl-Neelsen method followed by Hemacolor staining. The number of cells engulfing bacilli was counted in more than 100 fields (original magnification  $\times 1,000$ ) under a light microscope and ranged from 4 to 47. The percentage of each cell type was calculated and represented as Neutrophil, Macrophage, and Epithelial Cell. Horizontal bars indicate mean value. The data among groups were analyzed using one-way analysis of variance, the differences between two groups were tested using the unpaired *t* test, and no significance was found among groups. See Figure 1 legend for expansion of the abbreviation.

and phagocytosis of apoptotic neutrophils by macrophages results in the decreased viability of intracellular *Mtb*,<sup>25</sup> suggesting a cooperative role of neutrophils in the hosts' defensive strategy against *Mtb* infection.

Many reports have demonstrated the presence of neutrophils in the sputum of patients suffering from mycobacterial infections (in fact, large numbers of neutrophils are considered a hallmark of high-quality sputum).<sup>29,30</sup> Previous reports in which researchers observed fewer neutrophils than we report here show a greater proportion of epithelial cells, indicating less

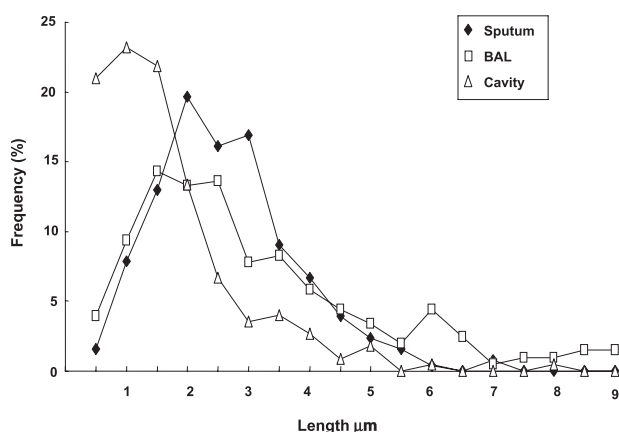


FIGURE 5. Frequency of *Mtb* bacilli lengths measured in sputum, in BAL, and in cavity caseum. More than 200 bacilli in each specimen were measured, grouped (as described in the "Materials and Methods" section), and plotted. Individual measurements were used for the calculation of average length and other statistical analyses. Extensive heterogeneity was observed in cell length in different specimens. The average values of *Mtb* cell lengths were  $2.9 \pm 0.1 \mu\text{m}$ ,  $3.6 \pm 0.2 \mu\text{m}$ , and  $1.9 \pm 0.1 \mu\text{m}$  for sputum, BAL and cavity caseum, respectively. A statistically significant difference between cavity caseum and both BAL and sputum was observed ( $P < .001$ ). See Figure 1 legend for expansion of the abbreviation.

productive sputum (the epithelial cells presumably originating from saliva).<sup>31</sup> Neutrophils are also persistently recruited to the sites of chronic mycobacterial infection.<sup>32,33</sup>

Of the many cell types present in the naive human lung that may mediate control of *Mtb*, most investigators have focused on alveolar macrophages or monocyte-derived macrophages. Yet the ability of human macrophages to kill *Mtb* in culture has never been convincing and consistent, suggesting that other cell types present at the site may also play important roles in innate resistance. Neutrophils are also essential for granuloma formation during chronic *Mtb* infection.<sup>34</sup> However, the ability of human neutrophils to solely mediate the outcome of pulmonary infection with *Mtb* is likewise questionable. Neutrophil depletion of mice was shown to have no impact on mycobacterial infections,<sup>35</sup> but reconstitution of mice with neutrophils increased resistance to *Mycobacterium avium*.<sup>36</sup> The ability of human neutrophils to kill virulent *Mtb in vitro* has been inconsistent at best,<sup>37-40</sup> but there have been consistent reports that circulating neutrophils from patients with active TB are highly activated upon stimulation with *Mtb*.<sup>41,42</sup>

The unit cell length of mycobacteria changes depending on the growth state *in vitro*, increasing during logarithmic growth and decreasing in the stationary phase.<sup>17</sup> In both sputum and BAL fluid, *Mtb* were found to be significantly longer than typical stationary phase cells. However, bacilli isolated from the cavities of resected lung tissue were significantly shorter than those from either of these two sources, averaging about  $1 \mu\text{m}$  in length (Fig 5). Thus, cavitory organisms appear to resemble in length stationary-phase organisms *in vitro*, whereas organisms in sputum

(and BAL) appear to resemble logarithmically replicating organisms *in vitro*.

The caseum of a liquefying cavity has long been thought to be the final site of the rapid replication of bacilli prior to their exit into the airways and subsequent transmission.<sup>43</sup> Surprisingly, our results suggest that this final burst of replication prior to transmission does not happen in the liquefying cavity, but rather upon the exit of the bacilli from that cavity into the sputum, and most likely in the context of an infected neutrophil.

There are, of course, several limitations to the current study, including the fact that the bacilli observed in the three different specimen types were not obtained from the same patient nor from patients with the same extent of disease. These patients have different histories of TB and treatment that could also affect the comparison. Nonetheless, the striking consistency in cell lengths and cell types observed across these samples suggests that this difference is meaningful.

Neutrophils may therefore play a much more profound role in the life cycle of this pathogen than has been previously appreciated. Neutrophils have been shown to participate in the transport of live mycobacteria from peripheral tissue to the lymphoid organ in mice,<sup>44</sup> and tissue neutrophils from TB-susceptible mouse strains have a significantly higher migration capacity, survive longer, and contain more intracellular mycobacteria.<sup>45</sup> There is also the fact that patients with noncavitary TB nonetheless produce sputum laden with AFB. Thus, neutrophils may act as transporters for bacilli from lesions to the pulmonary surface. Neutrophils may be short-lived, but they nonetheless appear to be an important microenvironment occupied by *Mtb* during two critical events in the life of a patient with TB: transmission and chemotherapy. Understanding the bacterial adaptations that occur during replication in neutrophils may suggest novel ways to interfere with transmission that could become an important adjunct to conventional chemotherapy.

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**Author contributions:** All of the authors have read and approved the final manuscript.

*Dr Eum:* contributed to the research conception, data acquisition and analysis, and drafting of the manuscript.

*Ms Kong:* performed the isolation of the cells from the specimens and the cytologic examinations.

*Ms Hong:* performed the isolation of the cells from the specimens and the cytologic examinations.

*Ms Lee:* performed the isolation of the cells from the specimens and the cytologic examinations.

*Dr Kim:* performed bronchial washings and surgeries and critically reviewed the manuscript.

*Dr Hwang:* contributed to the review and the final revision of the manuscript, and provided guidance.

*Dr Cho:* contributed to research conception and critically revised the manuscript.

*Dr Via:* contributed to research conception and analysis of the data and critically revised the manuscript.

*Dr Barry:* contributed to research conception and analysis of the data, and critically revised the manuscript.

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