Review Article

Marina Cañadas-Ortega, Clara Gómez-Cruz, Juan José Vaquero, and Arrate Muñoz-Barrutia*

The contribution of microfluidics to the fight against tuberculosis

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Abstract: The high mortality associated with tuberculosis brings forward the urgency of developing new therapies and strategies against the disease. With the advance of drug-resistant strains, traditional techniques have proven insufficient to manage the disease appropriately. Microfluidic devices have characteristics that can enhance treatment prescription and significantly advance our knowledge about the disease and its interaction within the human body. In addition, microfluidic systems provide advantages in terms of time and costs, which are particularly important in countries with low income and resources. This review will highlight how microdevices can help bridge the gaps in disease management, including their use for drug testing and development, drug susceptibility, basic research, and novel approaches to anti-TB vaccines and organ-on-chip studies.

Keywords: microfluidics, tuberculosis, microdevices, infectious diseases, respiratory diseases

1 Motivation

Infectious diseases of the respiratory airways are the third leading cause of death worldwide. Among them, tuberculosis (TB) constituted the sixth most deadly disease in 2000–2016. Around one-fourth of the global population is infected by Mycobacterium tuberculosis (Mtbb), making TB a global disease. Approximately 10 million new cases are diagnosed every year, and the number of deaths associated with TB per year exceeds a million. These cases mainly occur in developing countries and low-income areas, such as South-East Asia, Africa, and the Western Pacific [1]. Although most infections remain latent, in which the bacterium is inactive and asymptomatic, around 10% of cases develop active TB, where the bacterium is multiplying and invading tissues, mainly the lungs [2].

Numerous factors hinder the advances of the scientific community in the fight against TB, including the appearance of new drug-resistant strains of Mtbb (DR-TB) and the unmet requirements for both diagnosis and treatment.

The most used method for diagnosing pulmonary TB is sputum smear microscopy because of its low price and quickness, at the expense of low sensitivity and specificity. Tuberculin skin tests or chest X-rays are also used, with the disadvantage of presenting a low potential for point-of-care (POC) testing. The detection of antibodies specific for mycobacterial antigens and lipoarabinomannan (LAM), a lipopolysaccharide on the Mtbb bacterial cell wall, can also be used for diagnosis. However, the former presents low cost-effectiveness and accuracy, while the second lacks sensitivity. Finally, the use of molecular genotypic analysis (primarily based on nucleic acid amplification tests) is widely spread. However, it presents some disadvantages since most of the tests require tight temperature control and complex steps. For this reason, their use is limited to laboratories and requires trained staff. These methodologies are more extensively covered in the numerous reviews regarding the issue [3–5].

The development of DR-TB adds complexity to the diagnosis. This resistance emerges from patients’ poor
adherence to the long therapies against TB, insufficient monitoring of disease development, and other factors such as improper use of antibiotics in the food industry [6]. These drug-resistant strains can be of different types: multidrug-resistant (MDR) – not responding to treatment with isoniazid and rifampicin, the two first-line more powerful anti-TB drugs – or extensively drug-resistant (XDR) – resistant to the two previously mentioned ones, as well as to any fluoroquinolone and at least one of the three main second-line injectable drugs (such as capreomycin, kanamycin, or amikacin) [7].

This reduced drug susceptibility limits the treatment options, forcing the clinicians to use complex combinations of different anti-TB drugs. In addition, they often include multiple second-line drugs, which usually entails more extended hospital stays resulting in increased medical costs and higher human costs due to the severe side-effects caused by these second-line drugs [8–10].

Furthermore, drug resistance is usually harder to diagnose. For example, it requires phenotypic assays (like culture and microscopy of the strains, requiring several weeks to enable proper guidance) or the genotyping of DNA regions connected with drug resistance, which needs laboratories with appropriate biosafety levels.

These demands are critical in areas with technical constraints, such as underdeveloped or developing countries, where most cases are localized. However, the need for expensive laboratory infrastructure and thoroughly trained healthcare staff is one of the most significant barriers preventing these communities from accessing proper diagnosis or treatment at the POC [8]. Therefore, there is an urgent need for new diagnostic tools for active and latent TB, as well as for testing the drug susceptibility of these strains. Moreover, it is necessary to advance the development of new anti-TB drugs to increase the therapeutic efficacy in terms of costs, time, and side effects.

Microfluidic devices allow the manipulation of liquid volumes in the microscale [11], covering these needs in terms of costs, portability, number, and time of assays. These miniaturized devices work based on the application of electrical or mechanical properties to cover biological demands. The reduced volume and precise manipulation also result in lower reagent use and a minimization of the costs [12]. Regarding the development of new drugs and their testing, this technology enables multiplexing, designing devices with different channels to perform several reactions simultaneously. Additionally, several microchips can be coupled and imaged using a single microscope. These advantages lead to shorter testing times of potential drug candidates during preclinical development by reducing the assay time compared with traditional drug susceptibility tests. Thus, this technology allows performing more trials in less time, leading to quicker prescriptions [13].

These devices can be classified into three different groups: (i) high-throughput, automated systems with precise fluid control, aimed to facilitate laboratory research; (ii) those for low-scale experiments, dedicated to 3D in vitro cell culture (for organ-on-a-chip or drug screening applications), and (iii) simple and disposable devices to be used without advanced training or facilities [14]. Most devices used for drug susceptibility and fundamental research purposes fall into the second class. Diagnostic devices tend to fall in the third category, and a detailed summary can be found elsewhere [15]. In this manuscript, we review the last decade’s advances regarding the use of microfluidic devices applied to the field of TB, focusing on their role in drug testing and development, basic research, and emerging approaches.

### 2 Drug testing and development

The increasing number of drug-resistant strains of *Mtb* urgently requires the development of new fast and reliable tools to identify this resistance. To guide treatment prescription, it is thus essential to identify the particular drug resistance of specific strains by carrying out drug susceptibility tests (DSTs). Typically, they evaluate the growth or metabolic inhibition of the bacterial cells in drug-free or drug-containing media. More recently, new techniques for detecting mutations in drug-resistance associated genes have been established [16,17].

Mutations that result in drug-resistant strains are usually detected alongside *Mtb* diagnosis itself. However, more knowledge about the drug susceptibility of these strains is required due to the differences between genotypic and phenotypic resistance of *Mtb* and to the inability of current methods to detect all mutations involved in resistance (as they only cover a limited number of genomic regions) [18].

Furthermore, besides determining whether a specific strain is resistant to the first- or second-line TB drugs, it is essential to find new medicines that can control and eliminate these resistant strains. Situated farther from the POC but still necessary for the pharmaceutical industry are the microdevices oriented to developing new drugs. The goal of these devices consists of analyzing death processes and studying the effect of the compounds (mostly at the phenotypic level or through morphological changes) rather than identifying the susceptibility of a particular strain. The same chips developed for drug
discovery can also be used for DST, basic research, or other purposes.

In the drug testing and development process, microfluidic devices provide a system that allows the monitoring of colonies in the long term. Microdevices coupled with a microscope for time-lapse imaging of the growing bacterial colonies enable researchers to understand how the antibiotics affect the bacterial cells, either at metabolic or physiological levels, and to elucidate their death mechanism [19]. It is worth reminding that these technologies offer benefits such as the low amounts of compounds required or the possibility of multiplexing the chips, testing several drugs simultaneously [20].

Microdevices must trap bacteria in the microscope's focal plane to image the colonies and prevent movement during image acquisition, which can take up to several weeks. Different strategies have been developed to accomplish this, either using matrices or culture chambers in microfluidic devices to confine the cells in a single plane.

### 2.1 Culture of bacterial cells embedded in matrices

Different matrices have been used to trap the cells. Golchin et al. [21] used a thin layer of alginate hydrogel matrix to immobilize bacteria and promote planar growth, preventing the cells from going out of focus during long incubations. Dividing cells were recorded using time-lapse confocal imaging to observe phenotypic variations and antibiotic killing in individual mycobacteria cells. The nutrients were provided through media perfusion across a cellulose dialysis membrane from polydimethylsiloxane (PDMS) and a glass flow cell, sealed with a mechanical clamp (Figure 1 left). The group of Choi et al. utilized 3D agarose matrices to immobilize *Mtb* for rapid DST, developing the disc agarose channel (DAC) chip [22]. *Mtb* cultures are loaded in 24-well plexiglass chips, mixed with the liquid agarose, which is allowed to solidify (Figure 1 right). The antibiotics are flooded with the medium and allowed to diffuse in the agarose matrix; then, the chip is sealed and incubated to analyze the drug susceptibility. The 3D agarose structure allowed the immobilization and tracking of specific cells for 13 days with an automated imaging system. The susceptibilities of different strains (non-resistant/control, MDR, and XDR-TB strains) were tested with the DAC and compared with the results of liquid culture DST to validate the system. The agreement rates of specificity and sensitivity of different drugs obtained were around or above 90% when compared with liquid culture, thus proving its capacity for DST. The time required for the DST was 9 days using the DAC system. However, *Mtb* cells used to assess the validity were obtained from pre-cultured colonies on a liquid medium, which requires one or 2 weeks of culture to get a significant number of cells. Although the system reduces the turnover of conventional methods, which require 4–12 weeks (1–6 of sputum culture and 2–6 of DST), the sample culture still needs a long time.

![Figure 1](image-url)
Two years later, in 2018, the same DAC chip was used to overcome complications regarding the inoculum effect [18] (increment in the minimal inhibitory concentration (MIC) of antibiotic required when the size of the inoculum, i.e., the number of organisms inoculated, increases [23]). This effect constitutes one of the main problems of culture-based DST methods, as a standardized inoculum size is required to obtain reproducible DST results; additionally, it can cause a reduction in the drug effectiveness due to the increased number of bacteria [23,24]. Therefore, the DAC platform was used to evaluate the consistency of MIC values obtained from different inoculum sizes compared to the liquid culture reference method. Moreover, the critical concentration of 12 anti-TB drugs was determined from 254 clinical isolates (including pan-susceptible and resistant strains) without measuring the inoculum size. The tested anti-TB drugs gave consistent results with 96.3% overall agreement within 7 days. However, the isolates required a lengthy subculture on a liquid medium before the test, as in the previous study. In a recent 2021 study, the DAC system improved with a modified channel to increment the antibiotic diffusion into media to reduce turnaround times for DSTs. This fully automated DST system was named QuantaMatrix microfluidic agarose channel (QMAC) and integrated mycobacteria growth indicator tubes [25].

2.2 Monitoring in culture chambers

Culture chambers within microdevices have been widely used to analyze drug effects on Mtb. However, to achieve this, bacterial cells need to be encapsulated in the chambers. To this respect, Delincé et al. [26] developed InfectChip, consisting of a cellulose semipermeable membrane between a micropatterned coverslip and a PDMS chip. The coverslip is patterned with circular microchambers where the cells are trapped, grow, and divide, whereas the medium, is continuously flowing through a microchannel in the PDMS chip (Figure 2 left).

This device was used to study the dynamics and interactions of single-cell behavior between Mtb and motile eukaryotic phagocytes in a host-cell model, providing a platform to visualize the different stages of infection. However, despite its advantages, it presents several limitations, including a limited lifespan of the macrophages, which constrains the visualization of the infection process.

The design shares similarities with the device developed by Wakamoto et al. [29], used for basic research (Figure 4). It can be used to test new antibiotics. For example, they tested faropenem [30], and it was also used to analyze the effects of inhibitors against drug resistance mutation for isoniazid [31].

Other systems have been developed to encapsulate bacterial cells in microfluidic devices. For example, Sala et al. [27] proposed a model to analyze the effect of different drugs on dormant cells using a previously developed microdevice [28]. The microdevice design includes a PDMS layer patterned with long grooves, covered by a transparent membrane, and sealed with a thick PDMS layer with flow channels. Cells get trapped in the grooves between the membrane and the PDMS layer (Figure 2, right). This system differs from the previous ones [26,29] as the cells are not culture chambers in the coverslip but directly growing on PDMS grooves.

Figure 2: Microfluidic devices that use PDMS channels to trap cells. Left: 3D model of the microfluidic device from ref. [26], showing the semipermeable membrane that covers the patterned microchambers in the coverslip. The PDMS chip diffuses media through the membrane. Right: Representation of the microdevice used in ref. [27] to culture cells in PDMS grooves covered by a membrane, fed by diffusion through a flow channel encapsulated in a PDMS chip. Images reproduced with permission from Lab Chip (left, [26]) and Science (right, [28]).
2.3 Other methods for drug development

Time-lapse microscopy combined with microdevices for bacterial culture is the primary technique used in drug testing applications and basic research. Still, other methods are used in the field of drug development. Instead of focusing on the quantification of death rates and characterization of the persistent cells, this research analyzed the transcriptome and other markers to determine the response of the bacterial cells to the antibiotics applied.

Murima et al. [32] utilized a high-throughput microfluidic array developed in ref. [33] to obtain biomarker genes that could improve and shorten the traditional anti-TB drug-discovery schemes. This device contains an integrated fluidic circuit chip containing valves and 2304 chambers where the real-time polymerase chain reaction (PCR) occurs (Figure 3 left). After the fluids are pressure-driven through valves to the integrated fluidic circuit, composed of fluid lines, valves, and microchambers in which the PCR takes place. The sample and reagents are introduced through the inlets, which are pressure-driven through valves to the integrated fluidic circuit, composed of fluid lines, valves, and microchambers in which the PCR takes place. The high number of wells allows performing many simultaneous reactions, which were used to analyze transcriptomics of M. tuberculosis and M. bovis in the presence and absence of several growth inhibitors.

Raman spectroscopy was used to analyze the effect of isoniazid in the lipidic expression of M. smegmatis [34]. The device consists of a flow cell with a trapping chamber, formed by laser-cutting the channel between quartz slides and sealed with transfer tape, as can be observed in Figure 3, right. The inlet and outlet are made with silicone tubes.

The transparent lithium transducer allowed the acoustic pressure required for the cells to levitate and aggregate in the channel and to be trapped in the lateral direction to obtain the wavelength-modulated Raman spectra. The live-cell monitoring was carried out by acoustic trapping after 7 days of preculture.

3 Microfluidic applications in basic research

Thus far, the focus has been primarily on developing microfluidic technologies for diverse and specific purposes, mainly motivated by the novelty of the techniques, which had to be tested and characterized before being integrated into the laboratory and the scientific know-how. However, different research groups and facilities have also applied these technologies to better understand disease mechanisms. This knowledge ranges from changes in morphology and cell interactions to death rates and antibiotic exposure patterns.

Research groups dedicated to investigating Mtb and TB disease often use either already characterized or commercial microfluidic platforms for their experiments without developing a device themselves. Additionally, the investigations tend to be more focused on understanding a particular phenomenon rather than testing the global feasibility of a technique or drug. This section...
Figure 4: Schematic of the PDMS microfluidic chip mounted on a cellulose membrane and a glass coverslip for time-lapse imaging. (a) Top view depicting inlet, outlet, and parallel channels. (b) Cross-sectional view of the channels in the PDMS chip. (c) Closeup to the cells growing in the microchambers, covered by a membrane to allow media diffusion from the channels. Image reproduced with permission from Science [29].

summarizes the main applications of commonly used imaging devices.

Many researchers use variations of different platforms developed in the School of Life Sciences of the Swiss Federal Institute of Technology, particularly the one presented by Wakamoto et al. [29]. This microdevice consists of 100 µm × 50 µm parallel microchannels patterned on a PDMS chip, covered by a semipermeable membrane, then mounted on a glass coverslip (Figure 4). Cells get trapped between the membrane and the coverslip, continuously supplied with nutrients by diffusion from the flow channels across the membrane. The whole system is sealed in a sandwich configuration by a base adapter with screws. This enclosure allows 2D growth while restraining cells in the plane of focus for long imaging periods. In combination with time-lapse microscopy, this setup was initially used to study the enzymatic expression of persistent mycobacterial cells and their effect on cell survival.

The research group used microfluidics and time-lapse setup to study different parameters, characteristics, and behaviors of Mycobacterium cells. For example, in ref. [35], they measured chromosome division and replication to gain insights into the dynamics of Mycobacterium cellular growth. Other studies used the platform to assess antibiotics’ effects on intracellular adenosine triphosphate (ATP) levels [36] or to study the role of ribosomal activity and its associated gene expression in the latent state after chemotherapy [37]. It was also used to study the response to isoniazid of M. smegmatis penicillin-binding protein mutant [38] and NADH pyrophosphatase mutant [39]. More recently, this combination of microfluidic devices and time-lapse optical microscopy was used to measure single-cell pole growing to better understand the Mtb cell division process [40].

The same group developed other chips with different microchannel configurations, as the protocol presented in ref. [41]. Similarities are shared, like the channel dimensions, using a semipermeable membrane between the cells and the PDMS chip, or the mechanical clamping with screws. However, this second configuration is a multiplexed version that allows testing different bacterial strains (Figure 5). With this platform, the cell responses to DNA damage and its relationship with drug resistance were analyzed [42] or the characterization of the cellular response against a new molecular drug target [43].

Another custom-made device was proposed by Aldridge et al. [44]. In this case, the PDMS chip with the patterned microchannels is attached to the glass coverslip and sealed by plasma bonding treatment, avoiding the difficulties entailed by the mechanical clamp presented in the previous devices. The other main difference resides in the lack of membranes between the cells and the PDMS chip. In this device, the cells are pumped and grown directly in the channels, not patterned chambers. These channels for cellular growth connect with the main channel through which media is flowed, transporting nutrients. Bacterial cells are thus fed by diffusion between the main channel and the shallow ones, as can be observed in Figure 6, left.

This microdevice was used to analyze the asymmetric division of Mtb cells and its associated antibiotic resistance (particularly to Rifampicin) in ref. [48]. The media and the antibiotic were mixed and injected into the viewing device for the morphological analysis of the polarized division. The Morphological Evaluation and Understanding of Stress (MorphEUS) project also used
this platform to classify tested drugs into different categories according to their modes of action, characterized in ref. [49], continuing with the idea of the utility of morphological profiling to elucidate bacterial response to drugs.

Another use of the device includes measuring the polar growth and response of single cells in the presence of stressors such as antibiotics. Remarkably, the link between the asymmetrical division of physiological sub-populations and resistance is analyzed and compared with other bacteria such as *E. coli*. The platform was also used to study the changes produced in the host–pathogen interaction and bacterial burden upon vaccination [50].

Studying the biological processes of *Mycobacteria* can also benefit from commercial microfluidic platforms. For example, the CellASIC ONIX2 Microfluidic Systems (Millipore Sigma) are plastic plates with multiple channels and microfluidic chambers for cell culture (Figure 6, right), allowing tight control of the flow, temperature, and gas exchange. These plates were used [45] to characterize and elucidate the role of transpeptidase...
crosslinks in cell shape and death and the possibilities for drug targeting against them. In ref. [46], the same microfluidic plate was used to follow the replication of bacteria and the chromosome dynamics after adding drugs altering the cell division mechanisms. It was also used more recently in ref. [47] to investigate ultra-short peptides and their effect on membrane activity of mycobacterial cells to exploit their antimicrobial properties.

Other methods to characterize and visualize bacteria to retrieve basic biological information include the dielectrophoresis technique proposed in ref. [51]. In this work, a microdevice containing a single-cell suspension of M. smegmatis in buffer was prepared and then high frequencies were applied. Cell movement was recorded to analyze cell death and drug tolerance mechanisms.

4 Other applications

Microfluidic technology has shown utility in the fight against Mtb for different diagnostic methodologies, the advancement of drug candidates’ selection in its various forms, or enhancing the general knowledge about the disease through basic research. This section will focus on other contributions of these devices to the management of TB.

4.1 Production of anti-TB vaccines

Vaccination programs can be essential for disease management by preventing active and latent TB. The first (and to the day, only) vaccine against TB is the Bacille Calmette Guérin (BCG), introduced in 1921 [52]. It prevents extrapulmonary TB in infants by newborn vaccination; however, it fails to stop the pulmonary form of the disease [53]. Due to these limitations, new vaccine candidates are continuously being proposed and reviewed. Some of the leading candidates and their clinical phases are evaluated in detail in refs [54,55].

Particularly, cationic liposomes were formulated as a gene delivery system that could be used in gene therapy and vaccine due to their ability to form complexes with DNahsp65 [56–58]. The low yield production obtained with conventional methods and the escalation required led to the use of microfluidic devices to generate high lipid concentrations. Two PDMS devices were proposed based on single and double hydrodynamic focusing (SHF and DHF, respectively) to enhance vesicle formation and mixing [59], as depicted in Figure 7. These devices facilitate optimizing parameters affecting liposome production for the development and production of anti-TB vaccines. This process led to the high yield production of liposomes, suitable for gene delivery and vaccination due to their unilamellar nature and particle size.

This study demonstrates the possibility of taking advantage of microfluidic devices not only for low-scale analysis or diagnostics but also for scaled-up production of pharmaceutical compounds, especially in the cases of vaccines or antibiotics.

4.2 Diagnosis of coinfection of TB and HIV

TB is the leading cause of death in HIV-co-infected patients, with a proportion of approximately 8.6% of TB patients co-infected with HIV in 2019 [1]. Additionally, HIV infection in TB patients increments the probability of activating a latent TB state between 80- and 100-fold [60]. Nevertheless, its diagnosis remains challenging in HIV co-infected individuals due to a high frequency of smear-negative disease, high rates of extrapulmonary TB, and reduced pulmonary load [18]. Traditional techniques do not provide rapid and accurate results to manage HIV co-infected patients effectively [61]. Despite its importance, there are almost no microfluidic detection methods to detect both diseases simultaneously.

Figure 7: Schematic and photography of the microfluidic devices developed in ref. [59] depicting the single (a) and double (b) hydrodynamic focusing. Lipids are diluted in ethanol and injected in the center channel with a flow velocity of 143 mm/s. Deionized water flows through the exterior channels to hydrodynamically compress the central flow. In the case of double focusing, another flow-focusing stream is added to each side. Image reproduced with permission from Chemical Engineering Journal [59].
A multiplexed nucleic acid amplification test (NAAT) was developed by the Fudan University of Shanghai, portraying a microdevice for simultaneous detection of HIV, TB, and *Pneumocystis carinii pneumonia* (PCP) using a loop-mediated isothermal amplification strategy [62]. The PDMS device consists of a multiple loop-mediated isothermal amplification (LAMP) with the shape of a five-pointed star (Figure 8). The sample is loaded in the middle of the chip and distributed to the pools located at the points of the star by capillary forces. Three of the amplification pools are coated with LAMP probes against HIV, *Mtb*, and PCP. The fourth one serves as a positive control (coated with β-actin LAMP probe) and the last one as negative control (uncoated). Once the amplification pools fill with the nucleic acid sample and LAMP buffer, the inlet is sealed using uncured PDMS that requires on-site incubation. Results can be checked within 90 min with the naked eye (supporting its utility in low resources areas) by the appearance of white residue generated during the LAMP reaction.

Additionally, green fluorescence is produced by SYBR green I and can be detected using agarose gel electrophoresis. The detection limit of the multiplexed chip is 50 copies/µL for HIV and 100/µL copies for *Mtb* and PCP, respectively. Moreover, no nonspecific amplification or cross-reactivity was detected when samples of the diseases were tested using nucleic acids of different viruses as templates.

This platform is presented as a proof-of-concept for resource-poor countries. However, the accuracy of the primers and the rates of false positives and negatives needs to be characterized.

### 5 Future perspectives

Although many significant contributions have been made to the field of TB regarding mechanisms of action and several detection strategies, further work is required to understand the behavior of the disease entirely. In particular, it is still necessary to comprehend and bridge the gaps that limit the use of these technologies at the POC. Moreover, different perspectives are always open for investigation since perfect devices (*i.e.*, affordable, rapid, specific, sensitive, user-friendly) have not been deciphered. Some fields could strongly benefit from the advantages offered by microfluidic devices, especially those with translational capabilities or related to comorbidities. Notably, the reproduction and development of *ex vivo* tissues can significantly benefit from the use of microfluidic technologies. During the last decade, there has been an important advancement in the encapsulation and recreation of living tissues inside chips through organ-on-chip technology.

#### 5.1 Organ-on-a-chip and organoids

Advancing towards the elimination of TB requires improving the diagnosis and identification of drug resistance and a deeper understanding of the interactions between the pathogen and the host cells in the pulmonary tract during infection and drug treatment [63]. *Mtb* presents a sustained interaction over time with host cells [64], which displays a spatial organization of the host–pathogen interaction [65]. Moreover, the extracellular matrix influences the survivability of the infected host cells [66]. Additionally, interactions with the host cells and environment affect bacterial physiology, inducing stress-mediated variations in their genetic expression [67]. These factors disclose the importance of developing novel models representing the diversity of microenvironments and elements that mediate the interaction between pulmonary cells and *Mtb* [68]. Animal models or human 2D cellular models have been primarily used for drug development and testing, pharmacokinetics, and research in general. Basic 3D structures of lung tissues have also served to model the spatiotemporal features of the diseases [69,70].

Organoids and organ-on-a-chip are more complex models featuring 3D cellular organization that has
emerged in the last decade to bridge this gap between 2D cell cultures, animal models, and the actual 3D microenvironments. These technologies and their potential applications in the pharmaceutical analysis have been extensively reviewed elsewhere [63,71–74]. However, despite their strong potential for TB research, barely any endeavors have been dedicated to this field yet. Most lung-on-chip models will require adaptation for infection and security requirements in general, particularly TB-specific conditions, while organoids have not been used to analyze *Mtb* infections to date [71]. On a related note, a microsphere system combined with a microfluidic plate has been developed to model the effect of dynamic antibiotic concentration on *Mtb* [75]. A suspended microfluidic platform has also been used as a granuloma infection model [76]. Another example is the use of a commercial lung-on-a-chip model to analyze the interactions between *Mtb* and lung at the early infection stage [77].

Regarding the former [75], 3D microspheres were prepared by bioelectrospray methodology [78] using primary human cells (monocytes and T cells), active *Mtb*, and type I collagen on the microfluidic device, leading to the formation of human granulomas after 14 days. The system modulated the drug concentrations over time to replicate the pharmacokinetics observed *in vivo* during patient treatment. The granuloma’s genetic expression and the pharmacokinetics of antibiotics were measured and compared with the liquid culture, demonstrating the differences in the treatment outcome. The microfluidic devices were manufactured from a plexiglass sheet by micromilling, with inlet and outlet channels, and were then introduced in the pits of a 24-well plate before microsphere seeding.

The granuloma environment was also mimicked in ref. [76] using an *in vitro* suspended microfluidic platform adapted from the previously described Stacks system [79]. This system is based on capillary pinning to enable the stacking of layers with different microenvironments and uses open culture wells, which also allow easy imaging. Furthermore, the system does not require pumps (as traditionally occurs in microfluidic devices) because it functions based on surface tension and capillary forces. In this study, the Stacks system was used to create an infection model to study signaling microenvironments, combining human monocyte-derived macrophages and *M. bovis* BCG. The stacks were suspended in a 3D collagen extracellular matrix (ECM) and cultured to initiate paracrine signaling (Figure 9 left).

Another model using spheroids is presented in ref. [80], generated first by the combination of human monocyte and macrophage spheroids, subjected to different Mycobacteria strains. Then, the complexity of the model increased by incorporating peripheral blood mononuclear cells and lung fibroblasts. Finally, these 3D models were used to analyze the co-culture response to drugs such as Isoniazid and Rifampicin.

**Figure 9:** 3D organ-on-a-chip models for *Mtb* interaction with lung tissue. Left: Workflow showing the setup of a microfluidic device containing granuloma microspheres grown in a suspended microfluidics “Stack system” from ref. [76]. Immune and Mycobacteria cells arrange in collagen matrix into a 3D granuloma model. Right: Schematic representation of the lung-on-a-chip from ref. [77]. Alveolar epithelial cells and macrophages cover the upper face of the membrane in the air-filled alveolar channel, while endothelial cells reside in the lower, liquid-filled channel. The tissues able to maintain this interface are infected with *Mtb* and imaged with confocal microscopy. Images adapted with permission from Front Bioeng Biotechnol. (left, [76]) and eLife (right, [77]).
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Concerning the lung-on-chip [77], commercial lung chips were manufactured in PDMS from Emulate (Boston, MA), then preactivated and coated with ECM solution according to the manufacturer’s instructions. Then, endothelial cells were seeded in the different channels on the bottom face of the porous membrane (the vascular channel) and alveolar epithelial cells (AECs) on the epithelial channel (Figure 9 right). The use of two different cell types allows recreating the air–liquid interface present at the alveoli. After organ maturation, macrophages are introduced to the epithelial channel and allowed to attach to the epithelium. The lung tissues seeded on the chip capable of maintaining the interface are then infected with Mtb, allowing the analysis of its development in macrophages and AECs. Time-lapse fluorescence microscopy is used to study the dynamics of early infection and quantify the intracellular progress of the bacteria. This lung chip was also used in ref. [81] with murine cells and time-lapse microscopy to elucidate the role of surfactants produced by the alveolar epithelial cells in reducing bacterial growth.

These platforms that are still being developed and optimized present a broad range of applications. In the field of TB, they offer the possibility to study pathogen behavior in a more realistic environment and strongly increase the capacities of DSTs. In this way, the effect of antibiotics on bacterial populations and pulmonary tissue grown in the chip or organoid can be quantified. Different research groups worldwide continue their research in these areas. Furthermore, these systems grant the opportunity to analyze and model the interactions between drugs, pathogens, and tissues. They could also be helpful for the study of coinfections, especially those of the respiratory airway.

Studying these coinfections has growing importance, as respiratory tract infections (i.e., avian influenza, coronavirus) have resulted in pandemic situations during the last decade. Although different meta-analyses have found connections between susceptibility to COVID-19 and the presence of TB [82,83], further studies will be required to investigate the correlations between the diseases to advance the fight against them. Thus, the development of these platforms (organ-on-a-chip and organoids) can significantly enhance the knowledge about the interactions between the diseases and speed up the research pipeline.

5.2 Conclusion

Microfluidic devices have been present in the biomedical field for some decades now, enabling the advance and development of multiple techniques and platforms due to their numerous benefits over traditional methods. They have roles in the basic research field and at the translational and clinical stage in diagnosis and drug testing and development.

In this manuscript, we have dissected the potential applications of microfluidics to improve treatment prescription or to increment our knowledge about TB disease. Microdevices can provide an advantage over traditional diagnosis methods due to their simplicity and ease of use without medical facilities. They can also be used to detect different resistant strains of the bacteria, allowing more personalized treatment for each patient.

Moreover, many of the reviewed platforms have POC testing potential, some of them optimized in terms of cost for their use in underdeveloped or resource-constrained countries. There is a growing tendency to reduce the prices, materials, and laboratory setting requirements, as can be observed in the development of lab-on-paper devices in combination with smartphones compared to more traditional PDMS chips. In this way, microdevices position themselves at the forefront of biomedical research, with promising potential for future developments and opportunities.

Several research groups over the globe have routinely incorporated microfluidic devices in their laboratories as a daily tool, constituting an important source of knowledge about the basic forms of the disease and their relationship with other illnesses that entail threats to humanity. Moreover, through the development of new organ-on-chip and other techniques, insights into the behavior of the bacteria within the human body and the host reactions can significantly improve our understanding of infection mechanisms. Microdevices also present new opportunities in therapeutic and drug development testing. They not only enhance the diagnosis and treatment guidance but also facilitate basic research and drug development (Table 1).

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References


