

Lab on a Chip

Accepted Manuscript



This is an *Accepted Manuscript*, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. We will replace this *Accepted Manuscript* with the edited and formatted *Advance Article* as soon as it is available.

You can find more information about *Accepted Manuscripts* in the [Information for Authors](#).

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard [Terms & Conditions](#) and the [Ethical guidelines](#) still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this *Accepted Manuscript* or any consequences arising from the use of any information it contains.

ARTICLE

Microfluidic device for real-time monitoring of *Bacillus subtilis* bacterial spores during germination based on non-specific physicochemical interactions at the nanoscale level

Cite this: DOI: 10.1039/x0xx00000x

Received 00th January 2012,
Accepted 00th January 2012

DOI: 10.1039/x0xx00000x

www.rsc.org/

L. Zabrocka^b, K. Langer^a, A. Michalski^b, J. Kocik^c, and J. J. Langer^{*a}

A microfluidic device for studies on the germination of bacterial spores (e.g. *Bacillus subtilis*) based on non-specific interactions in nanoscale is presented. A decrease in the population of spores during germination, followed by the appearance of transition forms and an increase in the number of vegetative cells can be registered directly and simultaneously by using the microfluidic device, which is equipped with a conductive polymer layer (polyaniline) in the form of a nano-network. The lab-on-a-chip-type device, operating in a continuous flow regime, allows to monitor germination of bacteria spores and to analyse the process in detail. The procedure is fast and accurate enough for quantitative real-time monitoring of the main steps of germination, including final transformation of the spores into vegetative cells. All of this is done without the use of biomarkers or any bio-specific materials, such as enzymes, antibodies and aptamers, and is simply based on an analysis of physicochemical interactions at the nanoscale level.

Introduction

Germination of bacterial spores is a complex and still mysterious process¹ which results in a breakdown of the structure of the spores and in a concomitant loss of their resistance against physical and biochemical factors. This process, i.e. the conversion of spores into vegetative cells, begins under favourable conditions and is initiated by small molecules (amino acids and nucleosides). After initiation of germination, dormant spores begin to hydrolyse and release monovalent cations, protons and zinc from their core. Also, molecules of dipicolinic acid (DPA) and calcium which is associated with the DPA are released by the spores and replaced by water molecules. This process continues until a sufficient level of hydration is reached, which results in restoring enzyme action and hydrolysis of the spore coat (peptidoglycan) as well as swelling of the germ cell. Only when sufficient nutrients for outgrowth of spore cells are available, small acid-soluble proteins (SASP) associated with spore DNA are degraded and the vegetative cell escapes from the spore coat^{1,2}.

In recent years, a number of analytical tools for examining germination of spores have been developed¹, e.g.: fluorometry³, flow cytometry⁴, laser tweezers Raman spectroscopy^{5,6}, phase-contrast and fluorescence microscopy^{6,7}, surface-enhanced

Raman scattering (SERS) microscopy⁸, and SERS spectroscopy⁹, as well as an impedance-based method, though not a very accurate one, to detect spores by electrically detecting their germination in real time using a microfluidic biochip¹⁰. Most of these methods are based on advanced (and expensive) instrumentation and complex procedures. Some of the methods as mentioned above are very accurate, however, they are limited in that only a few individual cells can be examined, which is not a representative sample, thus it is difficult to characterise a whole colony of 10^6 - 10^9 (or more) cells when using such a method.

Fluorometry-based methods can use the process of the activation of esterases as a practical parameter to quantitatively measure spore germination since activation of proteases and cortex-lytic enzymes plays a key role during the initial stages of spore germination.

Expression of esterase activity is particularly suitable for measuring the early responses of spores since it displays rapid kinetics as measured with diacetyl fluorescein (DAF), a fluorogenic substrate which is hydrolysed nonspecifically by esterases, lipases, and proteases.

Therefore, this method allows to monitor the germination process only directly in the first 14 minutes after the germinants are added to dormant spores, since there is a 2-minute lag followed by a 2-minute burst of esterase activity during which activity increases more than 150-fold above the baseline³.

The microfluidic biochip can be used as a method for automatic and rapid electrical detection of germination of viable spores. The device includes special design features that facilitate spore capture and electrodes for impedance measurements. With a detection limit of fewer than 100 spores in a 0.1 nL chamber¹⁰, the spore detection time is similar to a conventional method based on the PCR (two hours from heat activation of a suspected organism⁹). This method, although very advanced and convenient, does not allow for real-time monitoring of spore transformation into vegetative cells.

Surface-Enhanced Raman Scattering Spectroscopy (SERS) has been used to detect the presence of DPAs, i.e. molecules which are released from spores during germination and can be used as a chemical signature for monitoring endospore germination of *Bacillus* species. Even more fascinating is the fact that only several hundred spores are sufficient to reliably measure the kinetics of germination at different concentrations of germinant and different temperatures using SERS⁹.

The downside of this method is observation of only a single factor, i.e. the release of DPAs from *Bacillus* spores, as a reference for such a complex germination process.

All methods that can detect spore physiology with limited manipulation and in a small time frame are valuable to public health and safety. Therefore, it is understandable that there is growing social emphasis on developing simple yet reliable techniques that will provide rapid information about a microbial sample without relying on culture techniques.

In this paper we present a new method for simultaneous monitoring of spores and vegetative cells during germination in a real time with no such limit and using for testing “reasonable” samples of 10^5 spores (or more). The method is based on measurements of time-dependent changes in the electrical conductivity of a polyaniline nanofibre network (Fig. 1a); the changes are induced thanks to the presence of spores and cells interacting directly on conductive polymer nanofibrils. Polyaniline is one of the best materials to use due to its stability, relatively high electrical conductivity and because it is electroactive and pH-sensitive.

A new nanodetector was designed as a lab-on-a-chip device operating in a continuous flow regime (Fig. 1a), analogously to our previous basic version of a continuous flow nanobiodetector, CFNBD, which has successfully been applied to detect bacteria vegetative cells^{11,12}. The system, based on polyaniline fibres (antibacterial), is self-cleaning, particularly when it operates in continuous flow mode, so it can be used to analyse multiple and different samples with practically no interference (the base line is stable between measurements).

After modification and improvement of sensitivity due to construction of a compact, microfluidic, mechanically stable lab-on-a-chip version of the nanodetector and after applying a more accurate measuring interface, we were able to monitor germination of bacteria spores in detail by using the same single device, which is not possible for other methods of comparable simplicity. When analysing the intensity and shape of the electrical signal registered thanks to changes in electrical conductivity of the nanofibre network in a function of time (time profile, Fig. 1f), one can obtain information on the number of spores and cells^{11,12,15} and on the processes in which they are involved. All that is necessary is the injection of a small sample with a suspension of germinating spores (usually

below 100 μL and ca. 10^5 spores) into the detecting device, done every 5 min or more often if necessary. This approach is useful for monitoring the germination process with an accuracy that is better than 5% in population, i.e. of 10^3 spores or cells and with a time resolution of process monitoring well below 1 min (in seconds in the case of an automatic sampling). These parameters are better (taking into account the representative character of the samples) or comparable to other methods which are applied in the case of complex systems, e.g. to analyse a mixture of spores and vegetative cells prepared *in vitro* or formed naturally during germination (please see the chapter *Comparison with other methods*). The ability to trace the transformation of spores into vegetative cells is essential not only for science (e.g. microbiology) but it is also of great importance from a practical point of view, e.g. for health, biotechnology, security and defence.

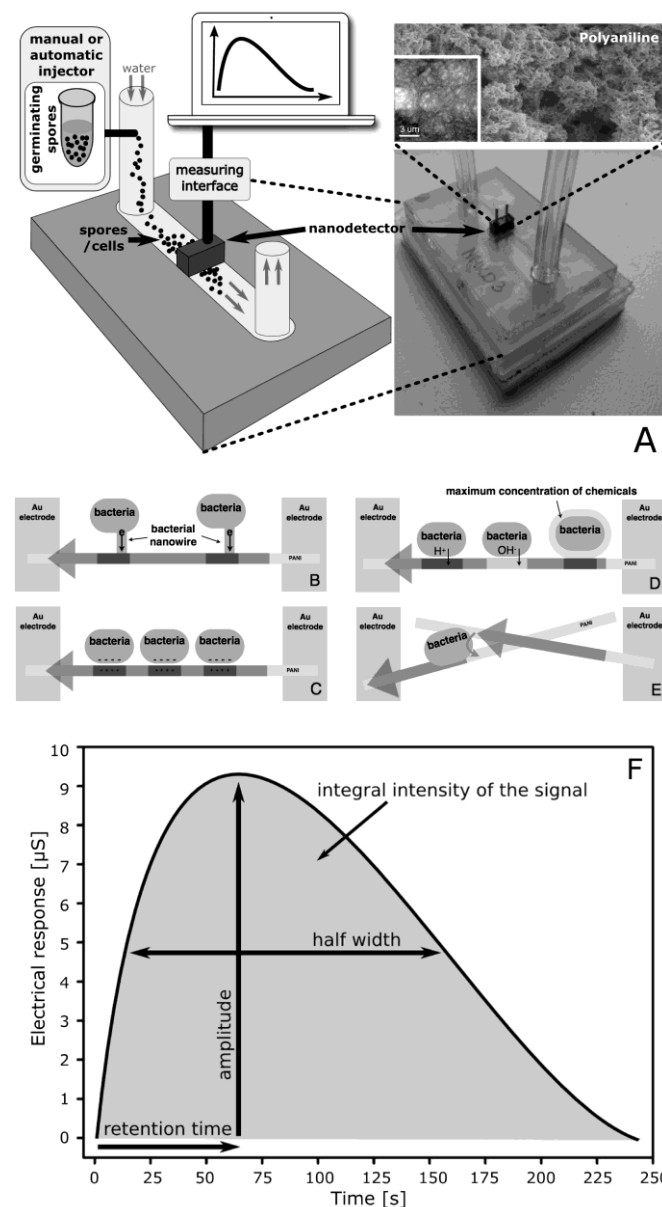


Fig 1. (a) Measurements are performed using a nanodetector which is designed as a lab-on-a-chip device operating in a continuous flow regime. The key component is a free-standing polyaniline (PANI)

nanofibre network which is electrochemically formed between two gold micro electrodes (inset: the initial stage of nanofibre formation, in a separate experiment which was done to help visualise the process). Modification of the nano-network's electrical conductivity, induced by the microorganism cells, is a source of the nanodetector's electrical response. The electrical conductivity of polyaniline nanofibres in contact with living cells is modified by: (b) the extracellular electron transfer (EET); (c) the presence of a cell surface charge and the FET mechanism; (d) the secretion of chemicals; (e) due to the quantum effects that occur in quasi-one-dimensional nanofibres (high electrical conductivity is possible only along PANI nanofibres and not perpendicularly, through sidewalls). The electrical conductivity is registered in a function of time (f). The result (time profile of the electrical response) is characteristic of different cells. The amplitude, and more precisely the integral intensity of the electrical response indicates the approximate number of cells in the sample. The retention time depends on the strength of interactions of cells and polyaniline nanofibres. The half-width depends on the number of modes of interactions.

Results and discussion

The basic, standard version of a continuous flow nanobiodetector (CFNBD) was successfully tested only against bacteria vegetative cells^{11,12,15}. This and similar systems have never been used before to detect spores, and particularly to monitor germination of bacterial spores, which is much more challenging as an analysis of a complex and dynamic process. Thus both the device and the method described here are unique.

Mechanism of detection

Modification of the electrical conductivity of the nano-network, induced by the spores and cells, is a source of the electrical response of the nanodetector. The detection mechanism is far from a routine approach. It is based on the non-specific (physicochemical) interactions of cells and spores which, when they touch the surface of the conducting polymer nanofibres, leads to a modification of the electrical conductivity of individual nanofibres and, consequently, of the whole network. These include direct charge injection, reversible chemical modifications of the nanofibrils, Coulomb interactions leading to FET-type features (FET – field-effect transistor), and quantum effects associated with the reduced dimensionality of individual nanofibrils (Fig. 1b,c,d,e).

The electric field generated by ions or the surface charge of cells is able to locally modify the charge carriers density in the nanofibres, which above the percolation threshold leads to an increase in the electrical conductivity due to the FET-type mechanism. Similarly, direct charge injection from the cells influences the electrical properties of the polyaniline nano-network. Quantum effects result in limiting the electrical conductivity only to one dimension, i.e. along the main axis of the nanofibres. Cells attaching to the surface of the nanofibres (even for a very short period of time) are responsible for breaking the limit, which causes short-circuiting between the fibrils and influences the electrical conductivity of the whole network. And, finally, chemical compounds released by cells in the closest vicinity of the nanofibres are active against the conducting polymer, such as polyaniline, which is electro-active and pH-sensitive (obviously other polymers of similar properties can also be used but have not yet been tested in our laboratory). Local chemical modifications induced by the cells result in a change of the electrical conductivity of the nanofibres above the percolation

threshold.

Owing to all of these effects, a considerable increase in the electrical conductivity is measured directly for the whole nano-network. The response is proportional to the density of the modifications, i.e. the number of cells. Because there are no stable links between the cells and the nanofibrils the processes are dynamic, that is why the response of the nanodetector, which is working in a continuous flow regime, is time-dependent.

Obtaining the signal of germinating spores

Injection of a small sample (100 μ l, approximately 10^5 spores) into a narrow channel of the lab-on-a-chip device, where a continuously flowing stream of water transports the sample into the detecting area of the NBD unit, results in generating a dynamic response that is registered as an electrical signal in a function of time (Fig. 1f). The time-profile of the response is characteristic of different types of cells, including spores and vegetative cells. This is caused by considerable differences in the size and morphology of the cells, in the surface structure and in all biochemical processes. The retention time of the sensor's electrical response depends on the strength of the interactions between cells and the polyaniline nanofibres. The effects measured are related to cell morphology and cell size, to the cell wall structure, cell surface hydrophobicity and the electrostatic surface charge of the cell. The half-width of the sensor's electrical response depends on the number of modes of interactions, which is related to the shape of the micro-organism, the presence of protrusions, tendrils, cilia on the outer surface of the bacteria, and on the symmetry of the electric charge distribution at the surface of the cells.

Germination of bacterial spores is associated with a breakdown of their structure, followed by dramatic changes in their chemical and physical properties – particularly of the surface layer, which can easily be detected and identified with the aid of a nanodetector due to the differences in the interactions of each type of cell (spores vs. transition forms vs. vegetative cell).

The data collected during spore germination are used to calculate the integral intensity, retention time and half-width of the registered signal (time profile). The integral intensity and signal amplitude are used to estimate the number of spores or vegetative cells in the sample analysed. The retention time, which is sensitive to the mobility of the cells, defines the type of analysed material (spores/transition forms/vegetative cells); the half-width is associated with interactions, including the influence of morphology of the analysed cells (shape and size). Thus, analysing the shape of the time profiles (Fig. 1f) allows us to distinguish between suspensions of spores, vegetative cells and their mixture, which also provides quantitative information about a mixture's composition.

Real-time monitoring of spore germination

The germination process is performed according to common procedure³ and monitored instantly by taking small samples (e.g. 100 μ l or less) every 5 min for injection into the lab-on-a-chip NBD system (Fig. 1a). The graph of integral intensity of the total signal measured versus the time of germination is sigmoid (Fig. 2a), which is typical of a transformation from one stage, i.e. the suspension of spores only, into the other, i.e. the suspension of vegetative cells, with a threshold at time 15 min after heat shock (65 °C for 30 min in

the presence of the germinates L-alanine and inosine). This corresponds to the dynamic transformation of *Bacillus subtilis* spores into vegetative cells. Analysing in detail the time profile of the signals, their half-width (Fig. 2b) and the retention time (Fig. 2c), it is possible to obtain direct insight into the interaction of spores and cells and the polyaniline nanofibrils. As can be concluded from the graphs (Fig. 2 a-c), the spores and vegetative cells interact with the polyaniline fibrils in a different way. The half-width of the signal measured for the cells (created from the germinating spores of *Bacillus subtilis*, is more than three times greater (68 seconds) in comparison to the spores of the bacteria (22 seconds). Similarly, the values of the retention time, measured at 10 and 60 min after administration of the germinates (mainly for the transition forms and vegetative cells, respectively), are 2.5 and 2.2 times longer as compared to those for the spores of *Bacillus subtilis* ATCC 6633. This is because of the differences in their surface properties, morphology and biological activity. Control experiments (calibration) using purified spores and vegetative cells, both having a defined population, showed that the electrical response of the nanodetector was proportional to the number of spores and vegetative cells in the sample being examined. However, the signal for vegetative cells was stronger, that is why we observed increasing integral intensity of the total signal measured with the time of germination (Fig. 2a). This can be explained by taking into account the difference in the morphology of the spores and cells and their different ability to interact with the nanofibrils. The conducting polymer (polyaniline) is in the form of positively charged macromolecules (poly-cation), which strongly interact with any negatively charged structures. Thus, having a polar-negatively charged surface, the vegetative cells are much more effective than spores¹⁶ in interactions with polyaniline nanofibrils, thus giving a stronger response of the NBD. In both cases the signal that was registered was proportional to the number of spores or cells, respectively, and in particular when taking into account its integral intensity. This has been proved in a series of calibration experiments. We were able to obtain even more information by analysing the time profile of the signals, i.e. the half-width and retention time (Fig. 2b,c), which gives direct insight into interactions between spores (or cells) and nanofibrils. One can clearly find that vegetative cells interact more strongly than spores, as expected when taking into account the difference in their surface properties, morphology and biological activity.

In fact, maximum interaction is observed for spores in transition form, 0-15 min after the heat shock. The signal of the transition form is even stronger than that generated by vegetative cells, thus a low population of the transition form in the steady state (below 500 000 ml⁻¹) can easily be detected. Interactions involving spore transition forms are the most intensive and effective. This is due to their unique surface structure¹⁷. Adhesion of such a form is stronger, thus it leads to a longer retention time and a broader signal (of a greater half-width) for spore transition forms than in any other case (Fig. 2b). Modification of the electrical conductivity of nanofibrils is stronger owing to the interaction of the charged surface of the transformed spores and also because of the direct chemical influence of “biomarkers” produced by the germinating spores², which can act onto the nanofibrils at the highest possible concentration due to close contact at the nano scale level.

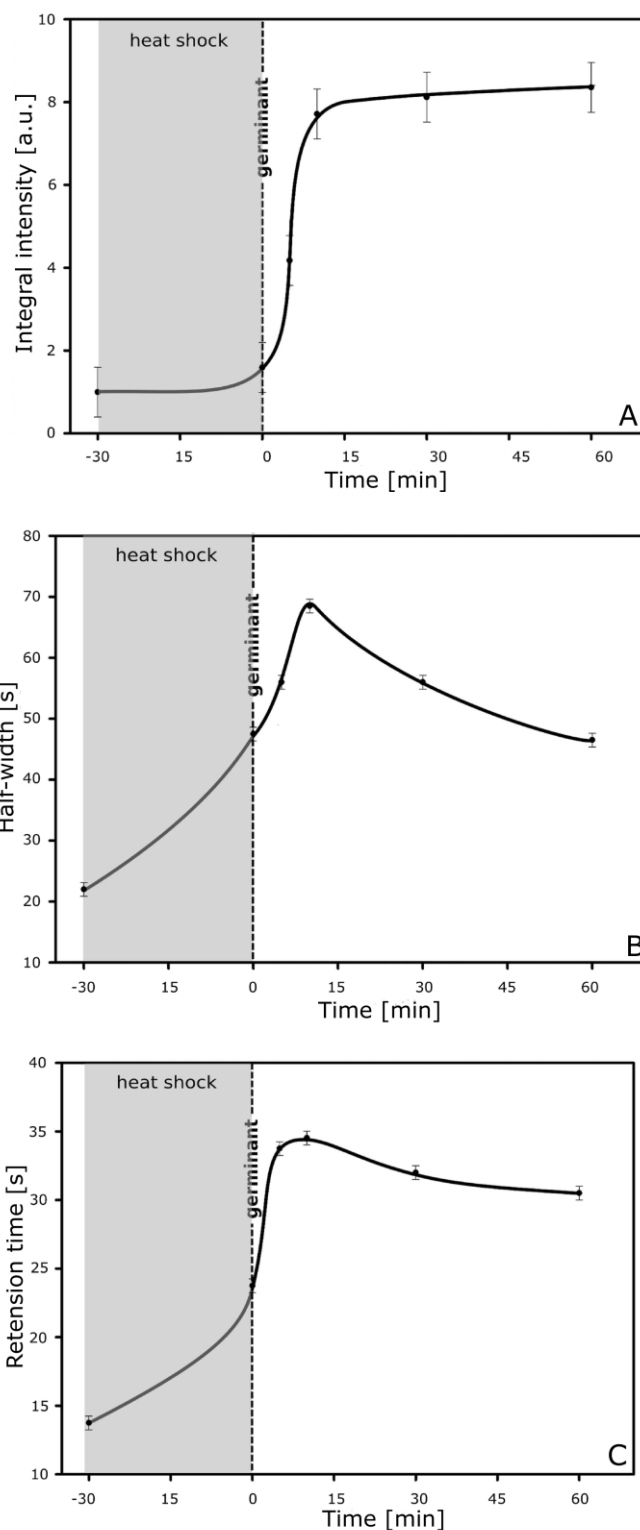


Fig 2. Graphs of (a) integral intensity, (b) half-width and (c) retention time for the signal measured during the whole germination experiment.

Defining specific signals for spores, vegetative cells and the transitional form

When analysing the amplitude of signals at the appropriate retention time (the amplitude is proportional to the number of spores or cells), i.e. 14 s for spores, 30 s for cells and 35 s for the transition form (the values defined from the plot of the RT vs. the duration time of the process, Fig. 2b), one can find information on the dynamics of the main processes induced during germination, including the formation of a transition form of spores before its final transformation into vegetative cells. The results are summarised in Fig. 3. The transition form of spores appears during heat shock, reaching a maximum concentration just after the thermal treatment within the next 15 min. By analysing the data collected in detail we can obtain more information about kinetics. Germination is a complex process which is composed of two main successive stages – [I] annihilation of spores (S) and formation of transition forms (TF), [II] transformation of transition forms into vegetative cells (C):



The process is described by a well-known formula:

$$-dP_S/dt = k_I P_S \quad (1)$$

$$dP_C/dt = k_{II} P_{TF} \quad (2)$$

$$dP_{TF}/dt = k_I P_S - k_{II} P_{TF} \quad (3)$$

and

$$P_S + P_{TF} + P_C = P_{S_0} \quad (4)$$

where P_{S_0} is the initial population of spores per unit volume (1 ml), P_S , P_{TF} and P_C are the population of spores, transition forms and cells per unit volume (1 ml) at time t , respectively; k_I and k_{II} are rate constants for process I and process II.

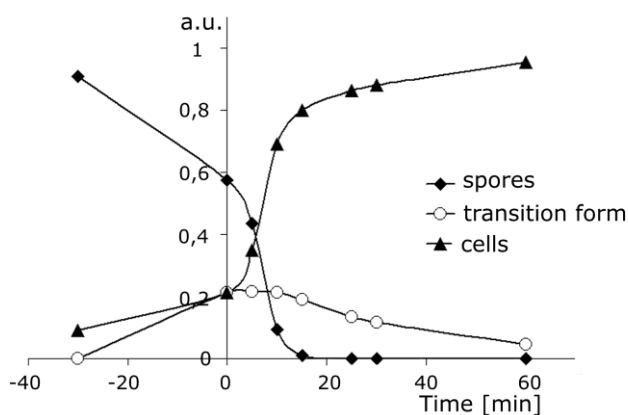


Fig 3. Changes in the population of spores (P_S , diamonds), transition forms (P_{TF} , circles) and vegetative cells (P_C , triangles) vs. time of germination, measured as the amplitude of signals at retention times of 14s, 30s and 35s, respectively (Fig. 2b). The values are normalised to 1 according to formula (4), where $P_{S_0} = 1$ [a.u.]. By analysing these data it is possible to obtain information on the kinetics of the germination process, including the formation of transition forms of spores before their final transformation into vegetative cells.

Based on calibration (an amplitude of 1 a.u. corresponds to 2×10^6 CFU/ml spores, or 2×10^5 spores injected) analysis of the experimental data presented in Fig. 3, one can obtain information on the rate of annihilation of spores (I) during heat shock (-30 to 0 min), which is 373 cell/s (in 1 ml). The rate increases dramatically up to 1600 cell/s (in 1 ml) under the influence of inducers of germination (inosine and L-alanine), just after their addition at time “0” (Fig. 3, Table 1). The rate of changes in the population of transition forms (TF) within 0-10 min is 0 because the rate of the transformation of spores in TF is equal to that of the formation of cells from TF. This is a quasi-stationary state. Then the population of TF decreases with a rate of 80 cell/s (in 1 ml) within a time range of 30-60 min. Formation of vegetative cells is most intensive (1600 cell/s in 1 ml) in the time range of 0-10 min. At the final stage the population of cells increases slowly with a rate of 80 cell/s (in 1 ml), which is equal to the value observed for the annihilation of the transition form (TF).

Table 1 Rate of transformation of *Bacillus subtilis* spores into their transition forms and, subsequently, the transition forms into vegetative cells (changes in the population) [cell / s*ml].

Time range [min]	-30 to 0	0 to 10	30 to 60
S, spores	-373	-1600	0
TF, transition forms	237	0	-80
C, cells	237	1600	80

Comparison with other methods

The results presented here are consistent with those obtained in parallel experiments performed in our laboratories using a conventional technique which is based on fluorescence measurements for detecting the activity of spore-derived esterase as an indicator of germination and a quantitative parameter of its progress. The plot presenting changes in the intensity of fluorescence of fluorescein in time, related to esterase activity from the germinating spores of *Bacillus subtilis* ATCC 6633 shows that the spores remain almost dormant in the absence of germination factors (an increase of fluorescence is low). These results correspond to those obtained using the bioluminometry method as reported elsewhere¹⁸. As expected, the addition of inosine and L-alanine changes the situation, as these compounds are well-known inducers of the germination of spores. They influence spore surface receptors which promote awakening from the dormant state. Then esterase biosynthesis leads to increased transformation of FDA-fluorescein and to more intensive fluorescence. The rate of increase of the fluorescence is 0.5 a.u./s at the beginning of the process (0-400 s after adding the germinants) and increases to 1.25 a.u./s within the next 600-800 s (Fig. 4). This corresponds exactly to the time-range where the most highly dynamic processes are observed with the use of a nanodetector (Fig. 3).

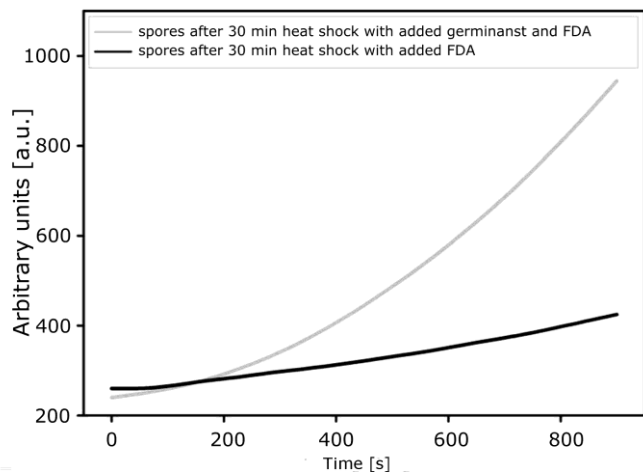


Fig 4. Changes in the intensity of fluorescence of fluorescein (generated from the biomarker FDA-fluorescein) during germination of *Bacillus subtilis* ATCC 6633 with and without germinants.

Experiments performed by Guiwen Wang et al. at the Yong-qing Li laboratory¹⁹, in which Raman spectroscopy and differential interference contrast (DIC) microscopy were used to analyse the kinetics of L-alanine-induced germination of wild-type *B. subtilis* spores, confirm the time intervals of the germination phases obtained with our method. A clearly visible change of DIC images between 10–20 minutes after addition of L-alanine corresponds to the most dynamic changes observed with our method (Fig. 3). This is concomitant with the completion of release of dipicolinic acid (CaDPA) by the spores, as confirmed by Raman spectroscopy with LTRS¹⁹. Pandey et al. examined the heterogeneity and germination of *Bacillus subtilis* spores with the help of a phase-contrast microscope. They found that most of the analysed *Bacillus subtilis* germinated within 3–5 minutes⁷. The 3–5 minute interval in the germination of *Bacillus* spores corresponds to the most dynamic changes and the maximum population of spore transition forms, which is clearly visible when using our method (Fig. 2b and Fig. 3).

The method presented here is adequate for the real-time monitoring of transformation of spores into vegetative cells (the germination process) by simultaneous observation of spores, spore transition forms and vegetative cells in a small but representative sample (10^5 spores). This is achieved by using one, single nanodetector which has been designed as an easy-to-handle lab-on-a-chip unit.

By analysing various parameters of the time profile of the electrical response, i.e. the amplitude, half-width and the retention time, which are specific for the analysed spores and vegetative cells, it is also possible to detect the presence of highly metabolically active transition forms of spores. The signals, obtained from the nanobiodetector between the fifth and twentieth minute after administration of a germinant, are dominated by a highly active form of *Bacillus subtilis* which is neither a spore nor a vegetative cell. The maximum activity of the transitional form falls on the 45th minute of the experiment, i.e. exactly 15 minutes after exposing the spores (thermally activated) to contact with L-alanine and inosine, which are used as the chemical activators of germination. Prior to that, the sample was dominated by spores of *Bacillus subtilis*, whose number decreased due to the progress of thermal activation. Hence, the amplitude of the signal registered at the retention time, which has been identified as characteristic of spores, was reduced by about half

during the 30 minutes of heating. The results are fully consistent with previous, independent studies of the process of transformation sequences of bacteria of the genus *Bacillus*⁷. Additionally, what is worth noting is that the compliance detail also applies to the time scale.

Experimental

The detecting system and procedure

The mechanism of the device is basically similar to that of a standard continuous flow system, CFNBD, which has been successfully applied for bacteria detection. However, now the detector was transformed to a compact lab-on-a-chip system (NBD) supported by a new, more accurate analytical procedure and data processing to reach improved time resolution (in seconds) and sensitivity (0.001 μ S), both of which are necessary to monitor the germination of spores.

The detecting device consists of two gold electrodes as simple bars (2 mm x 0.5 mm x 0.001mm) which are deposited on a plastic substrate with the use of a Quorum Technologies Q150R sputter coater. The electrodes are in good contact with the polyaniline nanofibrils formed over a narrow gap (5–10 μ m) between the Au microelectrodes so that the current flows through the dense conducting network.

A freestanding polyaniline nanofibril network (Fig. 1a) was prepared by direct synthesis onto Au microelectrodes by electrochemical oxidation of aniline hydrochloride (10% in water, pH \sim 1) at a potential of 0.8–1.2 V vs. Ag/AgCl with a controlled charge flow (using Potentiostat Autolab PGSTAT). The morphology of the PANI nanofibrils was examined with a scanning electron microscope (ZEISS EVO 40): the average length was 3–5 μ m (maximum above 10 μ m) and the thickness of single fibrils was about 100 nm or less (Fig. 1a).

To limit the ion current, gold microelectrodes were protected by a self-assembled monomolecular insulating layer formed of benzylthiol (after deposition of PANI nanofibrils). A solution of benzylthiol in methanol (10 mg/ml) was used. The reaction was carried out for 180 s at room temperature and then the electrodes were carefully washed with methanol and water. The insulating monomolecular layer considerably improved the electrical response of the nanodetector in the presence of the microorganisms by a significant reduction of the ion current.

The detecting unit was then assembled inside a micro-fluidic platform which consisted of three elements of poly(methyl methacrylate) (32 mm x 50 mm). The outermost element consisted of two inlets for silicone tubing with an internal diameter of 0.2 mm at a distance of 32 mm apart (Fig. 1a). In between there was a hole (3 mm x 6.5 mm) to insert the polyaniline detecting unit which was sealed with a silicone seal. In the middle element a microfluidic channel was formed (3 mm x 3 mm x 32 mm, or 3 mm x 100 μ m x 32 mm) that connected both inlets. All of the elements were joined using Tesa sticky tape (double-sided and waterproof).

The system was equipped with a peristaltic pump (GILSON MINIPULS 3), a measuring interface (e.g. conductivity meter ELMETRON CC 505, which is a tool that is safe for a nanodetector, effective and convenient, despite being used in a non-traditional manner to measure electrical conductivity and temperature at the same time and with very good accuracy) and a PC computer. Bi-distilled (BiD) or sterile deionised water was used as a medium to

transport the samples analysed through the detecting area (flow rate about 1 ml min^{-1}). Generally, a small volume ($100 \mu\text{l}$ or less) of the suspension of germinating spores of bacteria in water was injected every 5 min into the detecting system through a septa. Then the electrical conductivity was measured in a function of time. Changes in the electrical conductivity of nanofibres caused by the presence of spores and cells were plotted in a function of time (a time profile of the electrical response) and analysed. Parameters of the time profile, such as retention time RT and half-width HW, were characteristic of different microorganisms, and the intensity (or amplitude) of the signal was proportional to the number of cells. By measuring the amplitude at the appropriate RT one can obtain information on the components of a complex mixture of microorganisms.

For monitoring the germination of the spores of *Bacillus subtilis* ATCC 6633 with the NBD system, samples of $100 \mu\text{l}$ of suspension containing about 2×10^6 spores/ml were injected with a micro syringe. The average number of spores injected was about 10^5 , however, the device was able to work efficiently also in the case of a lower and higher number of spores, from 10^4 to 10^9 , respectively. After injection the electrical conductivity of the nanofibril network was measured with an accuracy of $0.001 \mu\text{S}$ and registered in a function of time within 300 s (every 1s with an accuracy of 0.1 s), as the time profile of the electrical response of NBD. Samples of $100 \mu\text{l}$ (about 10^5 spores) were collected and injected into the device through the septa according to the scheme: at the beginning of the experiment (time: 30 min), at the end of heat shock (time: 0 min), then at 5 min, 10 min, 15 min, 25 min, 30 min and 60 min after heat shock. This is reasonable because the process is most dynamic at the time of 0-10 min after heat shock, although another sampling time is also possible.

In order to compare the results, measurements were performed with:

- washed suspension of intact spores,
- water solutions of L-alanine and inosine (blank test),
- spore suspension after heat shock
- and spore suspension after heat shock with the addition of L-alanine and adenine (0, 5, 10, 15, 25, 30 and 60 minutes after the germinators had been added).

The measurements are triplicate.

Bacterial cell preparation

Growth and preparation of spores

Bacillus subtilis ATCC 6633 was grown on a standard agar plate with 5% sheep blood (MAST DIAGNOSTICA) in $37 \text{ }^\circ\text{C}$ (aerobic conditions). After 24 h, single colonies were transferred to flasks (100 ml) with TSB medium (50 ml). Liquid cultures were grown in a shaking incubator (24 h, $37 \text{ }^\circ\text{C}$, 115 rpm). Afterwards, 10 ml of liquid culture was transferred to a 1-litre Roux bottle containing 200 cm^3 sporulating agar (Sporulating Agar AK ,BTL). Roux bottles were incubated in $37 \text{ }^\circ\text{C}$ for 24 h and subsequently for 4 weeks in room temperature. After 28 days, $10\text{-}25 \text{ ml}$ of sterile distilled water was added to each bottle (depending on the density of suspension) and incubated for 1 h in $37 \text{ }^\circ\text{C}$. Then the bottles were washed by very careful and gentle shaking (additionally, $10\text{-}20$ of sterile medium-sized ($3\text{-}4 \text{ mm}$) glass beads were added to intensify the washing). This step was repeated if a significant amount of material was left on the agar surface. Then the obtained suspension was filtered with gauze (triple-folded) and centrifuged ($10\,000 \text{ RCF}$, 20 min , $4 \text{ }^\circ\text{C}$). The supernatant was removed and the pellet was suspended in 10 ml of sterile saline (in this step, additionally $10\text{-}20$ sterile medium-sized glass beads were added to intensify the mixing). The procedure was repeated. Then the pellet was suspended

in 10 ml of 65% isopropanol and incubated for 3 h in $20 \text{ }^\circ\text{C}$ in order to destroy any remaining vegetative cells. Afterwards the suspension was centrifuged ($10\,000 \text{ RCF}$, 20 min , $4 \text{ }^\circ\text{C}$), the supernatant was removed and the pellet was suspended in 10 ml of sterile deionised water at $4 \text{ }^\circ\text{C}$. This step was repeated five times. The quality of the spores was determined by microscopic examination to confirm a lack of debris from cell lysis and rods. After washing the pellet containing spores was suspended in sterile deionised water. The number of spores (cfu/ml) was estimated by 10-fold serial dilutions (triplicate) which were plated on nutrient agar Petri dishes. After 24 h, plates with a countable number of colonies were selected, colonies were counted and the final concentration was calculated.

Growth and preparation of vegetative cells

Bacillus subtilis ATCC 6633 was grown at $37 \text{ }^\circ\text{C}$ in a nutrient broth (BTL) with shaking (24 h and 4 h, 115 rpm). Vegetative cells were washed four times with cold ($4 \text{ }^\circ\text{C}$) deionised sterile water and then suspended in cold sterile deionised water. The quality of the vegetative cells was examined by microscopic observation, looking for the absence of spores. Fresh vegetative cells were prepared each time when required. The concentration of the vegetative cells was estimated by preparing serial log-dilutions (the method described above for spores) and expressed as colony-forming unit (CFU) per ml.

Parallel conventional monitoring of spore germination.

In order to compare the results, spore germination was tested in parallel at the same time by using a method in macro scale³. Spores derived from esterase activity were used as an indicator and a quantitative parameter of germination progress. For this purpose the spores were washed four times with cold ($4 \text{ }^\circ\text{C}$) deionised sterile water by centrifugation and were subsequently resuspended in water. The final pellet was suspended in cold ($4 \text{ }^\circ\text{C}$) sterile water at a concentration of about $2 \times 10^6 \text{ cfu ml}^{-1}$. The suspension of spores (1 ml) was incubated at $65 \text{ }^\circ\text{C}$ for 30 min for induction of germination. To boost germination by receptor interactions, after heat shock the solution of L-alanine and inosine (Calbiochem) was added at a concentration of 10 mM and 2 mM , respectively. Non-fluorescent fluorescein diacetate (FDA) (SIGMA-Aldrich), which is a substrate for the cellular esterase enzyme, was used as a germination indicator. The germination process stimulated the enzyme release from waking spores and thus started enzymatic hydrolysis of the FDA. The product of hydrolysis was fluorescein, which is a well-known fluorescent compound. The concentration of the fluorescent product was measured using the F-4500 HITACHI fluorescence spectrophotometer. Solid FDA (Sigma-Aldrich) was dissolved in acetone (POCh) to reach a concentration of 2 mg ml^{-1} and stored as a stock solution at $-20 \text{ }^\circ\text{C}$. Prior to each test the FDA was diluted to a concentration of 0.5 mg ml^{-1} , and $10 \mu\text{l}$ was added to 1 ml of each medium or each media component (FDA final concentration was 0.005 mg/ml). Measurements were conducted in quartz cuvettes containing 1 ml of spores suspension (about 2×10^6 spores/ml) in 0.1 M phosphate buffer (pH 7.6). For the measurements the following conditions were applied – excitation wavelength 488 nm , emission wavelength 526 nm , time-scan period 15 minutes, time accuracy 1 second, photo multiplier voltage 750 V , EX/EM slit 10 nm . The results are presented in Fig. 4.

Conclusions

The new lab-on-a-chip type device with a freestanding polyaniline nano-network allows to monitor the germination process in real time. The method is based on non-specific physicochemical interactions of cells and spores with conducting polymer nanofibrils, which results in a modification of the electrical conductivity of individual nanofibrils and, consequently, of the whole network. This generates a dynamic-type electrical response which is characteristic of different types of cells, including spores and vegetative cells. This is caused by the considerable differences in the size and morphology of spores and vegetative cells as well as in the surface structure and biochemistry of all processes influencing modification of electrical conductivity of polyaniline. All of these differences in interactions between each type of cells and the polyaniline conducting network, spores, transition forms and vegetative cells, can be detected with the aid of the described here device. Our results are fully consistent with previous independent studies of the germination process of bacteria of *Bacillus* genus. The method, based on non-specific physicochemical interactions in a nano-scale, is universal and could be useful for advanced studies of the kinetics and the mechanism of other microbiological processes. In a microfluidic version, it is a fast analytical technique, accurate enough for many applications in microbiology, biotechnology, health, security and defence, where spores are in use (proven sensitivity is of 10^4 spores injected, and the range of application exceeds a value of 10^{10}). Other possible applications are briefly presented in Supplement.

Acknowledgements

The work was supported by DARPA (USA) within project no. N10PC20099 and in part by the Srem Borough Council.

Notes and references

^a Faculty of Chemistry, Laboratory for Materials Physicochemistry and Nanotechnology, A. Mickiewicz University in Poznan, Srem, 63100, Poland.

^b Biological Threats Identification and Countermeasure Centre of Military Institute of Hygiene and Epidemiology, Pulawy, 24100, Poland.

^c Military Institute of Hygiene and Epidemiology, Warszawa, 01163, Poland.

- 1 E. Abel-Santos, *Bacterial Spores: Current Research and Applications* (UK: Caister Academic Press), 2012
- 2 P. Setlow, *Curr Opin Microbiol*, 2003, **6**, 550
- 3 L. Ferencsko, M.A. Cote and B. Rotman, *Biochem Biophys Res Commun*, 2004, **3**, 854
- 4 C. Laflamme, J. Ho, M. Veillette, M. C. de Latrémouille, D. Verreault, A. Mériaux and C. Duchaine, *Arch Microbiol*, 2005, **2**, 107
- 5 D. Chen, S.S. Huang and Y. Q. Li, *Anal Chem.*, 2006, **19**, 6936
- 6 L. Kong, P. Zhang, G. Wang, J. Yu, P. Setlow and Y. Q. Li, *Nat Protoc*, 2011, **5**, 625
- 7 R. Pandey, A. Ter Beek, N. O. Vischer, J. P. Smelt, S. Brul and E. M. Manders, *PLoS One*, 2013, **3**, e58972
- 8 D. D. Jr Evanoff, J. Heckel, T. P. Caldwell, K. A. Christensen and G. Chumanov, *J Am Chem Soc*, 2006, **39**, 12618
- 9 J. K. Daniels, T. P. Caldwell, K. A. Christensen and G. Chumanov, *Anal Chem*, 2006, **5**, 1724

- 10 Y. S. Liu, T. M. Walter, W. J. Chang, K. S. Lim, L. Yang, S. W. Lee, A. Aronson and R. Bashir, *Lab Chip*, 2007, **5**, 603
- 11 J. J. Langer, K. Langer, *Rev Adv Mater Sci*, 2005, **10**, 434
- 12 K. Langer, P. Barczyński, K. Baksalary, M. Filipiak, S. Golczak, and J. J. Langer, *Mikrochim Acta*, 2007, **159**, 201
- 13 S. Pal, E. C. Alocilja and F. P. Downes, *Biosens Bioelectron*, 2007, **9-10**, 2329
- 14 Y. Liu, S. Chakrabarty and E. C. Alocilja, *Nanotechnology*, 2007, **18**, 424017
- 15 J. J. Langer, P. Barczyński, J. Warchoń, K. H. Bartkowiak and K. Langer, *Biosens Bioelectron*, 2009, **9**, 2947
- 16 E. M. Sonnenfeld, T. J. Beveridge, A. L. Koch and R. J. Doyle, *J Bacteriol*, 1985, **3**, 1167
- 17 A. Driks, *Microbiol Mol Biol Rev*, 1999, **1**, 1
- 18 A. Bielawska-Drozd, *The ASA Newsletter*, 2005, **2**, 18
- 19 G. Wang, P. Zhang, P. Setlow and Y-Q. Li, *Appl Environ Microbiol*, 2011, **10**, 3368