This article is licensed under a Creative Commons Attribution-NonCommercial 3.0 Unported Licence.

Lab on a Chip



View Article Online **PAPER**



Cite this: Lab Chip, 2023, 23, 5165

Advancing in situ single-cell microbiological analysis through a microwell droplet array with a gradual open sidewall†

Jie Wang, ta Lin Du, ta ta Yuwei Han, Dawei Zhang ** and Dalei Jing**

The utilization of microfluidic analysis technology has resulted in the advancement of fast pathogenic bacteria detection, which can accurately provide information on biochemical reactions in a single cell and enhance detection efficiency. Nevertheless, the achievement of rapid and effective in situ detection of single-bacteria arrays remains a challenge due to the complexity of bacterial populations and low Reynolds coefficient fluid, resulting in insufficient diffusion. We develop microwell droplet array chips from the lateral hydrodynamic wetting approach to address this issue. The sidewall of the microwell gradually opens which aids in advancing the liquid-air interface and facilitates the impregnation of the solid microwells, preserving the Wenzel state and assisting in resisting the liquid force to separation from the drop. The feasibility of preparing cell arrays and identifying them inside the microwells was demonstrated through the simulated streamlined distribution of gradual and traditional microwells with different sizes. The water-based ink diffusion experiment examined the relationship between diffusion efficiency and flow velocity, as well as the position of the microwell relative to the channel. It showed that the smaller gradual microwell still has a good diffusion efficiency rate at a flow velocity of $2.1~\mu L$ min⁻¹ and that the infiltration state is easier to adjust. With this platform, we successfully isolated a mixed population containing E. coli and S. aureus, obtained single-bacteria arrays, and performed Gram assays after in situ propagation. After 20 hours of culture, single bacteria reproduced demonstrating the capability of this platform to isolate, cultivate, and detect pathogenic bacteria.

Received 6th July 2023, Accepted 3rd November 2023

DOI: 10.1039/d3lc00590a

rsc.li/loc

Introduction

Infectious diseases pose a significant risk to human health, and the spread of pathogenic microorganisms can lead to widespread disease outbreaks. Patients with pre-existing chronic conditions or weakened immune systems are particularly vulnerable to infections caused by pathogenic bacteria.^{1,2} The development of precise and effective pathogen detection technologies is essential for the successful diagnosis and treatment of infectious illnesses since existing detection methods are sometimes time-consuming and difficult.^{3,4} The single-cell analysis (SCA) technique based on microfluidics provides detailed information intracellular substances and biochemical reactions. This makes it an invaluable tool for obtaining individual differences among pathogenic microorganisms at the singlecell level, thus facilitating a more comprehensive understanding of complex life processes.5-7 In the field of microbial physiology, cell phenotyping and histological analysis, the SCA technique has played a crucial role in improved pathogen detection tools and processes depending on the advantage in pathogen isolation and detection systems in recent years.8-10

The preferred method for bacterial single-cell analysis in SCA technology is droplet microfluidics, which is effective manipulation on a chip and enables multiple analyses of individual droplets for ultra-high throughput acquisition of single cells. 11-14 Among them, microwell droplet arrays have inherent advantages over continuous flow droplets in that they allow for in situ detection by fixed physical microwells, which create a more stable microenvironment, prevent sample contamination, and allow for in situ culture and colony analysis of single bacterial cells through cell attachment and diffusion.15-19 The capture of a single bacterium in a microwell to create a single bacterial array has led to major improvements in the creation and manufacture of bioapplications. With the aid of supporting instruments, it

^a School of Optical-Electrical and Computer Engineering, University of Shanghai for Science and Technology, Shanghai, China200093. E-mail: dwzhang@usst.edu.cn

^b School of Mechanical Engineering, University of Shanghai for Science and Technology, Shanghai, China200093. E-mail: jingdalei@usst.edu.cn

^c State Key Laboratory of Genetic Engineering, School of Life Science, Fudan University, Shanghai, China200433

[†] Electronic supplementary information (ESI) available. See DOI: https://doi.org/ 10.1039/d3lc00590a

[‡] These authors contributed equally to this work.

has been demonstrated that the morphological form and hydrodynamic properties of microwells permit the capture of a single bacterium. 20,21 However, due to the complexity of the microbial community and the low diffusion efficiency of low Reynolds number laminar flow, the droplet array used for single-cell analysis has certain limitations when it comes to effectively isolating and culturing pathogenic bacteria. These limitations include inadequate representation of cells and the inability to verify the activity of individual bacterial cells.²² With lateral contact flow²³ and wetting transition criteria, 24 it is possible to achieve single-cell separation inside the microwell by using a physical barrier to separate the target from the general flora in transitional methods of surface modification or valve control. 25,26 The energy barrier created at the liquid-air interface of a hydrophobic surface during single-cell separation controls the wetting transition from the Cassie to the Wenzel state, which is usually metastable within the constraints of a thin microstructure. Filling the voids of the rough surface morphology with liquid can result in a reversible change to the unstable transition state, causing non-intimate contact between solid and liquid and the introduction of air. 27-29 This significantly decreased the ability to identify pathogenic bacteria by separating them into microwells and staining them. As a result, it is necessary to create microwell droplet array chips for extensive pathogenic bacteriology research that are easy to use and affordable and limit droplet fusion while preserving analytes in tiny volumes. 30,31

In this study, we develop a microwell droplet array chip that expedites wetting transitions into stable droplet arrays in microwells, enabling more effective in situ detection of single-bacteria arrays. By using the lateral hydrodynamic approach, we were able to capture individual bacterial cells and provide ample growth space inside the microwell arrays. Our design employs a gradual opening trend, which is inspired by the surface form of a lotus leaf, to enhance the movement of the liquid-air interface and promote the preembedding of bacterial cells to the bottom of the well. Unlike previous methods, we focused on the stability of the liquidair interface rather than surface modification or valve control. The channel above the microwell provides a confined environment and allows for the diversion of reagents. Through our investigations of microwell and channel dimensions, relative positions, and flow rate variations, we have discovered a high diffusion efficiency within the system. Our chip has been successfully used for single-cell isolation and in situ propagation culture experiments of mixed pathogenic bacteria. The microwell droplet array chip makes it simple to speed up wetting transitions for in situ detection of single bacteria, as demonstrated in Fig. 1. A conceptual representation of the channels and microwells can be seen in Fig. 1(a). Four different Gram reagents were sequentially pumped onto the chip after it had dried to stain the bacteria. The cross-section of the microwells is shown in the diagram in A-A. The assembled device is shown in Fig. 1(b). The gradually opening shaped microwells are displayed at the

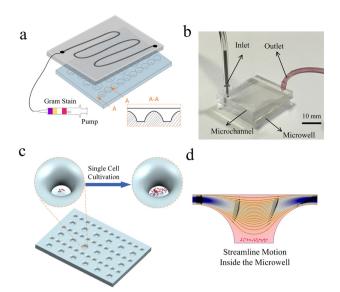


Fig. 1 Expediting wetting transitions of a microwell droplet array chip for *in situ* detection of single-bacteria arrays. (a) Conceptual diagram of the chip used for Gram stain. (b) Assembled device of the chip. (c) Detection and *in situ* cultivation of individual bacteria in microwell arrays. (d) Distribution of streamline within the microwell.

bottom, and the vertical sidewall microchannels are shown at the top. Fig. 1(c) shows a 3D schematic of single bacterium isolation and *in situ* cultivation in a microwell. The simulated streamlined motion of the reagents as they pass is shown in Fig. 1(d).

Experimental

Materials and equipment

Wet etching is used to create a glass-positive mould (Dongcheng Microfluidics Co., Ltd., Zhenjiang, China) for microwell array chip preparation. An SU-8 (2025, Kayaku Advanced Materials, Inc., MA) positive mould with a 50 µm width and a 25 µm thickness for a channel is created on silicon wafers (MJB4, SUSS Micro-Tec., Germany) by using the UV photolithography technique (see Fig. S1 and S2 in the ESI† for more details about the preparation process). The polydimethylsiloxane (PDMS) chips with microwells and channels are made of a Sylgard prepolymer and curing agent (184, Sigma-Aldrich, MO) in accordance with 10:1 from the glass positive mould and SU-8 positive mould, respectively. The inlet and outlet PDMS chips are punched with a 0.7 mm punch (iBiochip, China). With plasma surface treatment equipment (PDC-32G-2, Harrick Plasma, USA), two PDMS chips with microwells and channels are bonded together to obtain the microwell droplet array chip. A single disposable medical sterile 1 mL syringe and a syringe pump (PHD2000, Harvard Apparatus) are used to load various amounts of reagents into the chip. A charge-coupled device (CCD) camera (DP73, Olympus, Japan) is used to monitor the isolation and in situ propagation of single bacterial cells in the microwell arrays under an Olympus IX71 microscope.

Lab on a Chip

Bioreagents and pathogenic bacteria

Bacterial single-cell isolation and in situ propagation validation were carried out using S. aureus (29213, ATCC, VA) and E. coli (25922, ATCC, VA). The bacteria are measured and expanded quickly in a Luria-Bertani broth medium (HBO128, Qingdao Haibo, China) and blood agar plates (Bkmam, China) at 37 °C. A Gram Stain Kit (G1060, Solarbio, China) is used to stain for the identification of colonized bacteria. The concentration of the bacterial solution is figured out by contrasting the turbidity of the bacterial solution with that of a Mackenzie turbidimetric tube. At enough concentration of 1.5×10^3 copies per mL in accordance with the McFarland Equivalence Turbidity Standard, we carried out single-cell isolation tests for S. aureus. The bacteria are first purified by overnight incubation on blood agar plates, and the colonized bacteria are then transferred into a Luria-Bertani broth medium to obtain an appropriate concentration of the bacterial solution with a Mackenzie turbidimetric tube. A mixed flora solution was made in the same way. A droplet array is created by loading the solution onto a microwell array chip. It is incubated for in situ growth for 20 hours at constant humidity and standard atmosphere. We pre-embedded the bacteria into the bottom of the microwell and applied the Gram stain. Unlike regular staining, the channel is vented with air after each staining to "blow away" the staining chemical and prevent harm to bacteria caused by washing. Finally, the chips can be sent for observation.

Simulation model of microwells

The multiphysics field simulation program (COMSOL Multiphysics 5.6) microfluidic module is used to model the flow in microwells. The entire system is sealed and the slow laminar flow within the channel can be analysed for bacteria due to the presence of a drainage channel above the microwell. The sidewall expansion type microwell is formed by two 1/4-like circular arcs. The depth of the narrow channel above the microwell is 25 µm. We investigated how the geometry of the microwell affects the flow pattern. Simulations were performed to observe the effect of flow lines generated near individual microwells on the distribution of bacteria. The working fluid is assumed to be water (a homogeneous, incompressible Newtonian fluid; density 998.2 kg m⁻³, dynamic viscosity 0.001 kg m⁻¹ s⁻¹) and the flow is assumed to be laminar steady flow. Based on the Navier-Stokes momentum conservation equation, the steadystate velocity field and streamline in the microwell are solved. No-slip conditions are used on the walls of channels and microwells. The inlet solution is an aqueous solution, and the inflow velocity is 0.1 m s⁻¹.

Results and discussion

Transition principle for droplets and diffusion

The hydrophobic effect refers to the phenomenon caused by the reduction of the solid-liquid contact area due to air pockets trapped in the rough, porous microstructure. When the droplet is stationary on an ideal solid surface, Young obtained the equilibrium equation based on the principle of mechanical equilibrium³²

$$\sigma_{\rm sl} + \sigma_{\rm la} \cos \theta_{\rm Y} = \sigma_{\rm sa}$$

where $\sigma_{\rm sl}$, $\sigma_{\rm la}$ and $\sigma_{\rm sa}$ are the interfacial tensions between solid-liquid, liquid-air, and solid-air, and $\theta_{\rm Y}$ is the contact angle formed by the tangent of the solid-liquid interface and the tangent of the liquid-air interface. A droplet is said to be in the Wenzel state when it fills the cavity within the microstructure of a rough surface and is in complete contact with the rough surface.³³ When the wetting state does not occur and the air pockets are stable in the microstructure of the rough surface, it is called the Cassie state.³⁴ The hydrostatic pressure at which a superhydrophobic surface departs from the Cassie state, whether it reaches the Wenzel state, is referred to as critical pressure. 35,36 Therefore, from the perspective of force balance, we search for the critical pressure at which the liquid-air interface is depinning from the top of the microstructure to judge the occurrence of the wetting state transition process. 37-39 And this process is also related to the cross-sectional shape of the well. 40 Fig. 2a is a sketch of the liquid-air menisci corresponding to rectangular and gradient expansion microwells. Applying balance of forces and using the Young-Laplace law, one obtains

$$P + P_{\infty} - P_{\alpha} - \sigma \nabla \cdot \vec{n} = 0$$

where P, P_{∞} , and $P_{\rm a}$ are, respectively, the hydrostatic, atmospheric, and entrapped air pressures, σ is the surface tension, and \vec{n} is the interface unit normal vector. The critical pressure mostly is obtained by force balance analysis and depends on the homogeneity of the meniscus slope near the well walls. When its shape becomes more circular, the resistance of the pores to hydrostatic pressure increases.

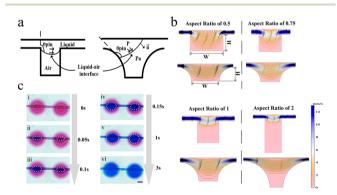


Fig. 2 Wetting transition on the microwell droplet array chip. (a) The liquid-air meniscus transition process occurs in vertical sidewalls and gradual opening trend sidewalls. (b) Visualized fluid flow paths in gradual opening trend microwells, for W/H of 200/100, 100/75, 100/100, and 100/200 μ m, respectively. The black solid lines with arrows indicate streamlines. (c) Diffusion tests on the microwell droplet array chip with gradual opening trend sidewalls. Where i–vi show the diffusion of blue ink at the corresponding moments. The scale bar is 50 μ m.

The movement of the forward liquid-air interface is enhanced by the gradual unfolding design of the sidewalls of the microwell, the bottom of the microwell is in contact with the liquid-air interface, and the liquid exhibiting the Cassie state on the hydrophobic surface is transformed into the Wenzel state. Although the Wenzel state is at a lower energy state, the energy potential barrier needs to be crossed for the system to change the permeability state. Ref. 41 discussed the effect of substrate surface topography on wettability, which was inspired from here. We changed the sidewall of the microwell from a vertical type structure to a 1/4 rounded type structure to enhance the transformation process of the depinning. The transition process of the two sidewall structures off the depinning is shown in Fig. 2a, where θ_{pin} is the contact angle of the liquid-air interface on the microwell sidewall, and the contact line is shifted when θ_{pin} is greater than the intrinsic contact angle of the material. In the Cassie state, the liquid-air interface is suspended between the vertical sidewalls (as shown on the left in Fig. 2a). The interface is curved due to the Laplace pressure inside the liquid. If the suspended interface does not remain fixed to the top of the sidewall, then it will be moved down into the microwell and wet on the surface. If the contact angle formed at the liquid-air interface is greater than the advance angle at that point, it will lead to depinning and the system will become the Wenzel state (as shown on the right in Fig. 2a). In contrast to vertical sidewalls, the pressure-volume energy of the liquid is modified by the gradually expanding sidewall design. The transition caused by depinning can also be considered as a process in which the work done by pressure is greater than the energy potential generated by the surface energy associated with the wetting roughness.

Fig. 2b is the steady-state flow velocity field and streamlined profile of the rectangular and gradient expansion microwells. We found that the wet state was affected by the aspect ratio of the microwell. The smaller the aspect ratio of the microwell, the easier it is to be wet. However, when the aspect ratio reaches 0.5 and below, although the microwell can be wetted, the bottom has a strong streamlined distribution, which means that the flowing liquid will affect the internal state of the microwell. Through the comparison of the streamlined distribution and velocity field, it is found that the gradient expansion microwells are more conducive to the entry of fluid. Due to the surface energy of the liquid and the pinning effect, the liquid is confined by the rectangular microwell and flows directly through the channel.⁴² The streamlines with gradient expansion microwells^{43,44} of the aspect ratio of 1 have a relatively uniform slope, which indicates smooth velocity changes within the microwells and that resistance to water pressure is enhanced at the bottom. The bottom of the gradient extension microwells is not disturbed by any strong flow and therefore does not encounter damage to bacteria caused by the high shear forces that can occur in lateral flow. We used microwells with an aspect ratio of 1 to perform diffusion tests in which blue ink aqueous solution was substituted for the red, as shown in Fig. 2c, to confirm the diffusion effect in this manner. Fig. 2c, i represents the state inside the system when the blue ink has not yet diffused. The white dotted line in Fig. 2c, ii-vi indicates the border between the two solutions, and it shows how quickly the blue liquid entered the microwells during the first 0.1 s of the experiment. The subsequent stage of 3 s begins with the liquid progressively diffusing, and the blue liquid fills the well until the red liquid completely fills the pores (as illustrated in Fig. 2c, v and vi).

Study of the staining procedure of Gram stain

To utilize microwell arrays in single-bacteria assays, we investigated the impact of the alignment, intake flow rate, and diffusion efficiency of the two chips on pinning and depinning. The PDMS chip with microwells could produce droplets of approximately 5 nL and 30 nL for small and large wells, respectively. The depth and breadth of a PDMS chip with channels were 25 µm and 50 µm, respectively. After bonding the two chips, we evaluated the relationship between microwells and channels (R is the radius of the microwell, and d for the distance between its centre and the centreline of the channel flow channel). In Fig. 3a, we observed that depinning occurs when R/d is larger for both microwell 1 and microwell 2. For microwell 1, the liquid exits the channel and enters the microwell when R/d is equal to 2. When the liquid-air interface touches the bottom of the wall, the microwell quickly enters the Wenzel wetting condition. The liquid-air contact is essentially fixed to the edge of the microwell as R/d slowly drops. As a result of the closed air inside the microstructure, the microwell is currently in the Cassie state, preventing liquid from entering. The liquid is depinning from the channel into the microwell for microwell 2 when R/d is equal to 1.3. The liquid is essentially trapped at

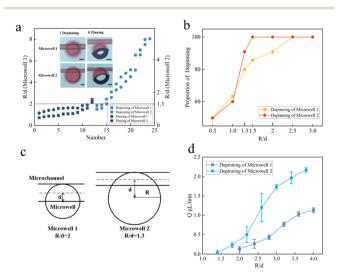


Fig. 3 Pinning and depinning based on the alignment and flow rate of the chips. (a) Influence of R/d. Micrographs depict typical depinning (i) and pinning (ii) outcomes. The scale bar is 50 µm. (b) Proportion of depinning at different R/d values. (c) Unique position of R/d in 100% depinning efficiency. (d) Intake flow rates on depinning. The error bars present standard deviations for 3 measurements.

Lab on a Chip Paper

the edge of the microwell as R/d steadily drops, preventing it from entering the interior of the microwell. This demonstrates that in the case of tiny R/d, the pinning of the liquid-air interface at the edge of the microwell results in the system being in the Cassie state, which is the non-wetting condition. As seen in Fig. 3b, depinning is more effective when the R/d is higher (i.e., the distance between the centreline of the channel and the centre of the circle is closer). When R/d is 2, the depinning efficiency for microwell 1 reaches 90.9%, and when it is increased to 2.5, it achieves 100%. This phenomenon was also seen in microwell 2, and when R/d was higher than 1.3, the depinning efficiency was 100%. The value of R/d at 100% depinning efficiency is hence referred to as a unique position. As seen in Fig. 3c, the location is currently tangent to both the upper border of the flow channel and the microwell. When R/d decreases, the channel is closer to the microwell centre from that particular point, and when it increases, the channel is farther from the microwell centre.

The condition involving depinning and pinning is influenced by the chips' alignment as well as the pace at which liquid flows through the channel. We looked at the impact that different intake flow rates had on depinning. The flow rate range of microwell 2 is higher than that of microwell 1, even when the flow rate exceeds 2.1 µL min⁻¹, depinning may be generated at a rate within the ranges of the error bars, as illustrated in Fig. 3d. The greatest flow rate of depinning of microwell 1 is just 1.1 μL min⁻¹, though.

The alignment of the chips and the rate of liquid flow through the channel both have an impact on the depinning and pinning state. We investigated the effects of various intake flow rates on depinning. The relationship between the flow rate at which it is capable of depinning and the relative positions of the microwell and microchannel are shown in Fig. 3d. In microwell 2, depinning may happen when the rate surpasses 2.1 μL min⁻¹. However, the maximum flow rate of depinning for microwell 1 is only 1.1 µL min⁻¹. The infiltration conditions within the microwell will change regardless of whether microwell 1 or microwell 2 is within a broad range of set flow rates, which decreases the reliance of the system on the flow rate, can simplify operations, and can speed up the process of biological detection.

To confirm that the microwell array chip is useful for Gram stain, we next investigated the diffusion ability of the two-phase liquid. We are all aware that different reagents and sample solutions at the microscale often flow laminarly with a low Reynolds coefficient, and the mixtures are frequently inadequate, which restricts the use of microfluidic technology. Our goal was to create a microfluidic system that could identify bacteria quickly using the Gram stain. Numerous substances, such as violet dyes, potassium iodide, and detergents, are used as Gram stain reagents. A crucial stage in the Gram stain procedure is washing. The cells will be treated with various staining agents at different stages of the staining procedure, and the unabsorbed agents must be removed by proper washing processes to guarantee that the final staining result is accurate. To produce the optimum Gram stain effect, it is specifically required to gently rinse with water after Gram stain to remove extra staining reagents and lingering detergent from the cell surface. To explore the diffusion replacement impact of the two-phase liquid, we first successively injected red/blue ink into the chip, as shown in Fig. 4a. The timing begins when the first droplet changes color, and within 3 seconds, all 5 droplets in the observation area shift from entirely red to completely blue, demonstrating that the microwell has undergone a complete solution diffusion replacement. This diffusion replacement takes around 9 seconds to complete on a 45-droplet array. Software called ImageJ was used to calculate the area of the blue/red portion of the five drops. The portion of the blue ink made up 28.9% of the red ink at time 0.05 s, and by the time 0.15 s, it made up 78.3%, an increase of 49.4% from the 0.05 s period. The portion of the blue ink to the red ink is 91.2% at 1 second, and even after the blue area reaches 100% at 3 s, it only grows by 8.8%. Without a change in pump speed, there is quite minimal change in the area ratio of the two inks in the image within 1-3 seconds, showing that post-diffusion is still occurring. The three-dimensional structure and the volume of the microwell, which is relatively large compared to the channel, are to blame for the extended diffusion substitution.

Then, as shown in Fig. 4b, the four reagent solutions included in the Gram kit were put through diffusion staining tests. The screenshots of crystal violet, iodine solution, 95% ethanol, and safranin solution flowing into the chip, respectively, are shown in Fig. 4b. The dyeing reagent solution entry microwells, the clear solution entry microwells, and the microwells following water washing are all shown in the screenshot as four rows of microwell arrays. See Video in the ESI† for further information on the dynamic staining and washing procedure.

Diffusion stain studies were then conducted using the four reagent solutions included in the Gram kit. Screenshots of crystal violet, iodine solution, 95% ethanol, and safranin solution, respectively, are presented in Fig. 4b, i-iv. Each screenshot shows four rows of microwell arrays that are empty, filled with the staining reagent solution, filled with a water solution, and finished washing. The results

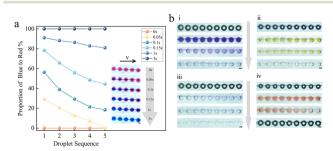


Fig. 4 Diffusion replacement in the microwell array chip. (a) Observation of diffusion in 5 ink droplets. (b) Gram stain, (i) crystal violet, (ii) iodine solution, (iii) 95% ethanol, and (iv) safranin solution. The scale bar is 100 μ m.

demonstrate that the Gram stain reagent solution can also effectively penetrate the microwell and diffuse, suggesting that the system has strong stability and a high efficiency of

diffusion and can be used to observe the identification of bacteria as Gram-positive or Gram-negative.

Single bacteria array and in situ detection

Then, we grew the extracted bacteria in situ, created a single bacterial array using the chip, and ultimately performed Gram stain to confirm the viability for biological applications. As is well known, the Gram stain procedure must penetrate the bacteria in the four reagent solutions for a few seconds to distinguish between positive and negative bacteria. There is a definite sequence to the reagent solutions that cannot be changed or skipped. As the chip for the bacterial array, we employed a PDMS chip with two kinds of microwells, and the channel leading to the reagent solution was 50 µm wide. In order for the bacteria to enter the microwells, droplets of the bacterial solution are first formed on the PDMS chip. We let the bacteria return to atmospheric pressure after turning off the pumping machinery. When the bacteria in the microwells are to be identified by Gram stain, the chip is put in an alcohol lamp to increase the temperature and let the solution evaporate, leaving the bacteria fixed in the microwells. A period of aqueous solution and air was necessary in the chip after each set of staining solutions colored the bacteria in order to isolate the effects of subsequent staining reagent solutions. Air must be passed between the two portions of solution in the capillary tube in order to prevent the fusion of each sequence of Gramstained and aqueous solutions. Several sets of dye

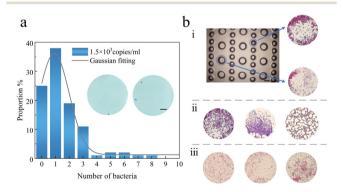


Fig. 5 (a) Statistical results of bacterial distribution in a single microwell. The blue histogram is the experimental result; the grey line represents the fitting of Gaussian distribution ($R^2 = 0.976$). The partial enlarged picture is the microscopy image of bacteria after staining; the scale bar is 5 μ m. (b) The morphology of mixed single cells cultured in situ in the microwell for 20 hours. (i) Localized top view of the microwell array (left) and mixed culture results of E. coli and S. aureus (right); (ii) only S. aureus; (iii) only E. coli. The entire image was captured using the $5\times$ microscope objective. At a $10\times$ magnification, the locally enlarged image of the microwell was taken. The scale bar is 200 μ m.

sequences were passed through for testing at the bacterial solution density of 1.5×10^3 copies per ml. The population distribution in Fig. 5a reveals that just one bacterium was present in around 38% of the microwells. The bacteria are more likely to be pre-buried towards the border of the microwell due to the lateral flow process in the microwell, which is also a major benefit for analyzing the development of adherent cells.

To undertake cellular phenotyping, cell-cell interaction research, and omics analysis, single-cell array in situ analysis is a crucial step. We combined S. aureus and E. coli solutions to create a mixed bacterial solution and then loaded it onto the microwell array based on studies on the production of single-cell arrays. After 20 hours, Gram stain allowed the bacteria in the microwells to be identified. The bacteria in the L-B medium solution reproduced freely independently. The bacteria in the microwells are stained as indicated in Fig. 5b, which depicts the outcome of in situ propagation. After propagation is finished, Fig. 5b(i) presents a partial top view of the microwell array and the mixture of E. coli and S. aureus that has been placed inside the microwells. Fig. 5b(ii) and (iii) show that only S. aureus and E. coli are present in the microwell, respectively. For further stained images of single bacteria and mixed bacteria, see Fig. S3 in the ESI.†

The build-up of metabolites and chemicals released by single cells or their offspring can occur more quickly when grown in arrays of microwells than when cultivated in bulk cultures. Numerous biological processes, including sample diagnostics, microbial genetic analysis, and antibacterial drug screening, depend on colony analysis at the single-cell The chips we developed enable cell-to-cell communication, the transport of some signalling molecules, such as peptides, that may contribute to cell growth, and can also be used to explain the pairing of taxa interactions or signal dependencies with other organisms.

Conclusions

In conclusion, we introduce, validate, and utilize microwell droplet array chips based on the lateral hydrodynamic wetting approach to in situ isolate and cultivate single-celled bacteria. The dynamics of bacterial populations, the stimulating impact at the gene level, and the clinical drug diagnostic analysis may all benefit greatly from the employment of SCA technology in the identification of harmful bacteria. The negative mould of the lotus leaf is where the concept of the sidewall gradually opening came from. This design is advantageous for controlling the wetting state of the liquid because it can overcome the energy barrier between the Cassie and Wenzel states. Due to the great diffusion efficiency of the liquid in the microwell array caused by pinning, it has several uses in amplification, sequencing, and metabolic processes. More notably, the chip can analyse at the single-cell level using just the lateral hydrodynamic wetting approach and can trap a variety of

bacteria in a specific region without the need for an external force field, which is a significant benefit for the omics research of cell co-culture.

Data availability

Lab on a Chip

The data that support the findings of this study are available from the corresponding authors upon reasonable request.

Author contributions

J. Wang and L. Du: conceptualisation, data curation, formal analysis, investigation, methodology, verification, visualisation, writing - original draft preparation. Y. Han: investigation, formal analysis, conceptualisation, methodology, supervision, visualisation. Jing: conceptualisation, data curation, investigation, methodology, project administration, supervision, verification, and writing - review and editing. D. Zhang: conceptualisation, methodology, funding acquisition, project administration, supervision.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

We thank Dr Enqi Wu and Mr Ming Yang for helping with the design parts of the experiment. This work was supported by the National Natural Science Foundation of China under Grant No. 62104148, No. 62274039 and No. 12272236.

References

- 1 G. Xing, W. Zhang, N. Li, Q. Pu and J.-M. Lin, *Chin. Chem. Lett.*, 2022, 33, 1743–1751.
- 2 Z. Xu, J. Wang, Z. Jia, Y.-X. Wu, N. Gan and S. Yu, *Analyst*, 2023, **148**, 1093–1101.
- 3 C. Mao, C. Xue, X. Wang, S. He, L. Wu and X. Yan, *Talanta*, 2020, 217, 121020.
- 4 L. A. Harris, S. Beik, P. M. M. Ozawa, L. Jimenez and A. M. Weaver, *Curr. Opin. Syst. Biol.*, 2019, 17, 24–34.
- 5 X. Xu, J. Wang, L. Wu, J. Guo, Y. Song, T. Tian, W. Wang, Z. Zhu and C. Yang, *Small*, 2020, 16, 1903905.
- 6 S. K. Govers, J. Mortier, A. Adam and A. Aertsen, *PLoS Biol.*, 2018, 16, 2003853.
- 7 C. Brasko, K. Smith, C. Molnar, N. Farago, L. Hegedus, A. Balind, T. Balassa, A. Szkalisity, F. Sukosd, K. Kocsis, B. Balint, L. Paavolainen, M.-Z. Enyedi, I. Nagy, L.-G. Puskas, L. Haracska, G. Tamas and P. Horvath, *Nat. Commun.*, 2018, 9, 226.
- 8 J. Shrestha, S. R. Bazaz, L. Ding, S. Vasilescu, S. Idrees, B. Söderström, P.-M. Hansbro, M. Ghadiri and M. E. Warkiani, *Lab Chip*, 2022, 23, 146–156.
- 9 C. H. Lin, Y. H. Hsiao, H. C. Chang, C. F. Yeh, C. K. He, E. M. Saln, C. Chen, L. M. Chiu and C. H. Hsu, *Lab Chip*, 2015, 15, 2928–2938.

- 10 J. Zhang, J. Xue, N. Luo, F. Chen, B. Chen and Y. Zhao, *Lab Chip*, 2023, 23, 1066–1079.
- 11 L. Cong, J. Wang, X. Li, Y. Tian, S. Xu, C. Liang, W. Xu, W. Wang and S. Xu, *Anal. Chem.*, 2022, **94**, 10375–10383.
- 12 F. Mi, C. Hu, Y. Wang, L. Wang, F. Peng, P. F. Geng and M. Guan, *Anal. Bioanal. Chem.*, 2022, **414**, 2883–2902.
- 13 X. Xu, L. Cai, S. Liang, Q. Zhang, S. Lin, M. Li, Q. Yang, C. Li, Z. Han and C. Yang, *Lab Chip*, 2023, 23, 1169–1191.
- 14 Y. Suo, W. Yin, W. Wu, W. Cao, Q. Zhu and Y. Mu, *Analyst*, 2022, 147, 3305–3314.
- 15 Y. Zhang, I. Naguro and A.-E. Herr, Angew. Chem., 2019, 131, 14067–14072.
- 16 X. L. Guo, Y. Wei, Q. Lou, Y. Zhu and Q. Fang, *Small*, 2022, **18**, 2107992.
- 17 Y. Huang, Z. Gao, C. Ma, Y. Sun, Y. Huang, C. Jia, J. Zhao and S. Feng, *Analyst*, 2023, **148**, 2758–2766.
- 18 J. Shemesh, T. B. Arye, J. Avesar, J. H. Kang, A. Fine, M. Super, A. Meller, D. E. Ingber and S. Levenberg, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, 111, 11293–11298.
- 19 M. Azizi, M. Zaferani, B. Dogan, S. Zhang, K. W. Simpson and A. Abbaspourrad, *Anal. Chem.*, 2018, **90**, 14137–14144.
- 20 Z. Gao, J. Yi, J. Zhao, H. Gu, H. Zhou and H. Xu, Sens. Actuators, B, 2020, 324, 128716.
- 21 L. Du, H. Liu and J. Zhou, Microsyst. Nanoeng., 2020, 6, 33.
- 22 E. Pope, C. Cartmell, B. Haltli, A. Ahmadi and R.-G. Kerr, Front. Microbiol., 2022, 13, 958660.
- 23 R. L. Lai and N. T. Huang, Microfluid. Nanofluid., 2019, 23, 121.
- 24 A. Tuteja, W. Choi, M. Ma, J. M. Mabry, S. A. Mazzella, G. C. Rutledge and R. E. Cohen, *Science*, 2007, 318, 1618–1622.
- 25 A. Karimi, S. Yazdi and A.-M. Ardekani, *Biomicrofluidics*, 2013, 7, 21501.
- 26 V. Narayanamurthy, S. Nagarajan, F. Samsuri and T. M. Sridhar, *Anal. Methods*, 2017, **9**, 3751–3772.
- 27 N. A. Patankar, Langmuir, 2010, 26, 8941–8945.
- 28 H. Kusumaatmaja, M.-L. Blow, A. Dupuis and J.-M. Yeomans, *Europhys. Lett.*, 2008, **81**, 36003.
- 29 R. Malinowski, I.-P. Parkin and G. Volpe, *Chem. Soc. Rev.*, 2020, **49**, 7879–7892.
- 30 Z. Gao, J. Yi, J. Zhao, H. Gu, H. Zhou and H. Xu, Sens. Actuators, B, 2020, 324, 128716.
- 31 D. C. Duffy, Lab Chip, 2023, 23, 818-847.
- 32 Y. Thomas, Philos. Trans. R. Soc. London, 1805, 95, 65-87.
- 33 R. N. Wenzel, Ind. Eng. Chem., 1936, 28, 988-994.
- 34 A. B. D. Cassie and S. Baxter, *Trans. Faraday Soc.*, 1944, 40, 546–551.
- 35 Y. Xue, S. Chu, P. Lv and H. Duan, *Langmuir*, 2012, **28**, 9440–9450.
- 36 W. Ren, Langmuir, 2014, 30, 2879-2885.
- 37 Q. S. Zheng, Y. Yu and Z. H. Zhao, *Langmuir*, 2005, 21, 12207–12212.
- 38 M. C. Salvadori, M. Cattani, M. R. S. Oliveira, F. S. Teixeira and I. G. Brown, *J. Appl. Phys.*, 2010, 108, 024908.
- 39 B. Emami, A. A. Hemeda and M. M. Amrei, *Phys. Fluids*, 2013, 25, 89–109.
- 40 B. Emami, H. V. Tafreshi, M. Gad-el-Hak and G. C. Tepper, *Appl. Phys. Lett.*, 2012, **100**, 13104.

- 41 G. Whyman and E. Bormashenko, *Langmuir*, 2011, 27, 8171–8176.
- 42 V. H. Lieu, T. A. House and D. T. Schwartz, *Anal. Chem.*, 2012, **84**, 1963–1968.
- 43 L. Du, Y. X. Li and X. L. Zhang, ACS Appl. Mater. Interfaces, 2023, 15, 17413–17420.
- 44 L. Du, A. Riaud and J. Zhou, *IEEE Trans. Nanotechnol.*, 2020, 19, 102–106.