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High-throughput microarray for antimicrobial susceptibility testing

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ABSTRACT

We describe the development of a novel, high-throughput, nano-scale microarray platform for antimicrobial susceptibility testing (AST). The platform allows to process 480 samples at 50 nL volume on a single chip, analyze by fluorescence read-out with an easy dunk-and-rinse step, and the ability to process multiple samples and chips simultaneously. We demonstrate the applicability of this chip for culturing community acquired methicillin resistant *Staphylococcus aureus* (CA-MRSA), and perform AST against clinical isolates of CA-MRSA. The chip platform holds promise for an impact in microbial biotechnology as an attractive high-throughput, lower sample volume and quicker alternative to conventional AST such as the traditional broth microdilution or the newer automated systems.

1. Introduction

Antimicrobial resistance remains a major threat to human health throughout the world, largely due to the indiscriminate use of antibiotics. The liberal use of antibiotics is driven by the need to quickly administer active antimicrobial therapy in the face of diagnostic uncertainty. AST is commonly performed using principles based on broth microdilution or disk diffusion assays, and takes 2-3 days, which includes the time for culturing the clinical isolates followed by antibiotic susceptibility testing by exposing the cultures to a panel of antibiotics over a range of concentrations [1]. Typically, clinical samples isolated from patients are cultured to increase the inoculum and isolate the pathogen; and then the isolates are exposed to a range of antibiotic concentrations either in 96-well plate suspension cultures or drug-filled cassettes (broth microdilution) or to a single concentration of the drugimpregnated disc placed on an agar plate containing the organism (disk diffusion). Susceptibility is then assessed by visual inspection either manually or by an automated instrument. Thus, it is conceivable that obviating relatively large culture volume or cell density may increase the throughput, and decrease the time required to obtain AST results. It is now well established that faster availability of AST data can enable the clinician to initiate or switch out of broad-spectrum treatment regimen to appropriate antimicrobial therapy sooner, and significantly

reduce health care costs and morbidity [2]. While there has been some progress in the recent years for rapid AST [3–5], the currently available tests are either limited to a few select organisms such as PCR-based methods [6,7], may not predict drug response reliably [8–10], limited in throughput or are cost-prohibitive such as MALDI-TOF MS or NMR [11] to most labs even in developed countries. To address some of the aforementioned issues, we have designed a novel microarray platform for high-throughput antimicrobial susceptibility testing (AST-Chip) and tested against select community-acquired methicillin-resistant *Staphylococcus aureus* (CA-MRSA) isolates.

2. Materials and methods

2.1. Culture conditions

Frozen stocks of *Staphylococcus aureus* strain UAMS 1 and clinical isolates were subcultured onto tryptic-soy-agar plates (TSA) (BD Difco, MD, USA), and propagated the in 10 ml of tryptic soy-broth (TSB) (BD Difco, MD, USA) in an orbital shaker at 37 °C. In order to capture cells in log-phase, 100 μ l from the overnight liquid culture were subcultured into 10 ml of TSB for 3 h.

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Fig. 1. High-throughput AST chip. (A) A schematic of high-throughput antimicrobial susceptibility testing using AST-Chip. Antimicrobial susceptibility testing using the chip can be conducted in 12 h; Microarray micrograph of *S. aureus* cultures formed on hydrogel spots of AST-Chip after 12–18 h. The spots are interspaced at 1 mm with a spot diameter of 0.5 mm; The nano-biofilms analyzed using FUN-1 is demonstrated in green color; (B) BacLight live/dead assay shows metabolically active cells in yellow-orange color, present as colonies. The scale bar is 100 μm; (C) SEM micrograph of *S. aureus* nano-biofilm colonies. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

2.2. Preparation of AST-Chip

The biofilm chip was prepared by modifying glass slides, and the growth conditions were optimized to obtain maximum biofilm yield of *S. aureus* using factorial design as described in [12,13]. The optimized media is a concoction of 2X Yeast Peptone Dextrose (YPD), 3X Brain Heart Infusion (BHI) along with 10% human serum. Antimicrobial stock solutions of doxycycline, vancomycin hydrochloride (Sigma, MO, USA), and clindamycin hydrochloride (RPI Corp., IL, USA), were diluted in phosphate buffered saline to a maximum concentration of 100 μ g/ml. Any subsequent dilutions needed for the antimicrobial susceptibility assays were made in PBS, mixed in a suspension of 0.5% alginate, and optimized media, and 50 nL of this mixture was spotted using a noncontact microarray spotter (Omnigrid Micro, Digilab Inc., Holliston, MA) on modified glass slides. An array of 40 rows and 12 columns was printed at room temperature with relative humidity of 100%. The AST-chip was stored at 4 °C for up to one week before use.

2.3. S. aureus nano-biofilms and viability assay

S. aureus cells were adjusted to a density of 1×10^7 cells/ml in modified growth medium, and printed on top of the drug spots. After

printing, the slides were then placed in a humidified hybridization cassette (Arrayit, Sunnyvale, CA, USA) to prevent evaporation of spots and incubated at 37 °C. All microarrayer functions such as sample loading, priming, printing, and spatial distribution of the array were controlled by AxSys programming (Digilab, MA, USA). The viability of nano-biofilms on the microarray was determined by staining with FUN-1 fluorescent dye as described in [13,14]. The fluorescence from the chips were measured using a microarray scanner (GenePix Personal 4100A; Axon Instruments, Union City, CA), and the images were analyzed with GenePix Pro V7 (Axon instruments, Union City, CA) to determine the antimicrobial susceptibility by normalizing the intensity between 0% and 100% for dead and live cells, respectively. The susceptibility values were obtained as average of 20-30 spots for each treatment from a single chip, and this experiment was performed in duplicate on two different days. The IC50 values were calculated using GraphPad Prism [15].

2.4. Microscopy

Fluorescence microscopy of *S. aureus* biofilms was performed after staining the slides with BacLight (Life Technologies) following manufacturer's protocol, and visualized at 40X (DMI6000, Leica). Scanning

electron microscopy was performed after fixing the slides and visualized using FEI Quanta 200 [12].

2.5. Antimicrobial susceptibility testing using Vitek-2 assays

Antimicrobial susceptibility testing was performed on a Vitek 2 compact instrument using AST-GP75 cards following the manufacturer's instructions. Briefly 3.0 ml of 0.45% saline was dispensed into two 12×75 mm tubes. Three to four well-isolated Staphylococcus aureus colonies on 5% sheep blood agar were selected with a rayon swab and transferred the first tube of saline. The suspension was homogenized by vigorously twirling the swab and vortexing. The inoculum density was adjusted to a 0.5 to 0.63 McFarland standard using a DensiCHEK instrument. The initial suspension was subsequently diluted by transferring 280 microliters into the second tube of saline. The Vitek AST-GP75 card and the second saline inoculum tube were then placed into a cassette and loaded into the instrument for filling and incubation. Accession numbers and organism identification data (S. aureus) were entered into the instrument database. A purity check plate was streaked from the second saline tube to assure a single morphotype was tested. Quality control as recommended by the manufacturer with appropriate ATCC strains was performed each day of testing. Results were printed and reviewed.

3. Results and discussion

The AST-Chip is a chemically modified microscope glass slide with nano-scale islands of hydrogels, consisting of antimicrobial agents in modified-growth medium. The clinical isolates were printed on the islands, incubated with drugs, and the cell density in each spot was measured by a fluorescence assay to obtain AST Fig. 1. The key steps in chip fabrication are described in detail elsewhere [13]. Briefly, the fabrication consists of three steps: (i) modification of the surface of glass slides with poly(styrene-co-maleic anhydride) (PSMA); (ii) printing 50 nl mixture of poly-L-lysine and barium chloride creating the tie-layer using a robotic microarrayer (Microsys, Digilab); and (iii) printing a 50 nl mixture of 0.5% alginate solution, optimized-growth medium and antimicrobial agents at various concentrations on top of the tie-layer. This process produces hydrophilic, drug-loaded islands on a hydrophobic background of the modified glass slide. In its final format, the AST-Chip contains of 480 spots of hydrogels of 50 nL loaded with different drugs at various concentrations Fig. 1A.

The growth medium in the mixture was optimized by factorial design to promote the growth of *S. aureus* [13]. We printed 50 nL spots of wild type (WT) *Staphylococcus aureus* (UAMS-1) in 0.5% alginate at 2500 cells/spot on top of alginate spots containing only optimized media Fig. 1B. We observed cell attachment to gel-spots in 30–45 min, following that the cells proliferated (1–8 h) and matured to a biofilm with the production of exopolymeric matrix material (8–16 h) [13]. After 12 h, the cell density at different spots was quantified using FUN-1, a vital cytoplasmic fluorescent dye compatible for analysis with a standard microarray scanner. A closer view of *S. aureus* biofilms is Fig. 2. Antimicrobial susceptibility profile of *S. aureus*. AST of methicillin against wild-type (UAMS-1WT) and community-acquired methicillin-resistant *Staphylococcus aureus* (CA-MRSA). (A) Microarray scanner image of AST-Chip with *S. aureus* cultures exposed to different concentrations of methicillin [MET]. The viability is indicated by the fluorescence intensity of the spots. (B) Scanning electron micrographs of S. aureus biofilms that are untreated (top) or treated with 10 μ g/ml MET (bottom); (C) Dose-response profile of *S. aureus* cultures obtained using AST-Chip.

shown by fluorescence intensity distribution from FUN 1, which confirms the uniformity of cell viability among different spots (Fig. 1A). We also visualized the biofilm colonies closer by viability dye BacLight (Fig. 1B), and by scanning electron microscopy (SEM), both of which show the formation of *S. aureus* biofilm colonies (Fig. 1C).

Having benchmarked the biofilm chip for the growth of S. aureus on top of hydrogel spots, we evaluated the antimicrobial-susceptibility of wild type (WT) Staphylococcus aureus (UAMS-1) and CA-MRSA (USA 300) strains against 0.1, 1.0, 10.0 and 100 µg/ml methicillin spotted on the AST-Chip. 50 nl of bacterial cell suspension in alginate was printed on top of the methicillin spots, and the chip was incubated for 12 h to allow for drug action, and stained with FUN1. The fluorescence intensities of control (no drug) and sodium hypochlorite-treated dead cells were arbitrarily set at 100% and 0% respectively. The inhibitory effects of the antibiotics were determined by the reduction in fluorescence intensity in comparison to the controls. As shown in Fig. 2A, the cell growth on the chip correlated with the drug concentration, and there was no cross-contamination across the spots despite their physical proximity and well-less culture. Consistent with the microarray scanner data, SEM micrographs confirmed that the untreated, control spots on the chip promoted S. aureus biofilm growth profusely, while methicillin completely inhibited biofilm growth (Fig. 2B). The susceptibility profile was deduced from the fluorescence intensities, and the data demonstrates that the AST-Chip accurately captures the decrease in susceptibility of CA-MRSA over WT strain (Fig. 2C).

Next, we performed AST for CA-MRSA clinical isolates collected from patients with skin and soft tissue infections (SSTIs) in primary care practices affiliated with the South Texas Ambulatory Research Network (STARNet) Practice-Based Research Network (PBRN). CA-MRSA accounts for a vast majority of skin abscesses, has high rates of clinical failure and disease recurrence, puts close contacts and health care workers at risk of infection, is difficult to eradicate, and can cause fatal bacteremia. To demonstrate the applicability of the chip platform, we compared the susceptibility of five clinical isolates against three commonly used antibiotics, clindamycin (CLI), doxycycline (DOX), and vancomycin (VAN), with the clinically used Vitek-2 assay. Table 1 shows the results typically obtained from the Vitek-2 assay. The 'R' and 'S', show the concentrations at which the isolates are resistance or susceptible, respectively, and thus we find that A32 is susceptible to CLI at 4 µg/ml, and other isolates are susceptible to all three drugs.

We tested the isolates, on the AST-chip, at their intermediate range of susceptibility of concentrations obtained from the Vitek-2 assay. The chip, consisting of 4 µg/ml of clindamycin (CLI), 4 µg/ml vancomycin (VAN), and 8 µg/ml of Doxycycline (DOX) with modified-media and alginate hydrogel was prepared, and the clinical isolates were tested. Table 1 shows the inhibition of bacterial growth in the presence of antibiotics using the biofilm chip, and it can be seen that the results are consistent with the Vitek-2 assay. The findings on the AST-Chip correlated well with the interpretive criteria of susceptible (S) and resistant (R), as determined by conventional AST from the Vitek 2 system. For instance, using Vitek 2, the isolate, A32, was classified as resistant to CLIN (MIC \geq 4 µg/ml) and susceptible to VAN (MIC < 0.5 µg/ml). The

Table 1

Effect of clindamycin, doxycycline and vancomycin in inhibiting the growth of *Staphylococcus aureus* isolates on Vitek 2 AST-GP75 cards and AST-Chip. The letters "R" and "S" refer to the respective resistance and susceptibility to drugs at the specified concentration.

Isolate	Age/Sex	Severity (Abcess)	Vitek			AST- Chip Inhibition (%)		
			CLI	DOX	VAN	CLI	DOX	VAN
A32	40/M	2	<u>> 4 R</u>	≤	<u><</u> 0.5 S	13.2	31.23	50.0
B2	52/F	Moderate-complicated (Yes)	\leq	<u>< 0.5 S</u>	<u>< 0.5 S</u>	64.8	30.1	51.1
I27	37/M	Moderate-complicated (No)	≤	<u>< 0.5 S</u>	<u>< 0.5 S</u>	27.8	36.6	50.7
K2	53/F	Moderate-complicated (Yes)	\leq	<u>< 0.5 S</u>	<u>< 0.5 S</u>	46.5	37.7	52.5
M2	38/M	Mild-Uncomplicated (Yes)	≤	<u><</u> 0.5 S	1 S	41.9	42.2	33.0

AST-Chip similarly demonstrated low percent inhibition (13%) to CLIN and high percent inhibition (50%) to VAN. In addition, the AST-Chip sensitively detected significant variation among the various clinical isolates, thus providing insights into the relative effectiveness of the drugs, which may not be available through conventional techniques. These results demonstrate concordance between conventional AST methods and the novel AST-Chip although future research will be needed to determine cut-off ('breakpoints') for this device. For instance, if we use an operational cut-off of < 20% inhibition as 'resistant' and > 20% as 'susceptible' strains for the AST-chip, as IC₈₀ is often used to interpret near complete inhibition, we observe that the A32 strain is can be classified as resistant to CLI [16].

In summary, we have successfully developed a rapid, robust, high density AST platform for CA-MRSA. We expect that the reduction in volume and an increase in throughput can be advantageous from a practical standpoint in clinical microbiology labs because: (i) each AST-Chip consisting of 1200 assays require a total of only a few microliters, and hence the assay can be directly performed on the clinical samples without a need for the initial expansion of culture density with a potential savings of 24 h; (ii) a 1000-fold scale down in volume may conceivably decrease the time required to obtain susceptibility data to less than 12 h [17]. In its current format, a single AST-Chip replaces up to twelve 96-well plates and several agar plates, and has sensitivity comparable to the current standard assays used in the microbiology labs but requiring much smaller sample volume and potentially shorter assay duration. The technology is flexible enough to be adapted to other bacterial and fungal organisms, integration into other novel assay platforms such as paper-based microfluidics, and along with other recent advances, will herald a new era in phenotypic susceptibility testing [18].

Conflict of interest

The authors have no conflict of interest to declare.

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