Direct Drug Cocktail Analyses Using Microscale Vortex-Assisted Electroporation

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Supporting Information

ABSTRACT: Combination therapy has become one of the leading approaches for treating complex diseases because it coadministers clinically proven drugs to concurrently target multiple signaling pathways of diseased cells. Identification of synergic drug combinations at their respective effective doses without unwanted accumulative side effects is the key to success for such therapy. In this work, we demonstrate the feasibility of the vortex-assisted microfluidic electroporation system for direct drug cocktail analyses where drug substances were individually delivered into cytosols in a sequential and dosage-controlled manner. Through quantitative analyses, the synergic combinational dosage ratios of the chemotherapeutic drug and the anticancer flavonoid were identified. When integrated with high-throughput label-free rare cell purification techniques, the presented system has the potential for development of personalized medicines as the system would be capable of comprehensively assessing drug combinations directly on patients’ cellular samples.

Combination drug therapy has been proposed as a treatment strategy for complex diseases such as cancer,¹ HIV,² cardiovascular disease,³ and type 2 diabetes.⁴ For such therapy, two or more already-licensed drugs are simultaneously administered to maximize therapeutic drug efficacy by targeting multiple signaling pathways while minimizing overlapping toxicity and inhibiting resistance-developing mechanisms.⁵−⁷ Due to its therapeutic benefits, extensive efforts have focused on the discovery of new drug combinations that work synergistically. For newly discovered drug combinations to rapidly progress to the clinic, systematic unbiased drug-screening strategies should complement existing hypothesis-driven approaches.⁸ However, identification of respective doses for individual agents that induce synergistic therapeutic outcomes is an essential but challenging task. Consequently, these drug combination screenings are very laborious; thus, the use of high-throughput liquid dispensing systems is inevitable for processing time reduction and precision improvement.⁹ However, not only do such systems have high fixed and operating costs,¹⁰ but their assay performances also are susceptible to noises and variations inherent in cell-based microplate assays.¹¹ Furthermore, a real dose–response curve and the cytotoxicity of drug combinations cannot be revealed if administered drugs have widely diverse physicochemical properties, yielding interdrug inconsistency in transport efficiency across the cell membrane.

Transient and reversible cellular membrane permeabilization utilizing electric pulses, namely, electroporation, can be an appealing means for drug-screening applications because it permits direct injections of newly developed cytotoxic molecules, which are inherently membrane-impermeable but can be subsequently modified to facilitate intracellular transport.¹²−¹⁴ In particular, it has been used to enhance chemotherapeutic drug efficacy for cutaneous cancers.¹⁵−¹⁷ Traditional electroporation drug-screening platforms have shown great promise in drug discovery; however, they have limited ability to perform cytotoxicity assays for combination therapies because their stochastic molecular delivery process precludes the determination of synergic drug dosage combinations. To overcome shortcomings of the bulk electroporation, many electroporation systems using microfluidic or nanomaterial platforms have successfully demonstrated reduction in operational costs and reagent consumptions with enhancement in molecular delivery efficiency and viability.¹⁸−²⁶ Still, very few systems allow for multimolecule delivery with controlled

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dosages besides nontrivial off-chip collection of processed biological samples for downstream analyses. Combining the advantages that microfluidics and electroporation offer, we demonstrated direct drug cocktail analysis using the vortex-assisted electroporation platform. The system enables sequential deliveries of precisely controlled amounts of multiple molecules into preselected cells, which can be subsequently released for downstream analyses. As a proof of concept, we examined the dosage-dependent cytotoxicity of chemotherapeutic drugs and an anticancerous flavonoid when those compounds were used as a single agent or in combination. The dose ratios of drug−flavonoid combinations that induce synergic and antagonistic effects were identified by systematically varying individual drug concentrations in the cytosol of cancer cells. The proposed platform has a great potential for facilitating comprehensive assessments of drug combinations directly on patients’ samples upon successful integration with the label-free cell purification system.

**MATERIALS AND METHODS**

**Cell Preparation.** Metastatic breast cancer cells, MDA-MB-231 (HTB-26, ATCC), were plated at a concentration of 1 × 10^5 cells/mL in a volume of 10 mL per T75 tissue culture flask (CELLSTAR, Greiner Bio-One, United States) in Leibovitz’s L-15 medium (Cellgro, Mediatech, Inc., United States) supplemented with 10% (v/v) fetal bovine serum (FBS; Gibco, Life Technologies, United States) and 1% penicillin–streptomycin (Sigma-Aldrich Co., United States). These cells were incubated in a humidified incubator at 37 °C with a 0% CO₂ environment. The cells were harvested for experiments 2 days after the seeding by treating them with 0.25% trypsin−EDTA for 2 min. The cells were pelleted by centrifugation for 5 min at 200g and resuspended in the growth medium to have a final concentration of 5 × 10^5 cells/mL.

**Device Design and Fabrication.** We have parallelized the previously reported microvortex-assisted electroporation platform to enhance the throughput without sacrificing any of its merits. The parallelized microfluidic electroporator consists of an inlet with multiple solution injection ports and coarse filters, two parallel inertial focusing channels (L = 7 mm, W = 40 μm, and H = 70 μm), and an outlet where two straight inertial focusing channels merge (Figure 1a). Individual straight channels consist of five electroporation chambers in series, and the electroporation chambers are placed 800 μm apart (W = 400 μm and L = 840 μm). Individual electroporation chambers have two holes for aluminum electrodes, and two transversally adjacent chambers share the hole for negative electrodes. 2-Dimensional projection of the device was designed using AutoCAD (Autodesk, Inc., United States), and the CAD file with micropatterns was converted to a GDSII file.
Electroporation of Target Cells for Drug Delivery. The system utilized the pneumatic flow control unit and the electrical equipment that we previously developed. Their detailed specifications28 and visualized operational protocols30 can be found elsewhere. In brief, the flow control unit independently and promptly pressurizes individual solution vials for rapid solution exchanges through the microfluidic system, while the electrical equipment generates high-voltage short pulses across the electroporation chambers on demand. Prior to electroporation experiments, a 15-pin aluminum electrode array60 and an outlet PEEK (poly(ether ether ketone)) tube were inserted into the microfluidic device, and the electrical equipment was connected to the electrode array. Solution vials, individually containing cells, drugs, and a growth medium for flushing, were mounted into the pneumatic flow control unit, and PEEK tubes from each vial were inserted into designated inlet ports in the microfluidic electroporator. The system was flushed with the growth medium at the operating pressure of 40 psi (equivalent to a flow rate of 400 μL/min) for 90 s prior to the target cell injection step to prime the flow speed required for cell trapping.34 Once the desired size and number distribution of trapped cells were attained in each cell-trapping vortex, nontrapped cells were removed from the device by switching the active solution port from the cell solution to the flush solution. Five short pulses with magnitude $V = 100$ V and pulse width $Δt = 30$ ms were applied promptly after the first drug solution was injected into the device. The resulting electric field strength across the electroporation chamber was $E = V/L_s = 0.7$ kV/cm. Here, $L_s = 1.5$ mm is the distance between the positive and negative electrodes that are in contact with the flowing solution. The magnitude, $V$, the width, $Δt$, and the frequency of applied electrical pulses were monitored in real time using an oscilloscope (Agilent, United States). Drug delivery doses were varied by exposing trapped cells to a single drug for incubation durations of 30, 60, and 120 s in continuous flow. For dual-drug delivery tests, trapped cells were sequentially incubated in individual drugs with various dosage combinations by systematically changing the treatment duration ratios of the two drugs (e.g., 30 s:30 s, 30 s:60 s, or 30 s:120 s). Upon completion of the drug delivery process, the treated cells were suspended in the growth medium and released from the device for downstream analysis by lowering the operating pressure to 30 psi. The collected cells were seeded in 96-well plates and cultured for 24-48 h prior to the cytotoxicity assays. The durations for which cells were cultured prior to the cytotoxicity assays were determined on the basis of conditions previously reported for the tested drugs (bleomycin,16 gemcitabine,32 topotecan and quercetin33).}

Real-Time Molecular Delivery Visualization. The nucleic acid fluorescent dyes propidium iodide (PI) and YOYO-1 (Life Technologies, United States) were used to visualize intracellular delivery of molecules in real time, allowing prompt determination of optimum electrical parameters and electroporation efficiencies. In addition, 1 μM bleomycin (BLM), which is fluorescently labeled with Alexa-Fluor 488 (Life Technologies), was used to identify electrical parameters required for successful drug delivery and dose control (see Figures S1 and S2, Supporting Information). For BLM–Alexa conjugation preparation, 25 μL of Alexa-Fluor 488 (8 mM) dissolved in DMSO was added to 100 μL of bleomycin (883 μM) dissolved in 0.9% saline to have a molar ratio of Alexa to BLM of approximately 2:1. Then sterile water was added to make 1 mL of total reaction volume, and the mixtures were stirred at room temperature for 19 h. Unreacted residual dye molecules were removed by performing high-performance liquid chromatography (HPLC) on a system equipped with an automated faction collector (Agilent 1100 series). The mixture was loaded onto the XTerra MS C4 column (Waters, United States) using a gradient of HPLC-grade water and acetonitrile as solvents A and B (2–10% B in 10 min and 10–95% B in 5 min), respectively, with ultraviolet (UV) detection at 292 and 495 nm (see Figure S2a). The purified conjugate was dried in a SpeedVac (Thermo Fisher Scientific) at 40 °C overnight, and the dried sample was dissolved in 1 mL of sterile Dulbecco’s phosphate-buffered saline (DPBS). The conjugate concentration was determined by UV–vis spectrophotometry (PerkinElmer, United States) on the basis of the BLM absorption spectrum. The molar extinction coefficient of BLM was taken as $ε = 1.5 × 10^4$ M$^{-1}$ cm$^{-1}$ at 292 nm,34 and the absorption signal of Alexa 488 did not interfere with drug concentration measurements.

Drug Preparation. Dose responses of metastatic breast cancer cells were tested using chemotherapeutic drugs, bleomycin, gemcitabine, and topotecan, and an anticancerous flavonoid, quercetin, to illustrate the drug-screening capabilities of the proposed platform (Sigma-Aldrich Co., United States). Conventional drug-screening assays utilizing well plates were conducted in parallel to evaluate the performance of the proposed microfluidic system. For the microfluidic electroporation drug-screening assays, the delivery dosages were varied by changing the incubation duration (e.g., 30, 60, and 120 s) at a flow rate of 400 μL/min where the concentrations of gemicitabine, bleomycin, topotecan, and quercetin in the injection vials were fixed at 210 μM, 57 nM, 240 nM, and 66 nM, respectively. For control experiments using the conventional method, the concentrations of the drugs were determined such that cells would be exposed in a well plate for 24 h to the drug solute quantity that is identical to that flown through the microfluidic device during the incubation period after the electroporation. That is, the viability of cells incubated in 210, 420, and 840 μM gemcitabine using the conventional assay was compared to that of cells treated with the drug for 30, 60, and 120 s, respectively, in the microfluidic device. Similarly, cells were incubated in bleomycin (57, 110, and 200 nM), topotecan (310, 620, and 1240 nM), and quercetin (83, 170, and 330 nM) for the control experiments.

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Cytotoxicity Determination. The CellTiter-Glo luminescent cell viability assay (Promega, United States) was used to determine the cytotoxicity of the tested drugs. First, the correlation between the luminescent intensity and the number of viable cells for the tested cell line was established from luminescence measurements from 96-well plates containing a known quantity of viable cells (LuMate, model 4400) (see Figure S3 (Supporting Information) for the calibration curve). For cytotoxicity studies on cells treated with the drugs, all cells present in the well, regardless of their viability, were manually counted prior to addition of the CellTiter-Glo reagent to the well. The number of viable cells was determined by identifying the corresponding number of viable cells represented by the measured luminescent intensity from the calibration curve. Viability was determined by taking the ratio between the number of viable cells and the total number of treated cells present in the well. The results from the combination treatments were analyzed to determine whether the combination was synergistic, additive, or antagonistic by calculating the combination index (CI) according to the Chou–Talalay method using the Compusyn software (Combosyn Inc., United States).

RESULTS AND DISCUSSION

In this work, a parallelized microfluidic electroporation platform was utilized for direct drug cocktail screenings. This platform renders a 10-fold enhancement in throughput in addition to all the merits that the previous single-chamber system provided. The system’s unique merits remaining uncompromised through parallelization include (i) superior viability of processed cells, (ii) on-demand injection of single substances, eliminating unforeseen adverse effects associated with drug–drug interaction prior to delivery, and (iii) sequential multimolecule delivery with highly efficient, precise,
and independent dosage controls. The system takes advantage of the microvortex-assisted cell-trapping mechanism to preselect subpopulations of cells with a uniform size distribution, thereby allowing less variations in electroporation efficiency and enhancement in viability per given electric field strength. Previously, we have shown that molecules ranging from small nucleic acid dyes to large naked plasmids could be introduced sequentially into preselected populations of cells as they orbit in the vortices. Parallelization of this platform did not interfere with the sequential molecular delivery process as we confirmed through sequential delivery of the nucleic acid dyes PI and YOYO-1 into cells trapped in the 10 electroporation chambers (Figure 1b). Moreover, the molecular amount transferred into the cytosol can be modulated by varying either the electrical field strengths or molecular injection times, and intracellular concentrations of delivered substances were probed using fluorescent drug surrogates (Figures S1 and S2, Supporting Information). In a similar manner, multiple drug compounds with varied concentration ratios were delivered into cells orbiting in the electroporation chambers.

To evaluate the system’s applicability toward drug-screening assays, we first assessed the dose–response characteristics of gemcitabine and bleomycin (Figure 2a,b), which have been studied extensively for electrochemotherapy for cutaneous metastasis. Consequently, these drugs are ideal candidates for comparative studies of the microfluidic and the conventional well-plate assay because their cytotoxic effects and optimum dosages have been well-documented. Cells treated with these drugs as single entities in well plates exhibited less significant dosing effects compared to those treated by the microfluidic device (Figure 2; Figure S2, Supporting Information). This could be attributed to the nonpermeant nature of the drugs. Correspondingly, higher IC_{50} values for both drugs were obtained from the conventional assays (Table S1, Supporting Information), suggesting that IC_{50} is strongly influenced by in-cell drug concentrations rather than accessible drug amounts in the incubation solution. This implies that the developed system would facilitate understanding of unidentified relationships between intracellular drug concentrations and pharmacological consequences, which presumably was ascribed to conventional assays’ stochastic and diffusive transmembrane transport mechanisms.

We further investigated the dose effects of a chemotherapeutic drug, topotecan, and a flavonoid with anticancer properties, quercetin, as single agents and in combination. Previously, the combination of these two substances was reported to have synergistic effects because quercetin selectively enhances cytotoxicity of chemotherapeutic drugs only in cancerous cells, suppressing off-target toxicity. As a single agent, topotecan dose-dependently reduced survival of cells processed using either drug-screening platform (Figure 2c). The fact that cells treated using the microfluidic platform responded more sensitively to the drug suggests that the proposed microfluidic assay can be used to identify the effectiveness of drugs with subtle variation in the concentrations. Quercetin alone, on the other hand, exhibited a negligible cytotoxic effect on cells processed with either system (Figure 2d) even at a high concentration, which is in good agreement with previous studies.

No apparent adverse effect on cells treated with atoxic quercetin using the current system further confirms that the dose responses of the tested chemotherapeutic drugs were not altered by the electroporation and solely represent the actual cytotoxic effects of the delivered drugs.

For sequential dual-drug screenings, cells were first exposed to topotecan and quercetin for identical durations (i.e., 30 s of incubation in solutions with 240 nM topotecan and 66 nM quercetin; see Figure 3a). More than 3-fold reductions in IC_{50} values obtained from both assays suggest that the addition of quercetin augments the cytotoxic effects of topotecan (Table S1, Supporting Information). These trends were evident at all combination ratios where a fixed concentration of topotecan was used (T:Q = 1:1, 1:2, and 1:4). The CI was computed for...
all tested combinations to quantitatively evaluate the drug combinatorial effects (Table S2, Supporting Information). In such an analysis, CI < 1 implies synergy, CI = 1 represents an additive effect, and CI > 1 corresponds to antagonism.\textsuperscript{35} Cytotoxicity assays of a fixed topotecan concentration assessed using both platforms illustrated that the drug combinations worked in synergy (CI < 1) at all tested ratios (Figure 3a). This trend of increased drug cytotoxicity continues as the dosage of topotecan increases while that of quercetin remains unchanged (T:Q = 2:1). The drug ratios T:Q = 1:1 and 2:1 were found to induce synergistic effects for both assays. Interestingly, the antagonistic effect (CI = 1.11) was detected only with the microfluidic assay at T:Q = 4:1, whereas the results from the conventional assay implied the synergistic effect of the combination (CI = 0.29). Such disparity could have originated from the fact that apparent drug amounts delivered into cytosols by the microfluidic system were much greater than those by the well-plate assay. Presumably, the maximal response of the combination, $E_{\text{max}}$, beyond which cytotoxicity is independent of drug concentrations\textsuperscript{39} has been attained for the microfluidic assay. It is also possible that the proposed system can identify elusive antagonistic drug combinations which are not detectable otherwise. Subsequent investigations are still required to reinforce this claim.

Owing to inherently small footprints of the microfluidic systems, the presented microfluidic electroporator can conduct cost-effective, single-cell-level multidrug screenings. As shown in the results, cells assayed using the presented system exhibited elevated cytotoxicity compared to those treated by the conventional well-plate method when identical amounts of drugs were used. Although the current system can screen a considerably lower number of cells than conventional methods, the channel geometry is simple and amendable to automation. Thus, the system can repeatedly process a greater number of cells to conduct an unbiased drug-screening approach to reveal serendipitous combinations of already-approved compounds. Moreover, the developed platform is anticipated to be used as a novel screening method to understand consequences of drug-target interactions for newly identified cytotoxic molecules whose transmembrane transport mechanisms are yet to be determined. Finally, the system can be readily integrated with the high-throughput vortex-assisted circulating tumor cell purification system whose clinical potentials have been demonstrated.\textsuperscript{27} Successful integration will allow performance of sophisticated drug-screening analyses of patient-derived cellular models to identify interpatient variability conferring factors in drug responses.

**CONCLUSION**

In this work, a microscale vortex-assisted electroporation platform was implemented for direct analysis of drug cocktails. Contrary to the bulk electroporation systems, this platform does not adversely affect cell viability, suggesting that cellular responses observed from cytotoxicity assays solely represent the actual effects of the drug cocktail. Through quantitative single-drug and drug combination analyses, we were able to identify the combinational dosages of the chemotherapeutic drug and the flavonoid inducing synergetic and antagonistic effects. The system has great potential to enable drug screenings for development of personalized medicines when it is integrated with the on-chip cell purification system.\textsuperscript{27}
Electronic Supporting Information (ESI)

Direct Drug Cocktail Analyses Using Microscale Vortex Assisted Electroporation

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List of Supplementary Information

Figure S1. Delivery dose control of fluorescent molecules by varying electric field strengths using parallel aluminum electrodes

Figure S2. Purification and delivery dose control of fluorescently labeled bleomycin

Figure S3. Luminescence calibration for viability analysis

Table S1. IC\textsubscript{50} values for microfluidic and conventional assay

Table S2. Combination index (CI) values for the combination therapy studies
Figure S1. Amount of transferred PI molecules into MDA-MB-231 cells could be controlled by varying the electric field strength with the fixed solution-injection and incubation time (40s) when aluminum electrodes were used. For drug screening assays, electrical field strength of 0.7 kV/cm (V = 100 V and Δt = 30 ms with 2 s intervals) was used to give a gentle electric field that does not have adverse effects on the cells. Scale bars are 100 µm.
Figure S2. (a) High performance liquid chromatography (HPLC) profiles used for the separation of the unbound Alexa Fluor 488 dye from the conjugated bleomycin. Successfully conjugated bleomycin exhibited a previously absent peak at 495 nm. The substance collected within the shaded elution time was used for drug delivery visualizations. (b) BLM-Alexa conjugate amounts delivered using the proposed microfluidic electroporator and the conventional well-plate assay were qualitatively compared by analyzing fluorescent intensities of processed cells. The percentage increase in fluorescent signals post drug delivery was computed using \[ \frac{\text{Signal}}{\text{Background}} \times 100\% \]. For the microfluidic assays, 5 pulses (\( V = 100\text{V} \) and \( \Delta t = 30 \) ms with 2 s intervals) were applied and cells were incubated with 1 \( \mu \text{M} \) conjugate for 30, 60, and 120 s, respectively, postelectroporation. Fluorescent signals from cells that were incubated in DPBS for 120 s postelectroporation were used as the control. For the conventional method, cells were incubated with 0, 1, 2, and 4 \( \mu \text{M} \) of BLM-Alexa conjugates in 96-well plates for 24 h under...
the culture condition. Fluorescent intensities of cells processed using the proposed technique linearly increased with incubation durations with the slope of regression line, \( S_e = 0.003 \), and the coefficient of determination, \( R^2 = 0.86 \). Cells assayed using the conventional method exhibited more gradual increase in intensity (\( S_c = 0.0014 \) and \( R^2 = 0.72 \)) than those tested by the microfluidic counterpart, suggesting that more efficient dosage controlled intracellular drug delivery accomplished by the proposed technique when the identical quantity of drug solutes was administered. Image contrast has been enhanced by applying the identical look-up table (LUT) to all images.

**Figure S3.** The number of viable MDA-MD-231 cells linearly correlates with luminescent intensity. Relatively larger standard deviation displayed at the last data point could be attributed by difficulties associated with manually counting a large population of cells.
**Table S1:** IC$_{50}$ values evaluated experimentally for both the microfluidic and conventional assay.

<table>
<thead>
<tr>
<th>Chemotherapeutic Drugs</th>
<th>Literature IC$_{50}$</th>
<th>Conventional IC$_{50}$</th>
<th>Proposed IC$_{50}$</th>
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</thead>
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<tr>
<td>Gemcitabine</td>
<td>340µM$^\dagger$</td>
<td>2000µM</td>
<td>302µM</td>
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<tr>
<td>Bleomycin</td>
<td>12µM$^\dagger$</td>
<td>20µM</td>
<td>0.12µM</td>
</tr>
<tr>
<td>Topotecan</td>
<td>1.1µM$^\dagger$</td>
<td>2µM</td>
<td>1.2µM</td>
</tr>
</tbody>
</table>

$^\dagger$ Previously reported IC$_{50}$ values for tested drugs on various cell types were included for comparison. IC$_{50}$ values for gemcitabine, bleomycin and topotecan have been reported for Ca-27 squamous carcinoma$^1$, MC38 colon cancer cell line$^2$, and MDA-MB-231$^3$, respectively.

**Table S2:** Combination index (CI) value for the combination therapy of topotecan (T) and quercetin (Q) at various ratios.

<table>
<thead>
<tr>
<th>Drug Combination Ratio [T:Q]</th>
<th>Conventional CI</th>
<th>Proposed CI$^*$</th>
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<tr>
<td>1:1</td>
<td>0.32</td>
<td>0.34</td>
</tr>
<tr>
<td>2:1</td>
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<td>0.32</td>
</tr>
<tr>
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<td>1:4</td>
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$^*$Note that the dose response value of quercetin at 120 s exposure was excluded from CI value determination because dose-independent and negligible cytotoxicity of quercetin resulted in illogical and negative coefficient of determination, R$^2$, for the dose response curve.

**References**