

1      **Electronic Detection of Apoptotic Cells on a Microchip**  
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14     **ABSTRACT**

15     Robust and rapid detection of apoptosis in cells is crucially needed for diagnostics, drug  
16     discovery, studying pathogenic mechanisms and tracking patient response to medical  
17     interventions and treatments. Traditionally, the methods employed to detect apoptosis rely on  
18     complex instrumentation like flow cytometers and fluorescence microscopes, which are both  
19     expensive and complex-to-operate except in centralized laboratories with trained labor. In this  
20     work, we introduce a microfluidic device that can screen cells in a suspension for apoptosis  
21     markers and report the assays results as electronic data. Specifically, our device identifies  
22     apoptotic cells by detecting externalized phosphatidylserine on a cell membrane – a well-  
23     established biomarker that is also targeted by fluorophore-based labeling in conventional assays.  
24     In our device, apoptotic cells are discriminated from others through biochemical capture  
25     followed by transduction of individual capture events into electrical signals via integrated  
26     electrical sensors. The developed technology was tested on simulated samples containing  
27     controlled amounts of cells with artificially-induced apoptosis and validated by benchmarking  
28     against conventional flow cytometry. Combining sample manipulation and electronic detection  
29     on a disposable microfluidic chip, our cell apoptosis assay is amenable to be implemented in a  
30     variety of settings and therefore has the potential to create new opportunities for cell-based  
31     diagnostics and therapeutics and contribute to healthcare outcomes on a large scale.

32     **Keywords:** apoptosis sensor, cell death detection, Annexin V assay, phosphatidylserine  
33     externalization, electronic cell viability assay, microchip-based apoptosis test.

35     **1. INTRODUCTION**  
36

37     Cell death is a fundamental, multi-step process that has profound effect on the function of  
38     multicellular organisms and occurs in nominal physiological states in addition to pathological  
39     conditions (Galluzzi et al., 2018; Kopeina & Zhivotovsky, 2022). Apoptosis is one of the most

1 widely recognized and studied type of cell death. The process is controlled and regulated by  
2 specific physiological signals or pathological factors (Taylor et al., 2008), allowing detailed  
3 study of its mechanics. Necrosis, in contrast, is an uncontrolled cell death typically characterized  
4 by cell swelling, followed by membrane rupture and release of intracellular contents (D'Arcy,  
5 2019).

6 A crucial event that occurs during apoptosis is the externalization of the phospholipid  
7 phosphatidylserine (PS), where PS is translocated from its normal inner leaflet placement on the  
8 cell membrane to the outer leaflet leaving it exposed to extracellular environment  
9 (Balasubramanian et al., 2007). This externalization process is an intrinsic component of tissue  
10 regeneration as it marks the cell with an “eat me” signal for neighboring cells and phagocytes  
11 whose function is the removal of apoptotic cells. Notably, PS externalization is one of the earlier  
12 events in apoptosis, preceding the failure of cell membrane integrity (Kerr et al., 1972). Given  
13 the point at which PS externalization occurs relative to the rest of the apoptotic process, detecting  
14 externalized PS with a high affinity biochemical probe like Annexin V makes such an assay  
15 crucial in studying apoptosis (Meers & Mealy, 1993).

16 When studying diverse cell populations that are undergoing apoptosis, itself an inherently  
17 stochastic process (Darzynkiewicz et al., 2004), it becomes crucial to leverage high-throughput  
18 platforms that can perform single cell interrogations. The gold standard for detecting apoptosis is  
19 through labeling cells under test with fluorophore-conjugated Annexin V and measuring  
20 fluorescence emission with a flow cytometer (Fischer et al., 2002; Vermes et al., n.d.; Worsley et  
21 al., 2022). However, this approach is limited in providing immediate interrogations to detect  
22 early apoptotic cells as the fluorescence staining procedures require a lead time before the  
23 sample is ready for analysis (Schutte et al., 1998). Moreover, fluorescence labeling has the  
24 potential to compromise protein activity due to the chemical cross-linking reactions present  
25 during the staining process (Koskinen et al., 2004). In addition, a flow cytometer is a complex  
26 instrument with a large footprint. Due to high maintenance costs and operational complexities,  
27 flow cytometers are often deployed in shared facilities and operated by skilled technicians,  
28 making it impractical to perform apoptosis assays at the point-of-care (PoC) or in mobile testing  
29 environments. Additionally, the extensive number of operational parameters that need to be  
30 manually selected, along with the necessary sample preparation, make the entire process and the  
31 interpretation of results dependent on the specific operator or laboratory conducting the analysis.

32 Several studies have leveraged recent technological advancements to develop apoptosis detection  
33 techniques, with the most common approaches utilizing fluorescence microscopy and image-  
34 based cytometry. One such study utilized a novel tetrahedral DNA probe to simultaneously  
35 image intracellular cytochrome c and telomerase to provide a comprehensive visualization of the  
36 apoptotic process (Dong et al., 2022). Another approach developed a microfluidic multicellular  
37 coculture array to integrate multiple cell-based assays, one of which was apoptosis detection,  
38 into a unified system (Chong et al., 2022). Alternatively, several electrochemical approaches  
39 have also emerged, including an Annexin V-modified biosensor that uses an impedance  
40 spectroscopy scheme to identify early apoptotic cells (Tong et al., 2009) and a differential pulse  
41 voltammetry (DPV) based measurement of caspase-3 activity that is used as an indicator of

1 apoptotic activity (Chen et al., 2015). All these sensing methods, however, suffer from low  
2 portability and/or high cost due to their instrumental and operational complexity.

3 In pursuit of assays that are more portable and affordable than conventional assays, significant  
4 advancements have been made using microfluidic microchip-based technologies. These solutions  
5 typically feature sensing elements integrated on-chip to allow compact sensing while  
6 maintaining direct contact with the cell sample (Arifuzzman et al., 2023; Civelekoglu et al.,  
7 2019; Civelekoglu, et al., 2022; Civelekoglu, et al., 2022; Civelekoglu, et al., 2022; Liu et al.,  
8 2019, 2020). These studies, however, adopt an immunoanalytical approach to target surface  
9 markers and do not directly target apoptotic cells.

10 Here, we introduce a microchip-based apoptosis assay that detects apoptosis by targeting  
11 externalized PS on cell membranes, eliminating the need for the user to pre-label the sample  
12 before performing the assay. This is a crucial advantage, since traditional testing practices such  
13 as flow cytometry require a multiple, labor-intensive sample preprocessing steps before running  
14 the assay. These steps typically include preparing various dilutions of the stock reagent, correctly  
15 mixing the diluted label solution with the sample, waiting for incubation, and performing  
16 centrifugation and resuspension. In contrast, our microchip-based apoptosis assay significantly  
17 simplifies this process by incorporating the sample preprocessing steps—such as labeling and  
18 mixing—directly within the microchip itself. This is made possible through our unique  
19 microchip design and manufacturing process, which chemically modifies specific regions within  
20 the microchip. These modifications allow the necessary chemical reactions to occur  
21 automatically when the sample is loaded, without any manual intervention. For the end user, this  
22 means that the sample can be directly loaded into the microchip without the need for dilution,  
23 mixing, incubation, or centrifugation. The entire process is automated within the microchip,  
24 resulting in a rapid, label-free apoptosis assay with a streamlined workflow. This not only  
25 reduces the time and effort required but also minimizes the potential for human error, making the  
26 assay more efficient and user-friendly. This capability makes our platform well-suited for  
27 integration with automated cell development processes, enabling efficient and flexible apoptosis  
28 assays as required. Additionally, the microchip-based assay allows it to be portable for  
29 deployment in point-of-care settings unlike a flow cytometer. This is particularly advantageous  
30 in remote or resource-limited environments where traditional laboratory facilities are  
31 inaccessible, enabling apoptosis assays to be performed efficiently and effectively outside of  
32 conventional laboratory settings. To develop an apoptosis assay platform with sample-to-answer  
33 capability, cells are internally labeled in flow on our microchip to transduce relatively weaker  
34 Annexin V-PS affinity into stronger avidin-biotin binding for reliable biochemical capture of  
35 apoptotic cells. A network of barcoded electrical sensors then quantifies the fraction of cells with  
36 externalized PS expression through capture statistics and presents the data as an electrical output  
37 signal. We applied our assay on studying human t-cell lymphocytes subjected to heat to induce  
38 apoptosis and validated it accuracy by benchmarking against the well-established flow  
39 cytometry-based Annexin V assay.

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1 **2. MATERIAL AND METHODS**

2 **2.1. Chemicals and materials**

3 We acquired the materials and chemicals used in this work from various sources. 3-  
4 aminopropyltriethoxysilane (APTES) was purchased from Gelest, Inc., while Neutravidin and  
5 BSA (Bovine Serum Albumin) were obtained from Thermo Scientific. Glutaraldehyde and  
6 trichloro(octyl)silane were sourced from Sigma-Aldrich, and 200 proof ethanol was obtained  
7 from Decon Labs, Inc. Phosphate-Buffered Saline (PBS) was purchased from Mediatech, and all  
8 chemicals used were of analytical grade. The experiment also utilized deionized (DI) water.  
9

10 The FITC Annexin V (Cat. No. 640906), Biotin Annexin V (Cat. No. 640939), and Annexin V  
11 Binding Buffer (Cat. No. 422201) were sourced from BioLegend (San Diego, CA). Cy5 biotin  
12 conjugate (Cat. No. 3100) was purchased from AAT Bioquest, Inc. (Pleasanton, CA). The Jurkat  
13 Clone E61 (ATCC® TIB152™) cell line was procured from the American Type Culture  
14 Collection (ATCC) (Manassas, VA). The Roswell Park Memorial Institute Medium (RPMI  
15 1640), fetal bovine serum (FBS), and 1% Penicillin/Streptomycin antibiotics were all acquired  
16 from Corning (Corning, NY).

17 For materials used during fabrication, 4-inch diameter silicon wafers were procured from  
18 UniversityWafer, Inc. SU-8 2000 series photoresist was obtained from MicroChem, while NR9-  
19 1500PY negative photoresist was acquired from Futurrex, Inc. Lastly, polydimethylsiloxane  
20 (PDMS) elastomer Sylgard 184 was purchased from Dow Corning.

21 **2.2. Microchip fabrication**

22 The microchip was fabricated using soft lithography and surface micromachining techniques.  
23 First, a 15  $\mu$ m thick layer of negative photoresist SU-8 (SU-8 2025, MicroChem) was spin-  
24 coated onto a silicon wafer. The design micropattern was transferred to the photoresist film by  
25 exposing it using a maskless aligner (MLA-1500, Heidelberg). The photoresist was then  
26 developed with a SU-8 developer (SU-8 Developer, MicroChem) to create the negative mold.  
27 The micropatterned wafer was then treated with trichloro(octyl)silane in a desiccator for 8 hours  
28 before a mixture of PDMS elastomer and its crosslinker (Sylgard 184 kit, Dow Corning) was  
29 prepared at a 10:1 weight ratio. This polymer mixture was poured onto the mold, degassed, and  
30 cured at 65 °C for 4 hours. The resulting PDMS was then peeled off and cut into individual  
31 chips.

32 To fabricate the electrical sensor network, a 1.5  $\mu$ m thick layer of negative photoresist (NR9-  
33 1500PY, Futurrex) was spin-coated onto a 2-inch by 3-inch glass microscope slide (6101,  
34 Premiere). Then, the microelectrode pattern was transferred to the resist layer using a maskless  
35 aligner (MLA-1500, Heidelberg). The photoresist layer was then developed using a developer  
36 (RD6 developer, Futurrex). To prepare the glass slides for the subsequent deposition process, a  
37 reactive ion etcher was used for descumming, ensuring optimal surface conditions. Using an  
38 electron beam evaporator, a 20 nm-thick chromium (Cr) film was deposited onto the glass slides,  
39 followed by the deposition of a 250 nm-thick (Au) film. This process ensured the formation of a  
40 reliable metal stack with strong adhesion to the glass surface. The sacrificial photoresist layer  
41

1 was removed by submerging the glass substrate in an acetone bath while employing mild  
2 sonication. Finally, the PDMS layer and the glass substrate were subjected to a one-minute  
3 oxygen plasma treatment to activate their surfaces. The activated PDMS layer and the treated  
4 glass substrate were aligned under a microscope and bonded together at a temperature of 65 °C  
5 to form the final structure.

6 **2.3. Measurement of micromixing efficiency for on-chip cell labeling**

7 Dye solutions with different colors (red and blue) were used in lieu of actual samples and  
8 reagents to evaluate different on-chip mixing strategies. Then, to quantify the efficiency of a  
9 specific microchip design, we calculated a mixing index (MI), which was derived based on the  
10 levels of primary color components within a 30 µm by 30 µm region of interest (ROI) in  
11 acquired images (São Pedro et al., 2023). Specifically, by measuring the presence of blue and red  
12 color content at distinct points throughout the mixer flow stream, we quantified the extent of  
13 mixing at various positions of interest. For each ROI, this measurement is done by first splitting  
14 the image into red, green and blue color channels. Omitting the green channel, the red and blue  
15 color channels of the ROI image are both converted into 8-bit grayscale, i.e., the intensity value  
16 of each pixel was converted into 8-bit levels—where 0 represents completely black and 255  
17 represents the highest color intensity. Splitting and normalizing the individual color channels this  
18 way allows the use of their intensity values in direct comparative analysis. Then, we determined  
19 the MI by computing the intensities of red and blue color channels within the ROI, at a given  
20 point on the mixer, relative to the intensities of red and blue color channels of the red and blue  
21 dyes before they enter the mixer. This normalization was deemed necessary as the stock red and  
22 blue dyes did not exhibit the same intensity values when converted to grayscale. Hence,  
23 assuming a value of 255 for both the red color channel intensity of the red dye and blue color  
24 channel intensity of the blue dye would lead to erroneous calculations. All in all, we calculated  
25 the MI as:

$$26 \quad MI = 100 \% - \left( \frac{|(R_{ROI} - B_{ROI})| - |(R_{unmix} - B_{unmix})|}{R_{unmix}} \times 100 \% \right)$$

27 , where  $R_{ROI}$  and  $B_{ROI}$  represents the amount of red and blue dye content measured in the ROI,  
28 respectively.  $R_{unmix}$  and  $B_{unmix}$  denote the red and blue channel intensities measured at the red and  
29 blue dye inputs of the mixer, which were used to normalize the measured red and blue channel  
30 intensities at the various points throughout the mixer.

31 **2.4. Chemical functionalization of the microchip for apoptotic cell capture**

32 We chemically functionalized inner surfaces of our fabricated microfluidic device to selectively  
33 capture the target cells out of a heterogenous mix. To utilize plasma-assisted surface activation  
34 for the subsequent chemical treatment, the process was initiated within the first 10 minutes after  
35 oxygen plasma-assisted PDMS-glass microchip bonding was performed. All chemical reagent  
36 infusions, including the washing step, were conducted at a flow rate of 500 µL/hr, with each  
37 infusion having a duration of 20 minutes. First, the device was wetted with ethanol followed by  
38 flushing the device with a solution of APTES in ethanol (3% v/v). Following an incubation of 1  
39 hour, the device was washed with ethanol and dried in a vacuum oven at 110 °C for 1 hour. The

1 device was then washed with deionized (DI) water, infused with a glutaraldehyde solution in DI  
2 water (3% v/v) and incubated at room temperature for 1 hour. We measured the cell speed in the  
3 microfluidic channel to ensure that the presence of glutaraldehyde within the microchip does not  
4 significantly affect it (Fig. S1 and Supplementary Video 1). Our findings indicate that the  
5 glutaraldehyde treatment does not impact the fluidity of the infused sample during operation  
6 when compared to an identical microchip without glutaraldehyde treatment. To continue the  
7 surface modification process after glutaraldehyde treatment, the device was flushed with DI  
8 water and then infused with a 1 mg/mL neutravidin solution in PBS. A total of 166.7  $\mu$ g of  
9 neutravidin was infused over a 20-minute period at a flow rate of 500  $\mu$ L/hr. Following this, the  
10 device was incubated for 4 hours at room temperature to facilitate glutaraldehyde-neutravidin  
11 binding and then flushed with PBS. To confirm consistent neutravidin immobilization, we  
12 conducted a characterization experiment. Three identical neutravidin-functionalized microchips  
13 were prepared alongside one control device without neutravidin. All four devices were incubated  
14 with Cy5-biotin for 1 hour, washed with PBS, and examined using fluorescence microscopy to  
15 measure fluorescence intensity. Cy5-biotin was selected because its presence after the labeling  
16 and washing steps indicates the presence of neutravidin, which would be absent if neutravidin  
17 had not been immobilized on the microfluidic walls and pillars. Fluorescence intensity was  
18 measured at three different locations on each device, and the average fluorescence intensity  
19 across the three neutravidin-functionalized devices was found to be similar (Fig. S2). This  
20 confirmed the consistent immobilization of neutravidin across different devices using our  
21 chemical immobilization process. Next, the device was flushed with PBS and incubated with a  
22 3% BSA blocking buffer for 1 hour for blocking the surface against non-specific cell capture.  
23 Finally, the device was thoroughly washed with PBS to remove unbound neutravidin, completing  
24 the functionalization process and producing a microchip ready for processing a sample.

## 25 **2.5. Preparation of human T lymphocyte samples**

26 Immortal T lymphocyte cells (Jurkat, Clone E6-1-TIB-152) were acquired from the American  
27 Type Culture Collection. After thawing, cells were cultured in RPMI media, supplemented with  
28 10% FBS and 1% Penicillin antibiotics to support optimal growth. The cells were cultured in  
29 media in a T25 flask within an incubator that was maintained at 37 °C and 5% CO<sub>2</sub>. Once the  
30 cells reach a confluence of 80%, they were collected and resuspended in 1X PBS before being  
31 placed in vial for use in experiments.

## 32 **2.6. Induction of controlled apoptosis on samples**

33 Jurkat cells were artificially induced into apoptosis using heat shock. Heat-induced apoptosis is a  
34 commonly observed phenomenon in heat-related illnesses, where irreversible cellular damage  
35 occurs at elevated temperatures ranging from 46 °C to 60 °C (Leber et al., 2012). In this work,  
36 Jurkat cells were exposed to 70 °C for 15 minutes to trigger the apoptotic processes. Following  
37 the heat exposure, the cells were immediately resuspended in fresh medium (RPMI 1640,  
38 Corning) and placed in a 5% CO<sub>2</sub> incubator set at 37 °C to maintain them in a stable  
39 environment for further analysis. Cells were then sampled for analysis at different time points  
40 over a 12-hour period as they underwent apoptosis. Sampled cells were suspended in an Annexin  
41 V binding buffer solution containing Ca<sup>2+</sup> ions to facilitate Annexin V-PS binding.

1    **2.7. Experimental setup**

2    The cell suspension and biotin-conjugated Annexin V solution were each loaded into separate 1  
3    mL syringes and infused into the microchip using a syringe pump running at a controlled flow  
4    rate of 100  $\mu$ L/hr. The on-chip electrical sensor network was excited by a 2 Vpp sinusoidal wave  
5    at 550 kHz through the input electrode pads. The current waveforms produced at the output pads  
6    were first amplified and converted into voltage signals using transimpedance amplifiers (TIA)  
7    and then demodulated by the lock-in amplifier (LIA) (HF2LI, Zurich Instruments). The baseband  
8    signal stream, time waveform recording of the barcoded sensor activity corresponding to  
9    detections of individual cells, was sampled at 57.57 kSa/s into a computer and processed by a  
10   custom-built algorithm suite that performed signal pre-processing and analytics.

11   

12    **2.8. Flow cytometry validation**

13    We labeled a sample being validated with FITC Annexin V (BioLegend). To label the sample,  
14    cells were first washed twice with PBS and then resuspended in Annexin V binding buffer  
15    (BioLegend) at a concentration of  $1 \times 10^6$  cells/mL. Next, 100  $\mu$ L of the cell suspension was  
16    transferred to a 5 ml test tube, and 5  $\mu$ L of FITC Annexin V was added. The cells were gently  
17    vortexed and incubated for 15 minutes at room temperature (25 °C) in the dark. Finally, the  
18    sample was diluted by adding 400  $\mu$ L of Annexin V binding buffer (BioLegend) to the tube.  
19    Once labeled, the sample was introduced into the flow cytometer (LSR-II, BD Biosciences) and  
20    subsequently, the laser power values were configured for optimal measurement (FSC: 275 V,  
21    SSC: 250 V, and FITC: 250 V). For every sample being validated, the processing was stopped  
22    once 5000 events were recorded. Finally, we analyzed the flow cytometry data using FlowJo  
23    (FlowJo, LLC).

24   

25    **3. RESULTS**

26   

27    **3.1. Microchip design and assay workflow**

28    We designed our microchip to (1) discriminate between apoptotic and non-apoptotic cells  
29    through biochemical capture of those presenting PS on their membrane and (2) convert cell  
30    capture statistics to electrical data for the number of apoptotic cells in the sample (Fig. 1a). For  
31    the biochemical capture stage, we have determined the Annexin V-PS affinity to be not sufficient  
32    to capture cells in a flow stream within our device in contrast to the avidin-biotin affinity, despite  
33    the fact that Annexin V is widely used to specifically label externalized PS on apoptotic cells in  
34    suspension (Abbady et al., 2017; Vermette et al., 2003) (Fig. S3). As a solution, we aimed to  
35    capture apoptotic cells on our device via stronger avidin-biotin binding, which required labeling  
36    of cells before they are introduced to the capture chambers. Therefore, we designed our  
37    microchip to label apoptotic cells with biotin while they flow within the device and specifically  
38    capture the labeled cells on surfaces functionalized with avidin (Fig. 1a). The microchip has two  
39    inlets that symmetrically feed a micromixer stage, where one inlet is used to introduce the cell  
40    sample to be interrogated while the other is used to introduce a buffer containing biotinylated  
41    Annexin V. In the mixer, the cell sample and Annexin V solution are mixed to ensure a

1 consistent distribution of Annexin V throughout the sample. Next, the mixed solution is  
2 delivered to the cell capture chamber functionalized with neutravidin, where the apoptotic cells  
3 labeled with biotinylated Annexin V are captured on the micropillars via the strong avidin-biotin  
4 interaction and the non-apoptotic cells are let to be discharged from the outlet. For the  
5 quantification of assay results, we integrated an electrical sensor network into the microchip,  
6 which logs individual cells as they enter and exit the capture chamber. Detected cells generate  
7 coded electrical signals, which are first sampled into a computer and are processed. The fraction  
8 of cells captured by the chamber are finally computed and scored as apoptotic (Fig. 1a).

9 The fabricated microchip consisted of two parts: a passive microfluidic layer and an integrated  
10 electrical sensor network (Fig. 1b). The microfluidic layer was created by molding PDMS via a  
11 standard soft lithography process. The dimensions of microfluidic structures were optimized to  
12 work with the target cell population, i.e., human T-cells, which have an average diameter of 11-  
13 14  $\mu\text{m}$ . The microfluidic channel was 20  $\mu\text{m}$  high, which allowed unobstructed flow of  
14 suspended cells while ensuring cells proximity to the electrical sensors on the channel floor. The  
15 capture chamber was designed to be a 145 mm-long 1.5 mm-wide serpentine channel that  
16 contained micropillars, which served both to increase the functional surface area and to  
17 structurally support ceiling of the wide microchannel against collapse. As expected, a higher  
18 density of micropillars was found to enhance the capture efficiency (Fig. S4). To optimize spatial  
19 distribution of captured cells, micropillars were arranged in three zones with increasing density.  
20 In the first zone, we intentionally maintain a lower pillar density to reduce the risk of cell  
21 clogging (Fig. S5), at the cost of a reduced capture rate (< 79%) (Fig. S4). Cell clogging  
22 promotes non-specific mechanical immobilization of cells regardless of whether or not they are  
23 apoptotic-increasing the false positive detection rate. The second and third zones, in contrast, are  
24 designed to progressively favor cell capture efficiency, ultimately achieving > 92%, by  
25 maintaining increasing pillar densities (Fig. S4). Since a significant portion of the cell population  
26 is captured in the first and second zones, the likelihood of cell clogging in the third zone, which  
27 has highly dense micropillars, is significantly reduced. This strategic heterogeneous micropillar  
28 design ensures a more effective and reliable cell capture across the entire device. The  
29 micropillars of first and second zones were arranged in a grid pattern and had 50  $\mu\text{m}$  lateral  
30 spacing. While the first zone kept a 160  $\mu\text{m}$  longitudinal spacing between micropillars, this value  
31 was reduced to 50  $\mu\text{m}$  in the second zone for a more densely packed section. The third zone  
32 featured a two-position staggered micropillar arrangement, where both lateral and diagonal  
33 spacings were 15  $\mu\text{m}$  (Fig. 1c and Fig. S5).

34 Besides the main fluidic pathway taken by the cells, the microfluidic layer contained auxiliary  
35 functionalization ports and channels that were located near the inlet and outlet of the capture  
36 chamber. These peripheral channels enabled exclusive delivery of reagents to the target area,  
37 ensuring cells to be exclusively captured in intended locations. Built-in particulate filters within  
38 these channels prevented coagulate products or precipitation from forming and entering the  
39 chamber during the introduction of the different functionalization reagents. The auxiliary ports  
40 were sealed following functionalization, leaving the microchip with two inlets and a single  
41 outlet.

1 The second major component of our microchip, the electrical sensor network, comprised of two  
2 barcoded sensors strategically positioned before and after the cell capture chamber, enabling a  
3 precise count of each cell captured in the chamber (Fig. 1b). We designed the sensor network  
4 based on the Microfluidic CODES platform (Liu et al., 2016, 2017), which utilized a code  
5 multiplexed architecture that allows spatiotemporal tagging for cells from an electrical signal.  
6 The barcoded sensors consisted of positive and negative output electrode fingers, acting as  
7 current sinks, while a common input electrode served as the current source. The fingers were  
8 arranged in an interdigitated manner such that the common electrode formed pairs with either the  
9 positive or negative electrodes at prescribed locations in the sensor. These pair arrangements  
10 determined the waveshape produced by the sensor and were hence designed to follow a known  
11 and desired sequence. Each sensor consisted of 15 pairs of electrodes (Fig. 1b), resulting in a  
12 bipolar waveform with 15 peaks, each representing a bit in the code sequence. The unique  
13 waveforms were processed through a correlation-based algorithm that matched each waveform  
14 with its corresponding sensor, thereby determining the time and location of the detection event.  
15 By aggregating detection events from the entry and exit sensors and using them in a mass  
16 balance equation, we determined the number of apoptotic cells captured in the chamber. It should  
17 also be noted that the barcoded sensor network was designed to detect cells at multiple locations  
18 simultaneously, all from a single electrical output within a single microchip. This eliminated the  
19 need for additional external electrical sensing and circuitry for each individual sensor, making it  
20 possible to potentially expand the platform to include additional chambers with different  
21 chemicals or antibodies without increasing the device complexity. Moreover, our on-chip  
22 barcoded sensors are passive, avoiding any additional complexity to the microfluidic device  
23 itself and relying on advanced computational algorithms to interpret the electronic  
24 measurements.

25 **3.2. On-chip labeling of apoptotic cells**

26 To efficiently discriminate between apoptotic and non-apoptotic cells, we evaluated different  
27 capture strategies for our assay. As mentioned previously, biochemical capture of PS-expressing  
28 cells via Annexin V immobilized on our microchip resulted in an inefficient capture (~16%) of  
29 the target cells (Fig. S3). In contrast, we found that when cells from a matched sample were pre-  
30 labeled with biotin, they were captured on an avidin-functionalized device with high (~93%)  
31 efficiency. Based on these results, we decided to target apoptotic cells via strong avidin-biotin  
32 binding in our assay. However, in order to create an assay that can directly operate on unlabeled  
33 samples, we aimed to perform the labeling of the cells within our microchip. In our device, cells  
34 were introduced into a biotinylated Annexin V solution before they reach the capture chamber so  
35 that biotin-expressing cells can be selectively captured on the avidin-coated micropillars.

36 To automatically label cells in flow, we built and tested passive micromixers with different  
37 geometries (Fig. 2a). Serpentine, triangular and linear mixers, each composed of 50 repeating  
38 units (Fig. S6), were tested by measuring their efficiency in mixing dye solutions with different  
39 (red and blue) colors when each were simultaneously driven at a flow rate of 100  $\mu$ L/hr.  
40 Processing the microscope images taken at the output of different mixers, we calculated a mixing  
41 index (MI) (Materials and methods), which served as a metric that represents the effectiveness of

1 each micromixer design. From our study, we found the triangular micromixer (MI, 93.9%) to be  
2 more efficient than the serpentine (MI, 86.5%) and linear (MI, 34.9%) micromixers and utilized  
3 this design in our assay.

4 Next, we optimized the triangular micromixer design and operating conditions to maximize the  
5 mixing efficiency. First, we measured the MI at various stages of the micromixer (Fig. 2b) and  
6 concluded that 50 mixing units was sufficient to thoroughly (MI, 94.3%) mix the fluids. Then,  
7 we studied the effect of turn angle on the mixing performance. We constructed several triangular  
8 micromixers with turn angles ranging from 5° to 20°, each with 50 units, and analyzed them  
9 (Fig. 2c). While, we observed the mixing performance increased with the turn angle, i.e., the 5°  
10 design produced an MI of ~79.3%, whereas a 20° design produced an MI of ~95.2%, larger turn  
11 angles also increased the micromixer footprint. Based on these results, we have settled on  
12 channels with 15° turn angle going forward. Finally, we investigated the impact of flow rate on  
13 the mixing efficiency (Fig. 2d). We tested a range of flow rates (50-200  $\mu$ L/hr), which were  
14 identical across the two inlets. We found that higher flow rates led to better mixing with the MI  
15 increasing from ~70.2% to ~94.8% as the flow increases from 50  $\mu$ L/hr to 200  $\mu$ L/hr.  
16 Considering the negative impact of higher flow rates on subsequent capture stage and the  
17 diminishing returns on mixing efficiency for flow rates >100  $\mu$ L/hr (Fig. 2d), we decided to  
18 operate the micromixer at a flow rate of 100  $\mu$ L/hr from each inlet.

19 Lastly, our on-chip labeling process eliminates the need for manual sample labeling, which  
20 traditionally involves several complex steps. In contrast, our on-chip micromixer-based apoptosis  
21 assay platform significantly simplifies the workflow for the end user. Users simply need to  
22 introduce an unlabeled sample, and the microchip autonomously handles the internal labeling  
23 necessary for the apoptosis assay. This enables our microchip-based apoptosis assay platform to  
24 provide a label-free experience for the user.

### 25 **3.3. Processing of cell capture signals**

26 To acquire electrical data from the sensor network tracking cells in our microchip, we built a  
27 system comprising of both hardware and software components (Fig. 3a). The hardware was  
28 responsible for (a) generating a sinusoidal waveform (550 kHz, 2Vpp) for driving the microchip  
29 to excite the embedded sensors and (b) collecting the cell detection-bearing signal output by the  
30 microchip and conditioning it for sampling. Due to the interdigitated arrangement of the drive  
31 and sense electrode fingers in each sensor, the injected sinusoidal waveform creates localized,  
32 alternating current electric fields in the volume above each sensor through which cells flow.  
33 When doing so, the flowing cells create intermittent changes in these electric fields that is  
34 reflected in the electrical current between the electrodes. The raw current (typically 1-10  $\mu$ A) is  
35 first amplified by a low-noise transimpedance amplifier (TIA) with a gain of  $\geq$ 1000x. The  
36 amplified signal is then demodulated within a lock-in amplifier (LIA) to separate the baseband  
37 cell signals from the carrier sinusoidal that was initially used to excite the microchip. This  
38 demodulated signal appears as a digital stream of barcoded sensor signals; all limited to  $\leq$ 5kHz  
39 bandwidth. Next, this cell signal stream is low-pass filtered ( $f_{3dB}$  10kHz) and sampled at 57.57  
40 kSa/s before being saved to computer memory. Regarding the sampling rate, since the cell  
41 signals' bandwidth does not exceed 5 kHz, a 20 kS/s rate meets the Nyquist criterion. However,  
42 we chose 57 kS/s to provide a spectral buffer. Finally, this sampled data is used in the subsequent

1 signal processing stages to produce a count of apoptotic and non-apoptotic subpopulations  
2 present in the cell sample flowed through the microchip.

3  
4 We developed a two-stage algorithm to interpret signals extracted from the microchip. The first  
5 stage screened the raw signal stream produced by the LIA and identified the time segments that  
6 contained sensor activity from cell detection (Fig. 3b). The segmentation was based on a sliding  
7 window that ran across the raw signal stream and saved the ones which had a power greater than  
8 a threshold. This threshold was set to be 34 dB above the noise floor, which was in turn  
9 determined by first aggregating signal windows during the initial 5 seconds of program and  
10 device operation. Then, 100 time-windows with the lowest powers were identified and averaged  
11 to set the noise floor. Any detection events with power below this threshold, primarily generated  
12 by debris, lysed cells, or small particles, were discarded.

13  
14 The second stage of our algorithm correlated the pre-constructed template library with  
15 segmented signals that only contained sensor activity to identify the specific sensor generating  
16 the signal. The template library was constructed by first generating digital code signals, i.e.,  
17 bipolar square waves that each followed the bit sequence encoded in each sensor. Using these  
18 digital waveforms for correlation, we first identified 20 low distortion, high SNR (>39 dB)  
19 sensor waveforms from the signal stream for each code. All sensor waveforms with the same  
20 code were normalized in power and time before being averaged to produce the template for that  
21 code. Similarly, any cell signals that could not be matched with any template were eliminated  
22 from further analysis. Then, we identified the sensor ID by running cross-correlation between the  
23 sensor waveform and each of the templates in the library. The correlating template that had a  
24 greater than 80% match was used to determine which sensor detected the event in question (Fig.  
25 3c). This was repeated for all waveforms in the signal stream to aggregate a count of every event  
26 detected by each sensor.

27  
28 In the final stage of signal analysis, the algorithm mapped the aggregated cell events to  
29 corresponding sensors on the microchip (Fig. 3d). Namely, since the positions of the sensors and  
30 their codes were known, each cell event was used to keep a running tally of the cells detected by  
31 each sensor. Once the location of all the cell events had been registered, the algorithm then  
32 computed the difference between the total number of cells registered at the entry sensor, which  
33 represented all the cells processed by the microchip, and the total number of cells registered at  
34 the exit sensor, which represented the uncaptured, non-apoptotic cells. Therefore, the differential  
35 count between the two sensors was interpreted as the number of apoptotic cells in the sample  
36 under test (Fig. 3d).

### 37 **3.4. Biochemical capture of apoptotic cells**

38 To ensure reliable cell capture, we optimized the process to chemically functionalize the  
39 microchip. We chemically treated the microchip to immobilize neutravidin on the inner surface  
40 (Materials and methods) to target the apoptotic cells already bound to biotinylated Annexin V  
41 from the micromixing stage. As a first step, we aimed to validate the functionalization process by  
42 investigating the neutravidin coating by introducing fluorophore-conjugated biotin in the device  
43 followed by imaging with fluorescence microscopy (Fig. 4a). The results confirmed successful  
44 immobilization of neutravidin uniformly across the device. It should be noted that, while all

1 inner surfaces of the microchip were coated with neutravidin, only capture chamber was  
2 designed to provide a combination of low flow speed and surface are to promote apoptotic cell  
3 capture.

4 Next, we investigated the cell capture efficiency as a function of biotin-labeling of target cells.  
5 For this analysis, we used biotinylated Annexin V solutions with different concentrations (2-10  
6  $\mu\text{g/ml}$ ) and measured the total number of apoptotic Jurkat cells captured on the device when  
7 matching cell populations were processed (Fig. 4b). We observed an increase in capture with  
8 biotinylated Annexin V concentration as increasing the concentration from 2  $\mu\text{g/ml}$  to 10  $\mu\text{g/ml}$   
9 improved the capture efficiency from ~36.6% to ~93.7%. Based on these results, we attributed  
10 the poor capture performance at lower concentrations to a shortage of available biotinylated  
11 Annexin V molecules to bind to the surfaces of apoptotic cells and set the labeling solution  
12 concentration to 10  $\mu\text{g/ml}$  for the remainder of this study.

13 While the abundance of biotinylated Annexin V ensured efficient on-chip labeling of target cells,  
14 excess unbound amount presented a challenge as they bonded with avidin on the device surface  
15 competing with the labeled target cells. We aimed to quantify the effect of unbound biotinylated  
16 Annexin V on the target cell capture efficiency by monitoring changes in the capture rate with  
17 increasing unbound biotinylated Annexin V in the microchip. In our measurements, apoptotic  
18 Jurkat cells at a concentration of  $4 \times 10^6$  cells/ml were driven through the microchip along with a  
19 biotinylated Annexin V solution at a concentration of 10  $\mu\text{g/ml}$  and the instantaneous capture  
20 rate was calculated as a function of time (and also the infused biotinylated Annexin V amount).  
21 We found that apoptotic cell capture remained efficient at ~97% when a total of <0.33  $\mu\text{g}$  of  
22 biotinylated Annexin V was infused into the device (Fig. 4c). We analyzed a total of ~18,000  
23 cells before the device became saturated with 0.33  $\mu\text{g}$  of biotinylated Annexin V infusion.  
24 Beyond this amount, the capture rate was found to steadily decline resulting in only ~3% of cells  
25 to be captured after 1.17  $\mu\text{g}$  of biotinylated Annexin V was input. While these measurements  
26 clearly showed a capacity limit for our assay due to saturation of avidin-coated surface, 0.33  $\mu\text{g}$   
27 biotinylated Annexin V is enough to analyze ~132K cells, a number that is orders of magnitude  
28 larger than the number of cells used for a typical cell viability assay with flow cytometry.

29 Finally, we studied the effect of flow rate on the cell capture performance (Fig. 4d). The flow  
30 rate is a critical parameter for assay - excessively high flow rates can result in false negative  
31 results due to inadequate interaction time between biotin-labeled apoptotic cells and the  
32 neutravidin-coated chamber surface, while very low flow rates can lead to false positive results  
33 due to non-specific binding of non-target cells to the capture chamber. To determine the  
34 optimum flow rate, we ran the microchip at different flow rates (50 - 600  $\mu\text{l/hr}$  per inlet) and  
35 measured the resulting cell capture efficiency (Fig. 4d). Virtually all (~95%) of the target cells  
36 could be captured on the microchip for flow rates <200  $\mu\text{l/hr}$ . At higher flow rates, the capture  
37 rate decreased as expected with only ~48% of cells captured at 600  $\mu\text{l/hr}$ . Based on these  
38 findings, we set 100  $\mu\text{l/hr}$  (per inlet) as the operational flow rate for the capture chamber. This  
39 flow rate allowed for ample avidin-biotin interaction time, facilitating effective capture while  
40 minimizing non-specific adhesion artifacts and sedimentation issues that were observed to be  
41 prevalent at slower flow rates. All capture rate measurements were conducted within the region

1 between the entry and exit sensors. We found that the capture rate outside this region, in the  
2 fluidic channels on either side of the inlet and outlet sensors, was negligible (<0.5%). This is  
3 attributed to the high flow speed of cells within such regions since the channel cross-section is  
4 significantly smaller than that of the micropillar-laden capture region. Given the constant  
5 volumetric flow rate throughout the device, regions with smaller cross sections experience high  
6 flow speeds, resulting in lower capture rate.

7 Each apoptosis assay we conducted on our microchip-based platform in this work employed a  
8 new device for two key reasons: First, at the optimal Annexin V concentration of 10  $\mu$ g/mL, the  
9 capture chamber saturates with unbound Annexin V once it exceeds a threshold amount (> 0.33  
10  $\mu$ g) infused into the microchip. This necessitates using the microchip for a single run, as its  
11 capacity is finite and Annexin V binding is irreversible. Second, our current protocol does not  
12 recover the captured cells. Once the capture chamber immobilizes the target cells, it might affect  
13 subsequent runs. Our approach of creating disposable devices ensures accurate and reliable assay  
14 results by preventing cross-talk between different assays.

### 15 **3.5. Testing the assay accuracy using control samples**

16 First, we tested the specificity of our optimized assay by processing control samples. Non-  
17 apoptotic (n=3) and heat-treated (n=3) apoptotic Jurkat cells were used as negative and positive  
18 controls, respectively. Our assay reported an average of ~12.3% of non-apoptotic Jurkat cells to  
19 be apoptotic versus ~93.6% of the heat-treated positive control (Fig. 5a). Once we confirmed our  
20 assay's ability to discriminate between non-apoptotic and apoptotic cell populations, we  
21 processed heterogenous control samples we prepared by mixing untreated and heat-treated cell  
22 populations at known ratios (1:1, 1:2, and 2:1) with our device (Fig. 5b). For each mixture  
23 processed, our assay determined the mix ratios of non-apoptotic and apoptotic cell  
24 subpopulations in close agreement to the nominal ratios used to prepare the samples. These  
25 results demonstrated that our assay can quantitatively and accurately report apoptotic cell  
26 fraction in a given cell suspension.

27 Next, we benchmarked our assay against the gold standard assay for cell apoptosis, i.e., flow  
28 cytometry-based Annexin V assay. First, we analyzed the untreated and heat-treated Jurkat cell  
29 populations using flow cytometry for an independent validation of their apoptotic cell content  
30 and confirmed the negative and positive controls we used in our experiments were  
31 predominantly composed of non-apoptotic (~93.8%) and apoptotic (~96.4%) cells, respectively  
32 (Fig. 5c). Next, we mixed the positive and negative control populations at a 1:1 ratio and  
33 processed matches samples using both our assay and a commercial flow cytometer (Materials  
34 and methods) (Fig. 5d). Using our microchip, we determined ~48.5% of the mixed cell  
35 population was apoptotic, while the flow cytometer reported ~52.2% of the matched population  
36 to be apoptotic. In repeated measurements (n=3) on matched samples, we found our assay results  
37 to be within <7% of those from flow cytometer. Treating the Annexin V assay run by a flow  
38 cytometer as the ground truth, we considered ~7% as the error rate for our assay. This error is  
39 likely to be due to several factors. While our microchip-based apoptosis assay and flow  
40 cytometry measurements were both based on Annexin V to PS binding, the transduction  
41 modalities are vastly different. Greater dynamic range provided by flow cytometer for measuring

1 surface expression is another factor that might have contributed to differences observed in our  
2 results. Having said that, these results demonstrated our microchip-based assay can achieve a  
3 level of performance close to a commercial flow cytometer and importantly, with no need for the  
4 user to label the sample.

5 **3.6. Serial monitoring of acute T-cell leukemia cell apoptosis**

6 As PS externalization is a time-varying phenomenon, accurately capturing this process  
7 throughout its progression would yield valuable insight into the kinetics of apoptosis especially  
8 in drug discovery and cell manufacturing applications. To demonstrate the utility of our  
9 microchip-based assay for monitoring a cell population as they go through apoptosis, we  
10 periodically analyzed samples of a Jurkat cell population briefly (15 mins) exposed to an  
11 elevated temperature (70 °C) (Materials and methods) (Fig. 6a). Before heat exposure, only a  
12 small fraction (~5.8%) of Jurkat cells were observed to go through apoptosis according to our  
13 assay results (Fig. 6a-i). However, at 30 minutes following the heat exposure, the majority cell  
14 population was observed to be apoptotic with ~65.9% of the cells expressing PS on their  
15 membrane (Fig. 6a-ii). After an hour-long interval in a CO<sub>2</sub> incubator at 37 °C, this time ~88.8%  
16 of Jurkat cells were determined to be apoptotic (Fig. 6a-iii). Finally, we analyzed the cells at 6.5  
17 hours after heat exposure and found that virtually all (~97.1%) cells were deemed apoptotic (Fig.  
18 6a-iv).

19 To independently monitor the apoptosis process, we processed matching samples via Annexin V-  
20 based flow cytometry. The flow cytometry results at different time points matched closely with  
21 our assay results (Fig. 6b i-iv). While the concordance between our assay and flow cytometry is  
22 critical to validate our assay for this longitudinal study, it should be noted that the flow  
23 cytometry measurements could only be performed with extensive sample preparation and  
24 labeling in contrast with ours that processed cells as sampled. As such, our assay could  
25 eventually be used to perform these measurements with higher temporal resolution. Taken  
26 together, these from this study demonstrated the potential of our microchip-based assay for  
27 longitudinal monitoring of cell populations for apoptosis.

28

29 **4. DISCUSSION**

30 While we utilized PS expression on cell membrane as biomarkers for apoptosis in this work, PS  
31 externalization has also been associated with several other biological events, including cell  
32 differentiation, development, signaling, and activation. Therefore, the quantitative detection of  
33 PS externalization on a low-cost and portable microchip has the potential to equip scientists with  
34 a practical tool to study mechanisms and cellular dynamics underlying various medical  
35 conditions such as cancer (Bucur et al., 2012; Carneiro & El-Deiry, 2020; Todaro et al., 2008),  
36 inflammation (Huynh et al., 2002; Martin, 2016), immune response (Niehues et al., 2001;  
37 Opferman, 2008), and neurodegenerative disorders (Chi et al., 2018; Erekat, 2022; Ma et al.,  
38 2022).

1 Here, we reported an integrated microchip-based assay that automates the Annexin V-based  
2 apoptosis assay for detecting programmed cell death by leveraging cell membrane-externalized  
3 PS. Benchmarking against the gold standard Annexin V assay using via flow cytometry validated  
4 the accuracy of our microchip-based assay with only small discrepancies. These deviations can  
5 mainly be attributed to fundamental differences in the sensing modalities. Furthermore, each  
6 technique responds differently to the presence of non-target particles, lysed cells and their residues  
7 that are produced during preparation, in addition to the inherent impurities within the sample.

8 Our all-electronic, multistage microfluidic scheme coupled with advanced signal processing and  
9 analytics software results in our platform offering distinct advantages over existing alternatives.  
10 First, unlike conventional immunocapture-based assays which require the cells in question to be  
11 pre-labeled with fluorophore-conjugated probes, our microchip performs all sample manipulation  
12 steps within the chip allowing the device to accommodate unlabeled cells. This elimination of  
13 the need for sample labeling by the operator not only reduces assay time and prevents sample  
14 loss during handling, but also creates the opportunity to run assays on samples where pre-  
15 labeling is infeasible or prohibitively challenging.

16 Second, our microchip transduces bioaffinity-based cell capture events into electrical signals.  
17 This capability allows the use of simple electrical hardware to generate the excitation signal and  
18 condition the information-bearing signal before it is digitized and processed by software.  
19 Notably, high integrated architecture extends the utility of our platform to applications, where  
20 minimizing the sample-to-answer time is of high importance. Optical systems, in contrast,  
21 involve a combination of complex optical and electrical components. The integrated nature of  
22 our system makes it viable in a wider range of settings than its more complex counterparts.

23 Third, the incorporation of a biochemical assay with an electronic sensing scheme allowed us to  
24 create a microchip that leverages the advantages of both detection modalities. In contrast to other  
25 electronic assays that rely on indirect cytometry via size and electrical properties of the cells  
26 within the sample, our approach directly interacts with the cell membrane antigens, utilizing the  
27 well-established biochemical markers to perform the measurement. This advancement not only  
28 enhances the platform's compatibility and reliability with existing apoptosis assay techniques but  
29 also ensures a heightened level of accuracy and specificity in the analysis.

30 Lastly, our microchip stands out for its affordability and versatility. Unlike traditional assays that  
31 require expensive and bulky systems such as flow cytometers, our microchip-based apoptosis  
32 assay provides a more cost-effective solution. Each microchip uses inexpensive surface  
33 modification reagents, costing less than \$5 per chip in research settings, and can be easily  
34 reconfigured for different assays and cell types. This flexibility extends to scalable cell  
35 processing, as the device's design and surface area can be adjusted to meet diverse bioanalytical  
36 needs. The device is intentionally designed for single use to minimize contamination risks, a  
37 crucial consideration in clinical point-of-care settings. Moreover, its portable form factor allows  
38 for on-demand assays, making critical testing possible outside of centralized laboratories. The  
39 all-electronic, label-free analysis ensures both convenience and reliability, making the microchip  
40 suitable for a wide range of scenarios and applications.

1    **5. CONCLUSION**

2    We have developed a microchip-based apoptosis assay that combines the specificity of  
3    biochemical assays with the practicality of electronic devices for data acquisition and  
4    transmission. Furthermore, the assay does not require sample labeling setting it apart from the  
5    conventional biochemical apoptosis assays. Given the cell viability is a widely employed  
6    endpoint in a variety of studies, an electronic device that can automatically quantify apoptotic  
7    cell population in a sample and achieve this with a specificity and sensitivity comparable to  
8    existing labor- and capital-intensive workflows could revolutionize variety of fields including  
9    basic cell and immunology research, drug discovery and cell manufacturing.

10    **CRediT AUTHORSHIP CONTRIBUTION STATEMENT**

11  
12    **A K M Arifuzzman:** Conceptualization, Methodology, Investigation, Validation, Writing –  
13    original draft, Writing – review & editing. **Norh Asmare:** Software, Investigation, Writing –  
14    original draft, review & editing. **Tevhide Ozkaya-Ahmadov:** Resources. **Aref Valipour:**  
15    Resources. **A. Fatih Sarioglu:** Conceptualization, Methodology, Supervision, Writing, – original  
16    draft, Writing – review & editing, Funding acquisition.

17  
18    **DECLARATION OF COMPETING INTEREST**

19  
20    The authors declare that they have no known competing financial interests or personal  
21    relationships that could have appeared to influence the work reported in this paper.

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28  
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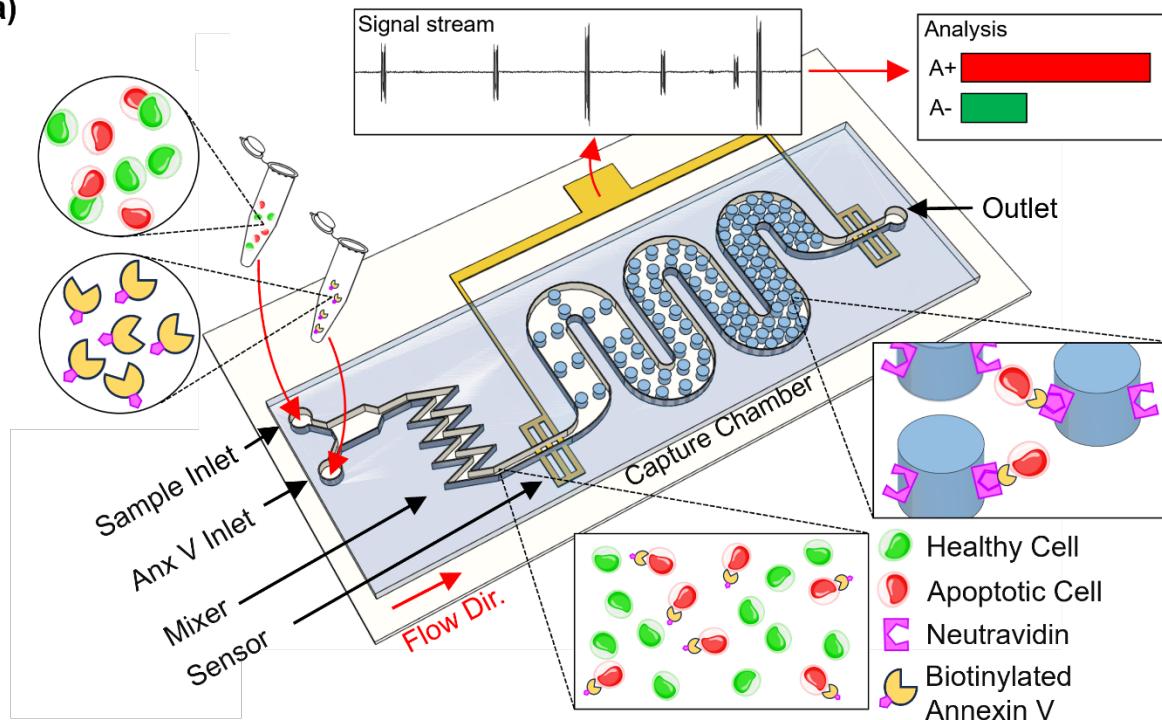
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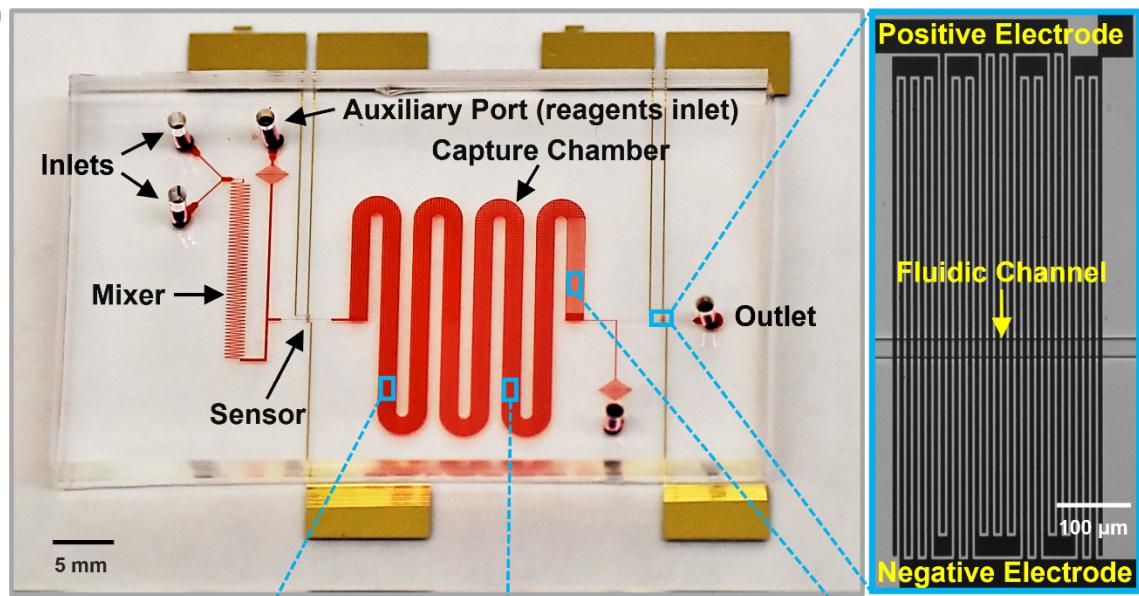
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1 Figures

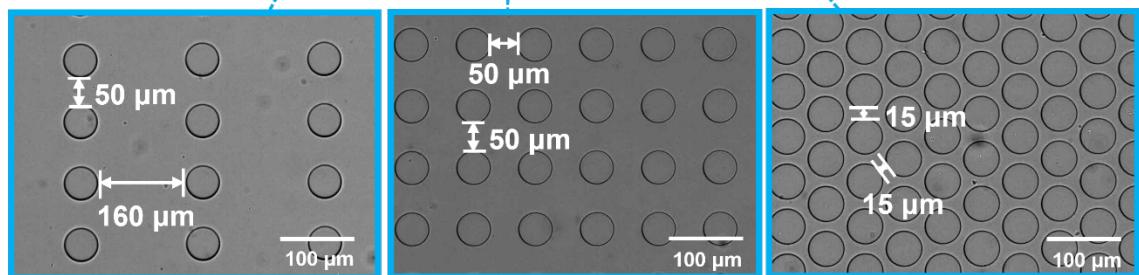
(a)



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1 **Fig. 1:** Workflow and microchip design for microchip-based apoptosis assay. (a) A schematic  
2 representation of the microchip-based apoptosis assay. The microchip automatically labels the  
3 cell population with Annexin V solution through a built-in micromixer and subsequently  
4 captures apoptotic cells that have undergone PS externalization in a functionalized chamber. An  
5 integrated barcoded sensor network transduces apoptotic cell capture events into an electrical  
6 signal, which is subsequently processed to compute the fraction of apoptotic cells in a sample.  
7 (b) A photograph of the fabricated microchip, whose microchannels were filled with a red dye  
8 for visualization purposes. Au electrode traces forming the on-chip electrical sensor network can  
9 also be seen sandwiched between the microfluidic channels and the glass substrate. Insets show  
10 close-up microscope images of the coded electrical sensors on the device, along with the  
11 microfluidic channels they monitor. (c) Microscope images of the micropillars and their spacing  
12 at the three different zones of the capture chamber, which maximizes interaction with cells as  
13 they flow through the device.

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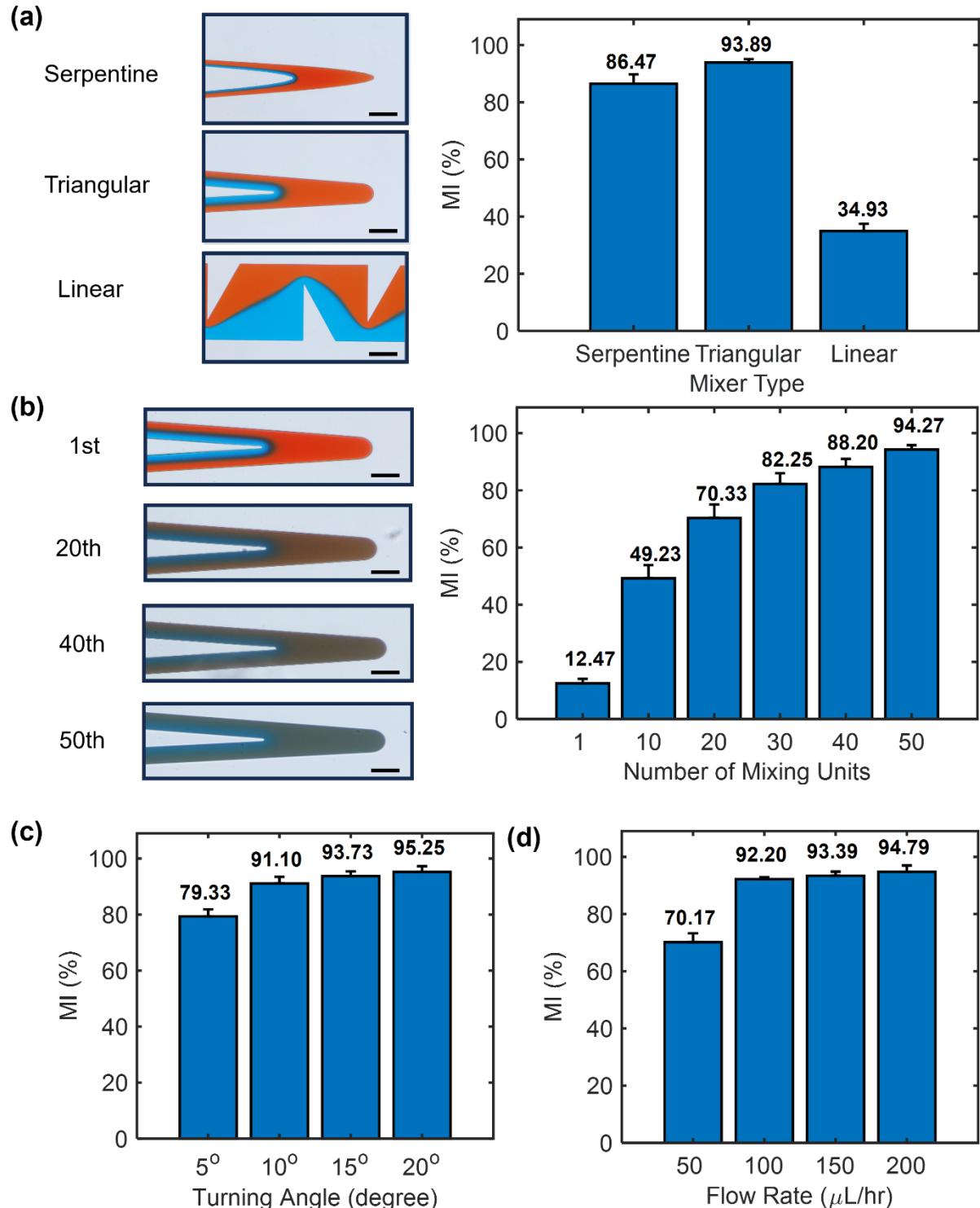
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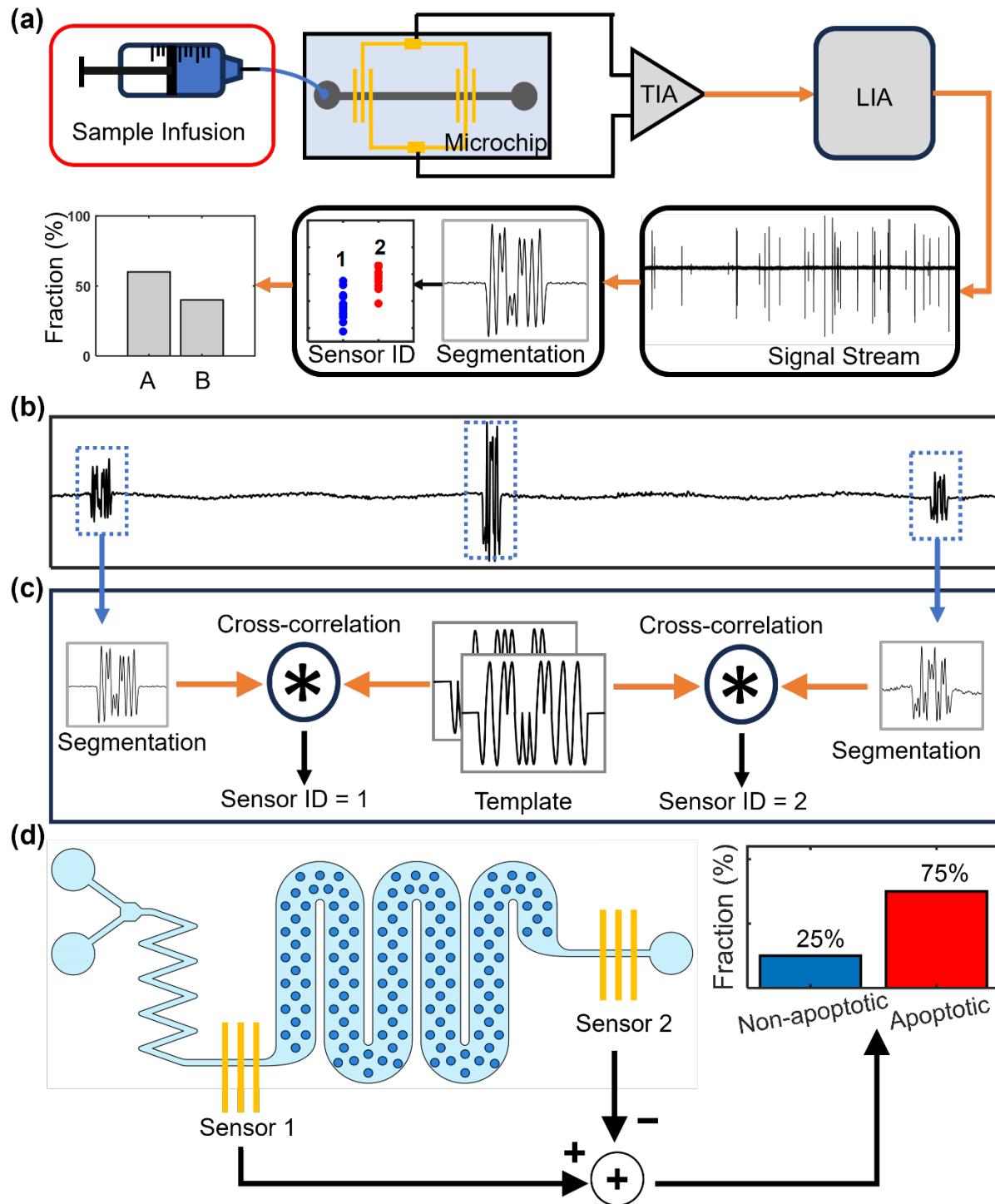
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**Fig. 2.** Characterization of the on-chip micromixer. (a) (Left) The bright field microscope images showing single mixing units of three different micromixer designs investigated in this study. (Right) Measured mixing performance for the three micromixer designs each comprising of 50 mixing units and driven at 100  $\mu\text{L}/\text{hr}$  from both inlets. (b) (Left) Colored microscope images of the triangular mixer in operation taken at different positions. The images show increasingly

1 mixed colors further along the micromixer. (Right) A plot showing the calculated MI values  
2 based on the color measurements taken at different points along the micromixer when driven at  
3 100  $\mu\text{L}/\text{hr}$  from both inlets. (c) A plot of calculated MI values for different micromixers designed  
4 to have different turn angles (5°, 10°, 15°, and 20°). The measurements were taken at the outlets  
5 of the micromixer designed with 50 mixing units, using a flow rate of 100  $\mu\text{L}/\text{hr}$  at both inlets.  
6 (d) Characterization of the optimized micromixer as a function of the flow rate. The plot shows  
7 the MI values calculated based on the measurements taken at the outlet of the micromixer. For  
8 all bar plots, the bar heights and error bars represent the mean (n=3) and standard error,  
9 respectively. Scale bars, 100  $\mu\text{m}$ .

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1 **Fig. 3.** Processing of electrical data from the developed apoptosis assay. (a) A schematic  
2 showing the individual steps in the operation of the developed microchip-based apoptosis assay  
3 system. (b) A representative section of the recorded signal stream after it was demodulated by  
4 the LIA. (c) A schematic representation of steps used in signal classification process. Waveforms  
5 from sensors are correlated with a pre-assembled template library for identification. (d) A  
6 schematic depicting how sensor signals were used to derive the number of captured cells (i.e., the  
7 apoptotic cells).

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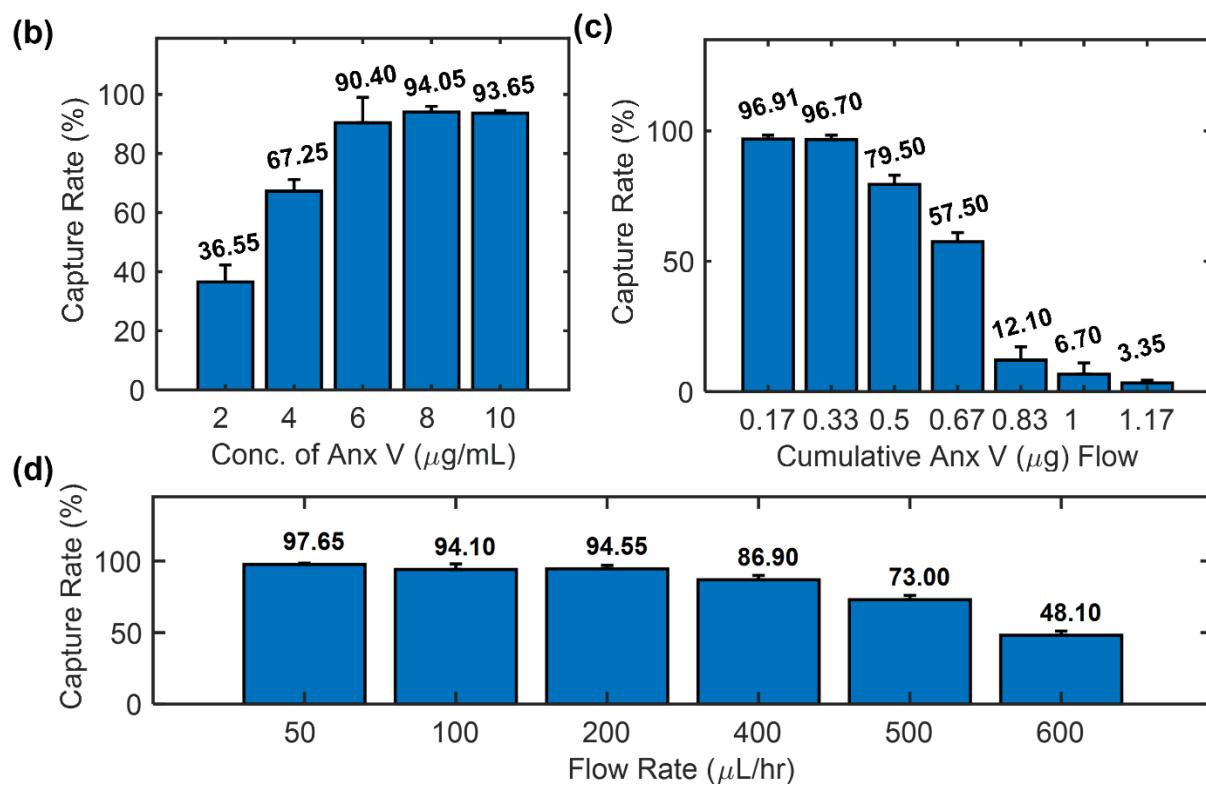
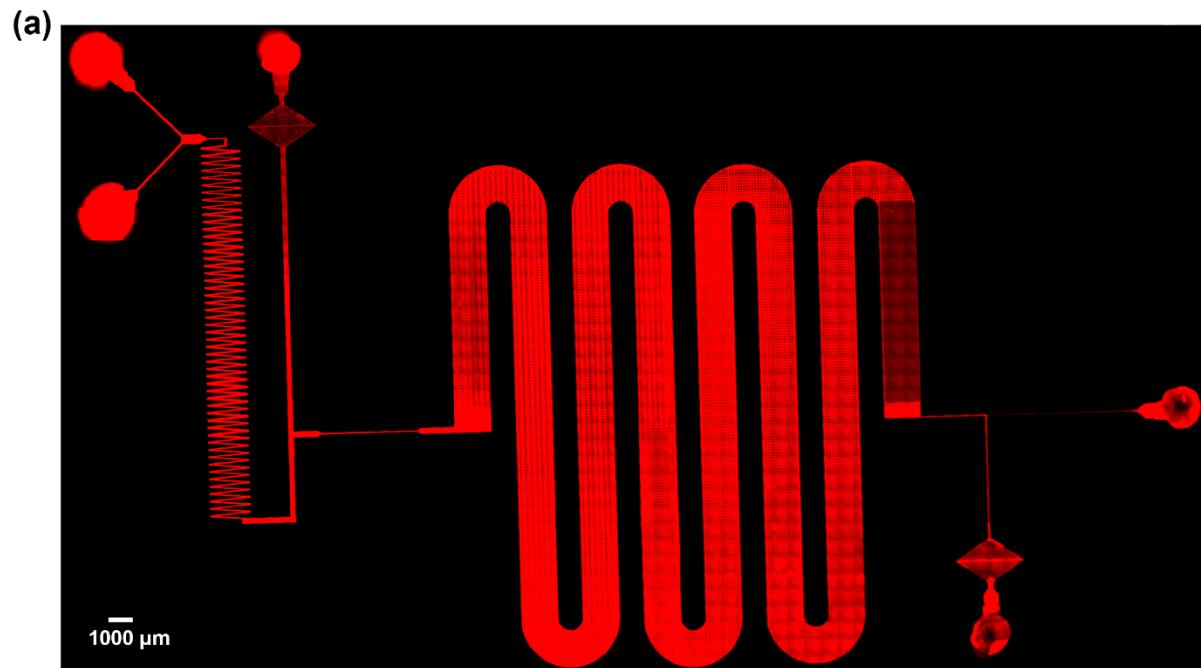
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1 **Fig. 4.** Optimization of the assay for apoptotic cell capture. (a) A fluorescence microscopy image  
2 of the microchip functionalized with neutravidin. Cy5 fluorophore-conjugated biotin molecules  
3 captured by the neutravidin-coated surface confirms the chemically active device surface. (b) A  
4 plot showing the apoptotic cell capture rates measured for varying concentrations of biotinylated  
5 Annexin V in the on-chip labeling solution. (c) Instantaneous apoptotic cell capture rate  
6 measured as the unbound biotinylated Annexin V accumulated on the inner surfaces of the  
7 microchip, whose functionalized surface was eventually neutralized leading to diminished cell  
8 capture rates. (d) Capture rates of the optimized assay measured for different sample flow rates.  
9 For all bar plots, the bar heights and error bars represent the mean (n=3) and standard error,  
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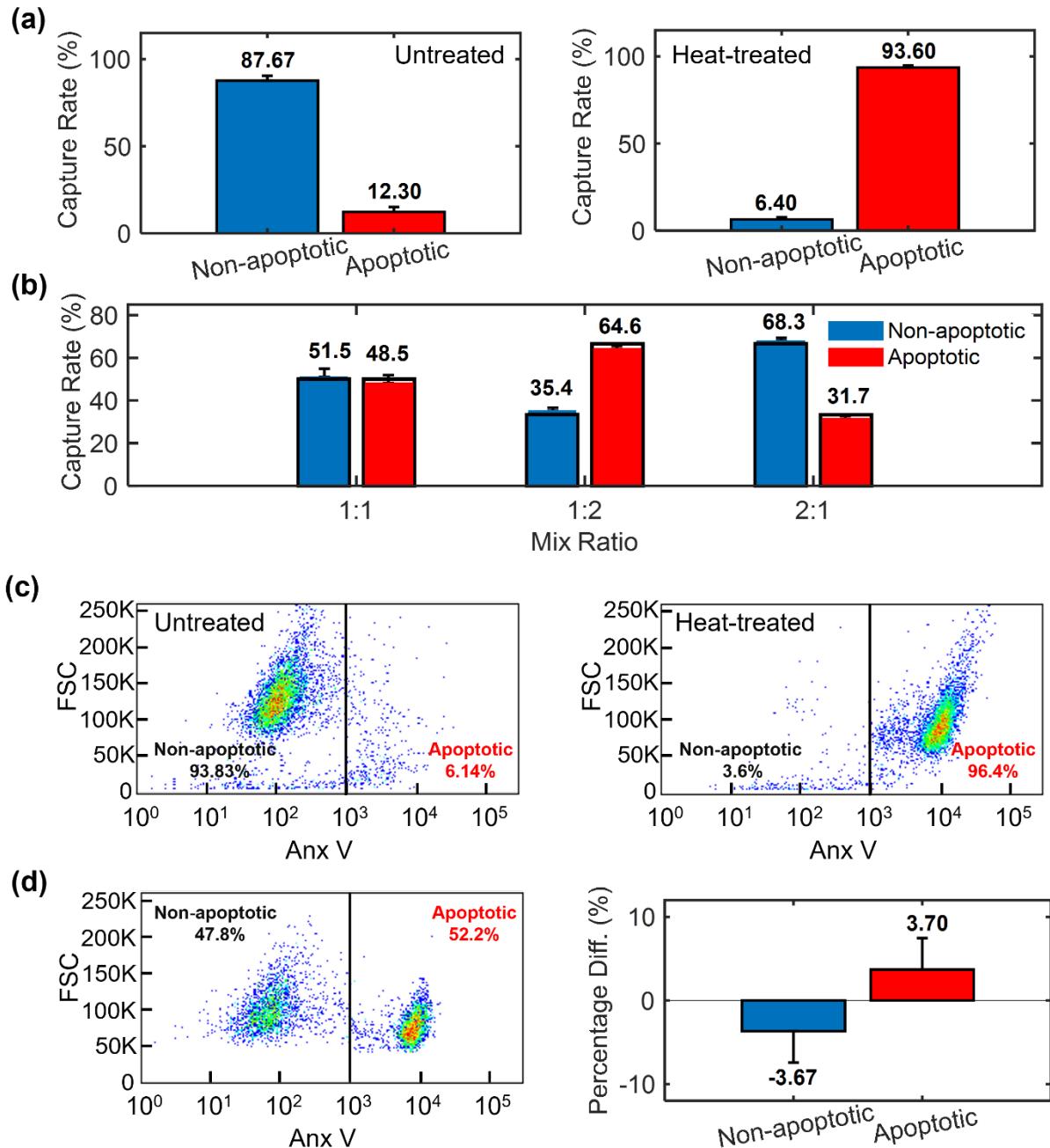
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2 **Fig. 5.** Evaluating the accuracy of our assay using control samples (a) Counts of non-apoptotic  
3 and apoptotic cells determined by our assay for untreated (left) and heat-treated (right) Jurkat cell  
4 population. (b) The figure shows the frequencies of non-apoptotic and apoptotic Jurkat cell  
5 subpopulations as measured by our system (colored bars) compared to the nominal mix ratios  
6 (1:1, 1:2, and 2:1) determined by a hemocytometer (unfilled bars). The agreement between our  
7 results and the nominal mixing ratios validates the assay accuracy. (c) Plots of PS expression of  
8 untreated (left) and heat-treated (right) Jurkat cell controls measured with through flow  
9 cytometry via Annexin V labeling. Cells were gated based on PS expression and were scored as  
10 either non-apoptotic or apoptotic (d) (Left) The plot illustrates PS expression measured in a

1 sample containing a 1:1 mix of non-apoptotic and apoptotic Jurkat cells using flow cytometry.  
2 Non-apoptotic and apoptotic cells were gated based on PS expression. (Right) The figure shows  
3 the differences in the measured non-apoptotic and apoptotic cell fractions in matched samples  
4 between flow cytometry and our microchip-based apoptosis assay. The flow cytometer results  
5 were taken as the reference. For all bar plots, the bar heights and error bars represent the mean  
6 (n=3) and standard error, respectively.

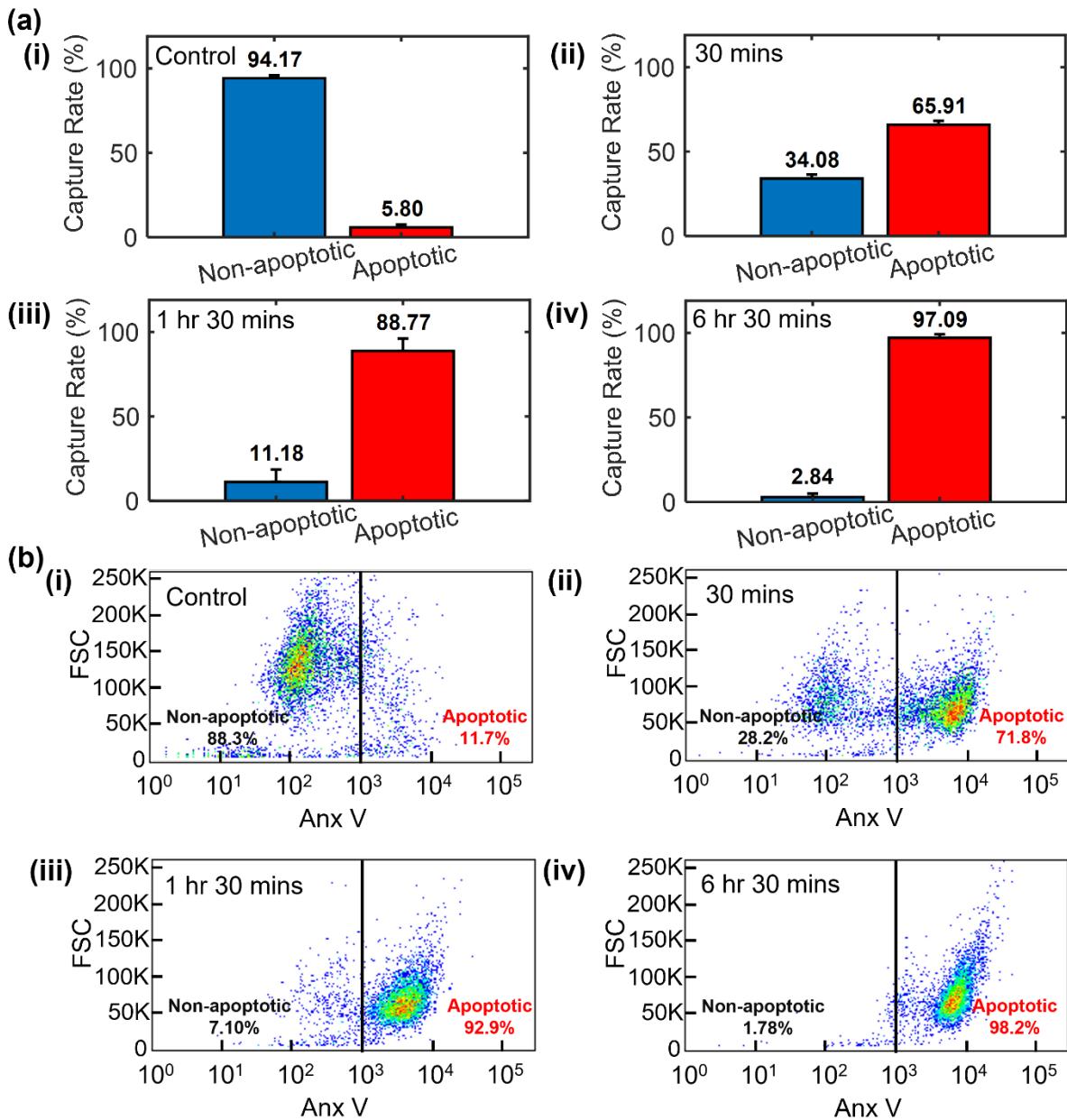
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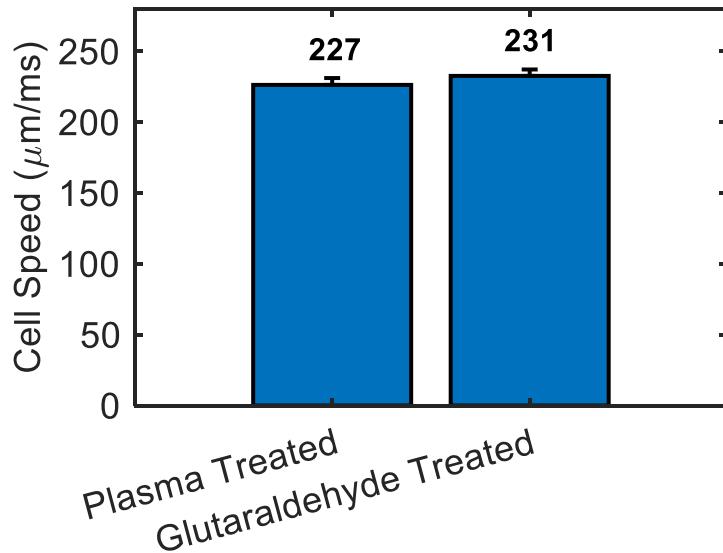
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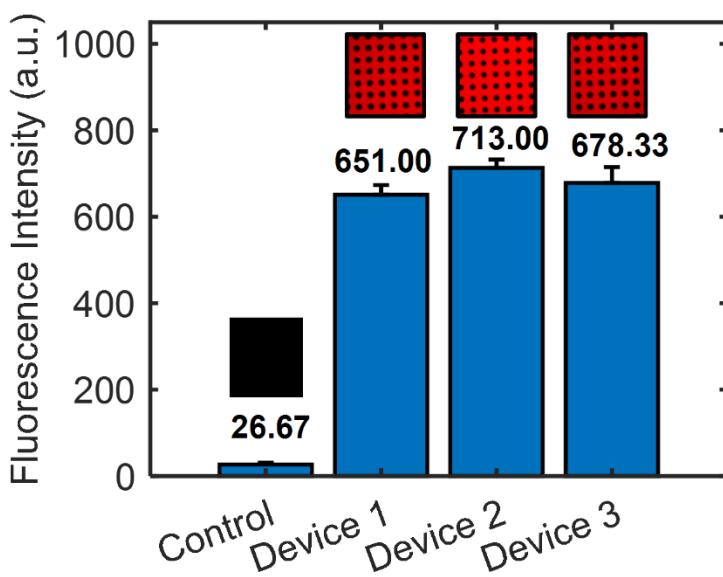


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2 **Fig. 6.** Serial monitoring of acute T-cell leukemia cell apoptosis. (a) Sequential measurement of  
3 PS externalization in Jurkat cells using our microchip-based apoptosis assay. The measurements  
4 were done (i) before any heat exposure and at (ii) 30 minutes, (iii) 1.5 hours, and (iv) 6.5 hours  
5 after the heat exposure. For all bar plots, the bar heights and error bars represent the mean (n=3)  
6 and standard error, respectively. (b) Plots show the PS expression measured in samples matched  
7 to those analyzed in (a) via flow cytometry for validation of our assay results.  
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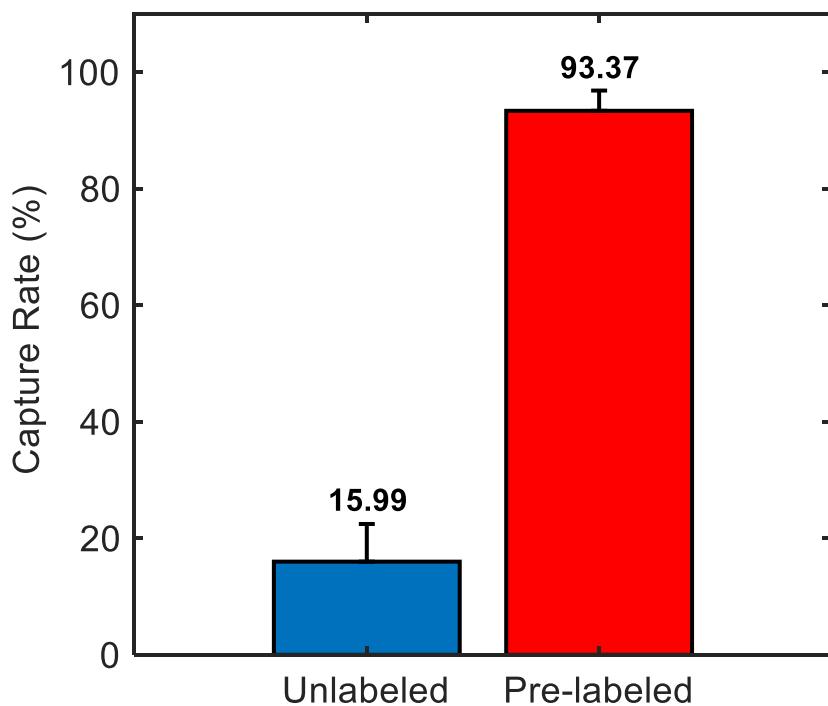
1 **SUPPLEMENTARY INFORMATION**



12 **Fig. S1.** Comparison of cell speed within plasma treated microchips with and without  
13 glutaraldehyde treatment. The samples were infused into the microchip using a syringe pump at a  
14 flow rate of 100  $\mu\text{L}/\text{hr}$ . The cell speed was measured within the microfluidic channel, which has  
15 dimensions of 20  $\mu\text{m}$  in width and 20  $\mu\text{m}$  in height. High speed camera footage (10,000 fps) was  
16 captured for both devices, and the field of view was placed at identical locations in both cases.  
17 An image processing program was developed to track each cell as it traversed the channel. The  
18 footage was denoised, background subtracted to isolate the movement of the cells within the  
19 frame and processed through a tracker that followed any visible cells throughout their time  
20 within the camera's view. To minimize error from any minute, irregular flow behavior of the  
21 cells, all the translations they made were collected over multiple frames ( $\geq 10$ ) before being  
22 averaged. This was done for all cells and repeated for both devices. The comparison shows there  
23 is no appreciable effect of glutaraldehyde treatment on cell flow speed within the microfluidic  
24 channel.

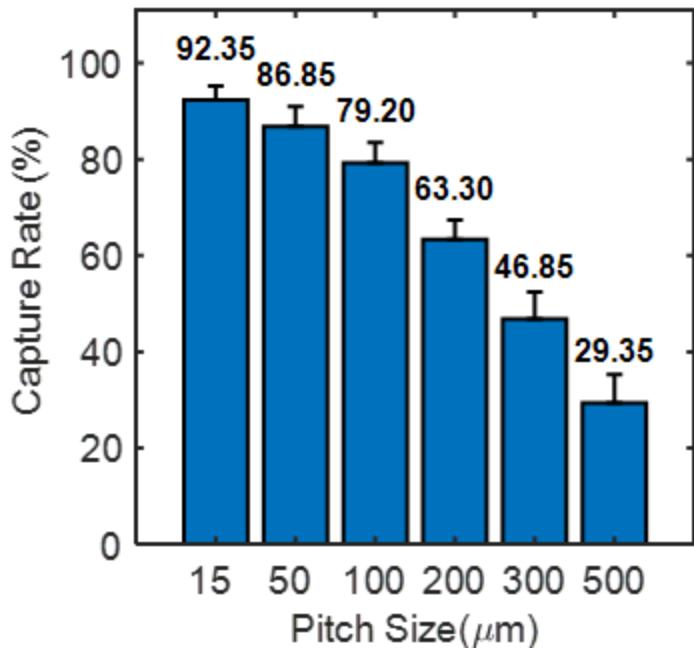


**Fig. S2.** Comparison of fluorescence intensity across four identical devices. Three devices were functionalized with the same amount of neutravidin (166.7  $\mu$ g), while the control device remained untreated. Each device was then incubated with Cy5 biotin. Fluorescence intensity measurements were taken at three different locations on each device using a fluorescence microscope. The average peak fluorescence intensity for each device was then calculated based on these three measurements. Insets show fluorescence images of the capture chambers from each device captured using a fluorescence microscope. The similarity found between the average fluorescence intensities across the three neutravidin-functionalized devices confirms that Cy5 biotin was conjugated to the immobilized neutravidin in a reproducible manner, thus validating the consistency of our functionalization process.



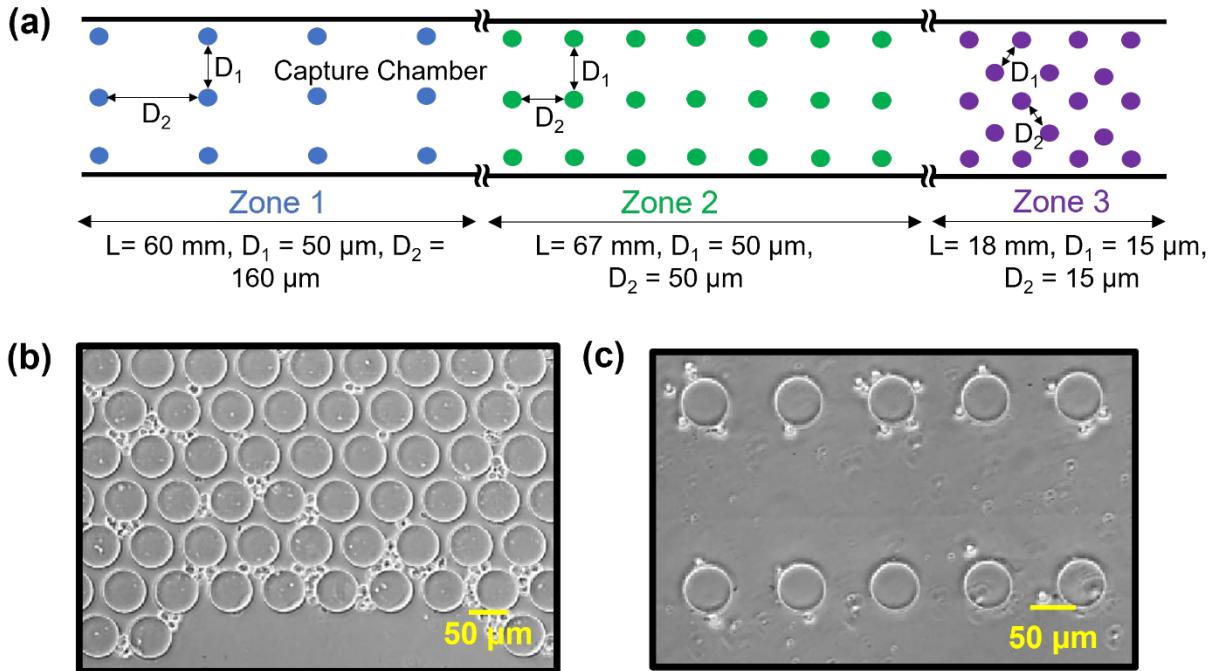
1 **Fig. S3.** The effect of pre-labeling of apoptotic cells on capture efficiency. The plot shows the  
2 measured rates of capture of apoptotic cells with Annexin V-PS and avidin-biotin binding. To  
3 utilize stronger avidin-biotin binding, cells were pre-labeled with biotinylated Annexin V and  
4 then captured on neutravidin-coated surfaces. For this plot, the bar heights and error bars  
5 represent the mean ( $n=3$ ) and standard error, respectively.

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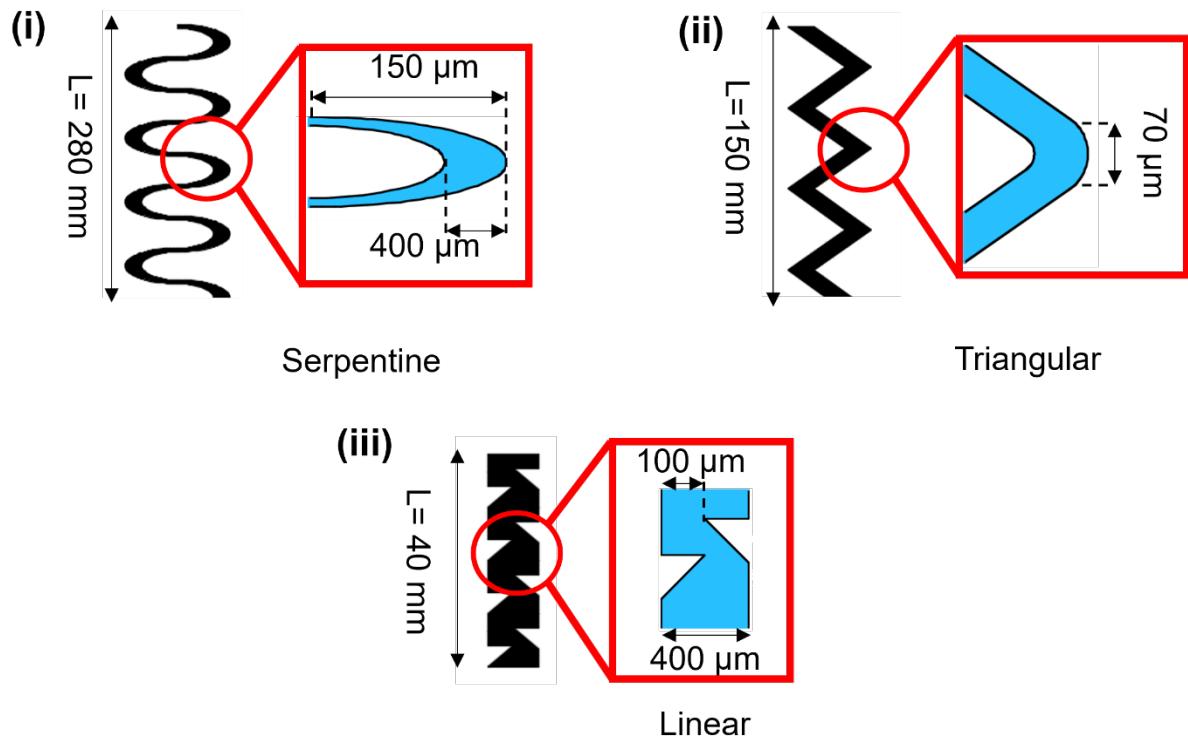


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2 **Fig. S4.** The plot illustrates the impact of micropillar density on the capture efficiency of  
3 apoptotic cells. It displays the measured capture rates of apoptotic cells with grid patterned pitch  
4 sizes ranging from 15  $\mu\text{m}$  to 500  $\mu\text{m}$ . Shorter pitch micropillars in the capture chamber result in  
5 higher capture efficiency due to increased interaction between the cells and the neutravidin-  
6 coated pillars. In contrast, longer pitch micropillars lead to a lower capture rate, approximately  
7 29%. In this plot, the bar heights and error bars represent the mean (n=3) and standard error,  
8 respectively.

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**Fig. S5.** Pillar Density Arrangement in the Device. (a) The device is segmented into three zones with varying densities of pillars (pillar diameter  $60 \mu\text{m}$ ). Zone 1 (length,  $L = 60 \text{ mm}$ ) features a lower density to prevent upstream cell clogging and unintended cell capture. Pillar density gradually increases in Zone 2 (length,  $L = 67 \text{ mm}$ ) to enhance capture efficiency, while Zone 3 (length,  $L = 18 \text{ mm}$ ) has the highest density to maximize the capture rate. (b) Initial high-density pillars at the onset of the capture chamber led to cell clogging and mechanical cell trapping, resulting in false-positive cell capture. (c) Lower density pillars in Zone 1 effectively mitigate cell clogging, ensuring accurate cell capture.



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2 **Fig. S6.** Schematic drawings showing different micromixer geometries investigated for this  
3 study, namely (i) a serpentine micromixer with a total mixing length of 280 mm, (ii) a triangular  
4 micromixer with a total mixing length of 150 mm and (iii) a linear micromixer with a total  
5 mixing length of 40 mm. Each micromixer tested comprised 50 of the mixing units shown in  
6 close-up.

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