


 Cite this: *Analyst*, 2016, **141**, 1888

High spatial and temporal resolution cell manipulation techniques in microchannels

Pedro Novo,* Margherita Dell'Aica, Dirk Janasek and René P. Zahedi

The advent of microfluidics has enabled thorough control of cell manipulation experiments in so called lab on chips. Lab on chips foster the integration of actuation and detection systems, and require minute sample and reagent amounts. Typically employed microfluidic structures have similar dimensions as cells, enabling precise spatial and temporal control of individual cells and their local environments. Several strategies for high spatio-temporal control of cells in microfluidics have been reported in recent years, namely methods relying on careful design of the microfluidic structures (e.g. pinched flow), by integration of actuators (e.g. electrodes or magnets for dielectro-, acousto- and magneto-phoresis), or integrations thereof. This review presents the recent developments of cell experiments in microfluidics divided into two parts: an introduction to spatial control of cells in microchannels followed by special emphasis in the high temporal control of cell-stimulus reaction and quenching. In the end, the present state of the art is discussed in line with future perspectives and challenges for translating these devices into routine applications.

 Received 5th January 2016,
Accepted 11th February 2016

DOI: 10.1039/c6an00027d

www.rsc.org/analyst

1. Introduction

Purification, concentration and counting of specific cell types from complex mixtures (e.g. blood, cell culture) is often required in clinical diagnostics but also fundamental research.

Well established methods include fluorescence or magnetic activated cell sorting (FACS and MACS). These techniques are readily available and widely used in laboratories as they allow effective counting and separation of cell populations. Besides routine applications, these techniques have been demonstrated to (i) detect rare cells,^{1–3} such as circulating tumor cells (CTC), and achieve high (ii) isolation/purification rate as well as (iii) throughput. However, FACS and MACS both lack of more sophisticated cell manipulation capabilities, e.g. to precisely control and adjust trajectories as required in elaborated

Protein Dynamics Group, Leibniz-Institut für Analytische Wissenschaften – ISAS – e.V., Otto-Hahn-Str. 6b, 44227 Dortmund, Germany. E-mail: pedro.novo@isas.de; <http://www.isas.de>



Pedro Novo

Pedro Novo received his MSc in Bioengineering and Nanosystems in 2010 and his PhD in Material Science in 2014, both from Instituto Superior Técnico (University of Lisbon, Portugal). During his graduation he conducted a multidisciplinary work aiming at developing a portable immunoassay readout by integrating microfluidics with micro-fabricated sensors. Currently a PostDoc at ISAS, his research interests consist on the develop-

ment of microfluidics for fast cell stimulation assays and lab on chips.



Margherita Dell'Aica

Margherita Dell'Aica received her diploma in Pharmacy in 2014 from the University of Trieste, Italy, where she focused on the HPLC-based separation of pharmaceuticals. In 2015 she joined the Protein Dynamics group at ISAS as a PhD student. The main focus of her research is studying the temporal dynamics of post-translational modifications such as protein phosphorylation after specific stimulation of cells.



cell stimulation experiments. To date, cell manipulation has been majorly performed using conventional techniques and labware such as pipettes, sample tubes, microwell plates and cell culture flasks. Notably, these strategies have only limited cell manipulation capabilities, are rather time and reagent consuming, lack sufficient precision and reproducibility and have only limited or no online monitoring capacities (e.g., integrated sensors for pH, fluorescence or magnetic field measurement).⁴ These limitations have motivated the development of microfluidic methods for enhanced spatial and temporal control of the neighboring cellular environment, as represented by numerous publications (>3.500 in the last two decades, according to a search report in ISI Web of Knowledge using the “cell microfluidics” keywords, October 2015).^{5–7}

Although different examples of microfluidic systems exist in nature (e.g., channel networks in paper), in literature the terms microfluidics or microchannels are generally employed for microfabricated fluidic networks that have characteristic dimensions below 100 μm . The possibility to tailor microfluidic networks enables the custom design of channels aiming at different manipulation methods. These may include a multitude of flow operations integrated in series or in parallel. As a consequence of the reduction of the characteristic dimensions in microfluidic channels, liquids flow in the laminar regime (low Reynolds numbers), allowing precise spatial and temporal control of flow patterns. Moreover, the conditions for laminar flow can be kept at high flow rate values (e.g., in the order of m s^{-1}), opening possibilities, e.g., for high-throughput cell manipulation^{8,9} and fast local environment switching times.¹⁰ Channel size can be fabricated to dimensions in the order of magnitude of cells, thus allowing precise control at the single cell level, but also requiring low volumes which is highly beneficial in many biological and clinical research scenarios which deal with limited amounts (e.g., cerebrospinal fluid) or numbers (e.g., research in rare disease) of samples. Control of the flow of cells by rational structure design, such as in pinched-flow *via* stream thinning elements (STE)s, or determi-

nistic lateral displacement (DLD), can be further complemented by integrated actuation methods such as those employing dielectro-, magneto- and acoustophoresis, as well as pressured valving.¹¹ Therefore, great attention has been paid to the development of microfluidic devices for cell manipulation, reflected by a vast number of publications that may be found in literature. Importantly, the overall reduction of chip size holds a great potential for portability as well as integration of sensors and actuators, which is an important factor in current lab-on-a-chip and point-of-care applications.

Integrating sensors or detection and quantification systems in lab on chips fall outside the scope of this review and therefore will not be discussed here. Suggested literature includes.^{12–16}

Owing to the large number of examples and broad spectrum of applications, this review will not cover the general state-of-the-art, but will rather focus on selected recent developments. We will first introduce microfluidic strategies used for fast cell manipulation, such as those for separation and concentration of CTCs. Then we will present the latest developments of microfluidic methods for fast cell stimulation and quenching reactions. Further examples such as for inertial,¹⁷ droplet,^{18,19} pneumatic,^{20,21} filter-based,^{22,23} electroosmotic,²⁴ immunoaffinity²⁵ and digital²⁶ microfluidics, cell traps²⁷ and optical tweezers,²⁸ are not covered here, but we recommend corresponding in-depth literature.^{29–35}

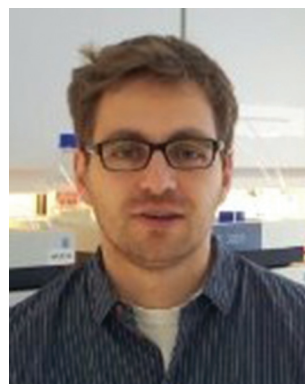
2. Biological relevance of cell manipulation studies

A cell is an extremely complex system comprising many different classes of biomolecules. Among those particularly proteins, lipids, and metabolites are highly dynamic and therefore key to the adaptability of cells to immediately respond to external stimuli. The human body comprises more than 100 different cell types and tissues, each evolutionary and



Dirk Janasek

Dirk Janasek studied Biochemistry at the Martin-Luther-University Halle-Wittenberg (Germany) and received his PhD degree in the field of chemiluminometric biosensors. After a PostDoc project on amperometric biosensors, he joined the Manz group in London in 2002 as a Leopoldina scholarship holder. Since 2003 he has worked at ISAS and is responsible for microfluidics and miniaturized separation systems.



René P. Zahedi

René Zahedi is head of the Protein Dynamics group at ISAS. He has received his PhD in Biochemistry at the Rudolf-Virchow Center for Experimental Biomedicine in Würzburg, Germany in 2008. He has been working in the field of mass spectrometry-based proteomics for more than 10 years, with a strong focus on post-translational modifications. He is currently speaker of the studies group Bioanalytics of the German Society for Biochemistry and Molecular Biology (GBM) and member of the Education Committee of the European Proteomics Association (EuPA).



individually tailored to fulfill specific tasks, individually or in concert with other cells and tissues. Indeed, this complexity is not limited to different cell types, but is also reflected in the heterogeneity among populations of the same cell type.

Even minor dysregulation of specific cells or cell–cell interactions can have dramatic effects on the homeostasis of a tissue or even the entire organism. Consequently, the detailed and quantitative study of cellular processes and their dynamics to understand causes and consequences of (dys)regulation is imperative to combat diseases such as cancer, cardiovascular and neurodegenerative disorders. It is furthermore one of the major goals of system biology research. To elucidate these processes and to reveal their underlying principles, one needs to investigate adaptations of cells on the level of proteins, lipids and metabolites under a large variety of conditions. Indeed, microfluidic applications hold a great potential to boost systems biology research, as they offer incomparable possibilities to manipulate and perturb cells, starting from minute sample amounts, in an automated, reproducible, fast and efficient way. Therefore, in the future microfluidics might be a cornerstone of sample preparation for disciplines such as (clinical) proteomics which aim at quantifying a large number of biomolecules from complex samples, *i.e.* body fluids, with high reproducibility, precision and accuracy. Moreover, microfluidics offers a higher throughput, as well as novel opportunities for temporal and spatial analysis of samples, which are otherwise not achievable.

In principle, manipulation studies have four major goals: (i) separation of cells and (ii) single cell analysis, (iii) subcellular analysis, as well as (iv) temporal analysis of cellular processes and signaling.

2.1. Separation of cells

Particularly for blood samples, which are often used in clinical diagnostics, there is a considerable need to separate and analyze specific cell types. This is not only limited to the classical blood cells erythrocytes, leukocytes or platelets and their subtypes, but also applies to microparticles, CTCs or even bacterial pathogens. Conventional biochemical methods based on differential and/or density centrifugation are rather slow and laborious, and even more importantly cannot be automated and therefore lack reproducibility. More modern methods such as FACS and MACS are more powerful, however, may require specific labeling of cells, can be time and cost intensive or have limited capacity. Here, microfluidics represents a most promising alternative that may allow a fast, comparably cheap, automated and high-throughput method to separate different cell types for subsequent analysis.

2.2. Single cell analysis

Since *in vitro* cell culture has been introduced in the early 1900's, many strategies have been developed to mimic natural cellular environments and to efficiently grow and differentiate cells. However, even in cell culture, there is a complex heterogeneity within cell populations and observed phenomena often represent average rather than individual responses.³⁶

The main interest for individual cell analysis derives from cancer research,^{37,38} where precise methods are needed to define cellular phenotypes from average population data. Starting with the first single cell-like proteomic approach in *Saccharomyces cerevisiae*,³⁹ many significant studies have followed and underline the importance of such focused assays.^{7,40} Particularly, in stem cell research, separate analyses of individual cells led to the characterization of varying levels of proteins abundance depending on time and differentiation of cells.⁴¹ The different analytical readouts used in single cell analysis comprise affinity based assays, such as flow cytometry and immune-cytochemistry, and non-affinity based assays such as mass spectrometry (MS). However, the biggest challenge in the field still remains the selective and robust isolation of single cells out of a tissue, cell culture or body fluid, often characterized by poor efficiency and reproducibility.

Indeed, recent advances in single cell analysis make use of microfluidics to overcome some of the limitations. Over the past decades a large number of high quality studies on single cell analysis using microfluidic chips have been published in the literature.^{42–48} Typically these studies aim either at (i) studying single cell's response to stimuli⁴⁹ or at (ii) detecting and concentrating specific cell types from a large population.⁵⁰ The most common microfluidic methodologies for single cell analysis involve (i) mechanical trapping, such as using posts^{27,49} and grooves,⁵¹ (ii) spatial control during fluid flow, such as by means of deterministic lateral displacement and inertial microfluidics,⁵⁰ or microdroplet encapsulation.⁵² For instance, Fernandes *et al.*²⁷ studied the modulation of alpha-synuclein toxicity in *S. cerevisiae* using a microfluidic gradient generator coupled to nine parallel chambers, each containing arrays of posts where cells were individually trapped (Fig. 1 – left). Then the number of cells with alpha-synuclein inclusions was investigated by subjecting cells to different levels of FeCl₃ and monitoring *via* fluorescence microscopy (Fig. 1 – right). Hence, this device enabled investigation of both single cell response and its magnitude upon exposure to stimuli, with applications in research on Parkinson's disease.

The field of single cell analysis, as can be assessed in the recommended literature, is vast and is starting to translate into successful industrial products, such as the case for the Fluidigm®C1 equipment,⁵³ an automated solution for single-cell genomics being capable of handling up to 800 cells individually in a single run, a considerable throughput capability.

2.3. Subcellular analysis

One of the most striking limitations of -omics technologies typically used for the in-depth analysis of cells is the lack of spatial information. Nevertheless, it is well established, that *e.g.* (i) many proteins have multiple localizations that may lead to different functions, or (ii) may translocate either upon specific cellular stimuli⁵⁴ or in disease conditions.⁵⁵ Such information, however, is lost in typical large scale proteome studies. Although these can provide quantitative information on expression levels for thousands of proteins, they cannot distinguish whether the distribution of proteins between



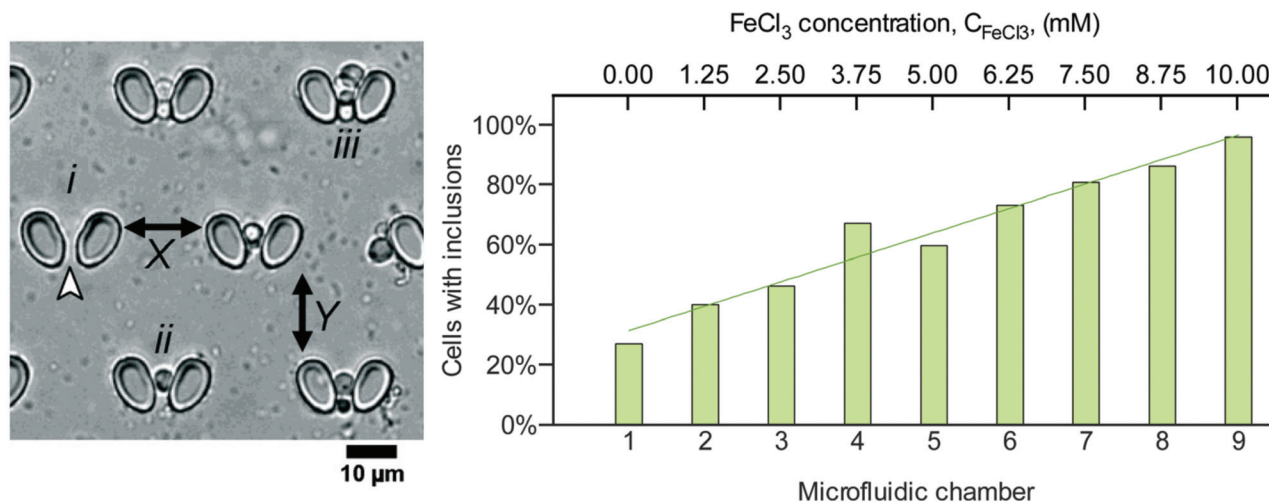


Fig. 1 Analysis of cell's response to different stimuli concentrations using a microfluidic chip combining a gradient generator with single cell trapping chambers. (Left) each trap holds a single *S. cerevisiae* cell. (Right) the number of cells per trapping chamber showing alpha-synuclein inclusions is a function of the concentration of the FeCl₃ stressor. Reproduced from ref. 27 with permission from the Royal Society of Chemistry.

different organelles or sub-compartments has changed. Therefore, the separate analysis of isolated organelles and their content is of major interest as it allows studying the subcellular localization and distribution of proteins, representing a crucial step towards a better understanding of their individual functions and dynamics. Thus, proteomic studies analyzing isolated organelles,^{56–58} or sub-compartments^{59–61} have been mainly based on classical, laborious and non-automated methods. Typically, they encompass the mechanical rupture of the plasma membrane, homogenization of cellular matrix, and separation of its components by differential/density centrifugation. Notably, free flow electrophoresis (FFE), which allows the continuous separation of a broad variety of analytes, including cells and organelles, represents an interesting alternative to the common centrifugation steps. For instance, Lu *et al.*⁶² developed an isoelectric focusing microfluidic device for subcellular organelle sorting from crude cell lysates from NR6 murine fibroblasts. Fig. 2 shows the separation of mitochondria from nuclei under free flow electrophoresis.

Alternative techniques, *e.g.* MACS, rely on the use of affinity tags or antibodies against specific epitopes present on the organelle's surface. Thus, microfluidics may play an important role and many recent publications have demonstrated different possibilities and prototypes for cell and organelle separations,^{62,63} some of which will be further described below.

2.4. Signaling dynamics

Recently, increasingly powerful mass spectrometry-based studies considerably improved our knowledge about cellular signaling, its complexity and dynamics.^{64–66} However, it concurrently became more and more evident, that the detailed elucidation of such dynamic processes as well as the underlying and intertwined networks requires more powerful sample preparation techniques. Notably, in cells different situations

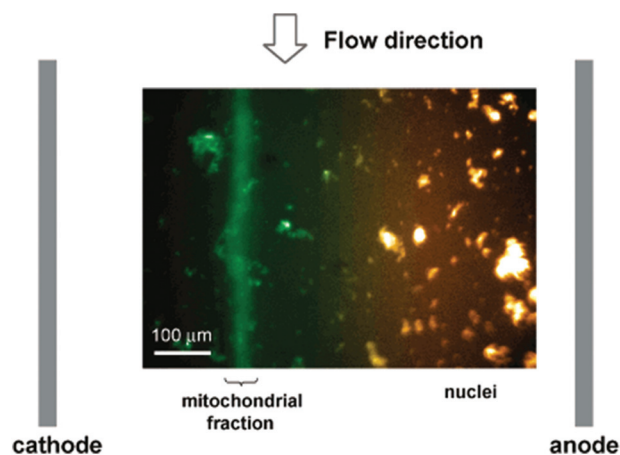


Fig. 2 Isoelectric focusing free flow electrophoresis in microfluidics for organelle sorting from a crude cell lysate using a pH gradient between 3 and 6 and under a 20 V cm^{−1} electric field. Mitochondria are focused in a band closer to the cathode whereas nuclei focus closer to the anode. Reprinted with permission from ref. 62. Copyright ©2004 American Chemical Society.

involve different timescale dimensions: processes such as regulatory protein phosphorylation upon kinase activation, or ion transport can occur within milliseconds to seconds, while others, such as changes in protein expression during cell cycle, require minutes to hours. Whereas rather slow processes can be easily followed in a time-resolved manner using conventional and mainly manual sample preparation methods, such strategies are not sufficiently reproducible and robust to study fast and short term processes. Initial signaling events are too fast to tolerate imprecise sampling and consequently measuring points; studying such fast processes demands for sub-second time resolution in combination with high accuracy and precision during sampling.



3. Fast free flow cell manipulation in microfluidic devices

Fundamental and technological developments in science progress hand-in-hand which also holds true for the use of microfluidics to study cell properties and/or cellular processes. Consequently, new reports of novel applications, designs and capabilities of microfluidic systems for cell manipulation emerge constantly. Optimizing existing protocols and/or developing original research are the main motivation drawing researchers' interest to the microfluidic technology. However, for many microfluidic systems handling issues have not been appropriately solved, and proper operation requires specialized professionals – often even the developers of the specific system – and well-controlled environments. This issue is relevant for future work and will be discussed later in this review.

Different microfluidic technologies for cell manipulation have been described in literature. These can be divided into passive, active or integrated strategies as represented in Fig. 3. The cells' position within microchannels can be controlled either by (i) rational design of the microfluidic structures, which results in deflection of cells' trajectories, as for the passive strategies, or (ii) as a result of a force generated by an actuator, taking into account the cells' electrical, magnetic and mechanical properties, as for active strategies. Often it is advantageous to combine different strategies to improve performance and functionalities within a single chip, referred to as integrated systems. However, integration is challenging, especially concerning handling simplicity, such as the need for alignment of micrometer-sized structures or fluidic and electronic connections.^{67–69} Firstly, we will revisit the aspects of different cell manipulation strategies and then highlight examples representing different integration designs thereof.

3.1. Geometry based manipulation techniques

Pinched flow (namely by using STEs), DLD and inertial microfluidics have been employed especially in cell sorting or controlled cell stimulation experiments. These are characterized by simplicity of fabrication and operation requiring only a flow control unit as external equipment.

The magnitude of the Reynolds number (Re) ($Re = \frac{\rho d v}{\eta}$, where d is the microchannel characteristic dimension, and ρ , η , ν and η are the fluid's density, velocity and viscosity, respectively) indicates whether the flow is laminar ($Re \ll 1$) or turbulent ($Re \gg 1$). In microfluidics, flow regimes are typically laminar due to very low Re as a consequence of the small characteristic dimensions (d). Hence, a particle per cell will tend to keep its flow trajectory coincident with the streamline in which it is suspended. However, when the cell's dimension becomes comparable to the microchannel cross section width, the cell's center can be deflected into a different flow lamina. This characteristic is explored analogously in STE (Fig. 3A) and DLD (Fig. 3B).

3.1.1. Stream thinning element. A schematic of the operation of a stream thinning element (STE) for changing a particle's suspension medium is demonstrated in Fig. 3A. Two particles with different radius flow through the upper inlet at a flow rate Q_1 which is much smaller than that of Q_2 . Hence, in the STE region, the flow width of the particle containing medium (σ) is much smaller than that of the second medium. Particle deflection occurs when the particle's radius (such as for r_{p1}) exceeds the width σ . An expansion channel placed at the STE output further guarantees that the particle is only submerged on the new medium. When considering the relatively small dimensions and high fluid speeds, this strategy has potential for the controlled cell deflection between media in the sub-millisecond range. For instance, Chiang *et al.*¹⁰ made use of a two-level STE microfluidic chip for sequential

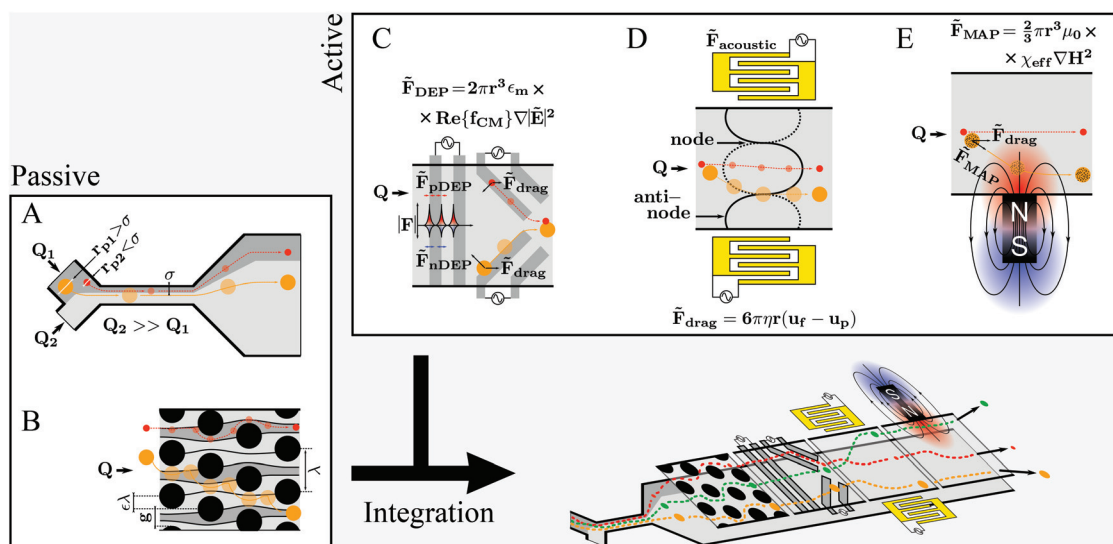


Fig. 3 Passive, active and integrated microfluidic strategies for cell manipulation. (A) STE controlled cell solution exchange. (B) DLD separation of cells with different diameters. (C) "Electrode cell highways" by DEP. (D) Separation of cells with different geometrical and mechanical properties by acoustophoresis. (E) Isolation of cells labeled with magnetic particles using magnetophoresis. Q indicates the inlet flow. In the drag force equation, $\vec{F}_{drag}$, η stands for the fluid viscosity, and u_f and u_p for fluid and particle velocity, respectively.



cell manipulation experiments, as demonstrated in Fig. 4, aiming at studying ligand-mediated cell surface signaling processes which are suspected to occur at the sub-second level. In that work, the cell is firstly deflected from its suspension medium (transparent) into a ligand containing medium to stimulate the insulin-like growth factor receptor. The reaction is arrested at the second STE in which the cell is deflected into a quenching medium. Their device showed an unprecedented time resolution control, yielding a cell switch (time it takes to completely submerge the cell membrane into the ligand medium) time of 1 ms and about 100 ms controlled cell stimulation prior to quenching.

3.1.2. Deterministic lateral displacement. The principle of cell deflection in deterministic lateral displacement (DLD) is identical to that in STE, but instead makes use of an array of pillars inter-spaced by a distance (g) larger than the cell's diameter (Fig. 3B). Typically, each successive pillar row is shifted by a distance, $e\lambda$, forcing the flow streamlines to bifurcate.⁷⁰ A cell is deflected when its radius is bigger than the width of the flow stream in which it was contained originally before it is bifurcated.⁷¹ On the other hand, small particles will maintain their trajectory at the post array's output. A critical particle size (D_c) which suffers net deflection can be calculated as (using a design constituted of an array of round posts distanced λ from center to center, with a gap g and an array to array post center offset of $e\lambda$):

$$D_c = 2\eta g e \quad (1)$$

in which η accounts for non-uniform parabolic flow through the pillar's gap.⁷² Hence, this technique is used for particle

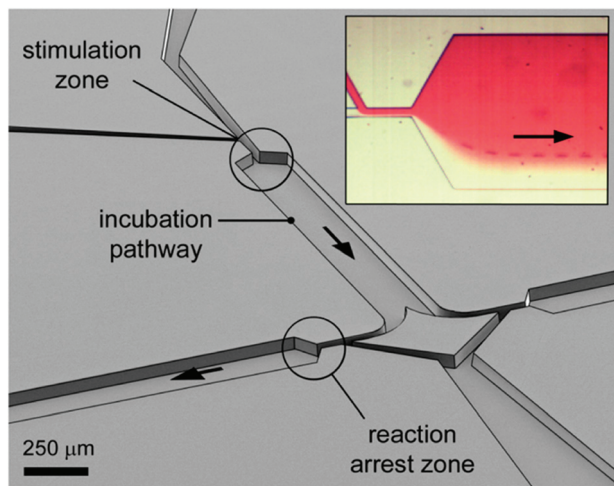


Fig. 4 Perspective open channel representation of a two-level pinched-flow microfluidic device for high temporal resolution of cell/ligand interaction. Insert figure: optical microscope image of the "stimulation zone". The cell, originally flowing in the transparent medium, is deflected into the red ligand containing medium at the first STE element. The time of exposure to the surface ligand is a function of fluid flow and length of the incubation path. The reaction is arrested at a second, down-stream located STE element. Reprinted with permission from ref. 10. Copyright ©2013 American Chemical Society.

separation depending on its dimensions and has been shown achieve high separation resolutions. For instance, Huang *et al.*⁷¹ demonstrated the use of DLD for the separation of sub-micrometer particles with a resolution of <20 nm. It is important to stress that this analysis is only valid for solid particles, whereas impacts against the posts may alter cell shape, thus changing the effective critical cell size deflected, D_c . Interestingly, Holmes *et al.*⁷² made use of the change of deformability in cells (by treating them with different concentration levels of glutaraldehyde, which makes them more rigid) and consequently differential cell separation results in a DLD microfluidic device. Cell deformation studies are of clinical relevance, such as in the case of sickle cell anemia which leads to a reduction of the red blood cells' deformability.⁷²

Notably, although simple and attractive, both DLD and STE impose shear stress on cells which might trigger unwanted biological processes or could even disrupt cells. As there is no general way for *a priori* simulation of potential cell damage, assumptions are based on empirical results and, generally, shear stress is considered qualitatively rather than quantitatively. Although it is possible to simulate shear stress induced on a particle (constituted by a defined material construction) upon flow in a microchannel, calculations become very complex for cells, owing to their erratic structural constitution (*e.g.* cell membrane, organelles, microtubules, *etc.*). Therefore, assessing cell viability after flow through microfluidic channels has been the gold standard for validating design and operation. Hence, additional experiments have to be performed every time a new or modified design is employed. Thus, Varma and Voldman⁷³ developed a cell-based shear stress sensor. They used NIH3T3 cells and created a transcriptional cell-sensor where transcription of Early Growth Factor-1 (a mechanosensitive protein) induces a fluorescent signal, achieving shear stress limits of detection down to 2 dyn cm⁻² or, in SI units, 0.2 Pa, which is well below the predicted physiologic range of 0.8–3 Pa to which osteoblasts and osteocytes are exposed,⁷⁴ for instance. Once biological changes induced by co-suspension of these cells with cells of interest can be excluded, it is possible to investigate simultaneously if a given shear stress value is overcome and whether this has an impact on the cells of interest.

3.2. Active manipulation techniques

Active manipulation techniques are based on the cells' physical properties. For instance, dielectrophoresis (DEP) and acoustophoresis (ACP) rely on the dielectric and mechanical properties, respectively. The overall physical properties of cells may be influenced, for instance, by controlled biological stimuli (*e.g.* response to membrane shear stress in osteocytes;⁷⁵ treatment of red blood cells with glutaraldehyde increases their membrane stiffness⁷²) or by surface-labeling using antibodies attached to tags or magnetic particles. The latter is often used for specific isolation/purification of cells from a mixture by magnetophoresis (MAP) since a vast selection of antibodies for different cell surface receptors and protocols for labeling are readily available. However, such



approaches depend heavily on antibody specificity as well as the availability of specific marker proteins that are uniquely expressed on one of the cell types at hand. Active manipulation techniques are not necessarily interchangeable and may actually be complementary at addressing complex cell manipulation procedures. Therefore, we will first present these techniques individually and later discuss integration developments. An advantage common to all these techniques when coupled to microfluidics is the possibility to control the position of cells in free-flow format. All following information will be related to free-flow strategies coupling DEP, ACP, MAP or combinations therefrom. For ease of reading we will not use the “free-flow” term.

3.2.1. Dielectrophoresis. Dielectrophoresis (DEP) is the motion of a dielectric particle caused by a force, generated due to polarization differences between cells and the medium under a non-uniform electric field.⁷⁶ The DEP force on a spherical cell considered to have uniform complex electrical permittivity is given by:

$$\vec{F}_{\text{DEP}} = 2\pi r^3 \epsilon_m \text{Re}\{f_{\text{CM}}\} \nabla |E|^2, \quad (2)$$

where r is the cell's radius, ϵ_m the medium's permittivity and $\text{Re}\{f_{\text{CM}}\}$ the real part of the Clausius-Mossotti factor, which is a function of the frequency of the applied electric signal and is limited to values between -0.5 and 1 . For a given set of fixed particle-medium properties, the DEP force magnitude is strongly dependent on the gradient of the square of the electric field. This value is greatly increased in microdevices as compared to bulk equivalents due to the small dimensions in microchannels.⁷⁷ Consequently, even low electric potential ($<30 \text{ V}_{\text{p-p}}$) values are sufficient to generate DEP forces above pN values,⁷⁸ sufficiently high to deflect cells in microfluidic devices. DEP has been very popular in biomedical applications for being a label free technique of relatively easy microfabrication and also for its ability to discriminate between cells of identical morphological aspect but with different electric footprint such as in the isolation of prostate tumor initiating cells as demonstrated by Salmanzadeh *et al.*⁷⁹ DEP can also be used in the particle size separation mode since the DEP force depends on the cube of the particle radius.

DEP may not be suitable for manipulating cells suspended in media of high electrical conductivity, which is typically the case for cell culture medium, and cause harm to cells by Joule heating.⁸ Therefore, cell medium exchange is required. Additionally, applied electric fields might be limited to avoid cell damage, concurrently limiting actuation force levels. Another concern regarding DEP is the lack of systematic studies, especially related to the use of high frequency signals (*e.g.* MHz) on the influence of the applied electric fields. These might cause changes observable only in the long term.

3.2.2. Acoustophoresis. Acoustophoresis (ACP) is another label-free technique⁸⁰ that makes use of sound waves to focus particles. Microfluidic ACP uses pressure resonators in microchannels to generate a standing wave field with pressure nodes and anti-nodes located at the channel cross-section. Thus, ACP

allows for free flow cell manipulation by placing the resonator in such a way that the standing wave is perpendicular to the fluid flow. Cells are focused on the pressure nodes or anti-nodes depending on the difference of their density (ρ) and compressability (β) as compared to the surrounding medium, and radius (r). As described by Laurell *et al.*,³⁵ the primary acoustophoretic force in the direction of the wave propagation (here defined along the x axis only) is defined as:

$$F_{\text{acoustic}}(x) = -\left(\frac{\pi p_0^2 V_c \beta_m}{2\lambda}\right) \phi(\beta, \rho) \sin(2kx), \quad (3)$$

where p_0 is the acoustic pressure amplitude, V_c is the cell's volume, β_m is the medium compressability, λ is the wavelength, k is the wavenumber defined as $k = \frac{2\pi}{\lambda}$, x is the distance from a pressure node and ϕ a measure of the mechanical differences between the cell and the medium defined as:

$$\phi(\beta, \rho) = \frac{5\rho_c - 2\rho_m}{2\rho_c + \rho_m} - \frac{\beta_c}{\beta_m}, \quad (4)$$

where the indices m and c correspond to the medium and cell properties, respectively. Particle aggregation at the pressure nodes or anti-nodes is further enhanced by a secondary acoustic force. This is a result of wave scattering by neighboring particles and is defined as:

$$F_{\text{secondary}}(x) = 4\pi r^6 \left[\frac{(p_c - p_m)^2 (3\cos^2\theta - 1)}{6p_m d^4} v^2(x) - \frac{\omega^2 p_m (\beta_c - \beta_m)^2}{9d^2} p^2(x) \right], \quad (5)$$

where d is the inter-particle distance, θ is the angle between the center line of the particle pair and the direction of the wave propagation, $v(x)$ is the particle velocity amplitude, ω the angular frequency and $p(x)$ is the acoustic pressure amplitude. Negative force values correspond to a inter-particle attraction situation and *vice-versa*. A direct observation of the above equation is that its value is highly dependent on the inter-particle distance, d , present in the denominator of both fractions. This indicates that the secondary acoustic force is only significant when the particles are very close to each other. Suggested literature for further insight into the physical concepts includes.^{35,81,82}

ACP offers an effective strategy for cell manipulation in microchannels, but contrary to DEP in which complex electrode “highways” can be fabricated for local cell manipulation, ACP techniques are only capable of focusing cells into pressure nodes or anti-nodes which are static along a channel cross section. On the other hand, the risk of affecting cell viability is much reduced as compared to DEP (in this case mainly due to Joule heating). For instance, Burguillos *et al.*⁸³ demonstrated that microfluidic ACP had no impact on survival or function of Microglia, Leukocytes or tumor cells under several acoustic pressure and suspension flow conditions. Li *et al.*⁸⁴ went further by claiming that acoustic-based separation offer excellent biocompatibility in terms of preserving both the cellular phenotype and genotype. They demonstrated the use of an ACP microfluidic device (Fig. 5) for the separation of CTCs



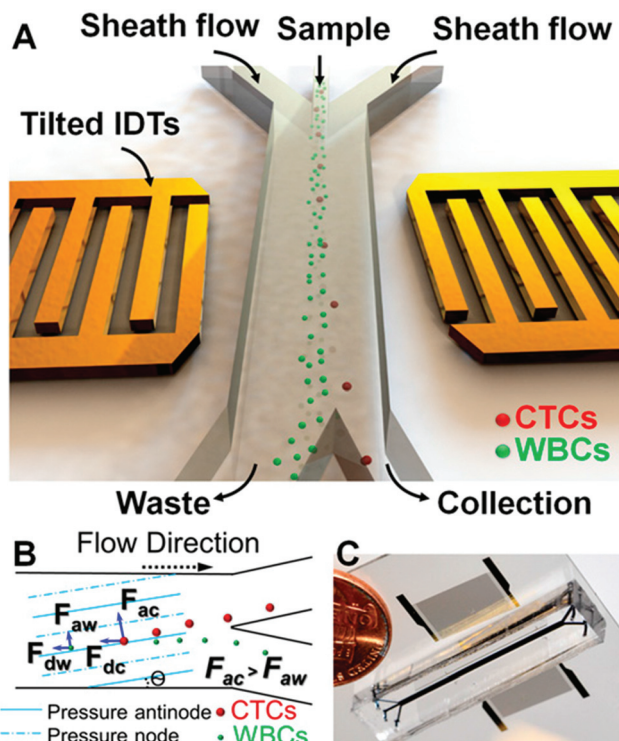


Fig. 5 ACP microfluidic device used by Li *et al.*⁸⁴ for isolation of CTCs from complex samples. (A) Tilted inter-digital transducers generate pressure nodes and anti-nodes within the microchannel cross-section, which force cells to separate. (B) CTCs and WBCs are separately collected into two outlets. (C) Photograph of the real microfluidic device next to a coin for size comparison. Reprinted with permission from ref. 84.

from white blood cells (WBC) containing solutions. Recovery rates above 83% were obtained for samples with CTC:WBC ratios of 1 : 100.000 and CTCs could be isolated from real clinical samples from breast cancer patients.

3.2.3. Magnetophoresis. In contrast to DEP and ACP, magnetophoresis (MAP) requires labeling of cells with magnetic or paramagnetic particles. MAP is a popular strategy since a vast selection of antibodies coupled to magnetic beads is commercially available and labeling of cells only requires a simple incubation procedure. Then, a bulk magnet, which ensures a force ($\vec{F}_{\text{MAP}}$)⁸⁵ (see eqn (6)) only in labeled cells, is used to deplete other cells and remove the solution. Since some antibodies bind antigens, such as cell membrane receptors, with very high specificity, MAP offers a rather simplified instrumental and proceedings for cell separation, purification and concentration. According to Karabacak *et al.*,⁵⁰ magnetophoresis has been popular because it “[...] requires simple and readily available tools”.

$$\vec{F}_{\text{MAP}} = \frac{2}{3}\pi r^3 \mu_0 \chi_{\text{eff}} \nabla H^2, \quad (6)$$

where μ_0 is the permeability of vacuum and χ_{eff} is the effective susceptibility of the cell which is related to the intrinsic susceptibility, χ_i , and given as $\chi_{\text{eff}} = \chi_i(1 + \chi_i/3)$.

Table 1 Cell manipulation techniques' details

Cell handling technique	Principle for cell manipulation	Force	Shear stress	Versatile cell control
STE	Dimension	—	++	✓
DLD	Dimension	—	+++	✓
DEP	Electric properties	5 pN ⁸⁶	++	✓
Acoustophoresis	Mechanical properties	2 pN ³⁵	+	—
MAP	Label cells with magnetic particles	6 pN ⁸⁷	+	—

The above advantages, together with aforementioned aspects of microfluidics, are the motivation behind developing microfluidic MAP equivalents. In parallel to the bulk equivalent, microfluidic MAP are characterized by simplicity of fabrication and magnet integration. This strategy, is very attractive for cell separation and focusing since low intensity magnetic fields are not known to harm cells. However it is “a one way” focusing strategy that lacks the versatility of DEP, for instance, in which the cell can be guided using dedicated electrode “highways”. Moreover, it does not offer simple solutions for multiple cell sorting, in contrast to multiplexed FACS techniques available in certain flow cytometers. Recently, Giudice *et al.*⁸⁵ reported the development of a particle sorting device employing MAP. First, a mixture of magnetic and non-magnetic beads are aligned along a streamline. Then, particles are separated into two different outlets *via* MAP force, yielding a deflection efficiency of 96%. See Table 1 for a summary of the above techniques' details.

3.3. Integration

The presented techniques for cell manipulation have inherent advantages and disadvantages which do not necessarily overlap, rendering them suitable only for some applications. Thus, cells that have identical dimensions, electrical and mechanical properties, and only differ in the expression (levels) of certain membrane surface proteins can only be differentiated in microfluidic devices by using appropriate labels, such as in MAP-based separation. Most studies use proof-of-principle examples of low complexity, such as cultured cell suspensions instead of complex mixtures such as blood, for development and verification of their devices. However, often such approaches are not sufficiently flexible to be successfully translated to specific real-world applications. For instance, size-based separation of blood cells does not necessarily allow distinguishing healthy from infected red blood cells. Hence, progressively researchers invest efforts to integrate different methodologies into a single lab-on-chip (LoC) (as represented in Fig. 3-bottom) in order to either merge advantages of different strategies while canceling out handicaps, or to enhance cell manipulation control by joint forces. In the next sub-sections, we will present recent examples of integration on microchannels for improved cell manipulation.

3.3.1. ACP and DEP. Collins *et al.*⁸⁸ developed an integrated DEP and ACP microfluidic device for the separation of particles with different dimensions and termed it virtual DLD. The device employs a novel fabrication scheme that made use



of interdigital electrodes, deposited over a piezoelectric lithium niobate, for both exerting ACP and DEP forces, as represented in Fig. 6. The device was capable of sorting particles with small dimension differences down to the sub micrometer range (*i.e.* 300 nm from 500 nm particles). The authors demonstrated that this integrated strategy yields similar overall particle separation results, even when using different microchannel heights, with DEP and ACP being dominant in small and tall microchannels, respectively. The reason is that due to the planar electrode fabrication the DEP force range is limited and therefore compatible with channel heights close to the dimension of cells. However, the limited DEP force range is compensated for by the ACP force which spans the overall channel height.

3.3.2. DEP and DLD. Beech *et al.*⁸⁹ developed an integrated device employing DEP to “tip the balance of DLD”. Here, the separation of particles by DLD is complemented by DEP forces in order to increase the critical particle diameter for separation by 50%. Furthermore, the use of DEP in this integrated construction allows (i) decreasing the shear stress applied to cells by avoiding or minimizing collision with posts, as well as (ii) discriminating between particles of the same dimension but different dielectric properties, as demonstrated by Shafiee *et al.*⁹⁰ who separated living from dead human Leukemia THP-1 cells.

3.3.3. ACP and MAP. Adams *et al.*⁹¹ showed the development of serially integrated ACP and MAP actuation in a microfluidic device in order to attain high enrichment of specific particles from a mixture. Their device had three individual outlets for separate collection of fluid fractions that ideally should only contain non-, acoustically or magnetically targeted particles. Therefore, they used a mixture of particles with different diameters: 0.9% 5 μm fluorescent green polystyrene particles (acoustic target), 0.7% 4.5 μm magnetic microspheres (magnetic target) and 98.4% 1 μm blue fluorescent polystyrene particles (non-targeted). The particle mixture distribution was chosen to mimic rare target separation and was flowed in the microchannels at a rate corresponding to 10^8 particles per h. Whereas they collected 94.8% magnetic target, 0.2% acoustic target and 5% non-target particles at the magnetic target outlet, the distribution at the acoustic target outlet was 89.2%

acoustic target, 2.4% magnetic target at 8.4% non-target particles. Additionally, 91% of all target particles were recovered from the initial mixture. This work demonstrates a simple integration yet efficient scheme for multi-parameter particle separation, which might be useful for biological applications. However, further development should be invested (as with most proof-of-principle designs employing particles) due to the considerable difference between synthetic particles and cells which might even compromise the device's proper operation, for instance, as discussed in the DLD section (*i.e.* regarding the different deformability of cells – not rigid particles).

3.3.4. DLD, inertial focusing and MAP. An almost transversal promise throughout LoC literature is that these devices perform at least comparable if not considerably better than traditional bench-top equivalents, while reducing complexity. Although this is generally true, one should note that often LoC devices only partially comply with the procedures required to attain the desired output (when compared to conventional methods) requiring third party processing by technicians. This is owing to the challenging integration of several independent functional units into a single all-in-one LoC device. As aforementioned, the combination of different strategies for particle or cell manipulation have increased microfluidic operation performance, however not without still leaving room for further development. Karabacak *et al.*⁵⁰ went beyond in the integration level and developed a two-level strategy for the isolation of CTCs from blood samples (Fig. 7A). The whole apparatus includes two devices: the first, named CTC-iChip1, performs DLD on a concurrent flow of whole blood and buffer, in order to isolate larger cells (WBCs and CTCs), while depleting abundant and undesired red blood cells and platelets. The first device is connected in series to a second, named CTC-iChip2, which further guarantees that only CTCs are collected at the output of interest. In a first stage, the cells are focused to a precise location of the channel cross section by inertial microfluidics. Then, in a second stage, WBCs (which were labeled with magnetic particles *via* anti-CD45 and anti-CD66b antibodies) are magnetically guided to a waste reservoir. Consequently, CTCs are collected into a second reservoir for counting purposes. The device was tested using real blood samples,

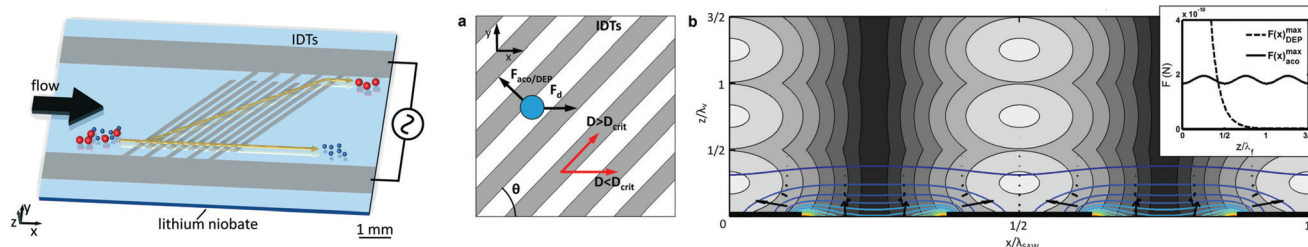
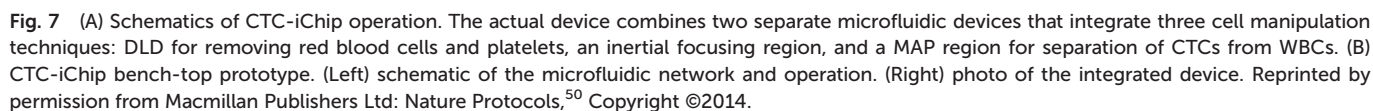


Fig. 6 Combination of DEP and ACP by integrating electrodes in a microchannel. (Left) schematics of the integrated microfluidic constitution and operation. (Middle) the electrodes make an angle θ toward the fluid flow direction. A particle, flowing from left to right will be subjected to a combination of drag force in the direction of the fluid flow, and DEP and ACP forces both perpendicular to the electrodes. Particles/cells with dimensions above the critical diameter will be deflected along the electrodes. (Right) representation of the acoustic pressure field magnitude in gray, and the first 10 DEP force potential contours in color and the DEP force vectors. Reproduced from ref. 88 with permission from the Royal Society of Chemistry.





and subject of many studies in the literature^{32,93,94} that will not be covered here. However, it is important to stress that, in contrast to traditional cell culture, microfluidic applications require more time and skills to successfully adhere/cultivate/maintain cells. On the other hand, microfluidics allows automation in a highly parallelized (high throughput) fashion under well-controlled conditions that are not achievable otherwise.^{7,9} Notably, adherent cells pose substantial limitations to the range of applications. When fast solution switching is required (*e.g.*, in the sub-second range) two important limitations have to be taken into account: (i) cells are shear stress sensitive and will not allow the use of high flow rates, and (ii) fast solution switching while avoiding contamination is a time limiting factor, because it usually requires physical separation of solutions (*e.g.*, using valves or air bubbles). In the next paragraphs we will present some work developed recently which is based on cell adhesion models.

4. Fast cell stimulation and quenching

4.1. Using perforated membranes

Chingozha *et al.*⁹⁵ developed a microfluidic device using a PDMS perforated membrane to allow stimulus from one side of the membrane to reach cells located on the other side (Fig. 8A). Hence, they created a physical barrier that controls the flow rate delivered to the cells in order to control the shear stress applied. At the same time, the hydraulic resistance was tuned to ensure rapid mass transport of the ligand. Cells were seeded into the cell's compartments by gravity-driven flow. As shown in Fig. 8B, an array of single cells (Jurkat E6-1 human acute T lymphoma) were compartmentalized within the cell microfluidic compartment, separated from the active flow layer *via* pores. With the cell layer inlet closed, the microfluidic device was flushed sequentially with buffer and stimulant with control of the frequency of switching. The authors used FITC and fluorescence microscopy in order to characterize the device's performance in terms of (i) homogeneity of the mass

Most recent developments on fast cell stimulation techniques make use of cell adhesion on specific parts of the microchannels. Then, cells can be subjected to a sequence of different solutions, such as growth medium exchange, stimulant containing solution and lysis or fixation buffer, simply by controlling the injection of solutions. Adsorbing, culturing and maintaining cell populations is *per se* not a trivial task

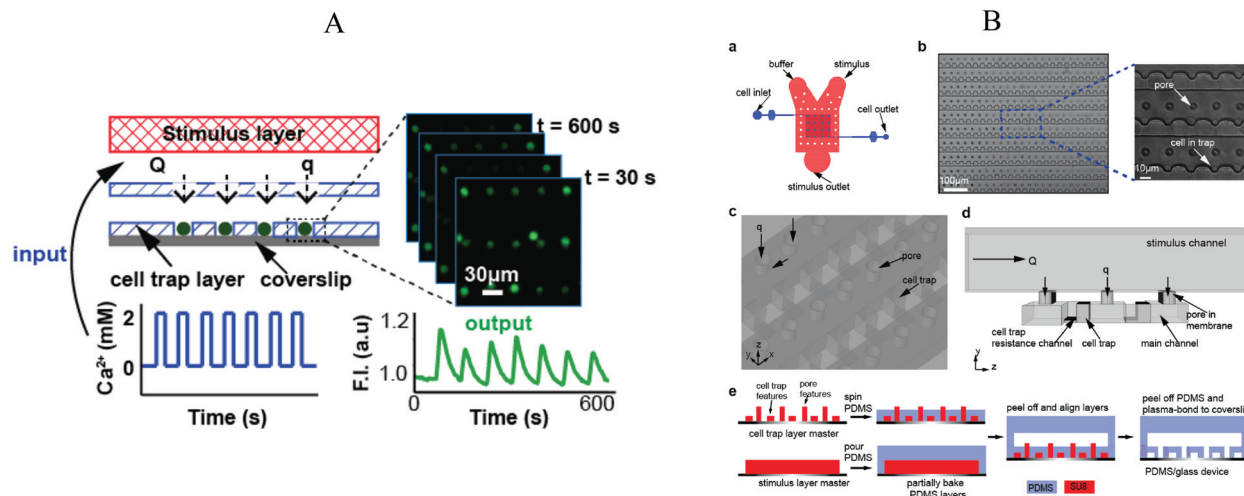


Fig. 8 (A) Illustration of the microfluidic device divided in three layers: (top) stimulus carrying layer, (middle) perforated membrane, (bottom) high density cell trap layer. Fluorescence measurement was performed under application of an oscillating stimulus. (B) Detailed representation of the microfluidic device, structures and microfabrication process. Adapted with permission from ref. 95. Copyright ©2013 American Chemical Society.

delivery to the different cell locations and (ii) required time (“rise-time”) to exchange the solution at the cell location. They found a negligible difference in the periodicity of the upstream and downstream flows (<0.01%) and a rise-time of 154 ms, both values in good quantitative agreement with their computer simulations.

The device was used to evaluate the response of intracellular Ca^{2+} depleted Jurkat to alternating flows of Ca^{2+} and EGTA solutions, using Fluo3-AM as signal. Different switching periods between 20 s and 2 min were tested, showing the potential of this device to perform controlled stimulation experiments. Moreover, the device offers a general strategy since cell adhesion to the microfluidic walls is not mandatory. However, ensuring that cells are individually immobilized, particularly in long term experiments, is not trivial, even more when using small cells such as bacteria.

4.2. Electro wetting – digital microfluidics

Recently, Ng *et al.*⁹⁶ described the use of digital microfluidics to perform immunocytochemistry in single cells. Digital microfluidics makes use of planar electrodes to change the contact angle of droplets in order to move them above a planar surface.⁹⁷ The device comprised two parallel plates, separated in about 180 μm distance, forming an “infinite” microfluidic chamber. An array of electrodes was deposited on one of the plates, covered with a Teflon film to avoid cell adhesion. In the second plate, specific areas for cell adhesion (*i.e.* hydrophilic areas) were patterned into a Teflon coating (Fig. 9). Fluids were moved around by sequential actuation of the planar electrodes. Except for the cell adhesion spots all areas of the microdevice were hydrophobic, allowing precise circulation of nl volumes of solution. By flushing a solution through the hydrophilic spots a “virtual microwell” was formed by trapping a given quantity of liquid between the two parallel plates. Its lateral footprint was defined by the hydrophilic area (Fig. 10).

Liquid exchange was performed by circulating a second solution through the virtual microwell area with an exchange efficiency of 93% in a single step and up to 99% in two steps.

The digital microfluidic device was used to investigate the activation dynamics of the phosphoinositide 3-kinase signaling pathway upon stimulation of fibroblasts with PDGF. Around 300–400 cells were immobilized in each virtual microwell. A series of 11 automated steps were conducted to perform cell starvation, ligand stimulation, fixation, permeabilization, blocking, staining and washing. Successive end-to-end solution exchanges were possible in about 1.3 s. Hence this device allows for fast cell stimulation assays down to the second time regime.

4.3. Chamber based microfluidics for sequential cell stimulation

Another concept to control the sequential flow of solutions to adherent cells on a chamber is by using pneumatic valving. Blazek *et al.*¹¹ developed a large scale integration platform containing 16 “logic blocks”, each addressing 8 chambers for conducting high throughput studies of fast protein phosphorylation kinetics in NIH3T3 fibroblasts (Fig. 11a). The chip was supplied with all solutions required using tubing and eventually flasks, thus avoiding any further disturbance during experiments. An automated pressure controller was used to ensure controlled flow rate delivery and timings of the different reagents used, with the full experiment taking about one and half days. The device’s theoretical time resolution of 1.5 s for cell-stimulation was measured by monitoring the time required to fully exchange the cell chamber buffer at different pressure values (see Fig. 11b and c). However, the experimental resolution time was 10 s, since the authors used stimulation pulses of 5 s followed by a cell fixation pulse. Full cell activation experiments were performed on chip and signals were quantified *via* proximity ligand assays. Cell counts between



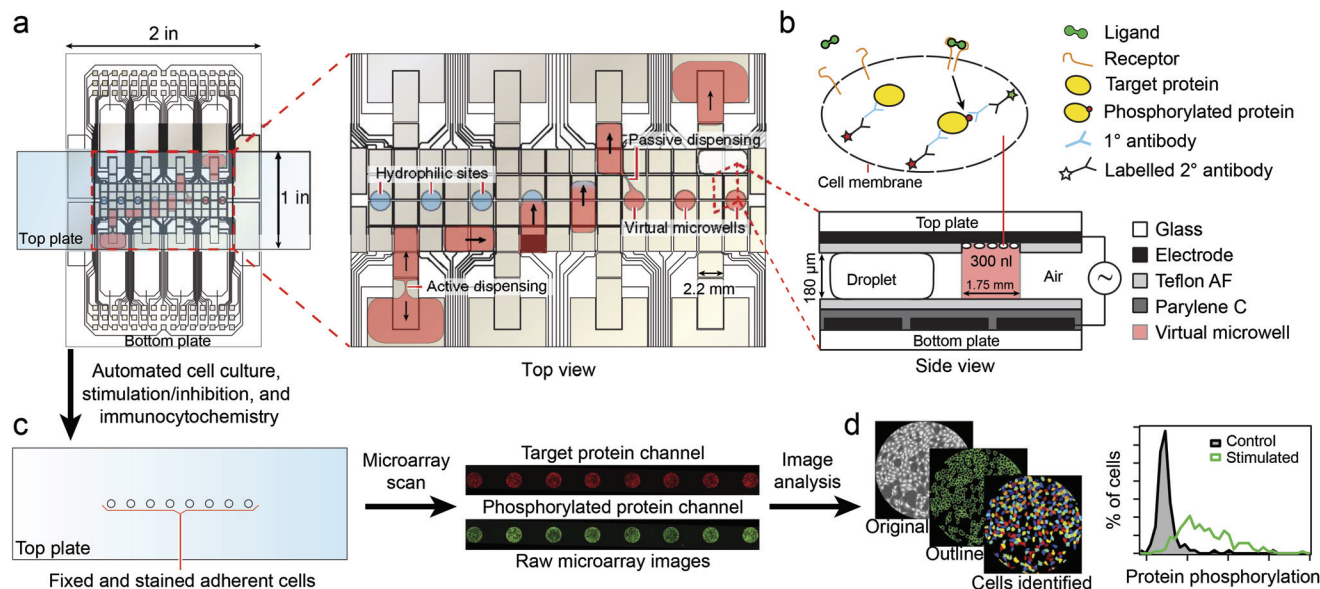


Fig. 9 Schematics of the digital microfluidics device. Fluid droplets are circulated by actuation of integrated planar electrodes passivated with a Teflon coating. In designated areas, the absence of hydrophobic coating allows the formation of a virtual microwell where cells are trapped for further stimulation experiments. Signal inspection was performed using a microarray scanner. Reprinted with permission from ref. 96. ©2015 Macmillan Publishers Limited.

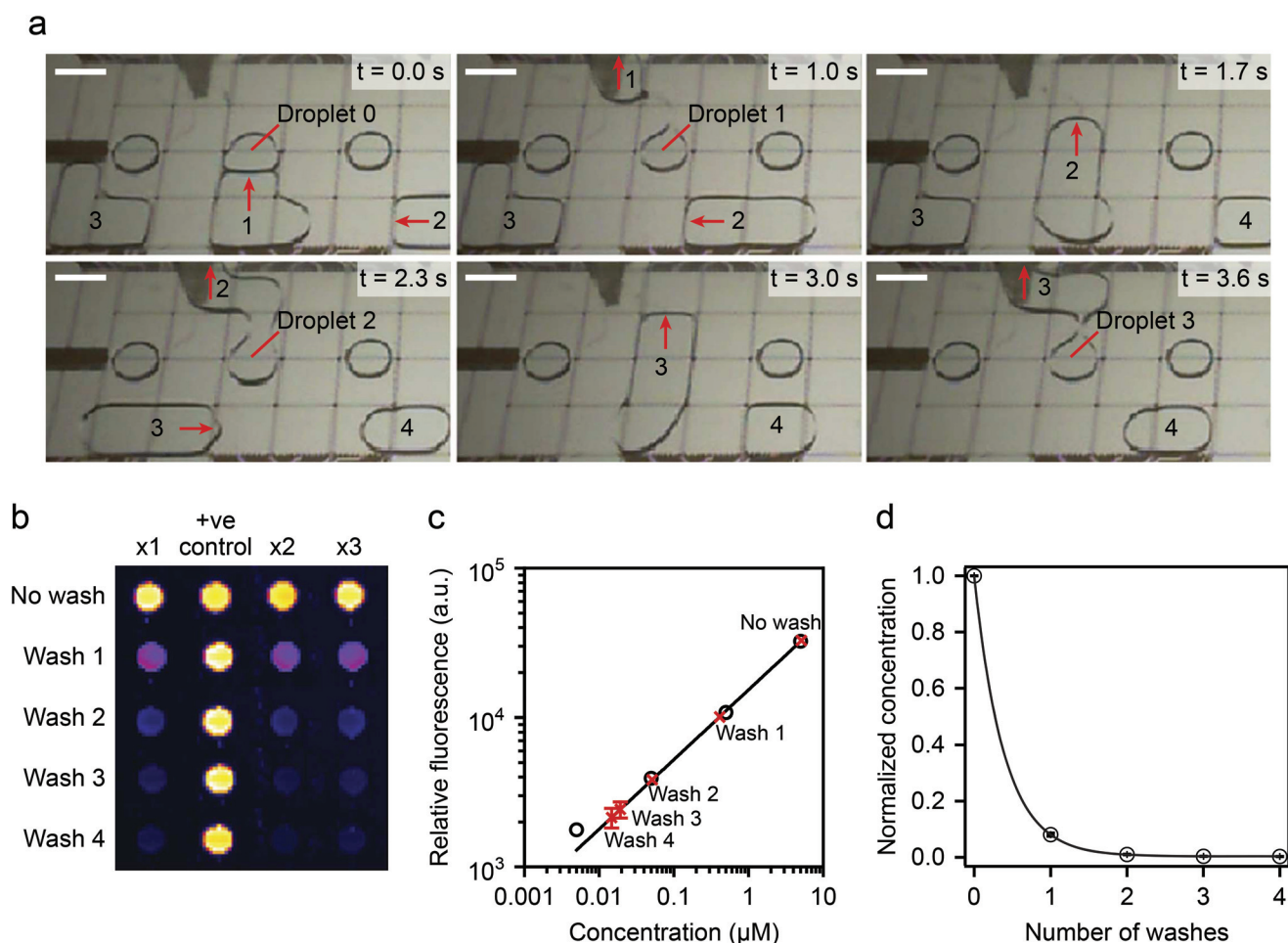


Fig. 10 (Top) device operation details. A virtual microwell containing 5 μl of fluoresceinamine is washed sequentially and the remaining fluorescence compared to calibrated data. (Bottom) washing efficiency experimental results: two washing steps displaced >99% of the original solution. Reprinted with permission from ref. 96. ©2015 Macmillan Publishers Limited.



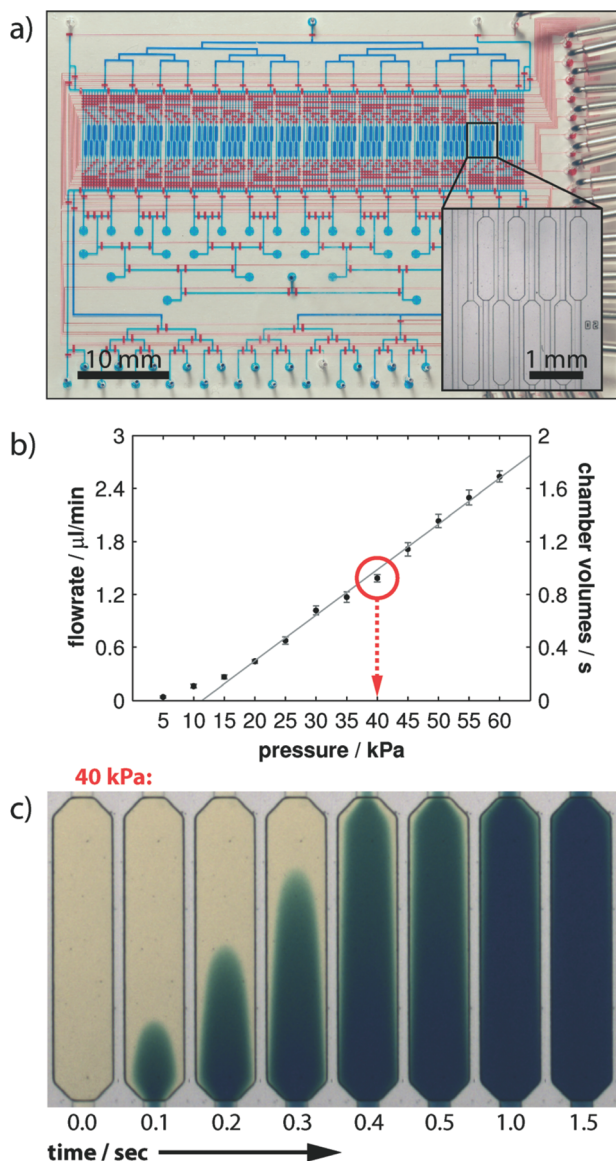


Fig. 11 (Top) microfluidic chip overall design. Flow and control micro-channel layers were filled with blue and red solutions for ease of visualization. Zoom of the cell culture chambers. (Middle) flow rate calibration results as a function of the applied pressure difference. (Bottom) time-lapsed images of buffer exchange in one cell culture chamber for an applied 40 kPa inlet pressure. Reproduced from ref. 98 with permission from the Royal Society of Chemistry.

350 and 500 were used to ensure statistically relevant data. The characteristic phosphorylation times for different cell receptors (PDGF and IGF-1) were between 13 and 35 seconds.

Despite reduced time resolution compared to the previous examples, this work demonstrates a highly compact and parallelized cell stimulation device that once properly interfaced with the outer world can perform complete cell stimulation experiments autonomously. However, interfacing complex tubing and pressure source connections in microfluidics is a challenging task.

4.3.1. Free-flow cell stimulation and quenching. “Whole cell quench flow” microfluidic systems can be used as alternative to adherence-based cell stimulation microfluidic methods, by using STE¹⁰ or micromixers.⁹⁸ Here, cells flow freely in microfluidic channels while pumping the cell suspension. Since cells will follow the solution flow, the fluid velocity at their membranes is lower than in the surface adherent version and therefore shear stress is reduced.⁹⁸ Once the cell is effectively transferred into the stimulation solution, stimulation time is a function of flow rate and dead volume, including how fast the cell is brought into contact with the stimulus and the reaction arrest. The same is true for aforementioned examples and the bottleneck in many others throughout literature, *i.e.* often methods lack of a precise quenching/lysis method. This is of particular relevance, as fast signaling events based on phosphorylation can proceed within milliseconds to seconds.⁹⁹ Therefore, analogous to stimulation, quenching/lysis should be performed in microfluidics (in opposition to off-chip) to ensure high temporal resolution, however requires additional care. An SDS containing cell lysate is highly viscous compared to typical microfluidic solutions and specific designs need to accommodate for this change. One example of such a strategy has already been presented in the “STE” section above. Briefly, a microfluidic device comprising an in series two-level STEs was used for whole cell quenched flow analysis. A cell suspension and stimulation reagent were input into the first STE level which ensures that cells (HeLa S3) migrate from the cell suspension into the stimulation media (IGF-1) in about 1 ms. Quenching was performed by merging the previous flow with quenching solution in a second STE stage. Although this strategy ensures high temporal resolution, it also requires high flow rate ratios to ensure that cells are properly deflected into different solutions. Taking a 1 : 7 flow

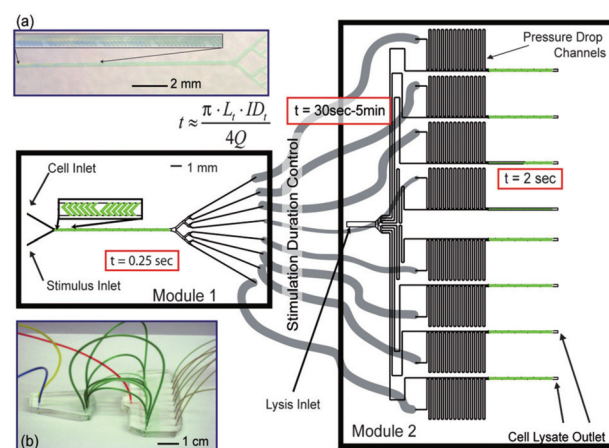


Fig. 12 A two module microfluidic application for cell stimulation and quenching. (Top left) photograph of module 1 showing mixing of colored solutions, achieved in 0.25 s. (Middle) the mixed solution is divided through 8 outlets which are connected to module 2 via different tubing lengths corresponding to different cell stimulation time points. (Right) the input from module 1 is mixed with lysis buffer in module 2. Reproduced from ref. 11 with permission from The Royal Society of Chemistry.



Table 2 Cell manipulation techniques details

Cell handling technique	Principle for cell manipulation	Time resolution	User friendly	Generalizable
Perforated membrane	Controlled solution exchange	++	++	+
Digital microfluidics	Electro-wetting	+	++	+
Chamber and valving	Solution exchange using valving	+	—	++
Pinched flow	Deviation of cells <i>via</i> STE	+++	++	++
Micromixers	Mixing cell and stimulus	++	+++	+++

ratio for instance, per ml of cell solution 7 and 49 ml of stimulation and quenching solutions are required, respectively, which may pose restrictive limitations to the concentration of the final lysate. Furthermore, for typical microfluidic device sizes and lysate collection requirements, single experiments may take hours of operation per time point experiment. Since in microfluidics liquids flow in the laminar regime, mixing of substances is controlled by diffusion. Diffusion constants are very low and especially reduced for larger molecules, hence mixing solutions in microfluidics is time inefficient. However, mixing times can be greatly reduced by increasing the number of interfaces between the liquids to mix within a microchannel cross section (*i.e.*, diffusion distances are decreased). Microfluidics especially designed for this effect have been termed micromixers.¹⁰⁰ Although micromixers might not offer such a high temporal resolution as STE elements, they offer the possibility to mix different solutions at comparative flow ratio values, hence avoiding dilution problems. For instance, Hirsch *et al.*⁹⁸ used a herringbone-based micromixer to activate T cells. The microfluidic device (Fig. 12) integrated several micromixers in parallel using a common source of cells, stimulus and lysis buffers. A mixed solution of cell and stimulus buffers, obtained in only 250 ms, was split into eight different time stimulation circuits (30 s to 5 min). Cell lysis was also performed in parallel and the experiments were carried out with an equivalent of 5% of material as compared to conventional methods (Table 2).

5. Discussion

As reviewed here, alongside high spatial resolution, high temporal control of cell manipulation in microfluidics has been performed at unprecedented levels. High temporal control is especially important when high time resolution is required and/or in characterization of fast occurring phenomena, both of which are of the interest of different fields such as research and clinics. However, key to success encompasses also technology transfer. The transfer of microfluidic technologies, such as those for cell manipulation, into real world applications faces two main obstacles, referred to in several publications in recent years.^{101,102} The first obstacle is to guarantee reproducibility of microfluidic operation in non-comfort zones, *i.e.* in the field of application rather than in the laboratory of development, without the need for specialists. Although, as demonstrated in this review, many studies have demonstrated high

potential and control regarding particle per cell manipulation, issues regarding interfacing microfluidic chips to up- and downstream applications of the external world are often omitted. Importantly, in many cases this may actually compromise the practical application of otherwise elaborate devices. The above mentioned issues are well-known and in the past years researchers have put efforts into both scaling integration levels of microfluidic systems and designing practical chip-world interfaces. This applies to microfluidic devices for particle or cell manipulation, but also to other microfluidic based applications, such as biosensing, which may actually be used complementary in cell studies; for instance, early cancer diagnosis facilitated by such technologies either by detection/quantification of cancer biomarkers and CTCs may improve treatment outcomes. The second obstacle is the development of a working prototype into industrialized products. This is mainly attributed to the inability to mass-scale produce microfluidics in their typical research environment form. Thus, PDMS has been the material-of-choice for microfluidic device fabrication, but casting is incompatible with established industrial fabrication techniques such as micro-injection and compression molding, and thermoforming. Thus, whereas microfluidic devices open up novel possibilities in fundamental research, there is still a gap to bridge for routine usage and applications, such as in clinical research.

6. Perspectives

Microfluidic devices, whether in their early development stages or as integrated systems, have increasingly received attention by the research community. Among those, particularly microfluidic devices for cell manipulation have attracted attention in the field of life science and (bio)medicine. Nowadays there is a huge demand, both economically and ethically, to miniaturize procedures and develop cell-based experiments that might replace traditionally performed animal models in accordance with the 3R (Replacement, Reduction, Refinement). Along this line, there is a general trend to reduce time and costs of analyses and increase efficiency. All these concepts may greatly benefit from migration to microfluidic formats. Indeed, microfluidic technology has opened new paths to perform original studies that had been impossible previously, and many tools are available. However, what is lacking is the development and fabrication of standardized microfluidic connections and ideally of-the-shelf parts, such



as in the case of electronics. Then, building a microfluidic circuit would be easily customizable, rendering it increasingly attractive to research in (bio)medicine and biology, among others. Furthermore, despite a vast research community of potential customers, the potential of microfluidics is seemingly still confined to experts. Breaking this invisible boundary by advertising its power to the general scientific public would greatly promote further interdisciplinary discussion, cooperation and development.

Acknowledgements

Financial support by the Ministerium für Innovation, Wissenschaft und Forschung des Landes Nordrhein-Westfalen, the Senatsverwaltung für Wirtschaft, Technologie und Forschung des Landes Berlin, and the Bundesministerium für Bildung und Forschung is gratefully acknowledged. Pedro Novo and Margherita Dell'Aica also thank for ISAS strategic funds.

References

- 1 Y. Hu, L. Fan, J. Zheng, R. Cui, W. Liu, Y. He, X. Li and S. Huang, Detection of circulating tumor cells in breast cancer patients utilizing multiparameter flow cytometry and assessment of the prognosis of patients in different CTCs levels, *Cytometry, Part A*, 2010, **77A**(3), 213–219.
- 2 T. Hristozova, R. Korschak, V. Budach and I. Tinhofer, A simple multicolor flow cytometry protocol for detection and molecular characterization of circulating tumor cells in epithelial cancers, *Cytometry, Part A*, 2012, **81A**(6), 489–495.
- 3 P. Li, Z. S. Stratton, M. Dao, J. Ritz and T. J. Huang, Probing circulating tumor cells in microfluidics, *Lab Chip*, 2013, **13**(4), 602–609.
- 4 A. Lenshof and T. Laurell, Continuous separation of cells and particles in microfluidic systems, *Chem. Soc. Rev.*, 2010, **39**(3), 1203–1217.
- 5 J. El-Ali, P. K. Sorger and K. F. Jensen, Cells on chips, *Nature*, 2006, **442**(7101), 403–411.
- 6 C. Wyatt Shields IV, C. D. Reyes and G. P. López, Microfluidic cell sorting: a review of the advances in the separation of cells from debulking to rare cell isolation, *Lab Chip*, 2015, **15**(5), 1230–1249.
- 7 A. J. Hughes, D. P. Spelke, Z. Xu, C.-C. Kang, D. V. Schaffer and A. E. Herr, Single-cell western blotting, *Nat. Methods*, 2014, **11**(7), 749–755.
- 8 S.-Y. Tang, W. Zhang, R. Soffe, S. Nahavandi, R. Shukla and K. Khoshmanesh, High resolution scanning electron microscopy of cells using dielectrophoresis, *PLoS One*, 2014, **9**(8), e104109.
- 9 N. Dénervaud, J. Becker, R. Delgado-Gonzalo, P. Damay, A. S. Rajkumar, M. Unser, D. Shore, F. Naef and S. J. Maerkl, A chemostat array enables the spatio-temporal analysis of the yeast proteome, *Proc. Natl. Acad. Sci. U. S. A.*, 2013, **110**(39), 15842–15847.
- 10 Y.-Y. Chiang, S. Haeri, C. Gizewski, J. D. Stewart, P. Ehrhard, J. Shrimpton, D. Janasek and J. West, Whole cell quenched flow analysis, *Anal. Chem.*, 2013, **85**(23), 11560–11567.
- 11 M. Blazek, T. S. Santisteban, R. Zengerle and M. Meier, Analysis of fast protein phosphorylation kinetics in single cells on a microfluidic chip, *Lab Chip*, 2015, **15**, 726–734.
- 12 A. van Reenen, A. M. de Jong, J. M. J. den Toonder and M. W. J. Prins, Integrated lab-on-chip biosensing systems based on magnetic particle actuation - a comprehensive review, *Lab Chip*, 2014, **14**, 1966–1986.
- 13 A. L. Washburn and R. C. Bailey, Photonics-on-a-chip: recent advances in integrated waveguides as enabling detection elements for real-world, lab-on-a-chip biosensing applications, *Analyst*, 2011, **136**, 227–236.
- 14 M. C. Estevez, M. Alvarez and L. M. Lechuga, Integrated optical devices for lab-on-a-chip biosensing applications, *Laser Photonics Rev.*, 2012, **6**(4), 463–487.
- 15 D. G. Rackus, M. H. Shamsi and A. R. Wheeler, Electrochemistry, biosensors and microfluidics: a convergence of fields, *Chem. Soc. Rev.*, 2015, **44**, 5320–5340.
- 16 D. King, M. O'Sullivan and J. Ducrée, Optical detection strategies for centrifugal microfluidic platforms, *J. Mod. Opt.*, 2014, **61**(2), 85–101.
- 17 H. Amini, W. Lee and D. Di Carlo, Inertial microfluidic physics, *Lab Chip*, 2014, **14**(15), 2739–2761.
- 18 R. Ramji, M. Wang, A. Asgar, S. Bhagat, D. T. S. Weng, N. V. Thakor, C. T. Lim and C.-H. Chen, Single cell kinase signaling assay using pinched flow coupled droplet microfluidics, *Biomicrofluidics*, 2014, **8**(3), 034104.
- 19 D. H. Yoon, A. Jamshaid, J. Ito, A. Nakahara, D. Tanaka, T. Akitsu, T. Sekiguchi and S. Shoji, Active microdroplet merging by hydrodynamic flow control using a pneumatic actuator-assisted pillar structure, *Lab Chip*, 2014, **14**(16), 3050–3055.
- 20 F. Volpetti, J. Garcia-Cordero and S. J. Maerkl, A microfluidic platform for high-throughput multiplexed protein quantitation, *PLoS One*, 2015, **10**(2), e0117744.
- 21 X. Qin, S. Park, S. P. Duffy, K. Matthews, R. R. Ang, T. Todenhöfer, H. Abdi, A. Azad, J. Bazov, K. N. Chi, P. C. Black and H. Ma, Size and deformability based separation of circulating tumor cells from castrate resistant prostate cancer patients using resettable cell traps, *Lab Chip*, 2015, **15**, 2278–2286.
- 22 P. Preira, V. Grandné, J.-M. Forel, S. Gabriele, M. Camara and O. Theodoly, Passive circulating cell sorting by deformability using a microfluidic gradual filter, *Lab Chip*, 2013, **13**(1), 161–170.
- 23 M. G. Lee, J. H. Shin, C. Y. Bae, S. Choi and J.-K. Park, Label-free cancer cell separation from human whole blood using inertial microfluidics at low shear stress, *Anal. Chem.*, 2013, **85**(13), 6213–6218.
- 24 Y. Ren, W. Liu, Y. Jia, Y. Tao, J. Shao, Y. Ding and H. Jiang, Induced-charge electroosmotic trapping of particles, *Lab Chip*, 2015, **15**, 2181–2191.



- 25 L. Diéguez, M. A. Winter, K. J. Pocock, K. E. Bremmell and B. Thierry, Efficient microfluidic negative enrichment of circulating tumor cells in blood using roughened PDMS, *Analyst*, 2015, **140**(10), 3565–3572.
- 26 S. C. C. Shih, H. Yang, M. J. Jebrail, R. Fobel, N. McIntosh, O. Y. Al-Dirbashi, P. Chakraborty and A. R. Wheeler, Dried blood spot analysis by digital microfluidics coupled to nanoelectrospray ionization mass spectrometry, *Anal. Chem.*, 2012, **84**(8), 3731–3738.
- 27 J. Tiago, S. Fernandes, S. Tenreiro, A. Gameiro, V. Chu, T. F. Outeiro and J. P. Conde, Modulation of alpha-synuclein toxicity in yeast using a novel microfluidic-based gradient generator, *Lab Chip*, 2014, **14**(20), 3949–3957.
- 28 X. Wang, S. Chen, M. Kong, Z. Wang, K. D. Costa, R. A. Li and D. Sun, Enhanced cell sorting and manipulation with combined optical tweezer and microfluidic chip technologies, *Lab Chip*, 2011, **11**(21), 3656–3662.
- 29 O. O. Saeed, R. Li and Y. Deng, Microfluidic approaches for cancer cell separation: Review, *J. Biomed. Sci. Eng.*, 2014, **07**(12), 1005–1018.
- 30 Y. Chen, P. Li, P.-H. Huang, Y. Xie, J. D. Mai, L. Wang, N.-T. Nguyen and T. J. Huang, Rare cell isolation and analysis in microfluidics, *Lab Chip*, 2014, **14**(4), 626–645.
- 31 H. Tsutsui and C.-M. Ho, Cell separation by non-inertial force fields in microfluidic systems, *Mech. Res. Commun.*, 2009, **36**(1), 92–103.
- 32 S. Halldorsson, E. Lucumi, R. Gómez-Sjöberg and R. M. T. Fleming, Advantages and challenges of microfluidic cell culture in polydimethylsiloxane devices, *Biosens. Bioelectron.*, 2015, **63**, 218–231.
- 33 E. K. Sackmann, A. L. Fulton and D. J. Beebe, The present and future role of microfluidics in biomedical research, *Nature*, 2014, **507**(7491), 181–189.
- 34 J. Nilsson, M. Evander, B. Hammarström and T. Laurell, Review of cell and particle trapping in microfluidic systems, *Anal. Chim. Acta*, 2009, **649**(2), 141–157.
- 35 T. Laurell, F. Petersson and A. Nilsson, Chip integrated strategies for acoustic separation and manipulation of cells and particles, *Chem. Soc. Rev.*, 2007, **36**, 492–506.
- 36 F. S. O. Fritzsche, C. Dusny, O. Frick and A. Schmid, Single-cell analysis in biotechnology, systems biology, and biocatalysis, *Annu. Rev. Chem. Biomol. Eng.*, 2012, **3**(1), 129–155. PMID: 22468600.
- 37 M. Niepel, S. L. Spencer and P. K. Sorger, Non-genetic cell-to-cell variability and the consequences for pharmacology, *Curr. Opin. Chem. Biol.*, 2009, **13**(5–6), 556–561.
- 38 A. A. Cohen, N. Geva-Zatorsky, E. Eden, M. Frenkel-Morgenstern, I. Issaeva, A. Sigal, R. Milo, C. Cohen-Saidon, Y. Liron, Z. Kam, L. Cohen, T. Danon, N. Perzov and U. Alon, Dynamic proteomics of individual cancer cells in response to a drug, *Science*, 2008, **322**(5907), 1511–1516.
- 39 J. R. S. Newman, S. Ghaemmaghami, J. Ihmels, D. K. Breslow, M. Noble, J. L. DeRisi and J. S. Weissman, Single-cell proteomic analysis of *S. cerevisiae* reveals the architecture of biological noise, *Nature*, 2006, **441**(7095), 840–846.
- 40 Y. T. Chong, J. L. Y. Koh, H. Frisen, S. K. Duffy, M. J. Cox, A. Moses, J. Moffat, C. Boone and B. J. Andrews, Yeast proteome dynamics from single cell imaging and automated analysis, *Cell*, 2015, **161**(6), 1413–1424.
- 41 H. H. Chang, M. Hemberg, M. Barahona, D. E. Ingber and S. Huang, Transcriptome-wide noise controls lineage choice in mammalian progenitor cells, *Nature*, 2008, **453**(7194), 544–547.
- 42 M. Junkin and S. Tay, Microfluidic single-cell analysis for systems immunology, *Lab Chip*, 2014, **14**, 1246–1260.
- 43 A. Rakszewska, J. Tel, V. Chokkalingam and W. T. S. Huck, One drop at a time: toward droplet microfluidics as a versatile tool for single-cell analysis, *NPG Asia Mater*, 2014, **6**, e133.
- 44 H. Yin and D. Marshall, Microfluidics for single cell analysis, *Curr. Opin. Biotechnol.*, 2012, **23**(1), 110–119. Analytical biotechnology.
- 45 T. A. Duncombe, A. M. Tentori and A. E. Herr, Microfluidics: reframing biological enquiry, *Nat. Rev. Mol. Cell Biol.*, 2015, **16**(9), 554–567.
- 46 P. K. Chattopadhyay, T. M. Gierahn, M. Roederer and J. Christopher Love, Single-cell technologies for monitoring immune systems, *Nat. Immunol.*, 2014, **15**(2), 128–135.
- 47 T. P. Lagus and J. F. Edd, A review of the theory, methods and recent applications of high-throughput single-cell droplet microfluidics, *J. Phys. D: Appl. Phys.*, 2013, **46**(11), 114005.
- 48 C. Gawad, W. Koh and S. R. Quake, Single-cell genome sequencing: current state of the science, *Nat. Rev. Genet.*, 2016, DOI: 10.1038/nrg.2015.16.
- 49 X. J. Li, Y. Chen and P. C. H. Li, A simple and fast microfluidic approach of same-single-cell analysis (sasca) for the study of multidrug resistance modulation in cancer cells, *Lab Chip*, 2011, **11**, 1378–1384.
- 50 N. M. Karabacak, P. S. Spuhler, F. Fachin, E. J. Lim, V. Pai, E. Ozkumur, J. M. Martel, N. Kojic, K. Smith, P.-i Chen, J. Yang, H. Hwang, B. Morgan, J. Trautwein, T. A. Barber, S. L. Stott, S. Maheswaran, R. Kapur, D. A. Haber and M. Toner, Microfluidic, marker-free isolation of circulating tumor cells from blood samples, *Nat. Protocols*, 2014, **9**(3), 694–710.
- 51 J.-P. Frimat, M. Becker, Y.-Y. Chiang, U. Marggraf, D. Janasek, J. G. Hengstler, J. Franzke and J. West, A microfluidic array with cellular valving for single cell co-culture, *Lab Chip*, 2011, **11**, 231–237.
- 52 L. Mazutis, J. Gilbert, W. Lloyd Ung, D. A. Weitz, A. D. Griffiths and J. A. Heyman, Single-cell analysis and sorting using droplet-based microfluidics, *Nat. Protocols*, 2013, **8**(5), 870–891.
- 53 See <https://www.fluidigm.com/products/c1-system> (consulted on the 1st february 2016).
- 54 P. E. Stenberg, R. P. McEver, M. A. Shuman, Y. V. Jacques and D. F. Bainton, A platelet alpha-granule membrane



- protein (GMP-140) is expressed on the plasma membrane after activation, *J. Cell Biol.*, 1985, **101**(3), 880–886.
- 55 E. D. Klootwijk, M. Reichold, A. Helip-Wooley, A. Tolaymat, C. Broeker, S. L. Robinette, J. Reinders, D. Peindl, K. Renner, K. Eberhart, N. Assmann, P. J. Oefner, K. Dettmer, C. Sterner, J. Schroeder, N. Zorger, R. Witzgall, S. W. Reinhold, H. C. Stanescu, D. Bockenbauer, G. Jaureguierry, H. Courtneidge, A. M. Hall, A. D. Wijeyesekera, E. Holmes, J. K. Nicholson, K. O'Brien, I. Bernardini, D. M. Krasnewich, M. Arcos-Burgos, Y. Izumi, H. Nonoguchi, Y. Jia, J. K. Reddy, M. Ilyas, R. J. Unwin, W. A. Gahl, R. Warth and R. Kleta, Mistargeting of peroxisomal EHHADH and inherited renal fanconi's syndrome, *N. Engl. J. Med.*, 2014, **370**(2), 129–138. PMID: 24401050.
 - 56 A. Sickmann, J. Reinders, Y. Wagner, C. Joppich, R. Zahedi, H. E. Meyer, B. Schönfisch, I. Perschil, A. Chacinska, B. Guiard, P. Rehling, N. Pfanner and C. Meisinger, The proteome of *Saccharomyces cerevisiae* mitochondria, *Proc. Natl. Acad. Sci. U. S. A.*, 2003, **100**(23), 13207–13212.
 - 57 S. W. Taylor, E. Fahy, B. Zhang, G. M. Glenn, D. E. Warnock, S. Wiley, A. N. Murphy, S. P. Gaucher, R. A. Capaldi, B. W. Gibson and S. S. Ghosh, Characterization of the human heart mitochondrial proteome, *Nat. Biotechnol.*, 2003, **21**(3), 281–286.
 - 58 J. R. Yates III, A. Gilchrist, K. E. Howell and J. J. M. Bergeron, Proteomics of organelles and large cellular structures, *Nat. Rev. Mol. Cell Biol.*, 2005, **6**(9), 702–714.
 - 59 M. Niemann, S. Wiese, J. Mani, A. Chanfon, C. Jackson, C. Meisinger, B. Warscheid and A. Schneider, Mitochondrial outer membrane proteome of *Trypanosoma brucei* reveals novel factors required to maintain mitochondrial morphology, *Mol. Cell. Proteomics*, 2012, **12**(2), 515–528.
 - 60 R. P. Zahedi, A. Sickmann, A. M. Boehm, C. Winkler, N. Zufall, B. Schönfisch, B. Guiard, N. Pfanner and C. Meisinger, Proteomic analysis of the yeast mitochondrial outer membrane reveals accumulation of a subclass of preproteins, *Mol. Biol. Cell*, 2006, **17**(3), 1436–1450.
 - 61 F.-N. Vögtle, J. M. Burkhart, S. Rao, C. Gerbeth, J. Hinrichs, J.-C. Martinou, A. Chacinska, A. Sickmann, R. P. Zahedi and C. Meisinger, Intermembrane space proteome of yeast mitochondria, *Mol. Cell. Proteomics*, 2012, **11**(12), 1840–1852.
 - 62 H. Lu, S. Gaudet, M. A. Schmidt and K. F. Jensen, A microfabricated device for subcellular organelle sorting, *Anal. Chem.*, 2004, **76**(19), 5705–5712. PMID: 15456289.
 - 63 M. Moschallski, M. Hausmann, A. Posch, A. Paulus, N. Kunz, T. T. Duong, B. Angres, K. Fuchsberger, H. Steuer, D. Stoll, S. Werner, B. Hagmeyer and M. Stelzle, MicroPrep: Chip-based dielectrophoretic purification of mitochondria, *Electrophoresis*, 2010, **31**(15), 2655–2663.
 - 64 K. Sharma, R. C. J. D'Souza, S. Tyanova, C. Schaab, J. R. Wiśniewski, J. Cox and M. Mann, Ultra-deep human phosphoproteome reveals a distinct regulatory nature of Tyr and Ser/Thr-based signaling, *Cell Rep.*, 2014, **8**(5), 1583–1594.
 - 65 Y. Zheng, C. Zhang, D. R. Croucher, M. A. Soliman, N. St-Denis, A. Pasculescu, L. Taylor, S. A. Tate, W. Rod Hardy, K. Colwill, A. Y. Dai, R. Bagshaw, J. W. Dennis, A.-C. Gingras, R. J. Daly and T. Pawson, Temporal regulation of EGF signalling networks by the scaffold protein Shc1, *Nature*, 2013, **499**(7457), 166–171.
 - 66 F. Beck, J. Geiger, S. Gambaryan, J. Veit, M. Vaudel, P. Nollau, O. Kohlbacher, L. Martens, U. Walter, A. Sickmann and R. P. Zahedi, Time-resolved characterization of cAMP/PKA-dependent signaling reveals that platelet inhibition is a concerted process involving multiple signaling pathways, *Blood*, 2013, **123**(5), e1–e10.
 - 67 P. Novo, F. Volpetti, V. Chu and J. P. Conde, Control of sequential fluid delivery in a fully autonomous capillary microfluidic device, *Lab Chip*, 2013, **13**, 641–645.
 - 68 P. Novo, V. Chu and J. P. Conde, Integrated optical detection of autonomous capillary microfluidic immunoassays: a hand-held point-of-care prototype, *Biosens. Bioelectron.*, 2014, **57**, 284–291.
 - 69 E. Tavares da Costa, M. F. Mora, P. A. Willis, C. L. do Lago, H. Jiao and C. D. Garcia, Getting started with open-hardware: Development and control of microfluidic devices, *Electrophoresis*, 2014, **35**(16), 2370–2377.
 - 70 J. McGrath, M. Jimenez and H. Bridle, Deterministic lateral displacement for particle separation: a review, *Lab Chip*, 2014, **14**(21), 4139–4158.
 - 71 L. R. Huang, E. C. Cox, R. H. Austin and J. C. Sturm, Continuous particle separation through deterministic lateral displacement, *Science*, 2004, **304**(5673), 987–990.
 - 72 D. Holmes, G. Whyte, J. Bailey, N. Vergara-Irigaray, A. Ekpenyong, J. Guck and T. Duke, Separation of blood cells with differing deformability using deterministic lateral displacement, *Interface Focus*, 2014, **4**(6), 20140011–20140011.
 - 73 S. Varma and J. Voldman, A cell-based sensor of fluid shear stress for microfluidics, *Lab Chip*, 2015, **15**(6), 1563–1573.
 - 74 S. M. Ponik, J. W. Triplett and F. M. Pavalko, Osteoblasts and osteocytes respond differently to oscillatory and unidirectional fluid flow profiles, *J. Cell. Biochem.*, 2007, **100**, 794–807.
 - 75 S. Weinbaum, S. C. Cowin and Y. Zeng, A model for the excitation of osteocytes by mechanical loading-induced bone fluid shear stresses, *J. Biomech.*, 1994, **27**, 339–360.
 - 76 H. A. Pohl and J. S. Crane, Dielectrophoresis of cells, *Biophys. J.*, 1971, **11**, 711–727.
 - 77 D. Das, K. Biswas and S. Das, A microfluidic device for continuous manipulation of biological cells using dielectrophoresis, *Med. Eng. Phys.*, 2014, **36**(6), 726–731.
 - 78 P. Cheng, M. J. Barrett, P. M. Oliver, D. Cetin and D. Vezenov, Dielectrophoretic tweezers as a platform for molecular force spectroscopy in a highly parallel format, *Lab Chip*, 2011, **11**(24), 4248–4259.



- 79 A. Salmanzadeh, L. Romero, H. Shafiee, R. C. Gallo-Villanueva, M. A. Stremmer, S. D. Cramer and R. V. Davalos, Isolation of prostate tumor initiating cells (TICs) through their dielectrophoretic signature, *Lab Chip*, 2012, **12**, 182–189.
- 80 X. Ding, Z. Peng, S.-C. S. Lin, M. Geri, S. Li, P. Li, Y. Chen, M. Dao, S. Suresh and T. J. Huang, Cell separation using tilted-angle standing surface acoustic waves, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, **111**(36), 12992–12997.
- 81 J. Nam, H. Lim and S. Shin, Manipulation of microparticles using surface acoustic wave in microfluidic systems: a brief review, *Korea-Aust. Rheol. J.*, 2011, **23**(4), 255–267.
- 82 H. Bruus, Acoustofluidics 10: Scaling laws in acoustophoresis, *Lab Chip*, 2012, **12**(9), 1578–1586.
- 83 M. A. Burguillos, C. Magnusson, M. Nordin, A. Lenshof, P. Augustsson, M. J. Hansson, E. Elmér, H. Lilja, P. Brundin, T. Laurell and T. Deierborg, Microchannel acoustophoresis does not impact survival or function of microglia, leukocytes or tumor cells, *PLoS One*, 2013, **8**(5), e64233.
- 84 P. Li, Z. Mao, Z. Peng, L. Zhou, Y. Chen, P.-H. Huang, C. I. Truica, J. J. Drabick, W. S. El-Deiry, M. Dao, S. Suresh and T. J. Huang, Acoustic separation of circulating tumor cells, *Proc. Natl. Acad. Sci. U. S. A.*, 2015, **112**(16), 4970–4975.
- 85 F. D. Giudice, H. Madadi, M. M. Villone, G. D'Avino, A. M. Cusano, R. Vecchione, M. Ventre, P. L. Maffettone and P. A. Netti, Magnetophoresis 'meets' viscoelasticity: deterministic separation of magnetic particles in a modular microfluidic device, *Lab Chip*, 2015, **15**, 1912–1922.
- 86 T. P. Hunt, D. Issadore and R. M. Westervelt, Integrated circuit/microfluidic chip to programmably trap and move cells and droplets with dielectrophoresis, *Lab Chip*, 2008, **8**(1), 81–87.
- 87 D. W. Inglis, R. Riehn, R. H. Austin and J. C. Sturm, Continuous microfluidic immunomagnetic cell separation, *Appl. Phys. Lett.*, 2004, **85**(21), 5093–5095.
- 88 D. J. Collins, T. Alan and A. Neild, Particle separation using virtual deterministic lateral displacement (vDLD), *Lab Chip*, 2014, **14**(9), 1595.
- 89 J. P. Beech, P. Jonsson and J. O. Tegenfeldt, Tipping the balance of deterministic lateral displacement devices using dielectrophoresis, *Lab Chip*, 2009, **9**, 2698–2706.
- 90 H. Shafiee, J. L. Caldwell and R. V. Davalos, Selective isolation of live/dead cells using contactless dielectrophoresis (cDEP), *Lab Chip*, 2010, **10**, 438–445.
- 91 J. D. Adams, P. Thévoz, H. Bruus and H. Tom Soh, Integrated acoustic and magnetic separation in microfluidic channels, *Appl. Phys. Lett.*, 2009, **95**(25), 254103.
- 92 See Fig. 1 of "chapter 4: Rates and duration" from "cell biology by the numbers" by ron milo and rob phillips, also available online on <http://book.bionumbers.org/rates-and-duration-introduction/> (consulted on the 3rd of december 2015).
- 93 M.-H. Wu, S.-B. Huang and G.-B. Lee, Microfluidic cell culture systems for drug research, *Lab Chip*, 2010, **10**, 939–956.
- 94 I. Meyvantsson and D. J. Beebe, Cell culture models in microfluidic systems, *Annu. Rev. Anal. Chem.*, 2008, **1**(1), 423–449. PMID: 20636085.
- 95 L. Chingozha, M. Zhan, C. Zhu and H. Lu, A generalizable, tunable microfluidic platform for delivering fast temporally varying chemical signals to probe single-cell response dynamics, *Anal. Chem.*, 2014, **86**(20), 10138–10147. PMID: 25254360.
- 96 A. H. C. Ng, M. Dean Chamberlain, H. Situ, V. Lee and A. R. Wheeler, Digital microfluidic immunocytochemistry in single cells, *Nat. Commun.*, 2015, **6**, 1–12.
- 97 R. B. Fair, Digital microfluidics: is a true lab-on-a-chip possible?, *Microfluid. Nanofluid.*, 2007, **3**(3), 245–281.
- 98 A. M. Hirsch, C. A. Rivet, B. Zhang, M. L. Kemp and H. Lu, Parallel multi-time point cell stimulation and lysis on-chip for studying early signaling events in T cell activation, *Lab Chip*, 2009, **9**, 536–544.
- 99 F. A. Solari, M. Dell'Aica, A. Sickmann and R. P. Zahedi, Why phosphoproteomics is still a challenge, *Mol. Biosyst.*, 2015, **11**, 1487–1493.
- 100 N.-T. Nguyen and Z. Wu, Micromixers - a review, *J. Micro-mech. Microeng.*, 2005, **15**(2), R1–R16.
- 101 B. D. Plouffe and S. K. Murthy, Perspective on microfluidic cell separation: A solved problem?, *Anal. Chem.*, 2014, **86**(23), 11481–11488.
- 102 G. M. Whitesides, Cool, or simple and cheap? Why not both?, *Lab Chip*, 2013, **13**, 11–13.

