Contents lists available at ScienceDirect

Journal of Chromatography A



journal homepage: www.elsevier.com/locate/chroma

Screen-printed digital microfluidics combined with surface acoustic wave nebulization for hydrogen-deuterium exchange measurements



Lucas Monkkonen^a, J. Scott Edgar^a, Daniel Winters^b, Scott R. Heron^{c,d}, C. Logan Mackay^e, Christophe D. Masselon^{f,g,h}, Adam A. Stokes^e, Patrick R.R. Langridge-Smith^e, David R. Goodlett^{c,d,*}

^a University of Washington, Seattle, WA, USA

^b Trioptics GmbH, Wedel, Germany

^c Deurion, LLC, Seattle, WA, USA

^d University of Maryland, Baltimore, MD, USA

^e University of Edinburgh, Edinburgh, UK

^f University Grenoble-Alpes, Grenoble, France

^g CEA, iRTSV, Biologie à Grande Echelle, Grenoble, France

^h INSERM, U1038 Grenoble, France

ARTICLE INFO

Article history: Received 2 July 2015 Received in revised form 16 November 2015 Accepted 17 December 2015 Available online 21 December 2015

Keywords:

Screen-printed digital microfluidics Surface acoustic wave nebulization Hydrogen-deuterium exchange Mass spectrometry

ABSTRACT

An inexpensive digital microfluidic (DMF) chip was fabricated by screen-printing electrodes on a sheet of polyimide. This device was manually integrated with surface acoustic wave nebulization (SAWN) MS to conduct hydrogen/deuterium exchange (HDX) of peptides. The HDX experiment was performed by DMF mixing of one aqueous droplet of angiotensin II with a second containing various concentrations of D₂O. Subsequently, the degree of HDX was measured immediately by SAWN-MS. As expected for a small peptide, the isotopically resolved mass spectrum for angiotensin revealed that maximum deuterium exchange was achieved using 50% D₂O. Additionally, using SAWN-MS alone, the global HDX kinetics of ubiquitin were found to be similar to published NMR data and back exchange rates for the uncooled apparatus using high inlet capillary temperatures was less than 6%.

© 2016 Published by Elsevier B.V.

1. Introduction

Hydrogen deuterium exchange (HDX) is a powerful technique for studying protein structure [1]. As evidenced by the rapidly growing body of HDX literature, which shows steady growth in the technique. For the most part, HDX workflows have deviated little from HPLC or a direct infusion apparatus coupled to electrospray ionization (ESI) MS. In this study, we investigate the use of two fundamental changes to the HDX workflow. First, we examine the use of an alternative ionization method called surface acoustic wave nebulization (SAWN) and second, we explore the use of microfluidics as an alternative to manual or an automated LC-type unit for sample preparation.

While ESI has been critical to HDX success and many other MS based assays [2,3], it is not without limitations. For instance,

dgoodlett@rx.umaryland.edu (D.R. Goodlett).

http://dx.doi.org/10.1016/j.chroma.2015.12.048 0021-9673/© 2016 Published by Elsevier B.V. while ESI is very sensitive, it can lead to in-source fragmentation and/or the oxidation of small molecules and proteins [4–6], and ESI requires its own charged, continuous flow apparatus. To address the analytical short-comings of ESI, many alternative ionization techniques have been developed, such as desorption ionization on silicon [7], desorption-ESI [8] and Laser Ablation-ESI [9].

In this study, we employed surface acoustic wave nebulization (SAWN) for HDX analysis. SAWN generates ions from a planar piezoelectric surface and delivers them to the inlet of a mass spectrometer [10]. To accomplish this, an alternating current is applied to interdigitated transducers (IDT, interlocking electrodes) on a piezoelectric LiNbO₃ wafer to generate a high frequency surface acoustic wave [11]. When the SAW reaches an area on the surface of the chip where an aqueous droplet of sample is located reflection of the wave within the droplet results in nebulization of the liquid sample within seconds. To date, a range of analytes of small molecules have been analyzed by SAWN-MS [5,6,10].

SAWN has several advantages over ESI and other ionization techniques for HDX. First, SAWN has been found to generate ions of lower energy than ESI [6], which has the potential to maintain



^{*} Corresponding author at: University of Maryland, Baltimore, MD, USA. *E-mail addresses*: lucasm7@uw.edu (L. Monkkonen),

the structural integrity of more analytes during the ionization process [5]. Additionally, given its relative "softness" compared to ESI, there is the potential advantage of reduced back exchange during HDX, which we test here. Second, in contrast to ESI, SAWN is very simple to operate with no possibility of clogging since it is a planar substrate. Specifically, SAWN operation only involves transferring a droplet of sample directly onto the chip and activation of the chip, which leads to the immediate nebulization of the sample. Here we validated the use of HDX via SAWN-MS using ubiquitin, a well-characterized protein [12–14].

In addition to the ionization source, we explored the use of a sophisticated fluid handling platform called digital microfluidics (DMF) for sample manipulation within the HDX workflow. The concept of DMF is representative of a class of techniques using the relatively weak interactions between electric fields and polarizable droplets of liquid on a planar surface. A typical DMF device consists of a flat substrate covered in a microelectrode pattern which is covered by subsequent layers of a suitable dielectric and a hydrophobic layer [15]. This technology has been referred to by many names during its development including; metal-insulator-solution-transport [16], digital microfluidic system [17], electrowetting on insulator coated electrodes [18], electrowetting on dielectric [19] and DMF [20], which we prefer.

Due to the promise of automated sample handling, small sample volumes, loss-less sample preparation, and miniaturized devices that can forgo large and costly LC systems, much effort has been placed into connecting microfluidics with ESI-MS as seen in the publication data presented in these review articles [21,22]. However, coupling ESI-MS to microfluidic systems remains a challenge due to the requirement of maintaining constant fluid flow for steady electrospray [23]. The most popular means for coupling microfluidics with electrospray is simply attaching a micro-electrospray emitter [21], such as the microfluidic bottom-up HDX device manufactured by Rob et al. [24]. This microfluidic device has successfully characterized HDX on very short timescales with low back exchange [25,26]. However, this device has several potential disadvantages such as time-consuming fabrication (due to attachment of a sprayer and incorporation of microchannels) and susceptibility to clogs (especially when using native buffer). Microfluidics has also been coupled with matrix assisted laser desorption ionization (MALDI) MS to circumvent these problems, but MALDI is susceptible to matrix effects [27]. Alternatively, integrating SAWN and DMF is simple and conserves sample since both are planar platforms which manipulate discrete droplets as low as half a microliter [20]. Other studies utilizing a combined DMF and SAWN approach for non-MS applications further supports the simplicity and minimal sample use of these two techniques [28–36].

Here, we demonstrate the ease of use of DMF-SAWN to perform HDX on an inexpensive DMF device. Briefly, the DMF device was made by screen-printing conductive ink on top of a flexible polyimide substrate and coating the device with hydrophobic materials. The motivation for utilizing an inexpensive DMF design was to minimize biofouling by making the device disposable. We report results obtained from fusing droplets of D₂O and angiotensin II on such a disposable DMF device and analyzing the sample immediately by SAWN-MS.

The significance of this report is three-fold. Firstly, it represents the first use of SAWN to ionize HDX samples for MS analysis. Secondly, we demonstrated that SAWN-MS can yield high-resolution, reproducible data for whole proteins with low back exchange. Thirdly, SAWN was successfully coupled with a disposable, screenprinted DMF device to provide a means for sample preparation at the mass spectrometer. This combination points the way toward monitoring more complex chemical reactions in real time by DMF-SAWN-MS with many different types of analytes for a new, sample conserving "lab on a chip" or micro total analysis system [22,37].

2. Experimental

2.1. Fabrication and operation of SAWN chips

Fabrication of SAWN chips has been reported in detail previously [6,10]. In summary, a SAW transducer consisting of 20 pairs of 100 µm interdigitated (IDT) electrodes (40 in total) with 100 µm spacing and 10 mm aperture, along with a secondary electrode to apply external electrical potential, were patterned onto the surface of 128 Y-cut X-propagating 3 in LiNbO₃ wafers purchased from Crystal Technology, Inc. (Palo Alto, Ca). The SAWN configuration was first designed in AutoCAD before being written into a chrome mask by a Heidelberg µPG 101 Laser Pattern Generator (Heidelberg Instruments Mikrotechnik GmbH Tullastrasse 2, D-69126Heidelberg, Germany) at the University of Washington Nanotech User Facility (https://depts.washington.edu/ntuf/). Wafers were then coated using AZ 1512 positive photoresist (AZ Electronic Materials, Somerville, NI) spun at 4000 rpm for 30 secondscreating a 1–1.2 µm thick resist layer. Exposure of the photo resist was done for 5 secondsusing an Oriel mask aligner (Newport Corporation, CA). The exposed wafers were placed in a development bath for 60 secondsin AZ 351 (AZ Electronic Materials, Somerville, NJ). To pattern conductive IDT electrodes a 20 nm chrome adhesion layer was deposited by heated vapor deposition followed by a 60 nm layer of gold, followed by lift-off in acetone for 30 min. The resulting SAWN IDT has a resonance frequency of 9.56 MHz. To operate the SAWN chip a MXG analog signal generator (Agilent N5181A, Santa Clara, CA) and a Mini Circuits ZHL-5W-1, 5-500 MHz RF amplifier (GwInstek GPS-2303, New York, NY) were used to generate and amplify the RF signal.

2.2. Fabrication of screen-printed DMF chips

Flexible, disposable, DMF chips were produced in a proprietary low-cost printing process using carbon-containing conductive ink on 50 μ m thick flexible polyimide foils (DuPont Kapton) to pattern electrodes with 100 μ m spacing, connection leads and external contact pads, followed by a 7 μ m layer of ink acting as dielectric layer. After the printing process, a fluoropolymer solution, either Teflon-AF (DuPont) or Cytop (Asahi Glass) solution was spin-coated onto the devices and left to dry in ambient condition to create a 150 nm thick hydrophobic layer. The finished devices could be stored in normal laboratory conditions over the time period of several months without influence on performance.

2.3. Operation of screen-printed DMF chips

A schematic showing the operation of the DMF is shown in Fig. 5. Droplets were moved, merged and transferred to the SAWN chip in AC mode [28,29,37,38] with a driving voltage of 500 Vpp and a frequency of 20 kHz. The devices performed flawlessly and without any noticeable degradation over several experimental cycles. The electrode geometry used was an arrangement of two parallel columns of square electrodes. The droplet was held between two adjacent electrodes from the respective columns by applying the driving voltage between them. To move the droplet to the next electrode pair, the driving voltage was first applied to the new pair, pulling the droplet in between the pairs, and then switching off the voltage between the first pair. DMF operation was controlled using control software running on a PC that was connected to custom drive electronics. For this, the 2.5 Vpp AC sine output of a 20 kHz signal generator was amplified by a high-voltage amplifier (Trek) to 500 Vpp and distributed to the DMF electrode pads using a distribution circuit based on solid-state relays. Connection to the DMF devices was through 2.54 mm pitch edge connectors.

2.4. HDX of ubiquitin

In an eppendorf tube, 1 µl of 200 µM ubiquitin, 40 mM ammonium bicarbonate (pH 7.4) was combined with 19μ l of D₂O for a final concentration of 10 µM ubiquitin, 2 mM ammonium bicarbonate. Exposure to deuterium was varied so as to obtain many time points to properly visualize the rate of deuterium uptake in ubiguitin over time. To quench the reaction, 1.5 µl of 10% formic acid was added to the mixture. Three spectra were obtained of three replicates of the same time point. To obtain a mass spectrum of the mixture, 1 µl of the quenched reaction mixture was placed on the SAWN chip surface in front of the interdigitated transducer. An ultrazoom scan of this droplet was recorded with a Thermo LTQ-Velos using the s-lens at 60% voltage and a capillary temperature of 200 °C after activating the SAWN chip with 25 dBm at 9.56 MHz. To calculate the overall percent deuteration, the percent deuteration of the +6, +7, and +8 ion was calculated as per the equation 1 reported in Zhang and Smith [39] with the centroided mass obtained using HX Express [40,41], and then these values were averaged for the final percent deuteration value. Estimated percent deuteration was achieved by using the method described in Pan et al. [13] and the rate constants from Bougalt et al. [12] adjusted for pH differences.

2.5. Measuring back exchange of ubiquitin

As before, 1 μ l of stock solution was pipetted into 19 μ l of D₂O. The reaction mixtures was boiled for 10 min and then dried by speed vac. Then, 19 μ l of D₂O and 1.5 μ l of 10% formic acid was added to the dried sample, mixed briefly, and then pipetted onto the SAWN chip for analysis by SAWN-MS. Back exchange was estimated according to the following formula:

$$\frac{MW_{measured} - MW_{undeuterated} - X_{fast} - X_{inexchangeable}}{X_{backbone}}$$

 $MW_{measured}$ is the centroid measured molecular weight (in Da), $MW_{undeuterated}$ is the average mass of ubiquitin (8564.8449 Da), X_{fast} refers to the number of side-chain exchanges in ubiquitin (72), $X_{inexchangeable}$ refers to the number of amides that exchange only extremely slowly in native conditions (21) [14], and $X_{backbone}$ is the total number of backbone exchanges possible in ubiquitin (73).

2.6. HDX mass spectrometry of angiotensin

 $4 \,\mu$ l droplet of angiotensin II (10 μ M, Proteomass) in H₂O and a $4 \,\mu$ l droplet of D₂O were pipetted onto separate regions of the flexible DMF chip. Then, an AC field to the electrodes adjacent to the droplets was applied to transport and fuse the two droplets. The fused droplet from the DMF chip was manually transferred onto the SAWN chip by touching the droplet on the hydrophobic DMF chip to hydrophilic SAWN chip. After transfer, SAWN was activated to aerosolize the droplet and HDX was monitored by a Thermo LTQ linear ion trap.

3. Results and discussion

3.1. HDX of ubiquitin

To demonstrate the utility of SAWN for HDX analysis of proteins, global HDX studies were carried out on ubiquitin, a well characterized protein. High-resolution mass spectra showing the isotopic envelope of deuterated ubiquitin were readily obtained by



Fig. 1. SAWN-MS monitored HDX of ubiquitin. The zero minute mass spectrum is obtained from ubiquitin exposed to quench solution. The five minute mass spectrum was obtained from ubiquitin exposed to D_2O for five minutes. The totally deuterated spectrum was obtained by boiling in D_2O for 10 min. All spectra were collected by an LTQ-Velos in ultrazoom scan mode.



Fig. 2. Time-course of deuterium uptake in ubiquitin. Time refers to the incubation time of ubiquitin in 95% D_2O at room temperature. The blue squares represent the experimentally derived percent deuteration. This was obtained by calculating the percent deuteration for each of the +6, +7, and +8 with the aid of equation #1 [39] and HX-express [40,41] and averaging these values. Error bars represent the standard deviation for the average of the three replicates of the percent deuteration levels found with each of the three charge states. The red triangles represent the estimated global percent deuteration based on combining NMR kinetic data of ubiquitin [12] with the equation by Pan et al. [13].

SAWN-MS on an ion trap mass spectrometer, and the mass spectra clearly demonstrate a shift of the ubiquitin ions toward higher m/z values with increased exposure to D₂O (Fig. 1). Due to these promising results using SAWN for HDX, we set out to record a full time course of the HDX reaction by SAWN to show that SAWN-HDX can be used to rigorously measure global HDX rates. These results are displayed in Fig. 2. Each data point represents triplicate measurements that highlight the reproducibility of this technique. Additionally, the experimental per-deuterated time-point of 94.6% is very close to the maximum possible 95% deuteration of ubiquitin under these conditions. Most importantly, however, the data from SAWN-MS correlates well with global estimated percent deuteration values based on HDX-NMR data obtained by Bougault, especially for the later time points [12] (Fig. 2). An additional point of validation is that the results of Pan et al. [13] compare well to the SAWN-MS data. Pan et al. [13] measured a global percent deuter-



Fig. 3. Fabrication of digital microfluidic (DMF) chips. DMF chips were fabricated by screening printing conductive ink onto 50 μ m thick flexible polyimide foils.

ation of 83% at 30 min of incubation for ubiquitin under similar conditions. When a simple natural log regression is fitted to the SAWN-MS data (r^2 value 0.9957), the estimated percent deuteration measured by SAWN-MS with 30 min incubation would be 80.4%, which agrees well with Pan et al. [13]. Finally, the back exchange of the SAWN-HDX setup was measured for HDX carried out by hand at room temperatures at three different inlet temperature values. At capillary temperatures of 100, 200, and 300 °C, the average exchange rates of three triplicates were 1.8%, 4.6%, and 6.0%, respectively. These measured back exchange rates for SAWN-HDX were either lower or close to the value reported by Rob et al. [26], depending on the inlet capillary temperature.

3.2. DMF chip fabrication

Though SAWN chips are simple to operate they remain rudimentary in their ability to accommodate fluid handling or chromatography. To address this issue we developed a technique for fabrication of low-cost screen-printed DMF chips on polyimide that can easily be coupled with SAWN and may be considered disposable (Fig. 3). These inexpensive DMF chips were prepared by screen-printing [42,43] on to flexible polyimide substrates. Specifically, screen-printed electrodes were developed using a carbon-containing conductive ink on top of which was printed a dielectric with a layer of hydrophobic Cytop on top of these first two layers.



Fig. 4. DMF Experimental Workflow. The basic HDX workflow is shown in (A). Droplets were pipetted onto the surface of the DMF chip (B) where they were moved by applying an AC field to adjacent electrodes until adjacent droplets combine (C). After being combined the droplet was transferred from the hydrophobic DMF chip to the natively hydrophilic lithium niobate SAWN wafer (D, E) by wicking action.

3.3. HDX via DMF-SAWN-MS

HDX was carried out on the signal peptide angiotensin II to demonstrate the functionality of DMF-SAWN-HDX. The movement of droplets on the DMF chip and transfer to SAWN is shown in Fig. 4. The design of the DMF chip used a series of electrodes spaced $100 \,\mu\text{m}$ apart (Fig. 4(A)). Samples were pipetted onto the surface of the DMF chip across from an equal volume of D2O as shown in Fig. 4(B). Once both droplets were in place on the DMF chip, the sample droplet consisting of a $4 \mu l$ droplet of angiotensin II in H2O was actuated toward the $4 \mu l$ droplet of D2O by applying a field on the adjacent electrode while simultaneously releasing the field on the initial electrode. In this way, one can actuate droplets back and forth across the DMF chip and combine droplets to initiate HDX as shown in Fig. 4(C). Transfer of the post-HDX reaction sample to the SAWN chip was performed by placing the droplet in contact with the surface of the SAWN wafer, which allowed the droplet to wick easily over to the natively hydrophilic lithium niobate SAWN chip from the more hydrophobic coated Cytop DMF chip (Fig. 4(D,E)).

The mass spectra of angiotensin II before (Fig. 5(A)) and after a one minute incubation (Fig. 5(B)) to allow HDX to occur are shown in Fig. 5. In addition to the expected $[M+H]^+$ ion of angiotensin II, the presence of a small amount of sodiated angiotensin II, $[M + Na]^+$, is observed. The post-HDX mass spectrum is shown in Fig. 5(B) which shows that eight deuteriums were found to be present on



Fig. 5. HDX of angiotensin II measured by DMF-SAWN-MS. An angiotensin II spectrum is shown prior to HDX (A) and one minute post-mixing (B) by DMF manipulation of a droplet of angiotensin in water with a droplet of D₂O. One minute of exposure of angiotensin II in 50% D₂O resulted in 8 deuterium exchanges.

angiotensin II, which is the maximum possible deuteration for angiotensin II in 50% D₂O. These results are expected for a small peptide like angiotensin II (1.046 KDa), since all the sites are highly solvent accessible.

4. Concluding remarks

In summary, we have demonstrated the use of SAWN-MS for monitoring HDX reactions of small proteins. Additionally, and importantly for future automation of the method, we have shown it is feasible to combine a DMF sample preparation chip with the SAWN chip to carry out HDX monitoring by MS. The DMF chips were produced by screen-printing onto a flexible, polyimide substrate making the chip effectively disposable. These HDX methods using SAWN-MS and DMF-SAWN-MS offer a new, facile means of elucidating protein structure. Given the simplicity of these methods, we envision them being used to carry out pilot studies rapidly at the MS interface.

Acknowledgements

Lucas Monkkonen thanks the NSF graduate research fellowship grant DGE-0718124 for support. Christophe D. Masselon acknowledges financial support from a CEA–Eurotalent outgoing fellowship (grant PCOFUND-GA-2008-228664). David R. Goodlett and Scott R. Heron thank the University of Maryland School of Pharmacy Mass Spectrometry Center (SOP1841-IQB2014), which in part supported this work. Daniel Winters, Adam A. Stokes, C. Logan Mackay, and Patrick R. R. Langrige-Smith, the United Kingdom's EPSRC for funding from the Radical Solutions for Researching the Proteome (RASOR) consortium (Grant from the Biotechnology and Biological Sciences Research Council (BB/C511599/1)).

References

- J.R. Engen, Analysis of protein conformation and dynamics by hydrogen/deuterium exchange MS, Anal. Chem. 81 (2009) 7870–7875, http:// dx.doi.org/10.1021/ac901154s.
- [2] J. Fenn, M. Mann, C. Meng, S. Wong, C. Whitehouse, Electrospray ionization for mass spectrometry of large biomolecules, Science 80 (246) (1989) 64–71, http://dx.doi.org/10.1126/science.2675315.
- [3] C.M. Whitehouse, R.N. Dreyer, M. Yamashita, J.B. Fenn, Electrospray interface for liquid chromatographs and mass spectrometers, Anal. Chem. 57 (1985) 675–679.
- [4] B.L. Boys, M.C. Kuprowski, J.J. Noël, L. Konermann, Protein oxidative modifications during electrospray ionization: solution phase electrochemistry or corona discharge-induced radical attack? Anal. Chem. 81 (2009) 4027–4034, http://dx.doi.org/10.1021/ac900243p.
- [5] S.H. Yoon, Y. Huang, J.S. Edgar, Y.S. Ting, S.R. Heron, Y. Kao, et al., Surface acoustic wave nebulization facilitating lipid mass spectrometric analysis, Anal. Chem. 84 (2012) 6530–6537, http://dx.doi.org/10.1021/ac300807p.
- [6] Y. Huang, S.H. Yoon, S.R. Heron, C.D. Masselon, J.S. Edgar, F. Tureček, et al., Surface acoustic wave nebulization produces ions with lower internal energy than electrospray ionization, J. Am. Soc. Mass Spectrom. 23 (2012) 1062–1070, http://dx.doi.org/10.1007/s13361-012-0352-8.
- [7] J. Wei, J. Buriak, G. Siuzdak, Desorption-ionization mass spectrometry on porous silicon, Nature 399 (1999) 243–246 (accessed 15.10.14) http://www. nature.com/nature/journal/v399/n6733/abs/399243a0.html.
- [8] Z. Takáts, J.M. Wiseman, B. Gologan, R.G. Cooks, Mass spectrometry sampling under ambient conditions with desorption electrospray ionization, Science 306 (2004) 471–473, http://dx.doi.org/10.1126/science.1104404.
- [9] P. Nemes, A. Vertes, Laser ablation electrospray ionization for atmospheric pressure, in vivo, and imaging mass spectrometry, Anal. Chem. 79 (2007) 8098–8106, http://dx.doi.org/10.1021/ac071181r.
- [10] S.R. Heron, R. Wilson, S.A. Shaffer, D.R. Goodlett, J.M. Cooper, Surface acoustic wave nebulization of peptides as a microfluidic interface for mass spectrometry, Anal. Chem. 82 (2010) 3985–3989, http://dx.doi.org/10.1021/ ac100372c.

- [11] Y. Tanaka, S. Tamura, Surface acoustic waves in two-dimensional periodic elastic structures, Phys. Rev. B 58 (1998) 7958–7965, http://dx.doi.org/10. 1103/physrevb.58.7958.
- [12] C. Bougault, L. Feng, J. Glushka, E. Kupce, J.H. Prestegard, Quantitation of rapid proton-deuteron amide exchange using hadamard spectroscopy, J. Biomol. NMR 28 (2004) 385–390 (accessed 16.10.14) http://link.springer.com/article/ 10.1023/B:JNMR.0000015406.66725.30.
- [13] J. Pan, J. Han, C.H. Borchers, L. Konermann, Electron capture dissociation of electrosprayed protein ions for spatially resolved hydrogen exchange measurements, J. Am. Chem. Soc, 130 (2008) 11574–11575.
- [14] Y. Pan, M. Briggs, Hydrogen exchange in native and alcohol forms of ubiquitin, Biochemistry 31 (1992) 11405–11412, http://dx.doi.org/10.1021/ bi00161a019 (accessed 16.10.14) http://pubs.acs.org//abs/.
- [15] M. Abdelgawad, A.R. Wheeler, The digital revolution: a new paradigm for microfluidics, Adv. Mater. 21 (2009) 920–925, http://dx.doi.org/10.1002/ adma.200802244.
- [16] R.B. Fair, M.G. Pollack, R. Woo, V.K. Pamula, R. Hong, T. Zhang, et al., A micro-watt metal-insulator-solution-transport (MIST) device for scalable digital bio-microfluidic systems, Int. Electron Devices Meet. Tech. Dig. (2001), http://dx.doi.org/10.1109/IEDM.2001.979513, 16.4. 1-16.4.4.
- [17] K.F. Bohringer, Towards optimal strategies for moving droplets in digital microfluidic systems, Proceedings. ICRA '04. 2004, IEEE, in: IEEE Int. Conf. Robot. Autom., vol. 2, 2004, pp. 1468–1474, http://dx.doi.org/10.1109/ROBOT. 2004.1308031.
- [18] C. Quilliet, B. Berge, Electrowetting: a recent outbreak, Curr. Opin. Colloid Interface Sci. 6 (2001) 34–39, http://dx.doi.org/10.1016/S1359-0294(00)00085-6.
- [19] Y. Li, R. Fu, D. Winters, Test structures for characterizing the integration of EWOD and SAW technologies for microfluidics, IEEE Trans. Semicond. Manuf. 25 (2012) 323–330 (accessed 15.10.14) http://ieeexplore.ieee.org/xpls/abs_all. jsp?arnumber=6212373.
- [20] A. Kirby, A. Wheeler, Digital microfluidics: an emerging sample preparation platform for mass spectrometry, Anal. Chem. (2013), http://dx.doi.org/10. 1021/ac401150q (accessed 15.10.14).
- [21] S. Koster, E. Verpoorte, A decade of microfluidic analysis coupled with electrospray mass spectrometry: an overview, Lab Chip 7 (2007) 1394–1412, http://dx.doi.org/10.1039/b709706a.
- [22] A. Oedit, P. Vulto, R. Ramautar, P.W. Lindenburg, T. Hankemeier, Lab-on-a-chip hyphenation with mass spectrometry: strategies for bioanalytical applications, Curr. Opin. Biotechnol. 31 (2015) 79–85, http://dx. doi.org/10.1016/j.copbio.2014.08.009.
- [23] P.A. Limbach, Z. Meng, Integrating micromachined devices with modern mass spectrometry, Analyst 127 (2002) 693–700, http://dx.doi.org/10.1039/ b200143h.
- [24] T. Rob, P.K. Gill, D. Golemi-Kotra, D.J. Wilson, An electrospray ms-coupled microfluidic device for sub-second hydrogen/deuterium exchange pulse-labelling reveals allosteric effects in enzyme inhibition, Lab Chip 13 (2013) 2528–2532, http://dx.doi.org/10.1039/c3lc00007a.
- [25] D. Resetca, D.J. Wilson, Characterizing rapid, activity-linked conformational transitions in proteins via sub-second hydrogen deuterium exchange mass spectrometry, FEBS J. 280 (2013) 5616–5625, http://dx.doi.org/10.1111/febs. 12332.
- [26] T. Rob, P. Liuni, P.K. Gill, S. Zhu, N. Balachandran, P.J. Berti, et al., Measuring dynamics in weakly structured regions of proteins using microfluidics-enabled subsecond H/D exchange mass spectrometry, Anal. Chem. 84 (2012) 3771–3779, http://dx.doi.org/10.1021/ac300365u.

- [27] S.L. Cohen, B.T. Chait, Influence of matrix solution conditions on the MALDI-MS analysis of peptides and proteins, Anal. Chem. 68 (1996) 31–37 http://www.ncbi.nlm.nih.gov/pubmed/8779435.
- [28] Y. Li, Y.Q. Fu, S.D. Brodie, M. Alghane, A.J. Walton, Integrated microfluidics system using surface acoustic wave and electrowetting on dielectrics technology, Biomicrofluidics 6 (2012) 12812, http://dx.doi.org/10.1063/1. 3660198.
- [29] A. Renaudin, P. Tabourier, V. Zhang, J.C. Camart, C. Druon, SAW nanopump for handling droplets in view of biological applications, Sens. Actuators B Chem. 113 (2006) 389–397, http://dx.doi.org/10.1016/j.snb.2005.03.100.
- [30] D. Beyssen, L. Le Brizoual, Ö. Elmazria, P. Alnot, Microfluidic device based on surface acoustic wave, Sens. Actuators B Chem. 118 (2006) 380–385, http:// dx.doi.org/10.1016/j.snb.2006.04.084.
- [31] Y.Q. Fu, J.K. Luo, X.Y. Du, A.J. Flewitt, Y. Li, G.H. Markx, et al., Recent developments on ZnO films for acoustic wave based bio-sensing and microfluidic applications: a review, Sens. Actuators B Chem. 143 (2010) 606–619, http://dx.doi.org/10.1016/j.snb.2009.10.010.
- [32] J. Zhou, M. DeMiguel-Ramos, L. Garcia-Gancedo, E. Iborra, J. Olivares, H. Jin, et al., Characterisation of aluminium nitride films and surface acoustic wave devices for microfluidic applications, Sens. Actuators B Chem. 202 (2014) 984–992, http://dx.doi.org/10.1016/j.snb.2014.05.066.
- [33] L.Y. Ye, H.C. Chang, P.P.Y. Chan, J.R. Friend, Microfluidic devices for bioapplications, Small 7 (2011) 12–48, http://dx.doi.org/10.1002/smll. 201000946.
- [34] L.Y. Yeo, J.R. Friend, Surface acoustic wave microfluidics, Annu. Rev. Fluid Mech. 46 (2014) 379–406, http://dx.doi.org/10.1146/annurev-fluid-010313-141418.
- [35] X. Ding, P. Li, S.S. Lin, Z.S. Stratton, N. Nama, F. Guo, et al., Surface acoustic wave microfluidics, Lab Chip 13 (2013) 3626–3649, http://dx.doi.org/10. 1039/c3lc50361e.
- [36] T. Franke, A.R. Abate, D.A. Weitz, A. Wixforth, Surface acoustic wave (SAW) directed droplet flow in microfluidics for PDMS devices, Lab Chip 9 (2009) 2625–2627, http://dx.doi.org/10.1039/b906819h.
- [37] S.K. Cho, H. Moon, C. Kim, Creating, transporting, cutting, and merging liquid droplets by electrowetting-based actuation for digital microfluidic circuits, J. Microelectromech. Syst. 12 (2003) 70–80, http://dx.doi.org/10.1109/JMEMS. 2002.807467.
- [38] K. Choi, A.H.C. Ng, R. Fobel, A.R. Wheeler, Digital microfluidics, Annu. Rev. Anal. Chem. 5 (2012) 413–440, http://dx.doi.org/10.1146/annurev-anchem-062011-143028.
- [39] Z. Zhang, D.L. Smith, Determination of amide hydrogen exchange by mass spectrometry: a new tool for protein structure elucidation, Protein Sci. 2 (1993) 522–531, http://dx.doi.org/10.1002/pro.5560020404.
- [40] M. Guttman, D.D. Weis, J.R. Engen, K.K. Lee, Analysis of overlapped and noisy hydrogen/deuterium exchange mass spectra, J. Am. Soc. Mass Spectrom. 24 (2013) 1906–1912, http://dx.doi.org/10.1007/s13361-013-0727-5.
- [41] D.D. Weis, J.R. Engen, I.J. Kass, Semi-automated data processing of hydrogen exchange mass spectra using HX-Express, J. Am. Soc. Mass Spectrom. 17 (2006) 1700–1703, http://dx.doi.org/10.1016/j.jasms.2006.07.025.
- M. Tudorache, C. Bala, Biosensors based on screen-printing technology, and their applications in environmental and food analysis, Anal. Bioanal. Chem. 388 (2007) 565–578, http://dx.doi.org/10.1007/s00216-007-1293-0.
 F.C. Krebs, M. Jørgensen, K. Norrman, O. Hagemann, J. Alstrup, T.D. Nielsen,
- [43] F.C. Krebs, M. Jørgensen, K. Norrman, O. Hagemann, J. Alstrup, T.D. Nielsen, et al., A complete process for production of flexible large area polymer solar cells entirely using screen printing-first public demonstration, Sol. Energy Mater. Sol. Cells 93 (2009) 422–441, http://dx.doi.org/10.1016/j.solmat.2008. 12.001.