

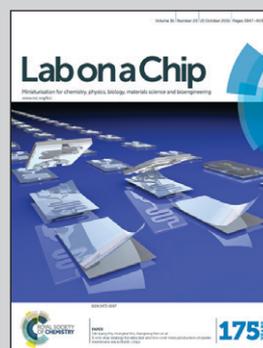


Featuring work from the BioNano Technology Lab,  
Dr. Christian D. Ahrberg, Prof. Andreas Manz and  
Prof. Bong Geun Chung, Sogang University, Korea

#### Polymerase chain reaction in microfluidic devices

We highlight recent developments of polymerase chain reaction (PCR) in microfluidic devices. Time and space domain microfluidic devices as well as isothermal nucleic acid amplification and digital PCR microfluidic devices are discussed.

#### As featured in:



See Bong Geun Chung *et al.*,  
*Lab Chip*, 2016, **16**, 3866.



CrossMark  
click for updates

Cite this: *Lab Chip*, 2016, 16, 3866

Received 2nd August 2016,  
Accepted 6th September 2016

DOI: 10.1039/c6lc00984k

www.rsc.org/loc

## Polymerase chain reaction in microfluidic devices

Christian D. Ahrberg,<sup>a</sup> Andreas Manz<sup>bc</sup> and Bong Geun Chung<sup>\*a</sup>

The invention of the polymerase chain reaction (PCR) has caused a revolution in molecular biology, giving access to a method of amplifying deoxyribonucleic acid (DNA) molecules across several orders of magnitude. Since the first application of PCR in a microfluidic device was developed in 1998, an increasing number of researchers have continued the development of microfluidic PCR systems. In this review, we introduce recent developments in microfluidic-based space and time domain devices as well as discuss various designs integrated with multiple functions for sample preparation and detection. The development of isothermal nucleic acid amplification and digital PCR microfluidic devices within the last five years is also highlighted. Furthermore, we introduce various commercial microfluidic PCR devices.

### Introduction

Since polymerase chain reaction (PCR) was invented by Kary Mullis over 30 years ago,<sup>1</sup> it has become a valuable tool for molecular biology as it can exponentially amplify deoxyribonucleic acid (DNA). Several million copies of DNA can be generated from a single molecule by only few cycles of PCR. A PCR cycle typically consists of three distinct steps, de-

naturation, annealing, and extension. In the denaturation step ( $\approx 95$  °C), double stranded DNA denaturates into two single strands. Afterwards the temperature is decreased for the annealing step ( $\approx 56$  °C). Primers, which are short, complementary sequences of DNA, can anneal to the single stranded target DNA. Furthermore, a polymerase attaches to the complex; the polymerase is often derived from thermophile organisms like *Thermus aquaticus*.<sup>2</sup> To increase activity of the polymerase, the temperature is raised for the extension step, typically to around 72 °C. The polymerase gains activity, synthesizing a second complementary strand of DNA from free nucleotides in solution. Therefore, an exponential increase of double stranded DNA concentration is obtained by repeating

<sup>a</sup> Department of Mechanical Engineering, Sogang University, Seoul, Korea.

E-mail: bchung@sogang.ac.kr

<sup>b</sup> Microfluidics group, KIST-Europe, Saarbrücken, Germany

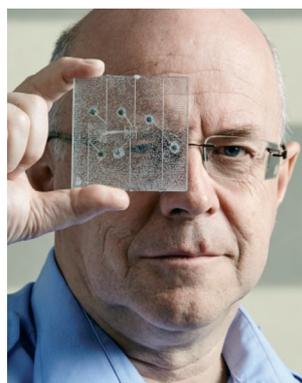
<sup>c</sup> Mechatronics Department, Universität des Saarlandes, Saarbrücken, Germany



Christian D. Ahrberg

Christian D. Ahrberg obtained his Bachelor and Master Degree in Chemical and Bioengineering from ETH Zürich. After graduating in 2012 he worked at Bayer Technology Services in Leverkusen before starting his PhD under supervision of Andreas Manz at the Mechatronics department of Saarland University in cooperation with KIST Europe, during which he worked on a portable real-time PCR device. Since graduating in 2016, currently he is a

postdoctoral researcher at the BioNano Technology Lab in Sogang University, Seoul, Korea.



Andreas Manz

Andreas Manz is currently professor at the Saarland University, and principal investigator at KIST Europe, Saarbrücken Germany. He graduated 1986 with PhD from Chemistry ETH Zurich Switzerland, was group leader at Ciba-Geigy Ltd. (Novartis) Basel Switzerland 1988–1995, professor at Imperial College London UK 1995–2004, head of the ISAS (Leibniz-Institute of Analytical Sciences) and professor at the Technical University Dortmund Germany 2003–2008. He was founder and chairman of the RSC journal *Lab on a Chip* 2001–2008, and scientific advisor to Caliper Technologies Inc. (now part of Perkin Elmer) 1995–2000.



three temperature steps. The method was extended to reverse transcription PCR.<sup>3</sup> Starting with a ribonucleic acid (RNA) molecule, a reverse transcription step is performed to obtain DNA complementary to the RNA strand before PCR. A further modification of PCR introduced fluorescent dyes into the reaction mixture, allowing the real-time monitoring of reaction progress.<sup>4</sup> This extension of the method allowed the determination of initial copies of DNA in the reaction mixture through standard curves. The method is referred to as quantitative real-time PCR (qPCR). Today, a number of PCR applications (e.g., diagnostics and medicine,<sup>5–7</sup> agricultural sciences,<sup>8</sup> and food sciences<sup>9</sup>) can be found.

Microfluidic devices offer small thermal mass, low thermal inertia, and rapid heat transfer. They present a number of advantages compared to macroscopic equivalents. The small volumes reduce sample and reagent consumption, leading to inexpensive operation of the systems. However, volumes in most cases are still large enough for bulk kinetics to apply. Furthermore, different functions can easily be integrated in microfluidic devices.<sup>10</sup> Diffusive mass transport and heat conduction scale with the characteristic length of the system. While in macroscale systems diffusion and heat conduction are slow for efficient transport, the smaller scales of microfluidic devices allow for utilization of these effects. Furthermore, these effects are easier to control than convective heat and mass transport used on larger scales instead. This allows for a better control of processes in microfluidic devices. A reduction in size also leads to a higher surface to volume ratio. This means that the exchange areas for mass and heat transport are larger, allowing for faster transport. On the other hand, surface and wall effects become more important compared to larger devices. As PCR requires many heating and cooling steps, total analysis time can be significantly reduced by miniaturization. Examples of PCR conducted in microfluidic chambers are among the first examples of PCR in microfluidic devices.<sup>11,12</sup> Soon after ligase chain reaction, a method closely related to PCR was shown in the same devices.<sup>13</sup> In 1998, the first flow through PCR in a micro-

fluidic device was demonstrated by Kopp *et al.*<sup>14</sup> A sample was flown through a microfluidic channel, repeatedly passing the three temperature zones for PCR. A 10  $\mu\text{L}$  sample could perform 20 cycles of PCR within 90 seconds. Numerous applications of PCR in microfluidic devices have been proposed.<sup>15,16</sup> Through the low sample and power consumption, these microfluidic devices are especially interesting for point-of-care applications. The possibility to combine sample preparation, PCR, and detection in the same automated microfluidic device also promises reduction of handling time as well as prevention of sample contamination. This review introduces and discusses various designs of PCR in microfluidic device developed within the last five years. First, space domain PCR microfluidic devices are discussed before moving on to their counterpart time domain PCR microfluidic devices. Second, isothermal nucleic acid amplification and digital PCR in microfluidic devices are discussed (Table 1).

## Space domain PCR

When the sample is moved through a microchannel featuring different temperatures along its length, the temperature is dependent on the position in the channel. Thus, this group of devices is referred to as space domain PCR device. Sample preparation can easily be integrated in the microfluidic device. The duration of the thermal steps of PCR and the number of cycles are predefined by the microfluidic design. The operation of space domain PCR devices requires movement of the sample along the channel.

### Serpentine channel designs

Since the first application of a microfluidic PCR device, a variety of devices improving and evolving the initial design have been developed. A polyimide device with three integrated resistive copper heaters has been presented (Fig. 1A).<sup>17</sup> The microfluidic device was designed for 30 cycles of PCR with a time ratio of 1:1:2 for denaturation, annealing, and extension step. Due to the low cost of the plastic device, it can be disposed after the experiment preventing cross-contamination that could arise through adsorption of DNA onto the channel walls. In another study, a similar design featuring 30 cycles of PCR was created in a polycarbonate substrate using a computer numerical controlled (CNC) vertical milling machine.<sup>18</sup> The temperatures were created using two copper heating blocks for denaturation and a combined annealing/extension step. Wu *et al.* reduced the flow resist of these devices by coating a 0.5 mm inner diameter steel capillary with Norland Optical Adhesive 61 (NOA61).<sup>19</sup> A smooth, hydrophobic inner surface was used to reduce the back pressure. While the previous examples experimented with different substrate materials reducing device costs and optimizing flow resistance, other researchers optimized the heaters or simplified device design by changing sample pumping methods and reducing the number of heaters. Harandi *et al.* optimized the heating of the sample using a metal filled composite material.<sup>20</sup> By employing copper comb structures



**Bong Geun Chung**

*Bong Geun Chung is currently Associate Professor at Department of Mechanical Engineering in Sogang University, Seoul, Korea. He received his Bachelor and Master degrees from Hanyang University, Seoul, Korea and his Ph.D. degree from University of California Irvine, USA in 2007. He was an Instructor at Department of Medicine in Harvard Medical School, Boston, USA. His research interests are in the areas of Lab on a Chip, nanomedicine, and tissue engineering. He is now a director of the BioNano Technology Lab in Sogang University, Seoul, Korea.*

*Bong Geun Chung is currently Associate Professor at Department of Mechanical Engineering in Sogang University, Seoul, Korea. He received his Bachelor and Master degrees from Hanyang University, Seoul, Korea and his Ph.D. degree from University of California Irvine, USA in 2007. He was an Instructor at Department of Medicine in Harvard Medical School, Boston, USA. His research interests are in the areas of Lab on a Chip,*



**Table 1** Summary table giving an overview of the various designs and approaches for microfluidic PCR devices including commercial PCR devices

Method	Design & system	Approach	Ref.
Space domain PCR	Serpentine channel designs	Disposable polyimide chip	17
		30 cycles of PCR	
		CNC fabricated polycarbonate device	18
		30 cycles of PCR	
		Steel capillary coated with NOA61	19
		Reducing back pressure	
		Metal-polymer composite material	20
		Improved temperature stability	
		Replacing active pumps with capillary forces	21
	Radial designs	Model of capillary forces in channel for PCR	22
		Reducing the number of heaters	
		Central heater, edges of chip cooled by heat pipes	
		Reducing the number of heaters 3D device heated at the bottom	23
	Digital and oscillating designs	Radial design	24
		Central heater, edges actively cooled	
		Radial design	25, 26
		Heated by focusing sunlight on center of device	
		Capillary wrapped around cylinder	27
		Cylinder with different temperature zones	
		Centrifugal microfluidic device	28
	Heating/cooling systems	Mounted on dual shaft centrifuge	
		Oscillating droplet	30
		Four parallel channels for multiplexing	
Oscillating droplet		31	
Nested PCR to increase specificity			
Digital microfluidic device		32	
Droplet moved between temperature zones			
Digital microfluidic device		33	
Real-time optical detection			
Digital microfluidic device		34	
Time domain PCR	Method for solvent replenishing		
	Capillaries in water bath	35	
	Forced convection inside capillaries		
Heating/cooling systems	Sample moved between water bath using servo motor	38, 39	
	Infrared heating through tungsten lamp	40, 41	
	Fan for active cooling		
	Infrared laser for heating	42	
	Silicon heating chip	44, 45	
	Portable real-time PCR device		
	Silicon heating chip	46	
	Melting curve analysis, multiplexing		
	Sample chamber on top of heat exchanger	47, 48	
	Heating and cooling through medium		
	Silica capillaries with thin film heaters	49	
	Cooling by fan		
	Capillary with bypass channel	50	
	Heating and cooling by Peltier elements		
	Integrated devices	Lysis and PCR in same chamber	54
		Removal of inhibitors through diffusion	
		Chemical lysis followed by DNA extraction with magnetic particles	55
Retention of DNA on silica gel membrane		57	
Release by master mix and amplification			
FTA membrane at entrance of amplification chamber retaining DNA		58	
Dielectrophoresis to capture pathogens on slip chip		60	
Followed by washing and PCR			
Separation of amplicons by CE after PCR		61, 62	
PCR in amplification chamber connected to CE	63		
Real-time detection of amplicons			



Table 1 (continued)

Method	Design & system	Approach	Ref.
	Centrifugal microfluidic designs	Centrifugal device with chambers	64, 65
		Counting of bacteria through Poisson distribution	
		PCR centrifugal device	66, 67
		Ice-valves next to PCR chamber	
		Centrifugal device dividing sample in 8 sub-samples	67, 68
		Centrifugal device subdividing sample	69, 70
	Arrays	Positive, negative control, and internal standards	
		LabDisc device with vapor diffusion barrier	71
		Preventing bubble formation in amplification chamber during PCR	
		LabDisc device including sample preparation	72
		Printed array of sample droplets in oil	73
		Dispensing of single cells onto hydrophilic spots for PCR	74
Single cell devices	Compartmentalization of sample by centrifugal device	75	
	Detection of amplicons by printed DNA probe array		
	Printing of single cells into individual reaction volumes	76	
	Trapping of single cells followed by lysis and RNA extraction	77	
	Encapsulation of single cells into droplets	78	
	Droplet sorting to recover genome of target species		
Isothermal nucleic acid amplification	Implementation of LAMP on microfluidic devices	PDMS valve to create 96 × 96 array of single cells	79, 80
		300 cell traps separated by valves	81
		PCR in chambers adjacent to cell traps	
		Continuous flow LAMP in droplets for DNA and microRNA	84, 85
		Multiplexed LAMP in device with 8 amplification	86
		LAMP in agarose beads	87
	LAMP for point-of-care devices	Extraction of RNA using magnetic beads followed by reverse transcription LAMP	88
		Centrifugal device with real-time detection of LAMP reaction	89
		Centrifugal device for multiplexed detection of 10 different species by LAMP	90
		Centrifugal LAMP device including sample preparation and detection of products	91
		LAMP in capillary heated by two pocket warmers for portable format	92
		Disposable polycarbonate cassette for LAMP	93
Digital PCR	Multiplexed digital PCR	LAMP followed by electrochemical detection	94
		Ten reaction chambers with different primers for multiplexed LAMP, detection by naked eye	95
		Duplexed digital PCR in device containing 38 panels each with 770 partitions	104
	Various designs	Duplex digital PCR in droplets utilizing TaqMan-probes with different emission wavelength	105, 106
		Cross-junction to generate water in oil emulsions for droplet digital PCR	107
		Creating sample array through PDMS membrane	108
		Megapixel digital PCR device	109, 111, 112
		Creating droplets at T-junction	110
		Creating sample array with negative pressure vapor proof-layer to stop sample evaporation	113
		Digital PCR in agarose beads	114
		Digital PCR to count nanoparticles tagged with DNA	115
		Integrated systems	Centrifugal device dividing samples into reaction volumes
Centrifugal device using step emulsification for creating droplets	117–119		
Device combining sample preparation and digital PCR	140, 141		
Slip chip for digital PCR	120–122		
Commercial PCR devices	Integrated devices	Centrifugal system with sample preparation and detection	123
		Automated systems including sample preparation, amplification and detection using prefilled reagent strips or blister-films	124–126
		Integrated, automated systems utilizing cartridges	127, 128
		Microfluidic silicon chip, with hybridization array for detection	129



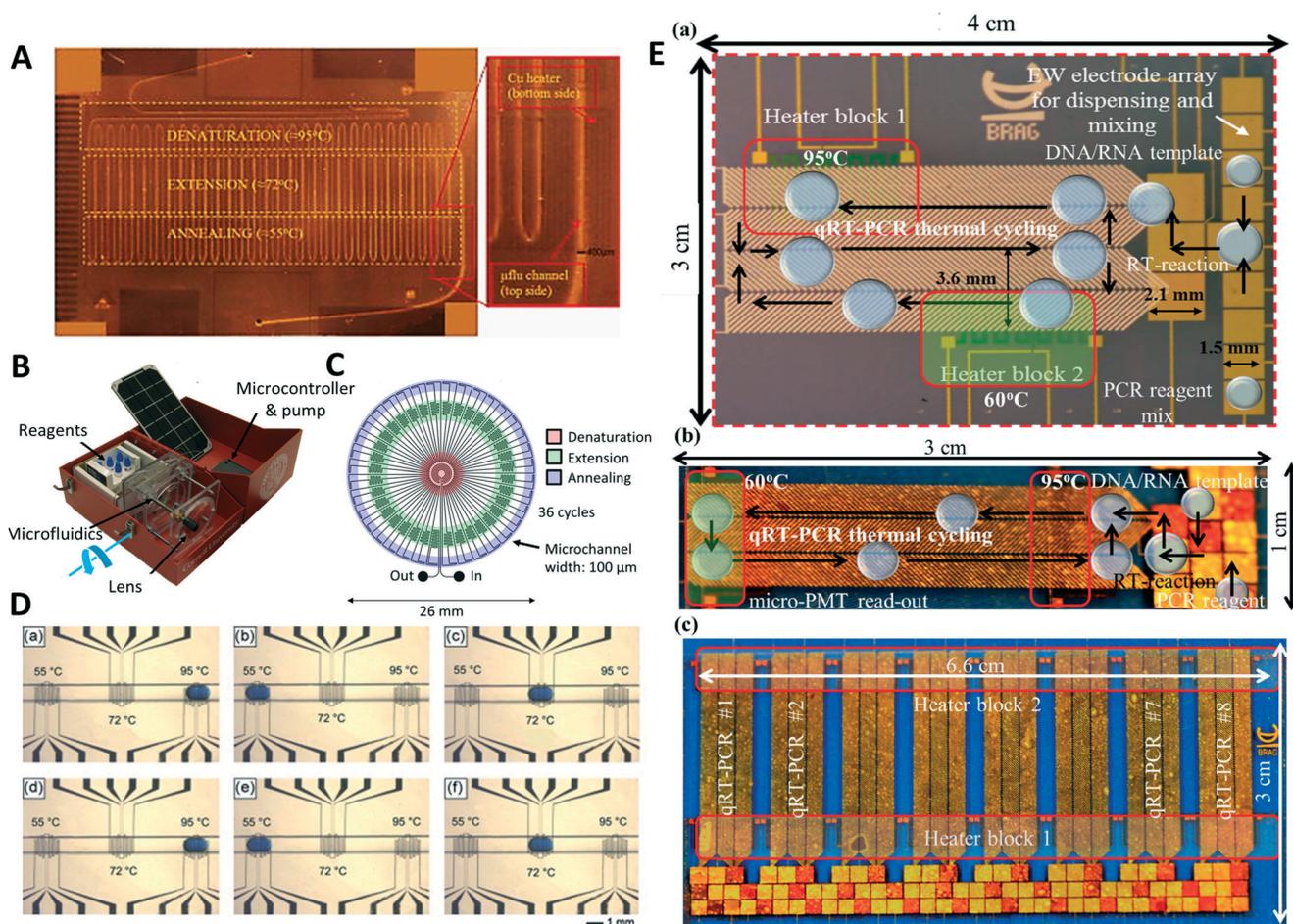
Table 1 (continued)

Method	Design & system	Approach	Ref.
	Digital PCR devices	Based on an array of chambers	133
		Droplet based	134, 135
		Integrated fluidic circuits for sample portioning	136

inside of a polyimide matrix, temperature control with a precision of 1 °C could be achieved. In 2014, Tachibana *et al.* suggested a method to reduce the complexity of devices by replacing active pumps by capillary forces.<sup>21</sup> The authors created a numerical model describing how the capillary force periodically changes. They developed a microfluidic device that could perform 50 cycles of PCR in 14 minutes. Capillary forces transported the sample through the microfluidic device, eliminating the need for an active pump.

Other publications reduce the number of heaters. One example uses a single heater mounted in the middle of the de-

vice to create an denaturation temperature, while the edges of the device are cooled using heat pipes cooled with a fan to create an annealing zone.<sup>22</sup> Another example used a three-dimensional (3D) poly(dimethylsiloxane) (PDMS) device on a single heater.<sup>23</sup> The PDMS microfluidic device consisted of a channel meandering between the top layer and bottom layer of the microfluidic device containing the heater. The heater was set to denaturation temperature creating a denaturation zone at the bottom of the microfluidic device, while the top layer acted as an annealing/extension zone. Most devices are close to the original serpentine channel PCR device. While



**Fig. 1** Space domain PCR devices. (A) A serpentine-based microfluidic PCR device containing three copper heaters (reprinted with permission from ref. 17, Copyright 2014 Elsevier). (B and C) A portable PCR device powered by solar energy with the radial design featuring a central denaturation zone. Data are transferred wirelessly to a tablet computer for analysis (reprinted with permission from ref. 26, Copyright 2016 PLoS One). (D) Example for an oscillating droplet inside of a microfluidic channel. The sample droplet is moved between three different heaters for denaturation (a), annealing (b), and extension (c), a second cycle of PCR (d–f). For nested PCR, the droplet is removed from the channel and mixed with the reagents for the nested reaction before being reintroduced to the channel (reprinted with permission from ref. 31, Copyright 2011 Elsevier). (E) Photographs of a spiral droplet-dielectrophoresis digital microfluidic device for PCR (a), the continuous, bi-directional droplet actuation scheme (b), and the eight-plex microfluidic device (c) (reprinted with permission from ref. 33, Copyright 2015 MDPI AG).



some advances make operation of these devices simpler and more robust, others might reduce the requirement of equipments. However, implementation of these simplifications often restricts the accessible operating conditions.

### Radial designs

The number of heaters can be reduced by creating symmetric devices. Schaerli *et al.* created a droplet PCR device with a circular geometry.<sup>24</sup> The center of the device was heated by a single copper rod to create a denaturation zone. Channels transported sample to the edge of the radial device where temperatures are suitable for annealing and extension due to the natural thermal gradient. A Peltier element was applied to the edge of the design to adjust the thermal gradients. A similar design was recently proposed utilizing the sun as a heat source, making the device independent of external power sources.<sup>25</sup> The authors used a radial device and focused the sun light on its center using a lens. An absorber layer on the device converted the sun light into heat, generating temperature zones for PCR. A portable PCR device was constructed by addition of a solar powered syringe pump and fluorescence measurements (Fig. 1B and C).<sup>26</sup> The portable device was used for the detection of Kaposi's sarcoma in pseudo-biopsy samples with a sensitivity of 83% and specificity of 70%.

In a different approach to radial designs, a capillary was wrapped around a cylinder machined out of three parts with a 1 mm gap in-between them.<sup>27</sup> Each part contained its own cartridge heater and provided denaturation, annealing, or extension temperature. The number of PCR cycles could be varied by changing the number of loops of the capillary around the cylinder. Furthermore, the system consisted of another heated cylinder before the amplification cylinder to perform a reverse transcription step and a fluorescence detection unit. Miao *et al.* used a radial centrifugal microfluidic PCR system.<sup>28</sup> The device employed a circular PCR containing four microfluidic chambers for denaturation, annealing, extension, and standby. The heaters were mounted underneath each chamber to provide the necessary temperature. The microfluidic device was mounted on a dual-shaft centrifugal device to allow for bidirectional pumping of the fluid using centrifugal forces. While only the number of PCR cycles can be modified in the capillary-based approach, the cycle number and duration of the single temperature steps can be changed by the centrifugal system.

### Digital and oscillating microfluidic designs

Electrowetting-mediated droplets are used in digital microfluidic devices. Cycle number and temperature step durations of PCR can be controlled by software manipulation using digital microfluidics. Oscillatory microfluidics is a simplification of digital microfluidics and samples are moved back and forth in a channel through a syringe pump. A first example of oscillating or bidirectional flow PCR was shown in 2007.<sup>29</sup> Zhang *et al.* used an array of independent parallel channels passing through three temperature zones to perform multiplexed

PCR.<sup>30</sup> The device used a syringe pump to generate the oscillation motion of sample between three temperature zones for the detection of different foodborne bacterial pathogens. In a similar manner, Sciancalepore *et al.* used a single straight channel with a sample oscillating to perform nested PCR, increasing the specificity of the reaction (Fig. 1D).<sup>31</sup> Droplets looping between three different heaters were employed for a digital microfluidic PCR device.<sup>32</sup> Optimization algorithms and simulations were used to increase the performance of the thermal looping and reduce sample cross-contamination. In a further use of digital microfluidics, a droplet was shuttled between two temperature zones for denaturation and annealing.<sup>33</sup> In Fig. 1E, a photograph of the device can be seen illustrating the movement of the droplets through the different temperature zones. Furthermore, it is illustrated how several devices can be operated in parallel using the same two heating blocks. Normal PCR and reverse transcription PCR were performed using this microfluidic setup. Through the addition of a fluorescence detection unit on top of the digital microfluidic device, end point detection of amplification and quantitative PCR could be performed. A further application added a geometry to replenish solvents to counter evaporation in air-matrix digital microfluidic applications of PCR.<sup>34</sup> A hole was drilled in the bottom layer of the digital microfluidic device. If the sample droplet was moved on top of the hole, solvent could be replenished using a syringe pump. In oscillatory and digital microfluidic devices, the number of cycles and duration of temperature steps can be changed unlike other space domain PCR devices.

### Heating and cooling systems

To regulate the temperature, a small water bath, heated by a tea candle, was used to perform PCR in a low power and portable manner.<sup>35</sup> The water bath contained a holder for 20  $\mu\text{L}$  capillaries for the sample. Due to the temperature gradient from the bottom to the top of the capillary, convective flow arose and the samples were cycled between different temperature zones. This demonstration is similar in function to internal convection devices shown previously, however the water bath allows a significant simplification of the heating system.<sup>36,37</sup> In a further operation mode using water baths, PCR samples are kept in vials for conventional PCR ( $\approx 20 \mu\text{L}$ ) and moved between different water bath for heating and cooling of the sample. A first example for these types of devices used 17  $\mu\text{L}$  glass vials for the PCR sample which was transported between 3–4 water baths using a sample holder on a pan/tilt servo motor.<sup>38</sup> The first water bath was used for performing a reverse transcription step when starting from RNA. The second and third were kept at temperatures for a denaturation and annealing/extension step. The last, optional bath remained at room temperature and was used to perform faster cooling steps. They were able to achieve 40 cycles of PCR in 5.5 minutes. By using smaller samples and a similar setup, Farrar *et al.* managed to perform 35 cycles of PCR in 14.7 seconds.<sup>39</sup> Next to the sample capillary, a temperature sensor was placed to trigger the stepper motor moving the sample into the next bath. Additionally, a fluorescence detection unit was



placed in the room temperature water bath, allowing real-time monitoring of PCR. These combinations of small sample volumes and large water baths for heating and cooling allow for fast cycle times. Water baths need significantly more energy to heat up compared to other heating equipments. Furthermore, integration with sample preparation or post amplification analysis is not possible.

## Time domain PCR

In a second class of devices, samples are stationary and the temperature is changed through active heating and cooling. The sample temperature is dependent on the time; hence it is referred to as time domain PCR device. As the temperature profile is not dependent on the channel design, changes to the PCR thermal cycling can be made by modification of the heating and cooling protocols. As no movement of the sample is required, there is no requirement for pumps. A disadvantage of this method of operation is that devices cannot be operated continuously limiting sample throughput.

### Different methods of heating and cooling

Time domain devices are reliant on performing a suitable temperature profile on a stationary sample. Active heating and cooling steps are required. Thus, more elaborate systems are required than used for most space domain PCR devices. To solve this problem, many different approaches (*e.g.*, fast heating or accurate temperature control) can be considered. A popular choice for heating is infrared light as the sample can be heated directly using this method without convection or conduction steps. One example used a tungsten lamp for infrared heating and a fan for forced convective cooling, achieving heating and cooling rates of about  $10\text{ }^{\circ}\text{C s}^{-1}$ .<sup>40</sup> In a similar approach, a tungsten lamp with a 780 nm long pass filter was used for heating.<sup>41</sup> Two modulated lasers with different frequencies were employed for the excitation of two dyes, EvaGreen and ROX. Emission was collected by a photomultiplier tube (PMT) in a color-blind manner and was demodulated by frequency analysis, showing that two amplicons could be quantified simultaneously. Pak *et al.* used a 1450 nm infrared laser for heating a 1.4  $\mu\text{L}$  sample inside of a microfluidic chamber.<sup>42</sup> Through the use of a laser instead of a tungsten lamp, heating rates up to  $60\text{ }^{\circ}\text{C s}^{-1}$  could be obtained. The small thermal masses of microfluidic systems allow fast heating rates. Plasmonic photothermal light-to-heat conversion was used in another demonstration by irradiating a thin gold layer inside of microfluidic PCR chambers with LEDs.<sup>43</sup> Photon-electron-phonon coupling in the gold membrane allowed heating the PCR solution up to  $150\text{ }^{\circ}\text{C}$  within 3 minutes with heating rates of  $12.79\text{ }^{\circ}\text{C s}^{-1}$ . In a further example, utilizing the small thermal masses, sample droplets ( $\sim 200\text{ nL}$ ) were placed on a hydrophobic glass surface and covered with small mineral oil droplets ( $\sim 1\text{ }\mu\text{L}$ ) to prevent evaporation.<sup>44,45</sup> The samples were heated using a silicon device with gold resistive heaters and temperature sensors to heat four samples in parallel. Combined with an

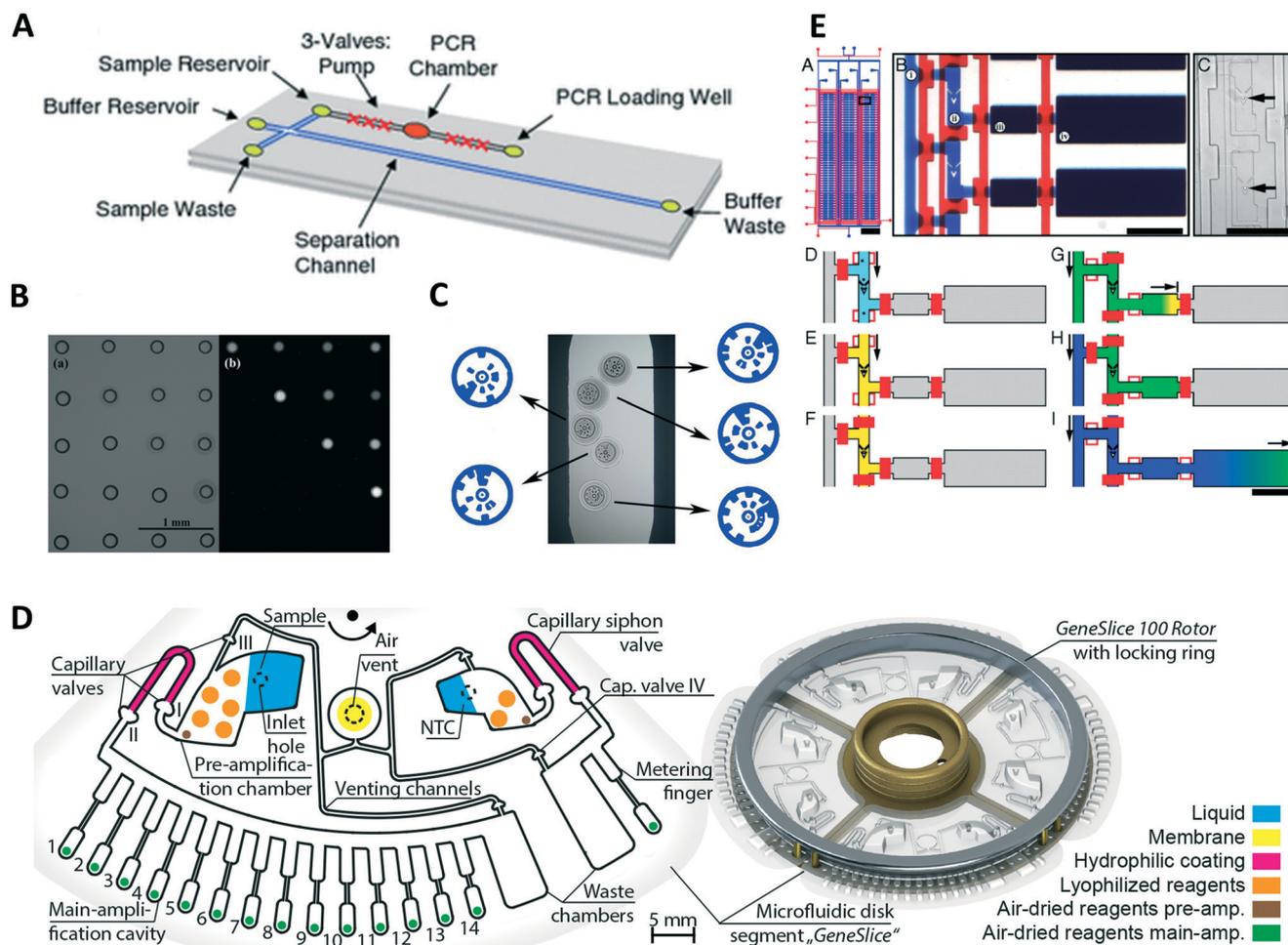
optical detection unit and an external 12 V power supply, the system formed a portable real-time PCR device. The small temperature gradients inside of the sample and fast heating rates were also utilized to perform fast melting curve analysis after each cycle of PCR,<sup>46</sup> showing that two targets could be amplified and quantified simultaneously using only a single intercalating dye.

Active heating and cooling can be accomplished by mounting a heat exchanger beneath the sample chamber for temperature control. To change from heating to cooling, the hot heating medium can be replaced by cooling medium through a valve. For example, Wheeler *et al.* used water as a medium flowing through a porous matrix as a heat exchanger.<sup>47</sup> 5  $\mu\text{L}$  sample could be heated and cooled with rates up to  $45\text{ }^{\circ}\text{C s}^{-1}$ , resulting in reaction times of under 3 minutes for 30 cycles. A similar approach used a mixture of propylene glycol and water flown through a second chamber on top of the 25  $\mu\text{L}$  sample chamber.<sup>48</sup> Melting curve analysis was conducted after amplification by applying a temperature gradient to the reservoir containing the cooling medium. Capillaries were also employed to conduct time domain PCR. Sundberg *et al.* flowed 50 nL PCR samples into one of eight parallel fused silica capillaries coated with thin film heaters.<sup>49</sup> Active cooling was provided by a fan, resulting in heating rates of  $50\text{ }^{\circ}\text{C s}^{-1}$  and cooling rates of  $20\text{ }^{\circ}\text{C s}^{-1}$ . Once PCR was completed after 12.5 minutes, flow could be continued and samples were collected from the end of the capillary. Another approach prepared a 14  $\mu\text{L}$  sample in a chamber with a bypass channel.<sup>50</sup> The sample chamber would be filled continuously, once the flow resistance increases significantly and sample flows through the bypass capillary. Heating and cooling were actively performed using two Peltier elements placed on each side of the sample chamber. The different methods of heating and cooling show different characteristics. While some offer fast heating and cooling rates, others have a higher precision or are able to handle larger sample volumes. The choice of heating method should be made depending on applications and requirements.

### Different approaches to PCR

In time domain PCR, the sample solution does not need to be moved and primers can be immobilized by binding them to a surface. This allows for a combination of PCR and microarray technology.<sup>51</sup> The authors conducted a nested PCR inside of a microfluidic chamber. Once the first PCR was completed, a nested PCR was carried out with some of the primers covalently attached to the glass surface. After complete reaction, the chamber was washed and filled with an aqueous solution containing an intercalating dye. From the pattern of fluorescence signals, the type of avian flu RNA could be identified. Jung *et al.* used optically encoded, hydrophilic porous polyethylene glycol microparticles to immobilize forward primers of several different micro RNAs (Fig. 2C).<sup>52</sup> Reverse transcription qPCR was performed in a channel. The fluorescence signal could be demultiplexed by decoding the code on the particles, leading to the parallel quantification of different





**Fig. 2** Time domain PCR devices. (A) Device capable of performing PCR from whole blood samples. Adjacent to the PCR chamber is microfluidic valves which can be used for pumping. After amplification targets can be separated using capillary electrophoresis on the same device (reprinted with permission from ref. 62, Copyright 2016 Springer). (B) Bright field image of a contact printed array of droplets (left) and fluorescence image of the same array after selective amplification of droplets using an infrared laser. This technique allows applying individual temperature profiles to each droplet as well as only reducing the testing to selected droplets (reprinted with permission from ref. 73, Copyright 2009 Royal Society of Chemistry). (C) Bright field image illustrating the barcoded hydrogel particles with primers for different microRNAs bound to them in microfluidic PCR device (reprinted with permission from ref. 52, Copyright 2016 Nature Publishing Group). (D) Example of a schematic of centrifugal microfluidic device for nested PCR (left). Image of the assembled centrifugal device fitting a commercial thermocycler (right) (reprinted with permission from ref. 70, Copyright 2015 PLoS One). (E) Device for the analysis of single cells via reverse transcription qPCR (a–c). Individual cells are trapped in cell traps (e) before a washing and cell lysis step (d–f). Next samples are flown into a reverse transcription chamber using active valves (g and h). Lastly qPCR is carried out in a chamber (i) (reprinted with permission from ref. 81, Copyright 2011 National Academy of Science).

micro RNAs. Qin *et al.* developed a microfluidic device from NOA81.<sup>53</sup> The device consisted of a microfluidic chamber on top of a Peltier element for heating and cooling. As NOA81 inhibits the PCR reaction, the authors coated the device with bovine serum albumin to prevent absorption of DNA and polymerase during the reaction.

### Integrated devices

While some applications focus on heating methods or modifications of PCR process, a number of publications consider the combination of time domain PCR, sample preparation, and post PCR analysis. Ma *et al.* performed a lysis step in a chamber connected to two side chambers.<sup>54</sup> Once cells were lysed, they replaced the lysing buffer in the reaction chamber with PCR solution by diffusion from the side chambers. In parallel, they uti-

lized the difference in diffusivity between long DNA strands and PCR inhibitors to remove the inhibitors from the solution in the same step. DNA was extracted from *Helicobacter pylori* by thermolysis in a chamber and bound to magnetic particles.<sup>55</sup> The magnetic beads remained in the chamber during several washing steps and a subsequent filling of the chamber with PCR master mix. After PCR, successful amplification was confirmed by fluorescence measurements using the intercalating dye SYBR green I. DNA was extracted with magnetic beads, before releasing the DNA into a microfluidic amplification chamber.<sup>56</sup> After amplification, targets were analyzed using capillary electrophoresis on the same device. Membranes can also be used to extract DNA from cell lysate. Cell lysate from human blood was prepared using a commercial product before introducing the lysate into a microfluidic device.<sup>57</sup> DNA was then



extracted from the lysate through a silica gel membrane and several washing steps before the DNA solution was flown into another chamber for PCR. Real-time detection allowed for the extraction of amplification curves for quantification followed by melting curve analysis. A membrane integrated to the entrance of the PCR chamber of a microfluidic device was used for DNA extraction.<sup>58</sup> After lysis, DNA retained in the membrane was washed and dried before filling the chamber with PCR master mix releasing the retained DNA. An aluminium oxide membrane was used to extract and concentrate DNA from saliva samples on a microfluidic device.<sup>59</sup> Samples added to the device and flown through the membrane, afterwards reaction mix was added to the wells and real-time PCR performed.

Dielectrophoresis was used to capture pathogens from diluted whole blood into individual grooves on a slip chip.<sup>60</sup> Once the pathogens were separated, a washing step was carried out to remove PCR inhibitors by flushing with water. The grooves were split from the feeding channel through the slip chip and cells were lysed through heating. The slip chip consisted of four parallel sample processing channels, allowing for the parallel detection of four different pathogens. Electrophoresis is the motion of charged particles relative to a fluid caused by an electric field. As particle velocities depend on particle net charge and size, it can be used for separation. Thus, electrophoresis is commonly applied for post amplification analysis in the form of slab gel electrophoresis. Miniaturization of slab gel electrophoresis is often performed in capillary electrophoresis (CE) which can achieve higher separation performance due to the higher area-to-volume ratio countering Joule heating. As CE is a powerful method of post amplification analysis, various integrated time domain PCR CE devices have recently been developed. Le Roux *et al.* used a PCR device with an infrared laser for heating to perform PCR.<sup>61</sup> After amplification samples injected into a microfluidic device, amplicons were separated by amplicon length. This combination of methods was used to quantify short tandem repeats. Similarly, Manage *et al.* used a CE microfluidic device after amplification to separate and identify amplicons (Fig. 2A).<sup>62</sup> As starting PCR from whole blood in this case, the CE separation step is necessary to perform fluorescence detection after the amplification reaction. This approach has the further advantages that intercalating dyes can be used and targets can be identified by length. The integration of valve pumps into the design eliminates the need for external pumps. An interesting variant of PCR combined a PCR chamber with a set of CE channels.<sup>63</sup> After each cycle of PCR, a small sample could be drawn from the sample chamber and analyzed by CE. This allows for real-time detection of PCR products as well as their quantification. As a separation is performed in each detection step, this device is capable of monitoring several amplification processes in parallel with the utilization of a single intercalating dye.

### Centrifugal microfluidic devices

In centrifugal microfluidics, fluids are distributed using centrifugal force and the Coriolis effect. The microfluidic devices are designed in the format of compact discs (CD) due to the large availability of CD-based centrifugal microfluidic device, which

could be a potential candidate for point-of-care applications. A silicon/glass sandwich disk containing 24 channels and 313 microchambers with volumes of 1.5 nL was used to count *Salmonella enterica*.<sup>64</sup> A solution containing the cells was placed in an entrance port and distributed over the microchambers using centrifugal force, leading to a Poisson distribution of bacteria in the microwells. PCR was initiated by thermocycling the entire disc. After PCR, the microwells showing amplification were counted to determine the initial numbers of cells. The same device was used for reverse transcription PCR.<sup>65</sup> Direct reverse transcription of RNA was possible by using a robust DNA polymerase immune to inhibition. The introduction of ice valves next to the PCR chamber opened access to other device materials like polycarbonate.<sup>66</sup> The liquid was frozen next to the PCR chamber, forming a plug sealing the chamber to prevent sample evaporation.

The “LabDisk” format developed at the University of Freiburg in Germany offers a range of different functions integrated on a disc ranging from sample preparation to analysis after amplification. The lab disc contains four units, showing that a sample can divide into eight sub-volumes (Fig. 2D).<sup>67,68</sup> Air-dried reagents for the PCR reaction are dissolved by the sample inside of the chambers. Thermocycling is conducted in a commercial thermocycler. This design was later modified to accommodate a positive control, no template control as well as standards.<sup>69</sup> The modifications were done without compromising the ability of the disc to fit a commercial thermocycler and without the addition of a significant number of pipetting steps. This design was later used for forensic animal family identification.<sup>70</sup> In a further modification, making the design compatible with the “LabDisk Player”, a vapor diffusion barrier added to the system to limit the pressure in the system that can arise through boiling of the PCR solution.<sup>71</sup> The vapor diffusion barrier was created by adding a pneumatic chamber filled with air closer to the center of rotation than PCR chamber. In case of gas formation in the PCR chamber, the pressure is relieved through the pneumatic chamber preventing undesired pumping of the solution. This “LabDisk” was extended with DNA extraction capabilities, allowing sample preparation steps to be carried out with the device.<sup>72</sup> Bacteria samples from culture could be pipetted into the inlets of the device. An automated program on the “LabDisk” player centrifuged the device at the right frequencies to carry out cell lysis, DNA extraction and washing, multiplexed PCR and detection. This allowed a complete sample to answer assay to be carried out with minimal requirements for manual handling of the device. The development of these centrifugal devices is particularly interesting as they provide integrated devices. Only few manual handling steps are required using these devices.

### Arrays

As samples are spatially fixed in space domain PCR devices, they can be encoded by their spatial position in arrays for multiplexing applications. For example, samples can be printed in a water-in-oil emulsion on a Petri dish with sample volumes in the nanoliter range (Fig. 2B).<sup>73</sup> The sample array was created



using contact printing in a custom device utilizing standard pipette tips. Single droplets can be heated by a focused infrared laser. Fast passive cooling is achieved through the surrounding oil. In another example, solution containing cells was dispensed by a pipetting robot on surface patterned with hydrophilic spots.<sup>74</sup> The droplets were covered with mineral oil to prevent evaporation. Cells captured in droplets were lysed before adding reagents for reverse transcription by the pipetting robot. Xu *et al.* first conducted PCR in a device in which the sample was compartmentalized using a centrifugal device.<sup>75</sup> After amplification, targets could be identified with an array consisting of printed DNA probes for different targets on aldehyde-modified glass slides.

### Single cell devices

A number of diseases are generated through abnormalities in a small number of cells. Microfluidics offers great opportunities for solving this task as cells can be compartmentalized into small volumes. Microfluidic devices combine the isolation of cells with PCR to analyze the single cells. Single cells were printed into commercial PCR tubes.<sup>76</sup> A microfluidic inkjet printing-inspired technique was utilized in combination with a vacuum shutter to ensure that droplets containing a cell were dispersed in the tubes. In a different approach, single cells were captured in a flow through cell trapping unit.<sup>77</sup> A chemical lysis step was carried out and the released messenger RNA captured by magnetic oligo(dT)<sub>25</sub> beads. After moving the beads to a PCR chamber, reverse transcription PCR was carried out. The effects of methyl methanesulfonate on a single human cancer cell could be tested. Lim *et al.* encapsulated cells into individual droplets.<sup>78</sup> The droplets were collected in a vial and PCR was carried out in a benchtop PCR device using TaqMan probes. Droplets were inserted into a droplet sorting device and droplets showing positive amplification were separated to recover the entire genome of the cells. Cells can also be isolated by pneumatically actuated valves in microfluidic systems. The system allows the simultaneous capturing and reverse transcription PCR in a 96 × 96 array.<sup>79,80</sup> A further valve-based design contained 300 cell traps in individual chambers separated by valves.<sup>81</sup> The reverse transcription chamber could be used for a chemical cell lysis step, when a “one pot” reverse transcription PCR was carried out in the PCR chamber. In Fig. 2E, the arrangement of cell traps and reaction chambers were used for reverse transcription PCR. The design of chambers was optimized to obtain a homogeneous distribution of cells, while preventing mechanical damages to them. Furthermore, background noise and effects of PCR inhibitors were reduced and the device was used to measure the expression rates of microRNA in individual leukemia cells.

## Isothermal nucleic acid amplification

An elimination of the temperature steps during PCR leads to a significant simplification of PCR by removing the requirements of heating and cooling systems. This can be achieved by isothermal nucleic acid amplification methods. Different methods, like nucleic acid sequence-based amplification and self-sustained se-

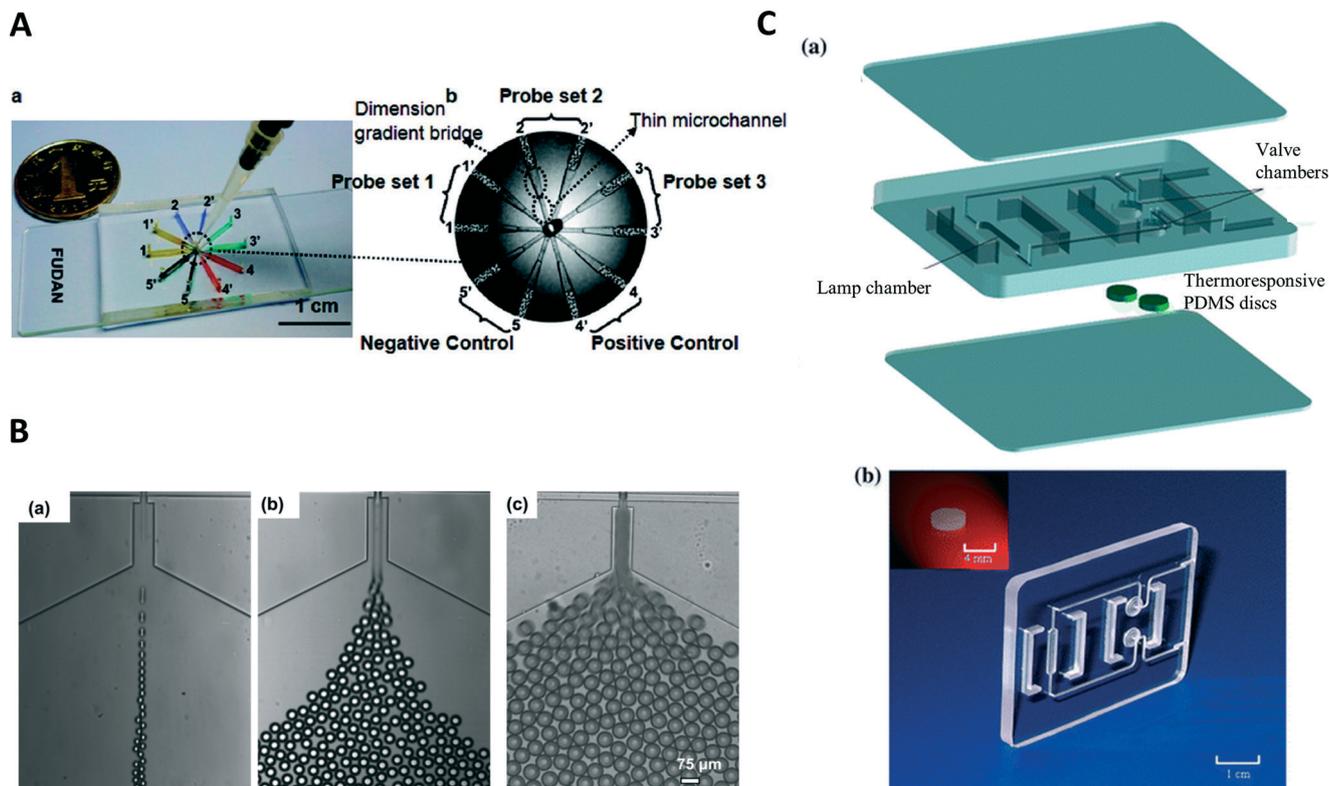
quence replication, are available for amplifying DNA at a single temperature. Another method is multiple displacement amplification, which has been demonstrated in microfluidic devices.<sup>82</sup> However, these methods often lack specificity, because they are conducted at relatively low temperatures ( $\approx 40$  °C). In 2000, Notomi *et al.* suggested loop-mediated isothermal amplification (LAMP), conducted at temperatures around 65 °C.<sup>83</sup> Four sets of primers and a DNA polymerase are employed. An inner primer complementary to the sense and antisense strand of the target DNA creates a double strand of DNA. The newly synthesized strand is released through strand replacement initiated by the outer primer. The released strand now serves as a target for the second set of inner and outer primer, resulting in a stem-loop structure. In following reactions, an inner primer initiates strand replacement polymerization, yielding the original stem-loop and a second stem loop. As a single temperature step is only required, devices are simpler and power consumption is lower. This makes LAMP an interesting option for point-of-care applications when combined with low sample volumes in microfluidic devices.

### Implementations of LAMP on microfluidic devices

Isothermal amplification methods have been used in combination with microfluidic devices. Applications in continuous flow in droplets can be found for the detection of DNA and microRNA.<sup>84</sup> A further application used droplets produced in a continuous manner.<sup>85</sup> The device was able to generate droplets, incubate the droplets at the temperature required for the LAMP reaction, and perform a fluorescence readout of droplets. A device with eight different sample chambers connected to a common central sample inlet was used for multiplexed LAMP reaction.<sup>86</sup> The device could be used to determine either three different human influenza A viruses or eight different swine flu viruses. Real-time detection of reaction progress was obtained through measurements of the redox current. Similar to solid phase PCR conducted in time domain devices, LAMP was carried out in agarose beads (Fig. 3B).<sup>87</sup> To prepare the beads, droplets were created from a solution containing agarose and the necessary reagents for LAMP. The droplets were collected for gelation initiated by cooling. Once the droplets had gellated into beads, the LAMP reaction was initiated by heating to 43 °C. Amplification was detected by monitoring fluorescence in a real-time manner using a camera equipped with matching fluorescence filters.

Integrated device are also proposed for this type of amplification. Wang *et al.* suggested a device combining extraction of RNA and reverse transcription LAMP reaction.<sup>88</sup> The samples are introduced to the device and RNA is extracted using magnetic beads. After washing steps, RNA transports to a LAMP chamber where a reverse transcription step is carried out. Afterwards products were analyzed using slab-gel electrophoresis of devices. The use of radial device design and centrifugal devices is also a popular choice for LAMP reactions. A radial device with four units containing five reaction chambers was designed.<sup>89</sup> Once the device was filled, it was transferred to a combined heating and fluorescence detection unit. The radial symmetry of the device was utilized as the device could be rotated to





**Fig. 3** (A) Microfluidic device for the parallel amplification of several genes. (a) Photograph of the device, the channels are filled with different dyes for visualization. (b) Schematic of the device illustrating placement of the probes for the different genes (reprinted with permission from ref. 95, Copyright 2010 American Chemical Society). (B) Generation of agarose precursor droplets with different flow conditions resulting in different droplet sizes. Beads are produced from these droplets and used for LAMP amplification in the solid agarose medium (reprinted with permission from ref. 87, Copyright 2012 AIP Publishing LLC). (C) Disposable polycarbonate cassette for LAMP reactions in point-of-care applications, exploded view of the PCR cassette (top) and photograph of the assembled cassette (bottom) including an image of the thermoresponsive PDMS disc (inset) (reprinted with permission from ref. 93, Copyright 2011 Springer).

measure fluorescence in the different reaction chambers using the confocal detection unit. Through this setup, real-time amplification curves could be extracted for the LAMP reaction. Other radial device designs utilize centrifugal forces to reduce the number of required handling steps. One disc-shaped device consisted of 24 reaction wells in which primers for the multiplexed detection of ten different pathogenic bacteria were pipetted and dried.<sup>90</sup> A solution containing the extracted DNA from fishes out of aquaculture was pipetted into the inlet of the device and divided between the reaction wells using centrifugal forces. LAMP reaction and real-time detection were performed on a device suitable for the temperature manipulation and fluorescence measurement of these disks. Sayad *et al.* used an integrated CD device for the detection of *Salmonella*.<sup>91</sup> The disc contained the steps required for reagent preparation, LAMP, and detection of amplification products. All of these steps could be automatically performed by a custom-made computer system.

#### LAMP for point-of-care devices

A simple point-of-care device was constructed by filling samples for LAMP amplification into a glass capillary.<sup>92</sup> The capillary was sealed on both ends using epoxy glue and was sandwiched between two pocket warmers to obtain an ampli-

fication temperature. Amplification was detected by naked eye using a dye and a portable UV light. This allows for performing a LAMP reaction completely independent of external power supplies. Liu *et al.* produced a disposable polycarbonate cassette with a size of 46 mm × 36 mm × 3.7 mm (Fig. 3C).<sup>93</sup> It contained two chambers for facilitating an isothermal amplification reaction and could be filled with only few pipetting steps due to thermoresponsive valves. An electrochemical DNA sensor was used by Hsieh *et al.* to detect the presence of a target DNA.<sup>94</sup> The reaction is performed before the electrochemical detection. Ten chambers connected to a central inlet were used for a multiplexed LAMP reaction.<sup>95</sup> Each chamber was functionalized with probes for a different target, allowing the multiplexed detection of ten different targets (Fig. 3A). After one hour of incubation at 63 °C, successful amplification could be detected by the naked eye through the precipitation of magnesium phosphate in the chambers, making this simple detection device suitable for point-of-care applications.

## Digital PCR

Digital PCR is an advanced refinement of PCR. While in PCR the process is carried out in a bulk solution, digital PCR



divides the sample into hundreds to thousands of sub-samples. The sample is partitioned and sub-samples contain less than one molecule of DNA on average with the molecule distribution following a Poisson distribution. After a digital PCR runs the number of partitions, showing that amplification can be counted and the starting number of molecules is estimated through the Poisson distribution. Digital PCR has the advantages over conventional qrtPCR that absolute quantification does not require the measurement of a standard curve. Furthermore, digital PCR offers advantages when working with minor differences in starting molecule count or target DNAs that are rare in the sample. A requirement for digital PCR is the partitioning of the sample in sufficiently small sub-samples. Microfluidics offers several options for performing this task, either by creating droplets or microwells, which can be filled through a variety of techniques.

### Applications

Droplet-based digital PCR has been used for several applications. Zhu *et al.* used droplet-based digital PCR for the detection of HER2 and CEP17.<sup>96</sup> The same samples were measured using alternative techniques, such as immunochemistry and fluorescence *in situ* hybridization to test patients with breast and gastric carcinomas. A high concordance of droplet digital PCR with the standard methods showed a high sensitivity. Another study compared the ability of digital PCR to amplify rare targets compared to qrtPCR.<sup>97</sup> In this study, KRAS mutation was detected in patient samples. While digital PCR was able to correctly identify 11 out of 14, adenomas qrtPCR only identified 9, demonstrating that the advantage of digital PCR has over conventional PCR in handling rare targets. These advantages were utilized to measure DNA contaminations of protein drugs produced by yeast cells.<sup>98</sup> For these experiments, the protein drug directly added to the digital PCR droplets. Thermal cycling carried out with primers for yeast, which produced the drug in this case. The higher amounts of sample could be tested in comparison to qrtPCR methods and the experiment protocol could be simplified.

Floren *et al.* used digital PCR to identify several different species.<sup>99</sup> Additionally they were able to determine from what tissue samples were taken due to minute differences in the amount of mitochondrial DNA, not detectable by qrtPCR. DNA from soil samples was quantified by Hoshino *et al.* using droplet digital PCR and standard qrtPCR.<sup>100</sup> While qrtPCR was prone to underestimating the DNA content of the samples due to inhibition, digital PCR showed to be more robust. While for some samples the inhibition effects could be overcome by dilution prior to amplification, other samples still displayed an underestimation of DNA content by qrtPCR compared to digital PCR. The robustness of digital PCR against inhibition was also confirmed by Sedlak *et al.* who looked into quantification of inhibition prone samples by qrtPCR and digital PCR.<sup>101</sup> The isothermal nucleic acid amplification is generally affected by inefficiencies in the reaction compared to normal PCR. Experiments were carried out to evaluate the suitability of isothermal amplification for digi-

tal PCR.<sup>102</sup> It found that the reaction speed of the isothermal reaction did not correlate with the digital efficiency. Furthermore, no correlation between reaction speed and specificity was found in the experiments. Digital PCR can also be used to gain into early cycles of PCR.<sup>103</sup> The ogive shape of digital PCR assays was decoded and information about the amplification reaction could be gained, especially about early cycle amplification.

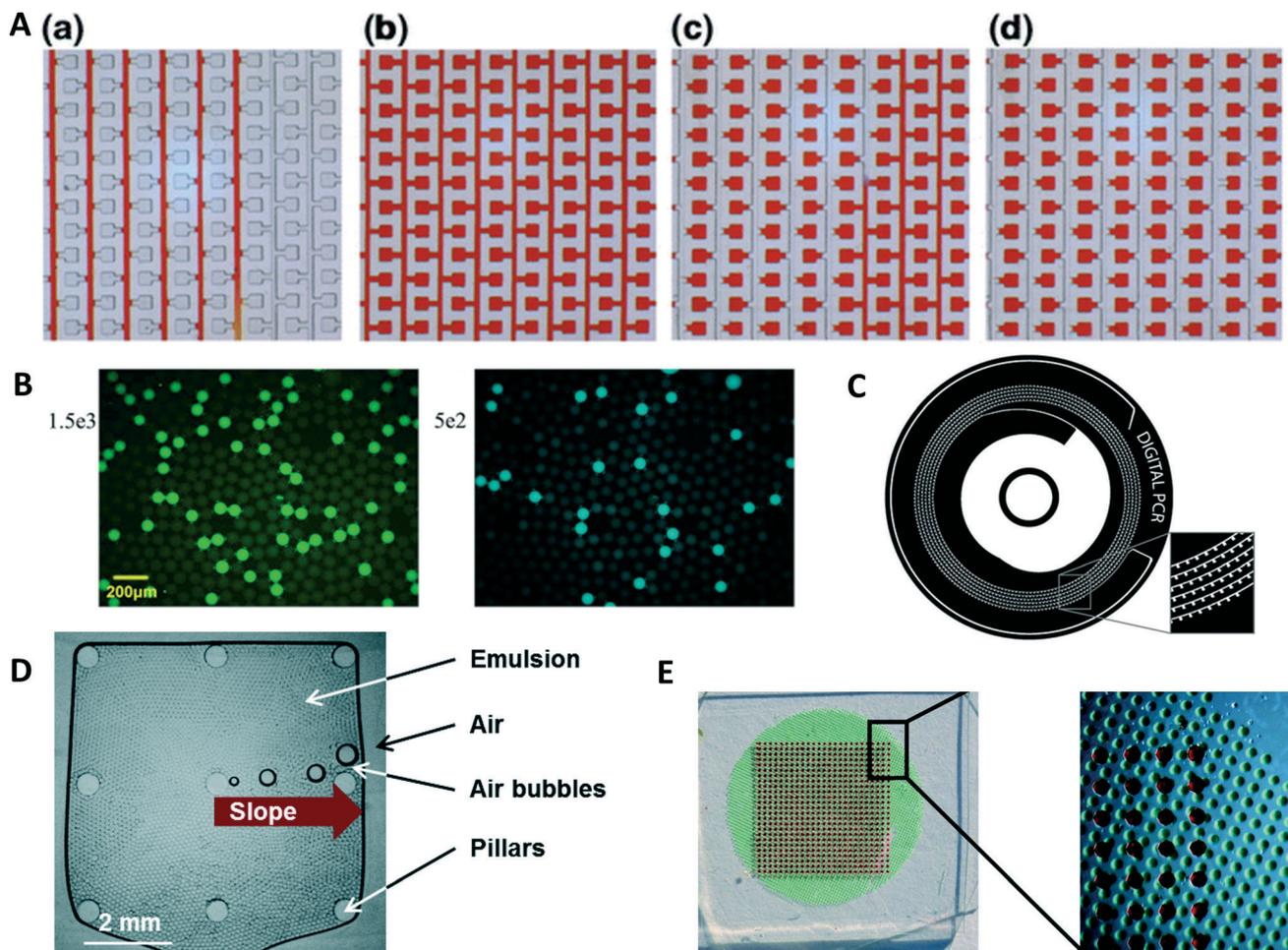
### Multiplexed digital PCR

A device with 48 individual panels, each panel containing 770 partitions, was used for duplexed digital PCR.<sup>104</sup> The device was used to detect genetically modified organisms by measuring the amount of modified genome in a sample. In another approach to droplet-duplexed digital PCR, *Escherichia coli* and *Listeria monocytogenes* were detected in parallel (Fig. 4B).<sup>105</sup> Droplets contained primers and TaqMan probes for both bacteria. To differentiate between the two species, colored reporters were used for the TaqMan probes. After amplification in a droplet array, two fluorescence images of the droplet array were taken. From the separate images, the number of amplified targets could be determined for each target. While the previous two methods employ spatially separated samples on the same device or different fluorescence wavelength for demultiplexing the results, Zhong *et al.* used a different approach.<sup>106</sup> In this assay, two different fluorophores were used to differentiate five different targets. To allow five different targets in the multiplexed reaction, linear combinations of the two dyes are formed using different brightness for the probes. This creates a unique fluorescent signature for each target. The same technique can also be used to run duplexed digital PCR using only one fluorescent reporter with different fluorescence intensities.

### Designs

Several different options have been suggested for the generation of reaction compartments using different microfluidic techniques. A cross-junction has been used to generate homogenous water-in-oil droplets in a high-frequency manner.<sup>107</sup> The droplets contained microRNA from lung cancer tissues and were incubated in a chamber, holding approximately  $2 \times 10^4$  sample droplets. The density of sample volumes per area was increased in microwell assays.<sup>108</sup> A device featuring 82 000 wells with a volume of 36 fL only occupied an area of  $2 \text{ mm} \times 2 \text{ mm}$ , giving a well density of 20 000 wells  $\text{mm}^{-2}$ . After removing the excess sample solution, wells were sealed by applying an overpressure to a second chamber above the microwell structure. A PDMS membrane was bent to close the wells preventing possible cross-contamination. Heating and cooling for PCR was performed by a Peltier heater. A megapixel digital PCR device was created with a density of 440 000 sample wells  $\text{cm}^{-2}$ .<sup>109</sup> The PDMS device consisted of channels from which several dead end chambers branched off. To fill the chambers, the sample solution was flown into the device under pressure, causing the air in the chambers to diffuse out through the PDMS. The channels





**Fig. 4** (A) A digital PCR device with a serpentine channel design with reaction chambers branching of the main channel. (a) In a first step, sample (replaced here with red dye) is flown through the degassed PDMS device. Once the reaction chambers are filled with sample (b) oil is flown through the channels (c) separating the sample into individual volumes (d) (reprinted with permission from ref. 112, Copyright 2015 Springer). (B) Multiplexed droplet digital PCR of *E. coli* and *L. monocytogenes*. Two different TaqMan probes are used, labelled with FAM *E. coli* (left) and VIC for *L. monocytogenes* (right) (reprinted with permission from ref. 105, Copyright 2015 Elsevier). (C) Centrifugal microfluidic device for the compartmentalization of sample. The sample is introduced in the central inlet chamber and then divided into the individual chambers by centrifugal force. Cross talk is prevented by filling the spiral channel with oil before amplification (reprinted with permission from ref. 116, Copyright 2014 Springer). (D) Photograph of an amplification chamber in a centrifugal droplet digital PCR device. A pyramid structure in the center of the chamber guides air bubbles developing during thermal cycling to the edge of the chamber (reprinted with permission from ref. 118, Copyright 2016 Royal Society of Chemistry). (E) Image of a device featuring an array of reaction chambers for amplification (filled with red dye). Below the array there is a vapor proof-layer preventing sample evaporation from the PDMS device (filled with green dye for visualization) (reprinted with permission from ref. 113, Copyright 2015 Royal Society of Chemistry).

were separated by flowing an immiscible oil through the channel to isolate the chambers. A microarray scanner was used for image analysis after the PCR run. A microfluidic device with a T-junction was used to generate droplets.<sup>110</sup> The device consisted of two inlets, one for an oil phase and another one for the sample and a reservoir to collect the droplets. A flow was created from the air in the reservoir diffusing into the degassed PDMS creating a negative pressure. Droplet dispensing frequencies of up to 12 Hz were achieved and droplet size was dependent on the channel size.

Thompson *et al.* proposed a device with a serpentine channel with 1020 reaction chambers.<sup>111</sup> The device was filled by applying a vacuum to the outlet, while pipetting sample solution in the inlet. Once chambers were filled, an

oil phase was flown through the channel to separate the chambers. A similar device had several parallel channels with 1040 chambers branching of them with volumes of 2.08 nL (Fig. 4A).<sup>112</sup> A fluorinated oil phase is filled into the channel. A device containing 650 chambers with volumes of 6.28 nL was filled by negative pressure in a chamber separated from the reaction chambers by a thin layer of PDMS (Fig. 4E).<sup>113</sup> Once chambers were filled, they were capped with silicon oil and an adhesive cover layer. To stop evaporation of sample during digital PCR, the vapor proof-layer consisting of an array of chambers containing water was placed instead of the negative pressure chamber. During PCR, a vapor proof-layer keeps the PDMS saturated with water to minimize sample evaporation which is can be a major problem when working



with small sample volumes. Agarose solution, forming nanoliter-sized beads, was used to perform digital PCR.<sup>114</sup> While forward primers were in free solution, the reverse primers conjugated to the agarose. As the agarose melts during PCR, the efficiency of the reaction is not influenced by the conjugation of the primers to the agarose matrix. Digital PCR in microliter wells was used to count nanoparticles.<sup>115</sup> The nanoparticles were tagged with DNA and diluted to a concentration of less than one particle per microliter. The digital PCR could be performed and the number of particles was counted by determining the number of positive microwells and the Poisson statistics.

### Integrated systems

The function of microfluidic PCR devices has been integrated into the design to simplify sample preparation and integrate detection of amplification products. A spinning disc for the enrichment and quantification of rare DNA variants was suggested by Sundberg *et al.* (Fig. 4C).<sup>116</sup> The disc consisted of a central loading chamber in which the sample and oil phase were injected. It demonstrated that compartments were generated through chambers rather than droplets on a centrifugal platform. The sample was separated over 1000 wells during rotation of the disc and crosstalk was eliminated through the oil phase. The digital PCR was termed quasi-digital as all chambers contained a number of copies of wild type DNA which was not the target of amplification. In another example of a spinning disc device, monodispersed droplets were created by centrifugal step emulsification.<sup>117</sup> The droplets could be created using a chamber filled with oil in which the aqueous sample injected through a nozzle over a step. Liquids were moved by centrifugal force which was created by the “LabDisc Player”. The player heated the sample to perform recombinase polymerase amplification, another method of isothermal nucleic acid amplification. Detection of successful amplification was also carried out by the “LabDisc Player” after amplification. To carry out conventional PCR protocols with the device, a method had to be developed to remove air bubbles during thermal cycling. Thus, the disk was extended with a pyramid structure in the middle of the amplification chamber and a capillary channel leading back to the sample inlet (Fig. 4D).<sup>118</sup> When gas bubbles developed, they were guided towards the edge of the amplification chamber through the pyramid structure. Once they arrived at the edge, they were removed from the chamber through the capillary channel enabling the conduction of thermal cycling digital PCR. This development allowed for the effective removal of gas bubbles, which could lead to issues in the analysis and droplet stability. This design was later modified to become independent of the “LabDisc Player”.<sup>119</sup> The smaller device contained the same features as the disk device was miniaturized to fit a mini-centrifuge. Isothermal digital PCR was conducted in an *in situ* cycler and transferred to a microarray scanner for fluorescence scanning.

Slip chips consist of two plates in contact with each other.<sup>120</sup> While the bottom plate contains reagent wells, the top plate acts as a cover. Furthermore, the bottom plate features ducts forming a fluidic path with wells in the top plate. A first reagent can be preloaded into the wells of the top device, while a second reagent can be filled into the wells of the bottom plate. Once the chambers in the top plate are filled, the top plate can be moved to align with the wells in the bottom plate. This slip chip technology was used to compartmentalize a sample for digital PCR to detect HIV RNA.<sup>121</sup> In further work, the robustness of digital isothermal amplification was compared against qrtPCR and was found favorable for point-of-care applications.<sup>122</sup> Thus, a fluorescence readout unit was developed for the slip chip using a smartphone-camera equipped with an objective lens and fluorescence filters. Digital PCR offers great advantages over conventional qrtPCR, such as robustness against inhibitors and quantification without the requirement for standard curves. Due to these advantages, digital PCR could be able to replace qrtPCR in a near future.

## Commercial PCR devices

The microfluidic PCR devices integrated into commercial thermal cyclers and real-time thermal cyclers. Furthermore, digital PCR instruments were introduced to the market, which requires microfluidics for sample partition.

### Integrated systems

A number of integrated, commercial thermal cyclers have been developed in the last years. They employ a variety of microfluidic techniques to integrate sample preparation, amplification and detection. Furthermore, integrated systems can allow for a test to be conducted in a doctor's office instead of requiring an external laboratory for testing. The GenePoc™ system uses a centrifugal platform for sample preparation as well as PCR on the same device.<sup>123</sup> The system is capable of testing eight different samples in parallel, while testing for up to 12 different targets per sample. Similarly, the cobas® Liat PCR system from Roche Diagnostics uses prefilled reagent strips to perform sample extraction, PCR, and subsequent detection.<sup>124</sup> Currently, three different assays are available for the system that can test for either Group A *Streptococcus* DNA or RNA of influenza A/B. The device can test one sample per run in 20 minutes due to the small sample volumes. In these small and low throughput systems, microfluidics is used to prepare samples for PCR followed by integrated amplification and detection on the same devices. The cross-contamination can be prevented and tests can be conducted easily. This allows medical doctors to conduct test, without requiring a specialised laboratory and trained laboratory staff. Other systems are aimed at high-throughput applications. The main advantage of microfluidics is that sample preparation can be automated to reduce the handling times. Two different approaches are most prominent. Reagents can be provided in reagent strips, like in the case of the BD



MAX™ system.<sup>125</sup> Another option is to provide reagents in blister films joined by channels as done in the FilmArray® system from bioMérieux.<sup>126</sup> Alternatively, cartridges can be provided for different assays as done in the case of Idylla™ system from Biocartis<sup>127</sup> or the GeneXpert® system from Cepheid.<sup>128</sup> All of these systems are able to run multiplexed PCR assays including automated sample preparation as well as detection.

The VerePLEX™ Biosystem from Veredus Laboratories does not offer automated sample preparation.<sup>129</sup> The system uses silicon-based microfluidic devices with a volume of 25 µL for end point PCR and is capable of performing up to 50 different tests on a single sample in a single reaction. After amplification, samples are hybridized on a microarray of up to 400 individual spots, allowing for multiplexed detection. The advantages of miniaturization are used to increase the number of detectable amplifications products per PCR run from up to eight in conventional TaqMan probe assays to up to 50. A study was carried out evaluating the performance of the cobas® Liat system for the detection of Influenza A and B viruses from throat swabs.<sup>130</sup> The study found a high sensitivity of 99.2% for Influenza A and 100% for Influenza B. Specificity for both targets was 100% in the study. A panel for testing 21 common viral and bacterial respiratory pathogens was developed for the FilmArray® platform.<sup>131</sup> The panel performed automated nested, multiplexed PCR on cultured organisms as well as nasal aspirants collected from patients, showing an analytical and clinical sensitivity as well as specificity comparable to other diagnostic platforms. A detection and identification of human influenza viruses was performed on the VerePLEX™ system.<sup>132</sup>

### Digital PCR systems

The microfluidics can achieve a division of samples into several thousand sub-volumes in the pico to nanoliter range. Three different microfluidic approaches can be found in commercial digital PCR devices. The QuantStudio® 3D Digital PCR system by Thermo Fisher Scientific uses an array of reaction chambers with volumes of 809 picoliter.<sup>133</sup> The array is filled using a doctor blading method, before thermal cycling is performed on a Peltier system. After amplification, arrays are analyzed on a separated array reader to measure fluorescence on two different wavelengths. Other methods use droplets, such as QX200™ Droplet Digital™ PCR system from Bio-Rad Laboratories<sup>134</sup> or the RainDrop® system from RainDance™ Technologies.<sup>135</sup> Both systems generate droplets on a microfluidic device. Droplets are collected in a vial and thermal cycling is performed on a conventional PCR machine. After amplification, droplets were introduced into a second microfluidic device to measure the fluorescence intensity of droplets. Both systems are capable of measuring two different colors of fluorescence probes, enabling multiplexing. While the QX200™ system can only differentiate two targets, the RainDance™ system can differentiate up to ten targets by applying different concentrations of

probes for targets. The Biomark HD system from Fluidigm® uses a system of integrated fluidic circuits (IFCs).<sup>136</sup> Reaction compartments are generated by separating the sample into sub-volumes using microfluidic valves. Samples are divided in 9180 partitions, each with a volume of 6 nL. The device has five fluorescence channels for detection. These commercial digital PCR devices are used for a variety of tasks. The droplet-based microfluidics from Bio-Rad was used for the differentiation of twelve different genetically modified maize lines,<sup>137</sup> while the droplet digital PCR device from RainDance™ was employed to detect and quantify of three cancer mutations.<sup>138</sup> The Fluidigm system was used for testing the suitability of digital RNA quantification.<sup>139</sup>

## Conclusions and perspectives

Since the first PCR in microfluidic devices was demonstrated, a number of techniques have been developed. They aim to improve the efficiency, reduce the complexity of devices, and integrate other functions, such as sample preparation and detection. However, a number of time and space domain PCR devices do not address the issue of providing simple user interfaces. While microfluidic devices offer great advantages over conventional bench-top systems, typical PCR users do not have access to microfluidic techniques, often lacking equipment or skill required for the operation. This provides a significant barrier for PCR users, preventing them from adapting microfluidic techniques. Additionally, the use of microfluidic devices can be difficult for users who do not have access to fabrication facilities. Only a few examples, like the “LabDisk”, went the additional step of not only designing the microfluidic device but also methods facilitating simple use of the microfluidic devices. Future research, especially in time and space domain PCR devices, should focus on making the microfluidic methods easily available for users. Through the origin of LAMP, technical requirements for devices were reduced. Furthermore, isothermal nucleic acid amplification methods show a higher robustness against temperature variations and inhibition compared to conventional PCR. This makes isothermal nucleic acid amplification an interesting candidate for point-of-care applications. However, like conventional PCR methods, access for end users should be as simple as possible. The higher requirements for primer design and advances in conventional PCR devices provide a further obstacle for isothermal devices. The microfluidic-based digital PCR has become a valuable alternative to conventional qPCR and has the potential to revolutionize the field. Digital PCR allows absolute quantification without the requirement for standard curves. Unlike conventional PCR, rare targets can be amplified and quantified with high precision and inhibition has less severe effects. Digital PCR depends on a compartmentalization of DNA molecules into individual volumes from bulk solution. Although digital PCR is already widely used, the impact of the technology could be extended and the technique might replace conventional qPCR. To achieve this, research should be conducted to allow for as



simple handling of digital PCR devices as for qrtPCR devices. Furthermore, automatic, high-throughput, and multiplexed digital PCR has to gain the same ease of use as currently offered by qrtPCR systems.

The commercial integrated PCR systems show that the automation possibilities and small fluidic handling are required. Thus, integrated microfluidic PCR devices have great advantages for high-throughput applications. Through the simple operation and high robustness against cross-contamination, integrated PCR devices allow for tests to be conducted in a medical doctor's office, eliminating the need for external laboratories. Although tests can be conducted in 20 minutes with the cobas® Liat PCR system, current applications do not seem to profit from further reductions in reaction time. Although assay costs could be decreased by a reduction of sample volumes, the cost advantages are not significant enough for most applications. Legal barriers can provide another hurdle for microfluidic systems; the commercial devices introduced here have less than 50 FDA-approved tests. This also prevents a wider application of portable point-of-care systems, in combination with the assay costs which are relatively high for developing countries. As microfluidics is the only way of creating digital PCR devices, a bigger variety of techniques is adapted in commercial digital PCR devices. Systems like RainDrop® or QX200™ require three different devices to perform the analysis. Utilizing the advantages of microfluidics has to offer integration of different functions and automation. Simplification of digital PCR processes and reduction of required equipment numbers could help to get more attention from end users.

Alternative methods of DNA and RNA detection, like DNA arrays and next generation sequencing that have been developed in the last years, offer another interesting perspective for development of microfluidic-based PCR. While PCR can detect single molecules of DNA or RNA in a rapid manner, its specificity is determined by the primer design. Furthermore, the detection of single nucleotide polymorphisms or repeats with PCR is difficult. Detection is significantly easier using DNA arrays. With sequencing methods, like pyrosequencing, the entire sequence of the target can be found. However, both DNA arrays and sequencing methods often need a minimum amount of DNA or RNA and thus require a sample preparation steps by PCR. Therefore, an integration of PCR and DNA arrays or sequencing methods on the same microfluidic device is an interesting option for future research.

## Acknowledgements

This work was also supported by BioNano Health-Guard Research Center funded by the Ministry of Science, ICT & Future Planning (MSIP) of Korea as Global Frontier Project (grant number H-GUARD\_2014M3A6B2060503), Republic of Korea. This research was supported by the National Research Foundation (NRF) of Korea grant funded by the MSIP (Grant number 2016R1A6A1A03012845). This work was also supported by

the NRF of Korea grant funded by the MSIP (Grant number 2015R1A2A1A15054236).

## References

- 1 K. Mullis, F. Faloona, S. Scharf, R. Saiki, G. Horn and H. Erlich, *Cold Spring Harbor Symp. Quant. Biol.*, 1986, **51**, 263–273.
- 2 R. Saiki, D. Gelfand, S. Stoffel, S. Scharf, R. Higuchi, G. Horn, K. Mullis and H. Erlich, *Science*, 1988, **239**, 487–491.
- 3 H. M. Temin, *Virology*, 1963, **20**, 577–582.
- 4 F. F. Chehab and Y. W. Kan, *Proc. Natl. Acad. Sci. U. S. A.*, 1989, **86**, 9178–9182.
- 5 F. Osman, E. Hodzic, A. Omanska-Klusek, T. Olineka and A. Rowhani, *J. Virol. Methods*, 2013, **194**, 138–145.
- 6 Y. Kim, Y. Choi, B.-Y. Jeon, H. Jin, S.-N. Cho and H. Lee, *Yonsei Med. J.*, 2013, **54**, 1220–1226.
- 7 J. F. Mehrabadi, P. Morsali, H. R. Nejad and A. A. Imani Fooladi, *J. Infect. Public Health*, 2012, **5**, 263–267.
- 8 A. Abd-Elmagid, P. A. Garrido, R. Hunger, J. L. Lyles, M. A. Mansfield, B. K. Gugino, D. L. Smith, H. A. Melouk and C. D. Garzon, *J. Microbiol. Methods*, 2013, **92**, 293–300.
- 9 M. Safdar and M. F. Abasiyanik, *Appl. Biochem. Biotechnol.*, 2013, **171**, 1855–1864.
- 10 A. Manz, N. Graber and H. M. Widmer, *Sens. Actuators, B*, 1990, **1**, 244–248.
- 11 P. Wilding, M. A. Shoffner and L. J. Kricka, *Clin. Chem.*, 1994, **40**, 1815–1818.
- 12 M. A. Northrup, C. Gonzalez, S. Lehwew and R. Hills, *Micro Total Analysis Systems: Proceedings of the  $\mu$ TAS '94 Workshop*, Springer, 1995, p. 139.
- 13 J. Cheng, M. A. Shoffner, K. R. Mitchelson, L. J. Kricka and P. Wilding, *J. Chromatogr. A*, 1996, **732**, 151–158.
- 14 M. U. Kopp, A. J. d. Mello and A. Manz, *Science*, 1998, **280**, 1046–1048.
- 15 P. A. Auroux, Y. Koc, A. deMello, A. Manz and P. J. R. Day, *Lab Chip*, 2004, **4**, 534–546.
- 16 L. Chen, A. Manz and P. J. R. Day, *Lab Chip*, 2007, **7**, 1413–1423.
- 17 D. Moschou, N. Vourdas, G. Kokkoris, G. Papadakis, J. Parthenios, S. Chatzandroulis and A. Tserepi, *Sens. Actuators, B*, 2014, **199**, 470–478.
- 18 M. L. Ha and N. Y. Lee, *Food Control*, 2015, **57**, 238–245.
- 19 J. Wu, W. Guo, C. Wang, K. Yu, Y. Ma, T. Chen and Y. Li, *Cell Biochem. Biophys.*, 2015, **72**, 605–610.
- 20 A. Harandi and T. Farquhar, *J. Micromech. Microeng.*, 2014, **24**, 115009.
- 21 H. Tachibana, M. Saito, K. Tsuji, K. Yamanaka, L. Q. Hoa and E. Tamiya, *Sens. Actuators, B*, 2015, **206**, 303–310.
- 22 J. J. Chen, M. H. Liao, K. T. Li and C. M. Shen, *Biomicrofluidics*, 2015, **9**, 014107.
- 23 W. Wu and N. Y. Lee, *Anal. Bioanal. Chem.*, 2011, **400**, 2053–2060.
- 24 Y. Schaerli, R. C. Wootton, T. Robinson, V. Stein, C. Dunsby, M. A. A. Neil, P. M. W. French, A. J. deMello, C. Abell and F. Hollfelder, *Anal. Chem.*, 2009, **81**, 302–306.



- 25 L. Jiang, M. Mancuso, Z. Lu, G. Akar, E. Cesarman and D. Erickson, *Sci. Rep.*, 2014, **4**, 4137.
- 26 R. Snodgrass, A. Gardner, L. Jiang, C. Fu, E. Cesarman and D. Erickson, *PLoS One*, 2016, **11**, e0147636.
- 27 Y. Li, C. Zhang and D. Xing, *Microfluid. Nanofluid.*, 2010, **10**, 367–380.
- 28 B. Miao, N. Peng, L. Li, Z. Li, F. Hu, Z. Zhang and C. Wang, *Sensors*, 2015, **15**, 27954.
- 29 L. Chen, J. West, P.-A. Auroux, A. Manz and P. J. R. Day, *Anal. Chem.*, 2007, **79**, 9185–9190.
- 30 C. Zhang, H. Wang and D. Xing, *Biomed. Microdevices*, 2011, **13**, 885–897.
- 31 A. G. Sciancalepore, A. Polini, E. Mele, S. Girardo, R. Cingolani and D. Pisignano, *Biosens. Bioelectron.*, 2011, **26**, 2711–2715.
- 32 Z. Li, T.-Y. Ho and K. Chakrabarty, *ACM Transactions on Design Automation of Electronic Systems*, 2016, **21**, 1–27.
- 33 R. Prakash, K. Pabbaraju, S. Wong, A. Wong, R. Tellier and K. Kaler, *Micromachines*, 2015, **6**, 63.
- 34 M. J. Jebrail, R. F. Renzi, A. Sinha, J. Van De Vreugde, C. Gondhalekar, C. Ambriz, R. J. Meagher and S. S. Branda, *Lab Chip*, 2015, **15**, 151–158.
- 35 Y.-F. Hsieh, E. Yonezawa, L.-S. Kuo, S.-H. Yeh, P.-J. Chen and P.-H. Chen, *Appl. Phys. Lett.*, 2013, **102**, 173701.
- 36 M. Krishnan, V. M. Ugaz and M. A. Burns, *Science*, 2002, **298**, 793.
- 37 N. Agrawal, Y. A. Hassan and V. M. Ugaz, *Angew. Chem., Int. Ed.*, 2007, **46**, 4316–4319.
- 38 K. Chan, P.-Y. Wong, P. Yu, J. Hardick, K.-Y. Wong, S. A. Wilson, T. Wu, Z. Hui, C. Gaydos and S. S. Wong, *PLoS One*, 2016, **11**, e0149150.
- 39 J. S. Farrar and C. T. Wittwer, *Clin. Chem.*, 2015, **61**, 145–153.
- 40 Y. Ouyang, G. R. M. Duarte, B. L. Poe, P. S. Riehl, F. M. dos Santos, C. C. G. Martin-Didonet, E. Carrilho and J. P. Landers, *Anal. Chim. Acta*, 2015, **901**, 59–67.
- 41 A. M. Schrell and M. G. Roper, *Analyst*, 2014, **139**, 2695–2701.
- 42 N. Pak, D. C. Saunders, C. R. Phaneuf and C. R. Forest, *Biomed. Microdevices*, 2012, **14**, 427–433.
- 43 J. H. Son, B. Cho, S. Hong, S. H. Lee, O. Hoxha, A. J. Haack and L. P. Lee, *Light: Sci. Appl.*, 2015, **4**, e280.
- 44 C. D. Ahrberg, B. R. Ilic, A. Manz and P. Neuzil, *Lab Chip*, 2016, **16**, 586–592.
- 45 C. D. Ahrberg, A. Manz and P. Neuzil, *Anal. Chem.*, 2016, **88**, 4803–4807.
- 46 C. D. Ahrberg, A. Manz and P. Neuzil, *Sci. Rep.*, 2015, **5**, 11479.
- 47 E. K. Wheeler, C. A. Hara, J. Frank, J. Deotte, S. B. Hall, W. Benett, C. Spadaccini and N. R. Beer, *Analyst*, 2011, **136**, 3707–3712.
- 48 T. Houssin, J. Cramer, R. Grojsman, L. Bellahsene, G. Colas, H. Moulet, W. Minnella, C. Pannetier, M. Leberre, A. Plecis and Y. Chen, *Lab Chip*, 2016, **16**, 1401–1411.
- 49 S. O. Sundberg, C. T. Wittwer, R. M. Howell, J. Huuskonen, R. J. Pryor, J. S. Farrar, H. M. Stiles, R. A. Palais and I. T. Knight, *Clin. Chem.*, 2014, **60**, 1306–1313.
- 50 C. Hurth, J. Yang, M. Barrett, C. Brooks, A. Nordquist, S. Smith and F. Zenhausern, *Biomed. Microdevices*, 2014, **16**, 905–914.
- 51 Y. Sun, R. Dhumpa, D. D. Bang, J. Hogberg, K. Handberg and A. Wolff, *Lab Chip*, 2011, **11**, 1457–1463.
- 52 S. Jung, J. Kim, D. J. Lee, E. H. Oh, H. Lim, K. P. Kim, N. Choi, T. S. Kim and S. K. Kim, *Sci. Rep.*, 2016, **6**, 22975.
- 53 K. Qin, X. Lv, Q. Xing, R. Li and Y. Deng, *Anal. Methods*, 2016, **8**, 2584–2591.
- 54 S. Ma, D. N. Loufakis, Z. Cao, Y. Chang, L. E. K. Achenie and C. Lu, *Lab Chip*, 2014, **14**, 2905–2909.
- 55 C.-Y. Chao, C.-H. Wang, Y.-J. Che, C.-Y. Kao, J.-J. Wu and G.-B. Lee, *Biosens. Bioelectron.*, 2016, **78**, 281–289.
- 56 P. Liu, X. Li, S. A. Greenspoon, J. R. Scherer and R. A. Mathies, *Lab Chip*, 2011, **11**, 1041–1048.
- 57 S.-M. Zhao, L. Zhu, C.-C. Zhu, Y. Li, H.-D. Wang, L. Zhang, D.-W. Du, G.-Q. Deng, A. Wang and Y. Liu, *Fenxi Huaxue*, 2014, **42**, 1393–1399.
- 58 X. Qiu and M. G. Mauk, *Microsyst. Technol.*, 2014, **21**, 841–850.
- 59 E. A. Oblath, W. H. Henley, J. P. Alarie and J. M. Ramsey, *Lab Chip*, 2013, **13**, 1325–1332.
- 60 D. Cai, M. Xiao, P. Xu, Y. Xu and W. Du, *Lab Chip*, 2014, **14**, 3917–3924.
- 61 D. Le Roux, B. E. Root, C. R. Reedy, J. A. Hickey, O. N. Scott, J. M. Bienvenue, J. P. Landers, L. Chassagne and P. de Mazancourt, *Anal. Chem.*, 2014, **86**, 8192–8199.
- 62 D. P. Manage, Y. C. Morrissey, A. J. Stickel, J. Lauzon, A. Atrazhev, J. P. Acker and L. M. Pilarski, *Microfluid. Nanofluid.*, 2010, **10**, 697–702.
- 63 Y. Liu, C. Li, Z. Li, S. D. Chan, D. Eto, W. Wu, J. P. Zhang, R.-L. Chien, H. G. Wada, M. Greenstein and S. Satomura, *Electrophoresis*, 2016, **37**, 545–552.
- 64 S. Furutani, H. Nagai, Y. Takamura and I. Kubo, *Anal. Bioanal. Chem.*, 2010, **398**, 2997–3004.
- 65 S. Furutani, H. Nagai, Y. Takamura, Y. Aoyama and I. Kubo, *Analyst*, 2012, **137**, 2951–2957.
- 66 M. Amasia, M. Cozzens and M. J. Madou, *Sens. Actuators, B*, 2012, **161**, 1191–1197.
- 67 O. Strohmeier, S. Laßmann, B. Riedel, D. Mark, G. Roth, M. Werner, R. Zengerle and F. Stetten, *Microchim. Acta*, 2013, **181**, 1681–1688.
- 68 M. Keller, S. Wadle, N. Paust, L. Dreesen, C. Nuese, O. Strohmeier, R. Zengerle and F. von Stetten, *RSC Adv.*, 2015, **5**, 89603–89611.
- 69 O. Strohmeier, N. Marquart, D. Mark, G. Roth, R. Zengerle and F. von Stetten, *Anal. Methods*, 2014, **6**, 2038–2046.
- 70 M. Keller, J. Naue, R. Zengerle, F. von Stetten and U. Schmidt, *PLoS One*, 2015, **10**, e0131845.
- 71 G. Czilwik, I. Schwarz, M. Keller, S. Wadle, S. Zehnle, F. von Stetten, D. Mark, R. Zengerle and N. Paust, *Lab Chip*, 2015, **15**, 1084–1091.
- 72 G. Czilwik, T. Messinger, O. Strohmeier, S. Wadle, F. von Stetten, N. Paust, G. Roth, R. Zengerle, P. Saarinen, J. Niittymaki, K. McAllister, O. Sheils, J. O’Leary and D. Mark, *Lab Chip*, 2015, **15**, 3749–3759.



- 73 H. Kim, S. Vishniakou and G. W. Faris, *Lab Chip*, 2009, **9**, 1230–1235.
- 74 Y. Zhu, Y.-X. Zhang, W.-W. Liu, Y. Ma, Q. Fang and B. Yao, *Sci. Rep.*, 2015, **5**, 9551.
- 75 Y. Xu, H. Yan, Y. Zhang, K. Jiang, Y. Lu, Y. Ren, H. Wang, S. Wang and W. Xing, *Lab Chip*, 2015, **15**, 2826–2834.
- 76 F. Stumpf, J. Schoendube, A. Gross, C. Rath, S. Niekrawietz, P. Koltay and G. Roth, *Biosens. Bioelectron.*, 2015, **69**, 301–306.
- 77 H. Sun, T. Olsen, J. Zhu, J. Tao, B. Ponnaiya, S. A. Amundson, D. J. Brenner and Q. Lin, *RSC Adv.*, 2015, **5**, 4886–4893.
- 78 S. W. Lim, T. M. Tran and A. R. Abate, *PLoS One*, 2015, **10**, e0113549.
- 79 V. Sanchez-Freire, A. D. Ebert, T. Kalisky, S. R. Quake and J. C. Wu, *Nat. Protoc.*, 2012, **7**, 829–838.
- 80 K. H. Narsinh, N. Sun, V. Sanchez-Freire, A. S. Lee, P. Almeida, S. Hu, T. Jan, K. D. Wilson, D. Leong, J. Rosenberg, M. Yao, R. C. Robbins and J. C. Wu, *J. Clin. Invest.*, 2011, **121**, 1217–1221.
- 81 A. K. White, M. VanInsberghe, O. I. Petriv, M. Hamidi, D. Sikorski, M. A. Marra, J. Piret, S. Aparicio and C. L. Hansen, *Proc. Natl. Acad. Sci. U. S. A.*, 2011, **108**, 13999–14004.
- 82 L. Chen, A. Manz and P. J. R. Day, *Anal. Biochem.*, 2008, **372**, 128–130.
- 83 T. Notomi, H. Okayama, H. Masubuchi, T. Yonekawa, K. Watanabe, N. Amino and T. Hase, *Nucleic Acids Res.*, 2000, **28**, e63.
- 84 M. C. Giuffrida, L. M. Zanolli, R. D'Agata, A. Finotti, R. Gambari and G. Spoto, *Anal. Bioanal. Chem.*, 2015, **407**, 1533–1543.
- 85 T. D. Rane, L. Chen, H. C. Zec and T.-H. Wang, *Lab Chip*, 2015, **15**, 776–782.
- 86 J. Luo, X. Fang, D. Ye, H. Li, H. Chen, S. Zhang and J. Kong, *Biosens. Bioelectron.*, 2014, **60**, 84–91.
- 87 L. Desbois, A. Padirac, S. Kaneda, A. J. Genot, Y. Rondelez, D. Hober, D. Collard and T. Fujii, *Biomicrofluidics*, 2012, **6**, 044101.
- 88 C.-H. Wang, K.-Y. Lien, T.-Y. Wang, T.-Y. Chen and G.-B. Lee, *Biosens. Bioelectron.*, 2011, **26**, 2045–2052.
- 89 T. Wang, Y. Zhang, G. Huang, C. Wang, L. Xie, L. Ma, Z. Li, X. Luo, H. Tian, Q. Li, X. Li, Z. Lv and X. Bao, *Sci. China: Chem.*, 2012, **55**, 508–514.
- 90 Q.-J. Zhou, L. Wang, J. Chen, R.-N. Wang, Y.-H. Shi, C.-H. Li, D.-M. Zhang, X.-J. Yan and Y.-J. Zhang, *J. Microbiol. Methods*, 2014, **104**, 26–35.
- 91 A. A. Sayad, F. Ibrahim, S. M. Uddin, K. X. Pei, M. S. Mohktar, M. Madou and K. L. Thong, *Sens. Actuators, B*, 2016, **227**, 600–609.
- 92 Y. Zhang, L. Zhang, J. Sun, Y. Liu, X. Ma, S. Cui, L. Ma, J. J. Xi and X. Jiang, *Anal. Chem.*, 2014, **86**, 7057–7062.
- 93 C. Liu, M. G. Mauk and H. H. Bau, *Microfluid. Nanofluid.*, 2011, **11**, 209–220.
- 94 K. Hsieh, B. S. Ferguson, M. Eisenstein, K. W. Plaxco and H. T. Soh, *Acc. Chem. Res.*, 2015, **48**, 911–920.
- 95 X. Fang, H. Chen, S. Yu, X. Jiang and J. Kong, *Anal. Chem.*, 2011, **83**, 690–695.
- 96 Y. Zhu, D. Lu, M. E. Lira, Q. Xu, Y. Du, J. Xiong, M. Mao, H. C. Chung and G. Zheng, *Exp. Mol. Pathol.*, 2016, **100**, 287–293.
- 97 D. Azuara, M. M. Ginesta, M. Gausachs, F. Rodriguez-Moranta, J. Fabregat, J. Busquets, N. Pelaez, J. Boadas, S. Galter, V. Moreno, J. Costa, J. de Oca and G. Capellá, *Clin. Chem.*, 2012, **58**, 1332–1341.
- 98 M. Hussain, R. Fantuzzo, S. Mercorelli and C. Cullen, *J. Pharm. Biomed. Anal.*, 2016, **123**, 128–131.
- 99 C. Floren, I. Wiedemann, B. Brenig, E. Schütz and J. Beck, *Food Chem.*, 2015, **173**, 1054–1058.
- 100 T. Hoshino and F. Inagaki, *Syst. Appl. Microbiol.*, 2012, **35**, 390–395.
- 101 R. H. Sedlak, J. Kuypers and K. R. Jerome, *Diagn. Microbiol. Infect. Dis.*, 2014, **80**, 285–286.
- 102 E. M. Khorosheva, M. A. Karymov, D. A. Selck and R. F. Ismagilov, *Nucleic Acids Res.*, 2016, **44**, e10.
- 103 D. L. Duewer, M. C. Kline and E. L. Romsos, *Anal. Bioanal. Chem.*, 2015, **407**, 9061–9069.
- 104 P. Zhu, W. Fu, C. Wang, Z. Du, K. Huang, S. Zhu and W. Xu, *Anal. Chim. Acta*, 2016, **916**, 60–66.
- 105 X. Bian, F. Jing, G. Li, X. Fan, C. Jia, H. Zhou, Q. Jin and J. Zhao, *Biosens. Bioelectron.*, 2015, **74**, 770–777.
- 106 Q. Zhong, S. Bhattacharya, S. Kotsopoulos, J. Olson, V. Taly, A. D. Griffiths, D. R. Link and J. W. Larson, *Lab Chip*, 2011, **11**, 2167–2174.
- 107 P. Wang, F. Jing, G. Li, Z. Wu, Z. Cheng, J. Zhang, H. Zhang, C. Jia, Q. Jin, H. Mao and J. Zhao, *Biosens. Bioelectron.*, 2015, **74**, 836–842.
- 108 Y. Men, Y. Fu, Z. Chen, P. A. Sims, W. J. Greenleaf and Y. Huang, *Anal. Chem.*, 2012, **84**, 4262–4266.
- 109 K. A. Heyries, C. Tropini, M. VanInsberghe, C. Doolin, O. I. Petriv, A. Singhal, K. Leung, C. B. Hughesman and C. L. Hansen, *Nat. Methods*, 2011, **8**, 649–651.
- 110 H. Tanaka, S. Yamamoto, A. Nakamura, Y. Nakashoji, N. Okura, N. Nakamoto, K. Tsukagoshi and M. Hashimoto, *Anal. Chem.*, 2015, **87**, 4134–4143.
- 111 A. M. Thompson, A. Gansen, A. L. Paguirigan, J. E. Kreutz, J. P. Radich and D. T. Chiu, *Anal. Chem.*, 2014, **86**, 12308–12314.
- 112 Q. Song, Y. Gao, Q. Zhu, Q. Tian, B. Yu, B. Song, Y. Xu, M. Yuan, C. Ma, W. Jin, T. Zhang, Y. Mu and Q. Jin, *Biomed. Microdevices*, 2015, **17**, 1–8.
- 113 Q. Tian, Q. Song, Y. Xu, Q. Zhu, B. Yu, W. Jin, Q. Jin and Y. Mu, *Anal. Methods*, 2015, **7**, 2006–2011.
- 114 X. Leng, W. Zhang, C. Wang, L. Cui and C. J. Yang, *Lab Chip*, 2010, **10**, 2841–2843.
- 115 D. Paunescu, C. A. Mora, L. Querci, R. Heckel, M. Puddu, B. Hattendorf, D. Günther and R. N. Grass, *ACS Nano*, 2015, **9**, 9564–9572.
- 116 S. O. Sundberg, C. T. Wittwer, L. Zhou, R. Palais, Z. Dwight and B. K. Gale, *Biomed. Microdevices*, 2014, **16**, 639–644.
- 117 F. Schuler, F. Schwemmer, M. Trotter, S. Wadle, R. Zengerle, F. von Stetten and N. Paust, *Lab Chip*, 2015, **15**, 2759–2766.



- 118 F. Schuler, M. Trotter, M. Geltman, F. Schwemmer, S. Wadle, E. Dominguez-Garrido, M. Lopez, C. Cervera-Acedo, P. Santibanez, F. von Stetten, R. Zengerle and N. Paust, *Lab Chip*, 2016, **16**, 208–216.
- 119 F. Schuler, C. Siber, S. Hin, S. Wadle, N. Paust, R. Zengerle and F. von Stetten, *Anal. Methods*, 2016, **8**, 2750–2755.
- 120 W. Du, L. Li, K. P. Nichols and R. F. Ismagilov, *Lab Chip*, 2009, **9**, 2286–2292.
- 121 B. Sun, F. Shen, S. E. McCalla, J. E. Kreutz, M. A. Karymov and R. F. Ismagilov, *Anal. Chem.*, 2013, **85**, 1540–1546.
- 122 D. A. Selck, M. A. Karymov, B. Sun and R. F. Ismagilov, *Anal. Chem.*, 2013, **85**, 11129–11136.
- 123 GenePOC, <http://www.genepoc-diagnostics.com/en/technology/>, (accessed 31.08.2016).
- 124 Roche Diagnostics, <https://usdiagnostics.roche.com/en/instrument/cobas-liat.html>, (accessed 31.08.2016).
- 125 BD Diagnostic Systems, <http://moleculardiagnosics.bd.com/product/max/>, (accessed 31.08.2016).
- 126 bioMérieux SA, <http://www.biomerieux-diagnostics.com/filmarray>, (accessed 31.08.2016).
- 127 Biocartis, <https://www.biocartis.com/>, (accessed 31.08.2016).
- 128 Cepheid, <http://www.cepheid.com/us/cepheid-solutions/systems/genexpert-systems/genexpert-i>, (accessed 31.08.2016).
- 129 Veredus Laboratories, <http://vereduslabs.com/system-and-software/vereplex-biosystem/>, (accessed 31.08.2016).
- 130 M. J. Binnicker, M. J. Espy, C. L. Irish and E. A. Vetter, *J. Clin. Microbiol.*, 2015, **53**, 2353–2354.
- 131 M. A. Poritz, A. J. Blaschke, C. L. Byington, L. Meyers, K. Nilsson, D. E. Jones, S. A. Thatcher, T. Robbins, B. Lingenfelter, E. Amiott, A. Herbener, J. Daly, S. F. Dobrowolski, D. H. F. Teng and K. M. Ririe, *PLoS One*, 2011, **6**, e26047.
- 132 J. Teo, P. D. Pietro, F. S. Biagio, M. Capozzoli, Y.-M. Deng, I. Barr, N. Caldwell, K.-L. Ong, M. Sato, R. Tan and R. Lin, *Arch. Virol.*, 2011, **156**, 1371–1378.
- 133 Thermo Fisher Scientific, <https://www.thermofisher.com/de/de/home/life-science/pcr/digital-pcr/quantstudio-3d-digital-pcr-system.html>, (accessed 01.09.2016).
- 134 Bio-Rad, <http://www.bio-rad.com/en-ch/product/qx200-droplet-digital-pcr-system>, (accessed 01.09.2016).
- 135 RainDance Technologies, <http://raindancetech.com/digital-pcr-tech/raindrop-digital-pcr-system/>, (accessed 01.09.2016).
- 136 Fluidigm, <https://www.fluidigm.com/products/biomark-hd-system>, (accessed 01.09.2016).
- 137 D. Dobnik, B. Spilsberg, A. Bogožalec Košir, A. Holst-Jensen and J. Žel, *Anal. Chem.*, 2015, **87**, 8218–8226.
- 138 E. Zonta, F. Garlan, N. Pécuchet, K. Perez-Toralla, O. Caen, C. Milbury, A. Didelot, E. Fabre, H. Blons, P. Laurent-Puig and V. Taly, *PLoS One*, 2016, **11**, e0159094.
- 139 R. Sanders, D. J. Mason, C. A. Foy and J. F. Huggett, *PLoS One*, 2013, **8**, e75296.
- 140 Q. Tian, B. Yu, Y. Mu, Y. Xu, C. Ma, T. Zhang, W. Jin and Q. Jin, *RSC Adv.*, 2015, **5**, 81889–81896.
- 141 Q. Tian, Y. Mu, Y. Xu, Q. Song, B. Yu, C. Ma, W. Jin and Q. Jin, *Anal. Biochem.*, 2015, **491**, 55–57.

