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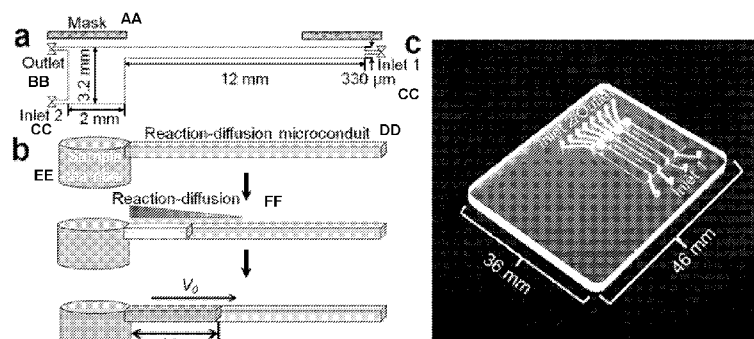


FIGURE 1

(57) Abstract: Provided herein are devices for monitoring nucleic acid amplification, nucleic acid amplification monitors, methods of quantifying nucleic acid amplification, methods of identifying an unknown nucleic acid molecule, and systems for monitoring nucleic acid amplification. The disclosed devices, methods, and systems comprise at least one nucleometer, comprising at least one sample chamber and at least one reaction-diffusion conduit in fluid communication with the chamber.

DEVICES AND METHODS FOR MONITORING AND QUANTIFYING NUCLEIC ACID AMPLIFICATION

CROSS REFERENCE TO RELATED APPLICATIONS

[0001] This application claims the benefit of U.S. Patent Application Nos. 62/020,135, filed July 2, 2014, and 62/020,484, filed July 3, 2014, the disclosures of which are incorporated herein by reference in their entirety.

GOVERNMENT RIGHTS

[0002] The subject matter disclosed herein was made with government support under grant numbers K25AI099160 and 1R41AI104418-01A1 awarded by the National Institutes of Health. The Government has certain rights in the herein disclosed subject matter.

TECHNICAL FIELD

[0003] Provided herein are devices, methods, and systems for monitoring and quantifying nucleic acid amplification.

BACKGROUND

[0004] Real-time amplification and quantification of nucleic acids has revolutionized genetic research, medical diagnostics, and environmental monitoring. In the case of infectious diseases, quantification of the pathogen-load in patient specimens is critical to assess disease progression, effectiveness of drug therapy, and emergence of drug-resistance. Currently, nucleic acid quantification requires expensive instruments, such as real-time PCR machines, for continuous monitoring of fluorescence emission from intercalating dye or molecular beacon probes. Although enzymatic amplification products can be detected without an instrument with lateral flow strips, this method suffers from low accuracy and low sensitivity.

[0005] Thus, there is a need for devices and methods providing low-cost, instrument-free, monitoring and end-point quantification of target nucleic acids undergoing enzymatic amplification. The invention is directed to these and other important needs.

SUMMARY

[0006] Disclosed are devices, methods, and systems for low-cost, reaction-diffusion based, instrument-free, monitoring and end-point quantification of target nucleic acids undergoing enzymatic amplification. The target concentration is estimated from a length

measurement along a reaction-diffusion-conduit. Exemplary uses of the disclosed devices, methods, and systems include on-site or high throughput applications and the identification of gene expression profiles.

[0007] Provided herein are devices for monitoring nucleic acid amplification comprising a nucleometer comprising a sample chamber and at least one reaction-diffusion conduit in fluid communication with the chamber. In some embodiments, each of the chamber and the conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the chamber and the conduit are capable of holding a blend of reactants for nucleic acid amplification.

[0008] Also disclosed are nucleic acid amplification monitors comprising a substrate and, on or forming a part of the substrate, a plurality of nucleometers, each nucleometer comprising a sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber; each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.

[0009] Methods of quantifying nucleic acid amplification are also provided herein, said methods comprising amplifying a sample comprising a nucleic acid molecule in any of the devices or the nucleic acid amplification monitors disclosed herein to generate an amplified nucleic acid molecule and measuring a reaction-diffusion length of said amplified nucleic acid molecule through the one or more conduits, wherein reaction-diffusion length is proportional to the number of nucleic acid copies in the sample and time.

[0010] Further disclosed are methods of identifying an unknown nucleic acid molecule, comprising amplifying a sample having an unknown nucleic acid molecule in any of the devices or nucleic acid amplification monitors disclosed herein, wherein each of the at least one conduits contain a plurality of primers specific for a different known nucleic acid molecule, and measuring a reaction-diffusion length within the reaction-diffusion conduit, wherein the presence of the reaction-diffusion length indicates that the sample having an unknown nucleic acid molecule comprises the known nucleic acid molecule to which the primers in the at least one conduit are specific and the extent of the reaction-diffusion length within the conduit indicates the number of the unknown nucleic acid molecules in the sample..

[0011] Further disclosed are systems for monitoring nucleic acid amplification comprising a nucleometer comprising a sample chamber, at least one reaction-diffusion conduit in fluid communication with the sample chamber, each of the sample chamber and the conduit

being capable of holding a blend of reactants for nucleic acid amplification, an optical imager, and a heat source.

[0012] The general description and the following detailed description are exemplary and explanatory only and are not restrictive of the invention, as defined in the appended claims. Other aspects of the present invention will be apparent to those skilled in the art in view of the detailed description as provided herein.

BRIEF DESCRIPTION OF THE DRAWINGS

[0013] The summary, as well as the following detailed description, is further understood when read in conjunction with the appended drawings. For the purpose of illustrating the disclosed devices, methods, and systems, there are shown in the drawings exemplary embodiments of the devices, methods, and systems; however, the invention is not limited to the specific devices, methods, and systems disclosed. In the drawings:

[0014] **FIG 1**, comprising FIGS. 1A-1E, represent the design and operation of an exemplary nucleometer for HIV viral load testing. (a) A schematic depiction of the cross-section of the nucleometer, consisting of a sample well and a reaction-diffusion microconduit. (b) An illustration of the nucleometer's operation. Initially, only the sample chamber contains the nucleic acid template (top). The template amplifies and diffuses into the reaction-conduit, where it continues to amplify at 62.5°C (middle). X_F indicates the position of the reaction front that propagates with a constant velocity (v_0) (bottom). (c) A photograph of a plastic chip housing four nucleometers. (d) Fluorescence images of the reaction-diffusion conduits of four nucleometers containing 10^4 , 10^3 , 10^2 , and 0 (negative control) HIV-1 RNA templates at various times after the start of incubation. (e) Four nucleometers each containing 100 HIV-1 RNA templates to illustrate reproducibility.

[0015] **FIG 2**, comprising FIGS. 2A-2I, represent exemplary experimental data and theoretical predictions of nucleometer's performance. (a) Normalized emission intensity $\hat{c} = c/c_{\max}$ as a function of position along the reaction-diffusion conduit at various times. The solid lines and symbols correspond, respectively, to predictions and experimental data. The number of target molecules is 10^3 copies. (b) Normalized emission intensity $\hat{c}$ as a function of time at positions $x=1.2$, 1.8, and 2.4 mm along the length of the conduit. The solid lines and symbols correspond, respectively, to the predictions and experimental data. The number of target

molecules is 10^3 copies. (c) The experimental rate of the reaction $(-\frac{\partial \hat{c}}{\partial t})_{\text{exp}}$ as a function of position (x) at various times. (d) The measured width of the reaction-rate peak at midheight Λ_{exp} as a function of time. (e) The measured position of the reaction front $X_{F, \text{exp}}$ as a function of time for various template concentrations (error bars = standard deviation (s.d.); $n = 3$; $R^2=0.998$). (f) The intercept ($t_{0, \text{exp}}$) of the line in FIG. 2e and the threshold time C_t of real-time RT-LAMP curves as functions of the number of templates (error bars = s.d.; $n = 3$; $R^2=0.99$). (g) $X_{F, \text{exp}} - X_{F, \text{exp}}^{(3)}$ as a function of the template number at various times t (error bars = s.d.; $R^2 = 0.99$, $n = 15$). (h) The predicted position of the reaction front ($X_{F, th}$) as a function of time for various numbers of templates. (i) The predicted intercept ($t_{0, th}$) of the asymptotes in FIG. 2h as a function of template number.

[0016] FIG. 3 represents an exploded view of an exemplary nucleometer chip consisting of three layers: a top PMMA film, a PMMA chip body, and a bottom PCR Sealers™ tape. The various features of the chip body were milled with a CNC machine.

[0017] FIG. 4 represents a micrograph of the reaction-diffusion micro-conduit's cross-section. The dashed line indicates the interface between the PMMA film and the PMMA chip body.

[0018] FIG. 5 represents an exemplary experimental setup for the nucleometer chip. The portable processor includes a USB-based, fluorescence microscope (AM4113T-GFBW Dino-Lite Premier, AnMo Electronics, Taipei, Taiwan). The processor can be powered either with four AA batteries or by grid power. The fluorescence image of the nucleometers can be directly displayed on the computer screen. The USB microscope can be replaced with LED illumination and a smartphone camera or other imaging devices.

[0019] FIG. 6, comprising FIGS. 6A-6C, represent an exemplary portable, processor for RT-LAMP amplification and detection. (a) A photograph of the processor. Inset: a flexible, polyimide-based, thin film heater (HK5572R7.5L23A, Minco Products, Inc., Minneapolis, MN). The heater can be replaced with an exothermic reaction chamber (b) A thermograph of the nucleometer chip's surface taken with an infrared camera T360 (FLIR Systems, Wilsonville, USA). The four reaction-diffusion reactors are located within the dashed square. (c) A mask made with black 3M Scotch electrical tape to block background emission. The mask is equipped with a ruler to assist in determining the position of the reaction front (X_F).

[0020] FIG. 7, comprising FIGS. 7A-7B represent an evaluation of the limits of detection of an exemplary nucleometer and benchtop, "tubed-based" LAMP. (a) A representative,

endpoint fluorescent image of the nucleometer chip with 50 copies (lanes 1 and 2) and 5 copies (lanes 3 and 4) of target HIV RNA molecules. (b) Real-time, benchtop monitoring of RT-LAMP amplification of HIV viral RNA with 50, 5, and 0 (negative control) target RNA copies per tube. Both the nucleometer and the real time machine detected successfully 50 target molecules and failed to detect 5 target molecules.

[0021] FIG. 8, represents real-time, benchtop monitoring of RT-LAMP amplification of HIV viral RNA with 10^4 , 10^3 , 10^2 , and 0 (negative control) target RNA molecules per tube on the benchtop PCR machine. The tubes include 0.04% HPMC to replicate the conditions in the nucleometer chip.

[0022] FIG. 9 represents the position of the reaction front (X_F) as a function of the number of target molecules ($n = 3$) at times (from bottom to top) 32, 40, 48 and 56 min after the start of incubation.

[0023] FIG. 10, comprising FIGS. 10A-10D, represent an exemplary nucleometer's sample well emission (a) before RT-LAMP amplification, (b) shortly after the onset of amplification, and (c) at saturation of amplification. The number of target molecules is 10^3 copies. (d) a fluorescent emission from the exemplary nucleometer's sample chamber as a function of the time. The solid line and the symbols correspond, respectively, to the best fit line based on equation 4 and the experimental data. The number of target molecules is 10^3 copies.

[0024] FIG. 11, comprising FIGS. 11A-11B, represents: (a) the concentration distribution of the labeled primers in the reaction-diffusion conduit at various times in the absence of an amplification reaction. (b) The emission intensity (normalized with the initial concentration) as a function of position at various times. The lines correspond to theoretical predictions and the symbols to the experimental data (with optimal estimate for the diffusion coefficient).

[0025] FIG. 12, comprising FIGS. 12A-12B, illustrates a schematic depiction of an exemplary nucleometer in which the sample chamber and the reaction-diffusion conduits are separated with a barrier consisting of a phase change material. (i) represents a top view with the barrier in place; (ii) represents a side-view with the barrier in place; (iii) represents a side view after the barrier had dissolved, melted, moved, been removed, etc. to allow hydraulic communications between the sample chamber and the reaction-diffusion conduit. Furthermore, the sample chamber contains an optional membrane for the isolation of nucleic acids. The device enables multi-stage amplification with one amplicon being amplified in the sample chamber and the other in the diffusion-reaction conduit.

[0026] FIG. 13, comprising FIGS. 13A-13B, illustrates a schematic depiction of an exemplary nucleometer comprising a single sample chamber and multiple reaction-diffusion conduits, each customized to amplify a different nucleic acid. The relative quantities of the various nucleic acids are determined from the relative lengths of the reacted regions, as indicated by the fluorescence emission, in the reaction-diffusion columns, enabling the identification of, among other things, a gene expression profile or bacterial flora.

[0027] FIG. 14 is an image of an exemplary nucleometer utilizing bioluminescent assay for real-time recording of the position of the reaction front. The luminescence blob propagates along the conduit and indicates the instantaneous position of the reaction front. The target in this particular assay is HIV.

DETAILED DESCRIPTION OF ILLUSTRATIVE EMBODIMENTS

[0028] The disclosed devices, methods, and systems may be understood more readily by reference to the following detailed description taken in connection with the accompanying figures, which form a part of this disclosure. It is to be understood that the disclosed devices, methods, and systems are not limited to the specific devices, methods, and systems described and/or shown herein, and that the terminology used herein is for the purpose of describing particular embodiments by way of example only and is not intended to be limiting of the claimed devices, methods, and systems.

[0029] Unless specifically stated otherwise, any description as to a possible mechanism or mode of action or reason for improvement is meant to be illustrative only, and the disclosed devices, methods, and systems are not to be constrained by the correctness or incorrectness of any such suggested mechanism or mode of action or reason for improvement.

[0030] Reference to a particular numerical value includes at least that particular value, unless the context clearly dictates otherwise. When a range of values is expressed, another embodiment includes from the one particular value and/or to the other particular value. Further, reference to values stated in ranges include each and every value within that range. All ranges are inclusive and combinable.

[0031] When values are expressed as approximations, by use of the antecedent “about,” it will be understood that the particular value forms another embodiment.

[0032] It is to be appreciated that certain features of the disclosed devices, methods, and systems which are, for clarity, described herein in the context of separate embodiments, may also be provided in combination in a single embodiment. Conversely, various features of the

disclosed devices, methods, and systems that are, for brevity, described in the context of a single embodiment, may also be provided separately or in any subcombination.

[0033] As used herein, the singular forms “a,” “an,” and “the” include the plural.

[0034] Various terms relating to aspects of the description are used throughout the specification and claims. Such terms are to be given their ordinary meaning in the art unless otherwise indicated. Other specifically defined terms are to be construed in a manner consistent with the definitions provided herein.

[0035] As used herein, the term “about” is meant to encompass variations of $\pm 20\%$ or less, variations of 10% or less, variations of $\pm 5\%$ or less, variations of $\pm 1\%$ or less, variations of $\pm 0.5\%$ or less, or variations of $\pm 0.1\%$ or less from the specified value.

[0036] As used herein, the term “fluid communication” refers to ability of liquids to diffuse between the sample chamber and the at least one reaction-diffusion conduit. For example, nucleic acid molecules diffuse from the sample chamber into the at least one reaction-diffusion conduit.

[0037] As used herein, “nucleic acid amplification” includes any technique used to detect nucleic acids by amplifying (generating numerous copies of) the target molecules in the sample. Suitable nucleic acid amplification techniques include, but are not limited to, loop-mediated isothermal amplification (LAMP), reverse transcription-loop mediated isothermal amplification (RT-LAMP), or polymerase chain reaction.

[0038] As used herein, the term “nucleometer” refers to a reaction-diffusion device for monitoring and quantitative detection of amplified nucleic acid molecules based on the position of the reaction front. Each nucleometer comprises at least one sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber.

[0039] “Target” refers to a sequence, or partial sequence, of a nucleic acid of interest. “Target” and “nucleic acid molecule” are used interchangeably herein.

Devices for monitoring nucleic acid amplification

[0040] Provided herein are devices for monitoring nucleic acid amplification comprising a nucleometer. The nucleometer comprises a sample chamber and at least one reaction-diffusion conduit in fluid communication with the chamber. In some embodiments, each of the chamber and the at least one reaction-diffusion conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the chamber and the at least one reaction-diffusion

conduit are capable of holding a blend of reactants for nucleic acid amplification. The sample chamber may also serve as a nucleic acid amplification reactor.

[0041] In some embodiments, the device for monitoring nucleic acid amplification can comprise a nucleometer comprising a sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber, each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.

[0042] In other embodiments, the device for monitoring nucleic acid amplification can comprise a nucleometer comprising a sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber, each of the sample chamber and the conduit holding a blend of reactants for nucleic acid amplification.

[0043] The sample chamber and diffusion-reaction conduit can be separated with a barrier. The barrier can be configured to block passage of reactants between the sample chamber and the reaction-diffusion conduit during sample introduction, nucleic acid isolation, etc. and can be configured to allow passage of reactants between the sample chamber and the reaction-diffusion conduit at a specified time or in response to an event. In some embodiments, the barrier can be removable. In some aspects, for example, the barrier can be physically removed from the nucleometer. In some embodiments, the barrier can be degradable. For example, the barrier can be composed of, or can comprise, a material that degrades. In some embodiments, the barrier can be dissolvable. For example, the barrier can be composed of a material that dissolves in solution. Exemplary materials that dissolve in solution include, for example, salt. In some embodiments, the barrier can be capable of being melted. For example, the barrier can be composed of, or can comprise, a phase change material that melts, or partially melts, at a certain temperature. For example, the barrier can melt and move through the action of surface tension (capillary) forces or buoyancy or combination thereof, thereby allowing diffusion between the sample chamber and the reaction diffusion conduit. In some embodiments, the barrier can be movable. For example, the barrier can be configured to open or otherwise allow passage of sample between the sample chamber and the reaction diffusion conduit in response to a mechanical force or various means of actuation. For example, the actuator may consist of bi-metal or shape memory alloy that alters shape in response to temperature variations.

[0044] The barrier can allow for a two-step or nested-like amplification. For example, two different amplification reactions can be run; one in the sample chamber and one in the reaction-diffusion conduit. The reaction in the sample chamber can, for example, amplify nucleic acid molecules non-specifically. Once the reaction in the sample chamber is complete,

the barrier can be removed, dissolved, degraded, melted, etc. allowing the amplified nucleic acid molecules to diffuse into the reaction-diffusion conduit, which can contain primers for a specific nucleic acid molecule. Within the reaction-diffusion conduit, specific nucleic acid molecules will then be amplified. The two reactions can operate with different enzymes, temperatures, etc.

[0045] The sample chamber can have a size and shape suitable for containing a nucleic acid amplification reaction. For example, the cross-section of the sample chamber can have a variety of suitable shapes, including, but not limited to, square, rectangular, triangular, pentagonal, hexagonal, heptagonal, octagonal, nonagonal, decagonal, round, or elliptical.

[0046] The bottom of the sample chamber can be flat, substantially flat, conical, round, substantially round, or pointed or contain depressions, for example, for the purpose of storing reagents needed for the amplification process.

[0047] In some embodiment the sample chamber can be a droplet, which can be manipulated to make contact with the reaction-diffusion conduit. Numerous techniques are known for manipulating droplets including, but not limited to, electrowetting, thermal gradients, solute gradients, conduit geometry such as taper, or any combination thereof. For example, in some embodiments, a droplet comprising the reactant mixture and nucleic acid sample can be placed onto a surface containing a plurality of electrodes. An electric field can be applied to the surface resulting in migration of the droplet towards (or away from) the reaction-diffusion conduit. The electric field can be applied until the droplet makes contact, such as a hydraulic connection, with the reaction-diffusion conduit. As amplification progresses, the amplified nucleic acid sample can diffuse into the reaction-diffusion conduit. In some embodiments, the droplet is the sample chamber. In other embodiments, the droplet is placed within the sample chamber. In some embodiments, the droplet can be encased in oil. In some embodiments, the droplet can be produced by a microfluidic device.

[0048] The sample chamber, when used for nucleic acid amplification, will have a size suitable for a containing a nucleic acid amplification reaction. Suitable sizes include sample chambers having a volume of from about 0.1 μl to about 300 μl . In one embodiment, the sample chamber has a volume of about 0.1 μl . In one embodiment, the sample chamber has a volume of about 0.5 μl . In one embodiment, the sample chamber has a volume of about 1 μl . In one embodiment, the sample chamber has a volume of about 10 μl . In one embodiment, the sample chamber has a volume of about 20 μl . In one embodiment, the sample chamber has a volume of about 50 μl . In one embodiment, the sample chamber has a volume of about 100 μl . In one embodiment, the sample chamber has a volume of about 150 μl . In one embodiment, the sample

chamber has a volume of about 200 μl . In one embodiment, the sample chamber has a volume of about 250 μl . In one embodiment, the sample chamber has a volume of about 300 μl . In some embodiments, the sample chamber has a volume greater than 300 μl .

[0049] The sample chamber can be from about 0.1 μl to about 300 μl in volume. In some aspects, the sample chamber can be from about 0.1 μl to about 250 μl in volume. The sample chamber can be from about 0.1 μl to about 200 μl in volume. The sample chamber can be from about 0.1 μl to about 150 μl in volume. The sample chamber can be from about 0.1 μl to about 100 μl in volume. The sample chamber can be from about 0.1 μl to about 50 μl in volume. The sample chamber can be from about 10 μl to about 300 μl in volume. The sample chamber can be from about 10 μl to about 200 μl in volume. The sample chamber can be from about 10 μl to about 150 μl in volume. The sample chamber can be from about 10 μl to about 100 μl in volume. The sample chamber can be from about 10 μl to about 50 μl in volume. In some aspects, the sample chamber can be from about 25 μl to about 300 μl in volume. The sample chamber can be from about 50 μl to about 300 μl in volume. The sample chamber can be from about 75 μl to about 300 μl in volume. The sample chamber can be from about 100 μl to about 300 μl in volume. The sample chamber can be from about 125 μl to about 300 μl in volume. The sample chamber can be from about 150 μl to about 300 μl in volume. The sample chamber can be from about 175 μl to about 300 μl in volume. The sample chamber can be from about 200 μl to about 300 μl in volume. The sample chamber can be from about 225 μl to about 300 μl in volume. The sample chamber can be from about 250 μl to about 300 μl in volume. The sample chamber can be from about 275 μl to about 300 μl in volume.

[0050] The height of the sample chamber can be from about 0.05 mm to about 10 mm. The height of the sample chamber can be from about 0.1 mm to about 9 mm. The height of the sample chamber can be from about 0.2 mm to about 8 mm. The height of the sample chamber can be from about 0.3 mm to about 7 mm. The height of the sample chamber can be from about 0.4 mm to about 6 mm. The height of the sample chamber can be from about 0.5 mm to about 5 mm. The height of the sample chamber can be from about 0.6 mm to about 4 mm. The height of the sample chamber can be from about 0.7 mm to about 3 mm. The height of the sample chamber can be from about 0.8 mm to about 2 mm. The height of the sample chamber can be from about 0.9 mm to about 1 mm.

[0051] In some aspects, the height of the sample chamber can be about 0.05 mm. In some aspects, the height of the sample chamber can be about 0.1 mm. In some aspects, the height of the sample chamber can be about 0.2 mm. In some aspects, the height of the sample chamber

can be about 0.3 mm. In some aspects, the height of the sample chamber can be about 0.4 mm. In some aspects, the height of the sample chamber can be about 0.5 mm. In some aspects, the height of the sample chamber can be about 0.6 mm. In some aspects, the height of the sample chamber can be about 0.7 mm. In some aspects, the height of the sample chamber can be about 0.8 mm. In some aspects, the height of the sample chamber can be about 0.9 mm. In some aspects, the height of the sample chamber can be about 1 mm. In some aspects, the height of the sample chamber can be about 2.5 mm. In some aspects, the height of the sample chamber can be about 5 mm. In some aspects, the height of the sample chamber can be about 7.5 mm. In some aspects, the height of the sample chamber can be about 10 mm.

[0052] The width of the sample chamber can be from about 0.05 mm to about 5 mm. The width of the sample chamber can be from about 0.1 mm to about 4 mm. The width of the sample chamber can be from about 0.2 mm to about 3 mm. The width of the sample chamber can be from about 0.3 mm to about 2 mm. The width of the sample chamber can be from about 0.4 mm to about 1 mm. The width of the sample chamber can be from about 0.5 mm to about 0.9 mm. The width of the sample chamber can be from about 0.6 mm to about 0.8 mm.

[0053] The width of the sample chamber can be about 0.05. The width of the sample chamber can be about 0.1 mm. The width of the sample chamber can be about 0.2 mm. The width of the sample chamber can be about 0.3 mm. The width of the sample chamber can be about 0.4 mm. The width of the sample chamber can be about 0.5 mm. The width of the sample chamber can be from 0.6 mm. The width of the sample chamber can be about 0.7. The width of the sample chamber can be about 0.8 mm. The width of the sample chamber can be about 0.9 mm. The width of the sample chamber can be about 1 mm. The width of the sample chamber can be about 2 mm. The width of the sample chamber can be about 3 mm. The width of the sample chamber can be about 4 mm. The width of the sample chamber can be about 2 mm. The width of the sample chamber can be about 5 mm.

[0054] The length of the reaction-diffusion conduit can be from about 0.05 mm to about 100 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 75 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 50 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 25 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 10 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 5 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 2.5 mm. The length of the

reaction-diffusion conduit can be from about 0.05 mm to about 1 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 0.5 mm.

[0055] The length of the reaction-diffusion conduit can be about 100 mm. The length of the reaction-diffusion conduit can be about 75 mm. The length of the reaction-diffusion conduit can be about 50 mm. The length of the reaction-diffusion conduit can be about 25 mm. The length of the reaction-diffusion conduit can be about 10 mm. The length of the reaction-diffusion conduit can be about 5 mm. The length of the reaction-diffusion conduit can be about 2.5 mm. The length of the reaction-diffusion conduit can be about 1 mm. The length of the reaction-diffusion conduit can be about 0.5 mm.

[0056] The height of the reaction-diffusion conduit can be from about 0.1 μm to about 1 mm. The height of the reaction-diffusion conduit can be from about 0.1 μm to about 750 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 500 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 250 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 100 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 50 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 25 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 10 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 5 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 1 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 0.5 μm .

[0057] The height of the reaction-diffusion conduit can be about 1 mm. The height of the reaction-diffusion conduit can be about 750 μm . The height of the reaction-diffusion conduit can be about 500 μm . The height of the reaction-diffusion conduit can be about 250 μm . The height of the reaction-diffusion conduit can be about 100 μm . The height of the reaction-diffusion conduit can be about 50 μm . The height of the reaction-diffusion conduit can be about 25 μm . The height of the reaction-diffusion conduit can be about 10 μm . The height of the reaction-diffusion conduit can be about 5 μm . The height of the reaction-diffusion conduit can be about 1 μm . The height of the reaction-diffusion conduit can be about 0.5 μm .

[0058] The width of the reaction-diffusion conduit can be from about 0.1 μm to about 1 mm. The width of the reaction-diffusion conduit can be from about 0.1 μm to about 750 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 500 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 250 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 100 μm . The width of the reaction-

diffusion conduit can be from about 0.1 μm to about 50 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 25 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 10 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 5 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 1 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 0.5 μm .

[0059] The width of the reaction-diffusion conduit can be about 1 mm. The width of the reaction-diffusion conduit can be about 750 μm . The width of the reaction-diffusion conduit can be about 500 μm . The width of the reaction-diffusion conduit can be about 250 μm . The width of the reaction-diffusion conduit can be about 100 μm . The width of the reaction-diffusion conduit can be about 50 μm . The width of the reaction-diffusion conduit can be about 25 μm . The width of the reaction-diffusion conduit can be about 10 μm . The width of the reaction-diffusion conduit can be about 5 μm . The width of the reaction-diffusion conduit can be about 1 μm . The width of the reaction-diffusion conduit can be about 0.5 μm .

[0060] The cross-section of the reaction-diffusion conduit may be, for example, a rectangle, a square, a circle, a semi-circle, a triangle, a trapezoid. The reaction-diffusion conduit may be, for example, straight, curved, spiral, or contain turns.

[0061] In some embodiments, the device can have one reaction-diffusion conduit in fluid communication with the sample chamber. In other embodiments, the device can have a plurality of reaction-diffusion conduits in fluid communication with the sample chamber. For example, the device can have between 2 to about 100 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the device can have 2 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the device can have 3 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the device can have 4 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the device can have 5 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the device can have 10 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the device can have 15 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the device can have 20 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the device can have more than 20 reaction-diffusion conduits in fluid communication with the sample chamber.

[0062] In some embodiments, the device can have 1 sample chamber, said sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. In some aspects, the device can have 1 sample chamber, said sample chamber having 1 reaction-diffusion conduit in fluid communication therewith. In some aspects, the device can have 1 sample chamber, said sample chamber having a plurality of reaction-diffusion conduits in fluid communication therewith. For example, in some aspects, the device can have 1 sample chamber, said sample chamber having about 2 to about 100 reaction-diffusion conduits in fluid communication therewith.

[0063] In other embodiments, the device can have a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. For example, the device can have 1 to 100 sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. In some aspects, the device can have 1 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 2 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 3 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 4 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 5 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 10 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 50 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 75 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the device can have 100 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith.

[0064] In some embodiments, the device can have a plurality of sample chambers, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. Thus, in some aspects, 100% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, less than 100% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 90% of said plurality of

about 90% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 80% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 70% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 60% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 50% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 40% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 30% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 20% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, about 1% to about 10% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.

[0066] The device can have at least one sample chamber having a known quantity of a preselected, nucleic acid molecule, which can serve as a “positive control” or “calibration” nucleometer.

[0067] The disclosed devices can be used in combination with porous membranes for nucleic acid extraction, concentration, and purification. Suitable porous membranes include, but are not limited to, silica membrane, Whatman FTA membrane, alumina membrane, cellulose membrane. Such porous membranes specifically bind nucleic acids allowing other non-nucleic acid material to pass through the membrane. For example, in some embodiments, a nucleic acid sample can be incubated with a porous membrane, the porous membrane can be washed to remove non-nucleic acid material, and the porous membrane can be added to the sample chamber. In other embodiments, the device can have a sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification. Suitable sample chambers include those disclosed in U.S. Application No. 14/001,347 (US2014/0162244). In some aspects, the porous membrane can be removably located within the sample chamber. In some aspects, the porous membrane can be attached to the inside of the sample chamber. In other aspects, all or a portion of the sample chamber can be made from a porous membrane.

[0068] The sample chamber and/or the reaction-diffusion conduit can be made of a porous polymer or cellulose material. The blend of reactants can be stored in the porous polymer or cellulose material until the device is ready for use.

[0069] In some embodiments, the device further comprises a waste conduit. The waste conduit can, for example, be used to empty the sample chamber, decrease the volume of the sample in the sample chamber, extract amplified material from the sample chamber, remove non-nucleic acid material from the sample chamber, or any combination thereof.

[0070] The waste conduit and reaction-diffusion conduits can have a valve to open and close the conduit, allowing and preventing, respectively, sample from flowing into the conduit. For example, in some embodiments, the sample chamber contains a porous membrane. To extract, concentrate, and/or purify the nucleic acid sample, the valves on both the waste conduit and reaction-diffusion conduits can be closed to allow the sample to interact with the membrane. To remove non-nucleic acid material, the valve on the waste conduit can be opened, while the valve on the reaction-diffusion conduit can remain closed. After the waste is removed, the valve on the waste conduit can be closed. Prior to amplification, the appropriate reactant mixture can be added to the sample chamber and the valve on the reaction-diffusion conduit can be opened, allowing diffusion of the amplified sample into the reaction-diffusion conduit. In some embodiments, the valve can be a passive valve. In other embodiments, the valve can be an active valve.

[0071] In some embodiments the device further comprises a light source. Suitable light sources include, but are not limited to, light-emitting diode (LED), compact fluorescent lamp (CFL), or incandescent.

[0072] The device can further comprise an optical imager. Suitable optical imagers include, but are not limited to, a fluorescent microscope or a camera. In some aspects, the camera can be from a cellular telephone, smart-phone camera, tablet computer, or computer.

[0073] The device can be operatively connected to a controller or computer.

[0074] In some embodiments, the device further comprises a ruler along the length of the reaction-diffusion conduit.

[0075] The device can further comprise a heat source. Suitable heat sources include a thermoelectric unit, an electric heater, an exothermic reactor, a laser, or any combination thereof. In some aspects, the heat source can provide a spatially varying temperature distribution along the length of the reaction-diffusion conduit. Such a heat source enables different temperatures to be applied to different regions of the reaction-diffusion conduit at different times. The

temperature can be regulated with a phase change material. In some embodiments, the temperature gradient along the reaction-diffusion conduit can be linear. The temperature in the sample chamber can be different from the temperature in the reaction-diffusion conduit. In other embodiments, the temperature in the sample chamber can be the same as the temperature in the reaction-diffusion conduit.

[0076] In some embodiments, each of the chamber and the at least one reaction-diffusion conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the chamber and the at least one reaction-diffusion conduit are capable of holding a blend of reactants for nucleic acid amplification. The blend of reactants can comprise a plurality of primers and enzymes. In some embodiments, the blend of reactants can further comprise one or more types of reporters. Those skilled in the art would understand that numerous primers, enzymes, and reporters are suitable for nucleic acid amplification procedures. Primers of any length suitable for nucleic acid amplification can be used herein. One skilled in the art would know that primers of nucleic acid amplification are used in pairs – a forward or sense primer and a reverse or antisense primer. “Plurality of primers” and “primer sets” are used interchangeably herein and refer to at least one pair of primers. In some embodiments, the plurality of primers can be a pair of primers. In some embodiments, the plurality of primers can be 2 pairs of primers. In some embodiments, plurality of primers can be more than 2 pairs of primers. Enzymes suitable for nucleic acid amplification include, but are not limited to, DNA polymerase, RNA-dependent DNA polymerase (reverse transcriptase), or DNA-dependent RNA polymerase (RNA polymerase). As used herein, the term “reporter” means any tag, label, or dye that can bind to, or intercalate within, the nucleic acid molecule or be activated by byproducts of the amplification process to enable visualization of the nucleic acid molecule or the amplification process. Suitable reporters include, but are not limited to, fluorescent labels/tags/dyes, intercalating agents, molecular beacon labels, and bioluminescent molecules. The sample chamber and reaction-diffusion conduit may contain a mixture of primers engineered to be specific to various targets, enabling the concurrent amplification and detection of multiple targets. The sample chamber may contain universal primers to amplify all targets, while the reaction-diffusion conduits may contain primers for specific targets. In some embodiments, the universal primers in the sample chamber can be immobilized to prevent passage of the primers into the reaction-diffusion conduit. Primers can be immobilized to, for example, the sample chamber or something within the sample chamber that cannot pass into the reaction-diffusion conduit. The reaction-diffusion conduit may also contain reporters (molecular beacons) each

engineered to bind to a specific target and each emitting at different range of the spectrum, allowing the detector, with appropriate filters, to image each target separately.

[0077] Other reactants include, but are not limited to, buffers and dNTPs.

[0078] The blend of reactants, or a portion thereof, can be pre-stored in the device. In some embodiments, the blend of reactants, or a portion thereof, are pre-stored in the device and are released upon an increase in temperature. For example, the blend of reactants can be released when the device is heated to its operating temperature.

[0079] The blend of reactants can be liquid or can be in a form that is capable of forming a liquid upon a change in temperature, such as a gel or solid. For example, the blend of reactants can be encased in, embedded in, or surrounded by, a material that melts with increasing temperatures, such as, for example, paraffin. In yet other embodiments, the primers and/or enzymes may be immobilized to the conduit walls or to a porous matrix that fills the reaction-diffusion conduit.

[0080] In some embodiments, the device can have at least two nucleometers, each containing different primers. The primers can recognize the same nucleic acid molecule but differ in length. Alternatively, the primers can recognize different nucleic acid molecules and be the same length. Alternatively, the primers can recognize different nucleic acid molecules and be different lengths.

[0081] A single nucleometer can have multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers. For example, one reaction-diffusion conduit can contain a first plurality of primers and another reaction-diffusion conduit can contain a second plurality of primers, the second plurality of primers being different from the first plurality of primers. In such a device, the sample chamber can contain: a mix of a first plurality of primers and second plurality of primers; only a first plurality of primers; only a second plurality of primers; or a third plurality of primers.

[0082] In some embodiments, the blend of reactants in the sample chamber and the blend of reactants in the conduit are capable of amplifying different nucleic acid molecules. In some aspects, for example, the blend of reactants in the sample chamber and the blend of reactants in the conduit are capable of amplifying at least two different nucleic acid molecules.

[0083] In some embodiments, at least one of the reaction-diffusion conduits can contain at least one polymer. In some embodiments, at least one of the reaction-diffusion conduit and the sample chamber can contain at least one polymer. The presence of the polymer in the reaction-diffusion conduit or reaction-diffusion conduit and sample chamber can slow the

diffusion of the amplified DNA and control the speed of propagation of the reaction front in the reaction-diffusion conduit, so that the diffusion may be easily monitored and/or quantified. In some embodiments, the at least one polymer can be a gel. In embodiments wherein the at least one polymer is a gel, the primers, enzymes, or both can be immobilized to the gel. In some embodiments, the polymer is a solid at an ambient temperature and a liquid at an amplification temperature. As used herein, "ambient temperature" refers to the temperature of the device prior to the amplification procedure. Thus, prior to the amplification procedure, the polymer can be a solid. As the temperature increases at the start of or during the amplification procedure, the polymer can turn into a liquid. In other embodiment, the reaction-diffusion conduit contains gel.

[0084] The disclosed devices can be used for a number of purposes including, but not limited to, melting curve analysis, reverse transcription, identifying a nucleic acid in a sample, diagnosis, quantifying the number of nucleic acid molecules in a sample, and testing amplification reactants. In some embodiments, the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.

Nucleic acid amplification monitors

[0085] Also disclosed herein are nucleic acid amplification monitors comprising a substrate and, on or forming a part of the substrate, and a plurality of nucleometers. Each nucleometer comprises an sample chamber and at least one reaction-diffusion conduit in fluid communication with the chamber. In some embodiments, each of the chamber and the at least one reaction-diffusion conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the chamber and the at least one reaction-diffusion conduit are capable of holding a blend of reactants for nucleic acid amplification.

[0086] Suitable substrates include, but are not limited to, glass, plastic, polymers, metal, silicon, aluminum oxide membrane, acrylic, paper, or any combination thereof. In some embodiments the substrate is a polymeric chip. Polymeric chips can be composed of numerous polymers including, but not limited to, polycarbonate (PC), poly-methyl methacrylate (PMMA), cyclic olefin co-polymers (COC), or any combination thereof. In one embodiment, the polymeric chip comprises polymethyl methacrylate (PMMA).

[0087] In some embodiments, the plurality of nucleometers can be on the substrate. For example, the nucleometer and the substrate can be separate components, and can be combined or attached such that the nucleometer is on a top surface of the substrate. In other aspects, the nucleometer and substrate can be from a single component wherein the nucleometer is located on a

top surface of the substrate. In other embodiments, the nucleometer can form a part of the substrate. For example, the nucleometer can be etched into the substrate. In other embodiments, the substrate can be molded to form the nucleometer.

[0088] The sample chamber can have a size and shape suitable for containing a nucleic acid amplification reaction. For example, the cross-section of the sample chamber can have a variety of suitable shapes, including, but not limited to, square, rectangular, triangular, pentagonal, hexagonal, heptagonal, octagonal, nonagonal, decagonal, round, or elliptical.

[0089] The bottom of the sample chamber can be flat, substantially flat, conical, round, substantially round, or pointed.

[0090] The sample chamber can be a droplet covered by oil, wherein the droplet is propelled by means such as electrowetting to make hydraulic connection with the reaction-diffusion conduit.

[0091] The sample chamber and diffusion-reaction conduit can be separated with a barrier. The barrier can be configured to block passage of reactants between the sample chamber and the reaction-diffusion conduit during sample introduction, nucleic acid isolation, etc. and can be configured to allow passage of reactants between the sample chamber and the reaction-diffusion conduit at a specified time or in response to an event. In some embodiments, the barrier can be removable. In some aspects, for example, the barrier can be physically removed from the nucleometer. In some embodiments, the barrier can be degradable. For example, the barrier can be composed of, or can comprise, a material that degrades. In some embodiments, the barrier can be dissolvable. For example, the barrier can be composed of a material that dissolves in solution. Exemplary materials that dissolve in solution include, for example, salt. In some embodiments, the barrier can be capable of being melted. For example, the barrier can be composed of, or can comprise, a phase change material that melts at a certain temperature. In some embodiments, the barrier can be movable. For example, the barrier can be configured to open or otherwise allow passage of sample between the sample chamber and the reaction diffusion conduit in response to a force, such as the amplification of the nucleic acid molecule or other surface tension.

[0092] The barrier can allow for a two-step or nested-like amplification. For example, two different amplification reactions can be run; one in the sample chamber and one in the reaction-diffusion conduit. The reaction in the sample chamber can, for example, amplify nucleic acid molecules non-specifically. Once the reaction in the sample chamber is complete, the barrier can be removed, dissolved, degraded, melted, etc. allowing the amplified nucleic acid

molecules to diffuse into the reaction-diffusion conduit, which can contain primers for a specific nucleic acid molecule. Within the reaction-diffusion conduit, specific nucleic acid molecules will then be amplified. The two reactions can operate with different enzymes, temperatures, etc.

[0093] The sample chamber has a size suitable for containing a nucleic acid amplification reaction. Suitable sizes include sample chambers having a volume of from about 0.1 μl to about 300 μl . In one embodiment, the sample chamber has a volume of about 0.1 μl . In one embodiment, the sample chamber has a volume of about 0.5 μl . In one embodiment, the sample chamber has a volume of about 1 μl . In one embodiment, the sample chamber has a volume of about 10 μl . In one embodiment, the sample chamber has a volume of about 20 μl . In one embodiment, the sample chamber has a volume of about 50 μl . In one embodiment, the sample chamber has a volume of about 100 μl . In one embodiment, the sample chamber has a volume of about 150 μl . In one embodiment, the sample chamber has a volume of about 200 μl . In one embodiment, the sample chamber has a volume of about 250 μl . In one embodiment, the sample chamber has a volume of about 300 μl . In some embodiments, the sample chamber has a volume greater than 300 μl .

[0094] The sample chamber can be from about 0.1 μl to about 300 μl in volume. In some aspects, the sample chamber can be from about 0.1 μl to about 250 μl in volume. The sample chamber can be from about 0.1 μl to about 200 μl in volume. The sample chamber can be from about 0.1 μl to about 150 μl in volume. The sample chamber can be from about 0.1 μl to about 100 μl in volume. The sample chamber can be from about 0.1 μl to about 50 μl in volume. The sample chamber can be from about 10 μl to about 300 μl in volume. The sample chamber can be from about 10 μl to about 200 μl in volume. The sample chamber can be from about 10 μl to about 150 μl in volume. The sample chamber can be from about 10 μl to about 100 μl in volume. The sample chamber can be from about 10 μl to about 50 μl in volume. In some aspects, the sample chamber can be from about 25 μl to about 300 μl in volume. The sample chamber can be from about 50 μl to about 300 μl in volume. The sample chamber can be from about 75 μl to about 300 μl in volume. The sample chamber can be from about 100 μl to about 300 μl in volume. The sample chamber can be from about 125 μl to about 300 μl in volume. The sample chamber can be from about 150 μl to about 300 μl in volume. The sample chamber can be from about 175 μl to about 300 μl in volume. The sample chamber can be from about 200 μl to about 300 μl in volume. The sample chamber can be from about 225 μl to about 300 μl in volume. The sample chamber can be from about 250 μl to about 300 μl in volume. The sample chamber can be from about 275 μl to about 300 μl in volume.

[0095] The height of the sample chamber can be from about 0.05 mm to about 10 mm. The height of the sample chamber can be from about 0.1 mm to about 9 mm. The height of the sample chamber can be from about 0.2 mm to about 8 mm. The height of the sample chamber can be from about 0.3 mm to about 7 mm. The height of the sample chamber can be from about 0.4 mm to about 6 mm. The height of the sample chamber can be from about 0.5 mm to about 5 mm. The height of the sample chamber can be from about 0.6 mm to about 4 mm. The height of the sample chamber can be from about 0.7 mm to about 3 mm. The height of the sample chamber can be from about 0.8 mm to about 2 mm. The height of the sample chamber can be from about 0.9 mm to about 1 mm.

[0096] In some aspects, the height of the sample chamber can be about 0.05 mm. In some aspects, the height of the sample chamber can be about 0.1 mm. In some aspects, the height of the sample chamber can be about 0.2 mm. In some aspects, the height of the sample chamber can be about 0.3 mm. In some aspects, the height of the sample chamber can be about 0.4 mm. In some aspects, the height of the sample chamber can be about 0.5 mm. In some aspects, the height of the sample chamber can be about 0.6 mm. In some aspects, the height of the sample chamber can be about 0.7 mm. In some aspects, the height of the sample chamber can be about 0.8 mm. In some aspects, the height of the sample chamber can be about 0.9 mm. In some aspects, the height of the sample chamber can be about 1 mm. In some aspects, the height of the sample chamber can be about 2.5 mm. In some aspects, the height of the sample chamber can be about 5 mm. In some aspects, the height of the sample chamber can be about 7.5 mm. In some aspects, the height of the sample chamber can be about 10 mm.

[0097] The width of the sample chamber can be from about 0.05 mm to about 5 mm. The width of the sample chamber can be from about 0.1 mm to about 4 mm. The width of the sample chamber can be from about 0.2 mm to about 3 mm. The width of the sample chamber can be from about 0.3 mm to about 2 mm. The width of the sample chamber can be from about 0.4 mm to about 1 mm. The width of the sample chamber can be from about 0.5 mm to about 0.9 mm. The width of the sample chamber can be from about 0.6 mm to about 0.8 mm.

[0098] The width of the sample chamber can be about 0.05. The width of the sample chamber can be about 0.1 mm. The width of the sample chamber can be about 0.2 mm. The width of the sample chamber can be about 0.3 mm. The width of the sample chamber can be about 0.4 mm. The width of the sample chamber can be about 0.5 mm. The width of the sample chamber can be from 0.6 mm. The width of the sample chamber can be about 0.7. The width of the sample chamber can be about 0.8 mm. The width of the sample chamber can be about 0.9

mm. The width of the sample chamber can be about 1 mm. The width of the sample chamber can be about 2 mm. The width of the sample chamber can be about 3 mm. The width of the sample chamber can be about 4 mm. The width of the sample chamber can be about 2 mm. The width of the sample chamber can be about 5 mm.

[0099] The length of the reaction-diffusion conduit can be from about 0.05 mm to about 100 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 75 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 50 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 25 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 10 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 5 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 2.5 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 1 mm. The length of the reaction-diffusion conduit can be from about 0.05 mm to about 0.5 mm.

[00100] The length of the reaction-diffusion conduit can be about 100 mm. The length of the reaction-diffusion conduit can be about 75 mm. The length of the reaction-diffusion conduit can be about 50 mm. The length of the reaction-diffusion conduit can be about 25 mm. The length of the reaction-diffusion conduit can be about 10 mm. The length of the reaction-diffusion conduit can be about 5 mm. The length of the reaction-diffusion conduit can be about 2.5 mm. The length of the reaction-diffusion conduit can be about 1 mm. The length of the reaction-diffusion conduit can be about 0.5 mm.

[00101] The height of the reaction-diffusion conduit can be from about 0.1 μm to about 1 mm. The height of the reaction-diffusion conduit can be from about 0.1 μm to about 750 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 500 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 250 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 100 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 50 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 25 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 10 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 5 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 1 μm . The height of the reaction-diffusion conduit can be from about 0.1 μm to about 0.5 μm .

[00102] The height of the reaction-diffusion conduit can be about 1 mm. The height of the reaction-diffusion conduit can be about 750 μm . The height of the reaction-diffusion conduit

can be about 500 μm . The height of the reaction-diffusion conduit can be about 250 μm . The height of the reaction-diffusion conduit can be about 100 μm . The height of the reaction-diffusion conduit can be about 50 μm . The height of the reaction-diffusion conduit can be about 25 μm . The height of the reaction-diffusion conduit can be about 10 μm . The height of the reaction-diffusion conduit can be about 5 μm . The height of the reaction-diffusion conduit can be about 1 μm . The height of the reaction-diffusion conduit can be about 0.5 μm .

[0103] The width of the reaction-diffusion conduit can be from about 0.1 μm to about 1 mm. The width of the reaction-diffusion conduit can be from about 0.1 μm to about 750 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 500 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 250 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 100 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 50 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 25 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 10 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 5 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 1 μm . The width of the reaction-diffusion conduit can be from about 0.1 μm to about 0.5 μm .

[0104] The width of the reaction-diffusion conduit can be about 1 mm. The width of the reaction-diffusion conduit can be about 750 μm . The width of the reaction-diffusion conduit can be about 500 μm . The width of the reaction-diffusion conduit can be about 250 μm . The width of the reaction-diffusion conduit can be about 100 μm . The width of the reaction-diffusion conduit can be about 50 μm . The width of the reaction-diffusion conduit can be about 25 μm . The width of the reaction-diffusion conduit can be about 10 μm . The width of the reaction-diffusion conduit can be about 5 μm . The width of the reaction-diffusion conduit can be about 1 μm . The width of the reaction-diffusion conduit can be about 0.5 μm .

[0105] The cross-section of the reaction-diffusion conduit may be, for example, a rectangle, a square, a circle, a semi-circle, a triangle, a trapezoid. The reaction-diffusion conduit may be, for example, straight, curved, spiral, or contain turns.

[0106] In some embodiments, the nucleic acid amplification monitor can have 1 reaction-diffusion conduit in fluid communication with the sample chamber. In other embodiments, the nucleic acid amplification monitor can have a plurality of reaction-diffusion conduits in fluid communication with the sample chamber. For example, the nucleic acid amplification monitor can have between 2 to about 100 reaction-diffusion conduits in fluid

communication with the sample chamber. In some aspects, the nucleic acid amplification monitor can have 2 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the nucleic acid amplification monitor can have 3 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the nucleic acid amplification monitor can have 4 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the nucleic acid amplification monitor can have 5 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the nucleic acid amplification monitor can have 10 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the nucleic acid amplification monitor can have 15 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects the nucleic acid amplification monitor can have 20 reaction-diffusion conduits in fluid communication with the sample chamber. In some aspects, the nucleic acid amplification monitor can have more than 20 reaction-diffusion conduits in fluid communication with the sample chamber.

[0107] In some embodiments, the nucleic acid amplification monitor can have 1 sample chamber, said sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 1 sample chamber, said sample chamber having 1 reaction-diffusion conduit in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 1 sample chamber, said sample chamber having a plurality of reaction-diffusion conduits in fluid communication therewith. For example, in some aspects, the nucleic acid amplification monitor can have 1 sample chamber, said sample chamber having about 2 to about 100 reaction-diffusion conduits in fluid communication therewith.

[0108] In other embodiments, the nucleic acid amplification monitor can have a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. For example, the nucleic acid amplification monitor can have 1 to 100 sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 1 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 2 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 3 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification

monitor can have 4 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 5 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 10 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 50 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 75 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith. In some aspects, the nucleic acid amplification monitor can have 100 sample chambers, each sample chamber having 2-100 reaction-diffusion conduits in fluid communication therewith.

[0109] In some embodiments, the nucleic acid amplification monitor can have a plurality of sample chambers, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. Thus, in some aspects, 100% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, less than 100% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 90% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 80% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 70% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 60% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 50% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 40% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 30% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 20% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 10% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some aspects, 5% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith. In some

aspects, 1% of said plurality of sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.

[illegible]

sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.

[0111] The nucleic acid amplification monitor can have at least one sample chamber having a known quantity of a preselected, nucleic acid molecule, which can serve as a “positive control” or “calibration” nucleometer.

[0112] The disclosed nucleic acid amplification monitors can be used in combination with porous membranes for nucleic acid extraction, concentration, and purification. Suitable porous membranes include, but are not limited to, silica membrane, Whatman FTA membrane, alumina membrane, cellulose membrane. Such porous membranes specifically bind nucleic acids allowing other non-nucleic acid material to pass through the membrane. For example, in some embodiments, a nucleic acid sample can be incubated with a porous membrane, the porous membrane can be washed to remove non-nucleic acid material, and the porous membrane can be added to the sample chamber. In other embodiments, the nucleic acid amplification monitor can have a sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification. Suitable sample chambers include those disclosed in U.S. Application No. 14/001,347 (US2014/0162244). In some aspects, the porous membrane can be removably located within the sample chamber. In some aspects, the porous membrane can be attached to the inside of the sample chamber. In other aspects, all or a portion of the sample chamber can be made from a porous membrane.

[0113] In some embodiments, the nucleic acid amplification monitor further comprises a waste conduit. The waste conduit can, for example, be used to empty the sample chamber, decrease the volume of the sample in the sample chamber, extract amplified material from the sample chamber, remove non-nucleic acid material from the sample chamber, or any combination thereof.

[0114] The waste conduit and reaction-diffusion conduits can have a valve to open and close the conduit, allowing and preventing, respectively, sample from flowing into the conduit. For example, in some embodiments, the sample chamber contains a porous membrane. To extract, concentrate, and/or purify the nucleic acid sample, the valves on both the waste conduit and reaction-diffusion conduits can be closed to allow the sample to interact with the membrane. To remove non-nucleic acid material, the valve on the waste conduit can be opened, while the valve on the reaction-diffusion conduit can remain closed. After the waste is removed, the valve on the waste conduit can be closed. Prior to amplification, the appropriate reactant mixture can be added to the sample chamber and the valve on the reaction-diffusion conduit can be opened,

allowing diffusion of the amplified sample into the reaction-diffusion conduit. In some embodiments, the valve can be a passive valve. In other embodiments, the valve can be an active valve.

[0115] In some embodiments the nucleic acid amplification monitor further comprises a light source. Suitable light sources include, but are not limited to, light-emitting diode (LED), compact fluorescent lamp (CFL), filtered flash light of a cell phone camera, or incandescent.

[0116] The nucleic acid amplification monitor can further comprise an optical imager. Suitable optical imagers include, but are not limited to, a fluorescent microscope or a camera. In some aspects, the camera can be from a cellular telephone, smart-phone camera, tablet computer, or computer.

[0117] The nucleic acid amplification monitor can be operatively connected to a controller or computer.

[0118] In some embodiments, the nucleic acid amplification monitor further comprises a ruler along the length of the reaction-diffusion conduit.

[0119] The nucleic acid amplification monitor can further comprise a heat source. Suitable heat sources include a thermoelectric unit, an electric heater, an exothermic reactor, a laser, or any combination thereof. In some aspects, the heat source can provide a spatially varying temperature distribution along the length of the reaction-diffusion conduit. Such a heat source enables different temperatures to be applied to different regions of the reaction-diffusion conduit at different times. The temperature can be regulated with a phase change material. In some embodiments, the temperature gradient along the reaction-diffusion conduit can be linear. The temperature in the sample chamber can be different from the temperature in the reaction-diffusion conduit. In other embodiments, the temperature in the sample chamber can be the same as the temperature in the reaction-diffusion conduit.

[0120] In some embodiments, each of the chamber and the at least one reaction-diffusion conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the chamber and the at least one reaction-diffusion conduit are capable of holding a blend of reactants for nucleic acid amplification. The blend of reactants can comprise a plurality of primers and enzymes. In some embodiments, the blend of reactants can further comprise one or more types of reporters. Those skilled in the art would understand that numerous primers, enzymes, and reporters are suitable for nucleic acid amplification procedures. Suitable primers include, for example, DNA or RNA primers. The length of the primer (i.e. number of nucleotides) depends, in part, on the nucleic acid molecule to be amplified. Primers of any

length suitable for nucleic acid amplification can be used herein. One skilled in the art would know that primers of nucleic acid amplification are used in sets or pairs – a forward or sense primer and a reverse or antisense primer. “Plurality of primers” and “primer sets” are used interchangeably herein and refer to at least one pair of primers. In some embodiments, the plurality of primers can be a pair of primers. In some embodiments, the plurality of primers can be 2 pairs of primers. In some embodiments, the plurality of primers can be more than 2 pairs of primers. Enzymes suitable for nucleic acid amplification include, but are not limited to, DNA polymerase, RNA-dependent DNA polymerase (reverse transcriptase), or DNA-dependent RNA polymerase (RNA polymerase). As used herein, the term “reporter” means any tag, label, or dye that can bind to, or intercalate within, the nucleic acid molecule to enable visualization of the nucleic acid molecule. Suitable reporters include, but are not limited to, fluorescent labels/tags/dyes and intercalating agents. The sample chamber and reaction-diffusion conduit may contain a mixture of primers engineered to be specific to various targets, enabling the concurrent amplification and detection of multiple targets. The sample chamber may contain universal primers to amplify all targets, while the reaction-diffusion conduits may contain primers for specific targets. In some embodiments, the universal primers in the sample chamber can be immobilized to prevent passage of the primers into the reaction-diffusion conduit. Primers can be immobilized to, for example, the sample chamber or something within the sample chamber that cannot pass into the reaction-diffusion conduit. The reaction-diffusion conduit may also contain reporters (molecular beacons) each engineered to bind to a specific target and each emitting at different range of the spectrum, allowing the detector, with appropriate filters, to image each target separately.

[0121] Other reactants include, but are not limited to, buffers and dNTPs.

[0122] The blend of reactants, or a portion thereof, can be pre-stored in the nucleic acid amplification monitor. In some embodiments, the blend of reactants, or a portion thereof, are pre-stored in the nucleic acid amplification monitor and are released upon an increase in temperature. For example, the blend of reactants can be released when the nucleic acid amplification monitor is heated to its operating temperature.

[0123] The blend of reactants can be liquid or can be in a form that is capable of forming a liquid upon a change in temperature, such as a gel or solid. For example, the blend of reactants can be encased in, embedded in, or surrounded by, a material that melts with increasing temperatures, such as, for example, paraffin.

[0124] In some embodiments, the nucleic acid amplification monitor can have at least two nucleometers, each containing different primers. The primers can recognize the same nucleic acid molecule but differ in length. Alternatively, the primers can recognize different nucleic acid molecules and be the same length. Alternatively, the primers can recognize different nucleic acid molecules and be different lengths.

[0125] A single nucleometer can have multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers. For example, one reaction-diffusion conduit can contain a first plurality of primers and another reaction-diffusion conduit can contain a second plurality of primers, the second plurality of primers being different from the first plurality of primers. In such a device, the sample chamber can contain: a mix of a first plurality of primers and second plurality of primers; only a first plurality of primers; only a second plurality of primers; or a third plurality of primers.

[0126] In some embodiments, the blend of reactants in the sample chamber and the blend of reactants in the conduit are capable of amplifying different nucleic acid molecules. In some aspects, for example, the blend of reactants in the sample chamber and the blend of reactants in the conduit are capable of amplifying at least two different nucleic acid molecules.

[0127] In some embodiments, at least one of the reaction-diffusion conduits can contain at least one polymer. In some embodiments, at least one of the reaction-diffusion conduits and the sample chamber can contain at least one polymer. The presence of the polymer in the reaction-diffusion conduit or reaction-diffusion conduit and sample chamber can slow the diffusion of the amplified DNA and control the speed of propagation of the reaction front in the reaction-diffusion conduit, so that the diffusion may be easily monitored and/or quantified. In some embodiments, the at least one polymer can be a gel. In embodiments wherein the at least one polymer is a gel, the primers, enzymes, or both can be immobilized to the gel. In some embodiments, the polymer is a solid at an ambient temperature and a liquid at an amplification temperature. As used herein, "ambient temperature" refers to the temperature of the nucleic acid amplification monitor prior to the amplification procedure. Thus, prior to the amplification procedure, the polymer can be a solid. As the temperature increases at the start of or during the amplification procedure, the polymer can turn into a liquid. In other embodiment, the reaction-diffusion conduit contains gel.

[0128] The disclosed nucleic acid amplification monitors can be used for a number of purposes including, but not limited to, melting curve analysis, reverse transcription, identifying a nucleic acid in a sample sample, diagnosis, quantifying the number of nucleic acid molecules in

a sample, and testing amplification reactants. In some embodiments, the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.

[0129] In some embodiments, the sample chamber and the reaction-diffusion conduit can be separate components.

Methods of quantifying nucleic acid amplification

[0130] Also disclosed herein are methods of quantifying nucleic acid amplification comprising amplifying a sample comprising a nucleic acid molecule in any of the devices or the nucleic acid amplification monitors disclosed herein to generate an amplified nucleic acid molecule and measuring a reaction-diffusion length of said amplified nucleic acid molecule through the one or more conduits, wherein reaction-diffusion length is proportional to the number of nucleic acid copies in the sample.

[0131] In some embodiments, amplifying the nucleic acid molecule comprises thermocycling. In other embodiments, amplifying the nucleic acid molecule comprises isothermal nucleic acid amplification.

[0132] Suitable methods for measuring a reaction-diffusion length of said amplified nucleic acid molecule include, but are not limiting to, measuring an electrical resistance, measuring a capacitance, measuring an absorption (for example, UV absorption), measuring an emission of a reporter, or any combination thereof. In some aspects, the reaction-diffusion length of said amplified nucleic acid molecule can be measured, for example, using a ruler. In some embodiments, the device or nucleic acid amplification monitor has a ruler along the length of the reaction-diffusion conduit. In other embodiments, the ruler is separate from the nucleometer.

[0133] In some embodiments, quantifying the length of the emitting column in the reaction-diffusion column can be a function of nucleic acid concentration and time. By reading the length of the emitting column and the time elapsed from the onset of the amplification reaction, for example, one can determine the number of nucleic acid molecules in the sample. In other embodiments, quantification can be independent of time. For example, one can monitor two or more nucleometers, at least one of which containing a known number of molecules (calibrator) and at least one containing an unknown number of nucleic acids (tester). The difference in the emission lengths of the tester and calibrator can enable one to determine the number of nucleic acid molecules in the tester, independent of the measurement time.

[0134] In some embodiments, the methods can further comprise comparing the reaction-diffusion length of the one or more reaction-diffusion conduits. For example, one or more of the reaction-diffusion conduits within the device can be compared to one or more of the reaction-diffusion conduits within the same device. Conversely, one or more of the reaction-diffusion conduits within the device can be compared to one or more reaction-diffusion conduits within a separate device. Comparison of the one or more reaction-diffusion conduits to reaction-diffusion conduits in the same device or in a separate device will allow one to evaluate, for example, the relative amount of one or more nucleic acid molecules. In other embodiments, the methods can further comprise comparing the reaction-diffusion length of the one or more reaction-diffusion conduits to a control. The control can be a reaction-diffusion conduit from the same device or from a separate device that contains a known quantity of a nucleic acid molecule, a nucleic acid molecule of a known identity, a sample from a subject having a known disease or condition, or any combination thereof. The quantity, identity, or sample from a subject may be indicative of the presence or absence of a disease or condition. The comparing can be used, for example, to determine: the presence or absence of a disease in a subject; the type or stage of a disease in a subject; or the effectiveness of therapy. For example, the one or more known nucleic acid molecules can be nucleic acid molecules that exhibit altered expression in the disease. In some aspects, the one or more nucleic acid molecules can be nucleic acid molecules whose expression is decreased in the disease. In some aspects, the one or more nucleic acid molecules can be nucleic acid molecules whose expression is increased in the disease. The reaction-diffusion length from the one or more known nucleic acid molecules can be indicative of a disease state. In some aspects, the reaction-diffusion length from the one or more known nucleic acid molecules can be indicative of a disease type. In some aspects, the reaction-diffusion length from the one or more known nucleic acid molecules can be indicative of a disease stage. In other embodiments, the reaction-diffusion length from the one or more known nucleic acid molecules is indicative of a normal (non-disease) state.

[0135] The one or more known nucleic acid molecules can be derived from cells, blood, serum, bacteria, or any other suitable source of a nucleic acid molecule. In some aspects, the nucleic acid molecules can be derived from cells from a subject suspected of having a disease. In some aspects, the nucleic acid molecules can be derived from the bacterial flora of a subject. Comparing the reaction-diffusion length of the sample to the reaction-diffusion length from one or more known nucleic acid molecules that are indicative of a disease state, a normal

(non-disease) state, or both, can be used to determine if a subject has the disease state.

Exemplary diseases include, for example, cancer.

[0136] Different reaction-diffusion conduits can store different primers specific to genes of interest. By comparing the reaction-diffusion lengths in different reaction-diffusion conduits, in which selected genes were amplified, with a library of gene expression profiles, one can determine, for example, the gene expression profile of the sample.

[0137] In some embodiments, the nucleic acid molecule is DNA. In some embodiments, the nucleic acid molecule is RNA. In embodiments wherein the nucleic acid molecule is RNA, the methods can further comprise, prior to the amplifying, reverse transcribing the RNA to generate cDNA.

[0138] The one or more known nucleic acid molecules can be markers of a disease. For example, the one or more known nucleic acid molecules can be nucleic acid molecules that exhibit altered expression in the disease. In some aspects, the one or more nucleic acid molecules can be nucleic acid molecules whose expression is decreased in the disease. In some aspects, the one or more nucleic acid molecules can be nucleic acid molecules whose expression is increased in the disease. For example, the one or more known nucleic acid molecules can be markers of cancer.

[0139] The results of the methods of obtaining a gene expression profile can be used, for example, to screen for a disease of interest, select an appropriate drug for a patient, or monitor therapy.

[0140] The disclosed methods can be used to determine the number of nucleic acid molecules in a sample by, for example, comparing the reaction-diffusion length in a reaction-diffusion conduit containing molecules from the sample to a reaction-diffusion length in a conduit having a known number of nucleic acid molecules. The control molecules may be pre-stored in the device or consist of house-keeping molecules in the sample. The disclosed methods can also be used to determine the relative expression of nucleic acid molecules in a sample by, for example, comparing the reaction-diffusion length to a reaction-diffusion conduit having a control sample and calculating the percentage or the reaction-diffusion length of the sample.

Methods of identifying an unknown nucleic acid molecule

[0141] Also disclosed herein are methods of identifying an unknown nucleic acid molecule, comprising amplifying a sample comprising an unknown nucleic acid molecule in any of the devices or nucleic acid amplification monitors disclosed herein, wherein each of the at

least one conduits contain a plurality of primers specific for a different known nucleic acid molecule and measuring a reaction-diffusion length within the reaction-diffusion conduit, wherein the presence of the reaction-diffusion length indicates that the sample having an unknown nucleic acid molecule comprises the known nucleic acid molecule to which the primers in the at least one conduit are specific and the extent of the reaction-diffusion length within the conduit indicates the number of the unknown nucleic acid molecules in the sample..

[0142] For example, the device or nucleic acid amplification monitor suitable for the disclosed methods of identifying an unknown nucleic acid molecule can contain one or more nucleometers. The nucleometers can have a plurality of reaction-diffusion conduits, each of said reaction-diffusion conduit containing a plurality of primers specific for a different known nucleic acid molecule. For example, each of the reaction-diffusion conduits can contain different primers for different viral nucleic acids. In some aspects, each of the reaction-diffusion conduits can contain different primers for different pathogens, such as bacteria, virus, and parasite nucleic acids. In other aspects, each of the reaction-diffusion conduits can contain different primers for different fungal nucleic acids. In yet other aspects, each of the reaction-diffusion conduits can contain different primers for different biomarkers which identify human disease. For example, each of the reaction-diffusion conduits can contain different primers for different cancer biomarkers. In some embodiments, a single nucleometer can comprise a plurality of reaction-diffusion conduits, each reaction-diffusion conduit containing different primers for a variety of different pathogens, viruses, bacteria, fungus, human, animal, and plant diseases, or any combination thereof.

[0143] The disclosed methods of identifying an unknown nucleic acid molecule can be used, for example, in diagnosing a patient with an unknown infection or disease. For example, a nucleic acid sample can be isolated from a human, an animal, or a plant and added to the sample chamber of any one of the devices or nucleic acid amplification monitors disclosed herein, wherein the device or nucleic acid amplification monitor comprises at least one nucleometer comprising at least one sample chamber and at least one reaction-diffusion conduit, said at least one conduit containing a plurality of primers specific for a different known nucleic acid molecule. Amplification will proceed in a reaction-diffusion conduit if that conduit contains primers specific for the unknown nucleic acid molecule. Amplification, which can be monitored by an emission, will indicate that the sample having an unknown nucleic acid molecule comprises at least a nucleic acid molecule for which the primers are specific.

[0144] In addition to identifying unknown nucleic acid samples, the disclosed methods can be used to determine a number of nucleic acid molecules in the unknown nucleic acid sample. For example, in some embodiments, the method comprises determining the difference in length of reaction-diffusion from a nucleometer having a sample having a known quantity of a nucleic acid and a nucleometer having a sample having an unknown nucleic acid molecule, wherein the difference in the length of the reaction-diffusion between the known quantity of a nucleic acid molecule and the sample having an unknown nucleic acid molecule indicates a number of nucleic acid molecules in the sample having the unknown nucleic acid molecule.

[0145] Suitable methods for measuring a reaction-diffusion length of said amplified nucleic acid molecule include, but are not limiting to, measuring an electrical resistance, measuring a capacitance, measuring an absorption (for example, UV absorption), measuring an emission of a reporter, or any combination thereof. In some aspects, for example, the length of the reaction-diffusion can be determined by using a ruler. In some embodiments, the device or nucleic acid amplification monitor has a ruler along the length of the reaction-diffusion conduit. In other embodiments, the ruler is separate from the nucleometer.

Systems for monitoring nucleic acid amplification

[0146] Further disclosed are systems for monitoring nucleic acid amplification comprising a nucleometer, an optical imager, and a heat source. The nucleometer comprises an sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber. In some embodiments, each of the sample chamber and the at least one reaction-diffusion conduit hold a blend of reactants for nucleic acid amplification. In other embodiments, each of the sample chamber and the at least one reaction-diffusion conduit are capable of holding a blend of reactants for nucleic acid amplification.

[0147] Any of the nucleometers disclosed herein are suitable for the disclosed system.

[0148] The system can further comprise a controller, an output device, or a combination thereof.

[0149] The sample chamber can have a size and shape suitable for containing a nucleic acid amplification reaction.

[0150] Each of the optical imager, heat source and output device can be operably controlled by the controller.

EXAMPLES

Methods

Nucleometer chip fabrication

[0151] The 46 mm long × 36 mm wide × 3.0 mm thick, Poly(methyl methacrylate) (PMMA, Acrylic glass) body of the chip was milled with a precision, computer-controlled (CNC, HAAS Automation Inc., Oxnard, CA) milling machine (FIG. 1C). An inlet port and an exit port were connected to the sample chamber and a third port was connected to the distal end of the microconduit (**FIG. 1A**). After milling, the chip body was sonicated in 100% ethanol for 15 minutes, rinsed with water, and air-dried at room temperature. To eliminate any RNase and DNase that could degrade nucleic acids or interfere with enzymatic reactions, the chip body was dipped in Decon™ ELIMINase™ decontaminant (Thermo Fisher Scientific Inc., Waltham, MA) for 2 minutes, rinsed twice with sterile molecular biology-grade water (Thermo Fisher Scientific Inc.), and air-dried at room temperature.

[0152] The chip body was ceiled and floored with PMMA film (top) and PCR Sealers™ tape (bottom), respectively (**FIG. 3**). Both were cut with a CO₂ laser (Universal Laser Systems). The top PMMA film was solvent-bonded to the chip body with acetonitrile (Sigma-Aldrich) at room temperature. The bonded chip was heated overnight (Isotemp Vacuum Oven Model 280A, Fisher Scientific Inc., Pittsburgh, PA) at 55 °C to remove any residual solvent. Finally, the PCR Sealers™ tape was used to seal the bottom of the chip.

HIV RNA purification from plasma samples

[0153] Viral RNA was extracted from HIV-1 standards (AcroMetrix® HIV-1 High Control, Benicia, CA) with QIAamp Viral RNA Mini Kit (Qiagen, Valencia, CA) according to the manufacturer's protocol. Briefly, 140 µL of virus suspension was lysed with 560 µL virus lysis buffer containing carrier RNA. 560 µL ethanol was added to the lysate and the mixture was centrifuged in a spin column (630 µL aliquots) at 10,000 rpm for 2 minutes. Prior to eluting the HIV viral RNA, wash buffers were loaded into the spin column and centrifuged at 14,000 rpm for 5 minutes. The RNA was eluted with 60 µL of elution buffer. Negative controls were prepared from a de-identified HIV-negative plasma sample (provided by the Penn Center for AIDS Research CFAR with the Institutional Review Board approval (protocol: 814752)) using the same extraction procedures as described above.

RT-LAMP reagents

[0154] The RT-LAMP primers were designed as described in Curtis, K.A., Rudolph, D.L. & Owen, S.M. J. Virol. Meth. 151, 264-270 (2008) at the Center for Disease Control and Prevention (CDC) and were synthesized by Sigma-Aldrich. The real-time benchtop RT-LAMP experiments were carried out with 15 μ L reaction volumes. The reaction mixture consisted of 0.2 μ M of F3 and B3, each; 0.8 μ M Loop F and Loop B, each; and 1.6 μ M of FIP and BIP, each, 1.25 U AMV reverse transcriptase (Life Technologies, Carlsbad, CA); $0.53 \times$ EvaGreen dye (Biotium, Hayward, CA); 0.04% (w/v) hydroxypropyl-methyl-cellulose (HPMC); and 9 μ L Isothermal Master Mix (ISO-001nd, OptiGene, Horsham, UK).

[0155] The HPMC was dissolved in Isothermal Master Mix, centrifuged at 10,000 rpm for 2 minutes, and filtered through a Corning Costar[®] Spin-X[®] centrifuge tube equipped with cellulose acetate membrane filters with a pore size of 0.45 μ m to remove any traces of insoluble HPMC.

[0156] A ten-fold dilution series of HIV viral RNA extracted from a HIV-1 standard panel and a negative control without template prepared from a HIV-negative plasma sample were tested in parallel. The real-time, “tubed-based” RT-LAMP was carried out in a Peltier Thermal Cycler PTC-200 (Bio-Rad DNA Engine, Hercules, CA). Reactions were carried out at 62.5 °C for 60 minutes with real-time fluorescence monitoring. Real-time RT-LAMP results were analyzed and the threshold time C_t (the time needed for the emission intensity to exceed a predetermined value) was obtained.

Device operation

[0157] 5 μ L of RT-LAMP master mixture, comprised of all the reagents necessary for the RT-LAMP and 0.04% HPMC (excluding the HIV RNA template), was inserted into each reaction-diffusion microconduit through inlet port 1 (FIG. 1A and 1C). Then, inlet ports 1 of all four nucleometers were sealed with PCR Sealers[™] tape. Next, 15 μ l of RT-LAMP master mixture and HIV RNA template of various concentrations were injected into the sample chambers through the inlet ports 2 (FIG. 1A and 1C). Subsequently, both the inlet ports 2 and outlet ports were sealed with PCR Sealers[™] tape to minimize evaporation during the amplification process. The nucleometer chip was placed on a custom, portable heater and incubated at 62.5 °C for about 60 minutes to enable isothermal amplification.

Portable processor for RT-LAMP

[0158] The custom made, portable processor (FIGS. 5 and 6) for the nucleometer consisted of a chip holder equipped with a flexible, polyimide-based, thin film heater (Model HK5572R7.5L23A, Minco Products, Inc., Minneapolis, MN) (inset in FIG. 6A), an electronic circuit board, and a thermocouple positioned at the interface between the thin film heater and the nucleometer chip. When the nucleometer chip, filled with LAMP master mixture, was inserted into the processor, the sample chambers and diffusion conduits were in thermal contact with the thin film heater.

[0159] To thermally calibrate the device, a calibration chip with a type-K thermocouple (Omega Engr., each wire 75 mm in diameter, and a junction diameter of 170 μm) was constructed in the reaction-diffusion conduit. The sample chambers and reaction-diffusion conduits were filled with water. The thermocouple reading was monitored with a HH506RA multilogger thermometer (Omega Engr., Stamford, CT, USA). In addition, an infrared image of the heated microfluidic chip was taken with an infrared thermography camera T360 (FLIR Systems, Wilsonville, USA) to evaluate temperature uniformity (FIG. 6B).

Endpoint, fluorescence image for quantitative detection

[0160] The fluorescence excitation and emission imaging were carried out with a handheld, USB-based, fluorescence microscope (AM4113T-GFBW Dino-Lite Premier, AnMo Electronics, Taipei, Taiwan) (FIGS. 5 and 6). The USB-based, fluorescence microscope has built-in, filtered blue LEDs for excitation, a 510 nm emission filter, and a CCD camera for fluorescence imaging. The microscope was interfaced with a computer through a USB interface. Images were acquired with a DinoCapture 2.0 software program. The images were processed with MatLabTM software to remove background noise and uneven illumination effects. A normalized and averaged fluorescence intensity signal for each lane was extracted from each processed image. The locations of the reaction fronts of different samples were directly read out by eye with the fluorescence ruler (FIG. 6C).

[0161] As an alternative to fluorescent dye to indicate the length of the reaction zone, bioluminescence can be used (FIG. 14). The bioluminescent real time reporter (BART) was included with the reagents. During DNA amplification in microconduits, the reporter emits visible light at the reaction front, providing an indication of the position of the reaction front. When bioluminescence is used, there is no need for excitation, eliminating any background

emission. The emission consists of white light that can be recorded directly by charged coupled device camera, cell phone camera or by eye (FIG. 14).

Image Processing and Analysis

[0162] Images were processed post-acquisition to facilitate comparison with numerical simulations. Initially, the noise was removed from all images using a low pass filter. Then, the images were corrected by applying the following scaling:

$$I_c = \frac{I_f - I_b}{I_f - I_b},$$

where I_c is the corrected emission intensity, I_f is the filtered intensity, I_b is the background intensity, and I_f is the flat maximum intensity. The background intensity is the average of the first five video frames when there are too few amplicons to generate a significant signal. The maximum intensity was obtained from the last frame acquired. The intensity signal at each position x was averaged along the width of the conduit. All image processing and mathematical calculations were performed with MatLab.

Results

[0163] The disclosed nucleometers enable monitoring and end-point quantitative detection of amplified nucleic acids based on the position of a reaction front. The nucleometer is comprised of a sample chamber and a reaction-diffusion conduit, containing all the reagents needed for enzymatic amplification, polymeric additive HPMC to slow diffusion, as well as intercalating dye reporter (FIG. 1A-C). A sample laden with target nucleic acids was introduced into the sample chamber and the amplification reaction was triggered thermally. As time progresses, amplicons diffused into the conduit, where they continue to react and amplify. After a certain time threshold, the conduit consisted of two distinct regions (FIG. 1B): the bright, left segment ($0 < x < X_F$), where the amplification reaction had already generated a sufficient number of amplicons to emit detectable fluorescence emission; and the dark, right section ($x > X_F$) into which amplicons had yet to diffuse. As time proceeded, the reaction front (X_F), separating between the bright and dark regions, propagated to the right. Without intent to be bound by theory, it is believed that the position of the reaction front indicates target analyte concentration.

Many nucleometers can be housed on a single chip and imaged simultaneously for concurrent monitoring of multiple amplification processes, calibration standards, and controls.

[0164] Polymethyl methacrylate (PMMA) chips consisting of four nucleometers (FIG. 1C and FIG. 3) were fabricated. Each nucleometer consisted of a 2mm diameter $\times$ 2.80mm deep sample well ($\sim 9\mu\text{L}$) connected to a $330\mu\text{m}$ wide $\times$ $330\mu\text{m}$ deep $\times$ 12mm long reaction-diffusion conduit (FIG. 1A and FIG. 4). Reverse transcription, loop-mediated isothermal amplification (RT-LAMP), as described in Notomi, T. et al. *Nucleic Acids Res.* 28, E63 (2000) and Tomita, N., Mori, Y., Kanda, H. & Notomi, T. *Nat. Protoc.* 3, 877-882 (2008), was used to quantify HIV viral load. Samples containing 0, 10^2 , 10^3 , and 10^4 HIV-1 RNA molecules were inserted into the sample wells (FIG. 1B) and incubated at 62.5°C using a custom, portable, processor (FIG. 5 and 6). The emission from the reaction-diffusion conduits was recorded using an USB fluorescent microscope (FIG. 1D). The position of the reaction front (X_F) was quantified with a ruler affixed on the processor (FIG. 6C). At any given time, the greater the number of target molecules, the larger X_F . Thus, with appropriate calibration, the number of initial target molecules can be inferred from X_F . Although X_F increases as time increases at any target concentration, the differences between X_F values associated with different concentrations are time-independent.

[0165] The reproducibility of the nucleometer was evaluated by introducing identical target concentrations (10^2 copies HIV-1 RNA) into all four sample wells (FIG. 1E). All four conduits exhibited nearly identical X_F ($\pm 4.5\%$) at any given time. Furthermore, the limits of detection were tested by reducing the number of target molecules. As few as 50 RNA copies were consistently detected, comparable to the performance of a benchtop RT-LAMP (FIG. 7).

[0166] FIG. 2 illustrates the experimental data and compares it with the predictions of a simple theoretical model. The emission intensity was taken to be proportional to the amplicons' concentration $c(x,t)$, assumed uniform in each cross-section of the conduit. FIG. 2A depicts $\hat{c} = c/c_{\text{max}}$ as a function of position x at various times t . The lines and symbols correspond, respectively, to predictions and experimental data. The location of the reaction front $X_F(t)$ was defined as the position at which $\hat{c}(X_F, t) = 0.5$. When $x < X_F$, $\hat{c} \sim 1$ and the amplification reaction is nearly complete (the regions with fluorescent emission in FIG. 1D). When $x > X_F$, $\hat{c} \sim 0$ and no amplification has yet occurred (the dark regions in FIG. 1d). FIG. 2B depicts $\hat{c}$ as a function of time at various positions x . An observer located at position x will not see a signal until after a

certain time delay. FIG. 2C depicts the experimentally-determined rate of the reaction $\left(\frac{\partial \hat{c}}{\partial t}\right)_{\text{exp}}$

as a function of position (x) at various times. The rate of the reaction resembles a propagating peak that travels at a fixed velocity v_0 . The peak's width at midheight (Λ) did not vary with time (FIG. 2D), i.e., the reaction front is non-dispersive.

[0167] The position of the reaction front $X_F(t)$ was depicted as a function of time when the number of target molecules is 10^2 , 10^3 , and 10^4 (FIG. 2E, $n = 3$). For sufficiently large times, $t > t_I > t_0$, the experimental data correlates well with straight lines ($R^2 = 0.998$).

$$X_F(t) = v_0(t - t_0) \quad , \quad (1)$$

Where t (s) is the observation time, t_0 (s) is the intercept with the horizontal axis, and t_I (s) is the delay time until a visible signal is observed. All the lines in FIG. 2E have nearly the same slope, indicating that the front propagates at a nearly constant speed of $v_0^{\text{exp}} = 1.73 \pm 0.15 \text{ } \mu\text{m/s}$ ($n=9$) independent of target concentration. In contrast, t_0 decreases as the target concentration increases (FIG. 2F), playing a similar role to the threshold time (C_t) in a standard real-time quantitative amplification. For comparison, C_t is also shown in FIG. 2F (see also FIG. 8).

$$t_0 = A - B \log(c_0) \quad , \quad (2)$$

where A and B are constants and c_0 is the number of target molecules ($R^2 = 0.996$).

[0168] Although the position of the front X_F is time-dependent, the distances between any two fronts associated with different numbers of target molecules are not (FIG. 9). Thus, time-dependence can be eliminated by subtracting the position of the front of a calibration lane ($X_F^{(c)}$) containing a known target concentration from that of the test lane. To demonstrate this, variables associated with lanes 1, 2, and 3 (FIG. 1D) were denoted with superscripts 1, 2, and 3. FIG. 2G depicts $X_F^{(i)} - X_F^{(3)}$ as a function of the number of target molecules $c_0^{(i)}$. It is observed that all the data collapses to a single straight line ($n = 15$, $R^2 = 0.99$), eliminating any explicit dependence on the time at which X_F was measured. Thus, in the presence of a calibration nucleometer (c),

$$\Delta X_F^{(i)} = X_F^{(i)}(t) - X_F^{(c)}(t) = v_0(t_0^{(c)} - t_0^{(i)}) \quad . \quad (3)$$

[0169] In the presence of one or more calibration nucleometers, one can rely on the differences among reaction front positions to determine target concentration, independent of measurement time. Although not essential, a device could include at least one calibration nucleometer. The calibration nucleometers can, of course, also serve as positive controls. With the aid of equation (2), equation (3) can be rewritten to express explicitly the dependence of ΔX_F on target analyte concentration.

$$\Delta X_F^{(i)} = v_0 B \log \left(\frac{c_0^{(i)}}{c_0^{(c)}} \right) \quad (4)$$

[0170] When a monitoring device includes two calibration nucleometers c1 and c2 with known target concentrations c0I(1) and c0I(2), respectively, the difference of the emission

$$\Delta X_F^{(c)} = X_F^{(c2)} - X_F^{(c1)} = v_0 B \log \left(\frac{c_0^{(c2)}}{c_0^{(c1)}} \right) \quad \text{and}$$

$$\Delta X_F^{(i)} = \Delta X_F^{(c)} \frac{\log(c_0^{(i)}) - \log(c_0^{(c1)})}{\log(c_0^{(c2)}) - \log(c_0^{(c1)})}$$

[0171] A simple reaction-diffusion mathematical model to simulate the experiment was proposed to gain further insight into the operation of the nucleometer. The amplicon production during enzymatic amplification was approximated with the production term $kc_{\max}\hat{c}(1-\hat{c})$, where the reaction rate constant $k \sim 0.008 \text{ s}^{-1}$ was determined empirically by fitting theoretical predictions with real time RT-LAMP amplification curves (See below – estimation of the reaction rate constant). $c_{\max}$ was estimated at $\sim 1.1 \times 10^{-10} \text{ mol/m}^3$. The variable c in the theory plays the same role as I in the experiment. The reaction diffusion process in the nucleometer was modeled with the dimensionless equation

$$\frac{\partial \hat{c}}{\partial \hat{t}} = \frac{\partial^2 \hat{c}}{\partial \hat{x}^2} + \hat{c}(1-\hat{c}) \quad (-\hat{d} < \hat{x} < \infty) \quad (5)$$

In the above, distance was scaled with $\sqrt{D/k}$ and time with k^{-1} . The diffusion coefficient $D \sim 10^{-10} \text{ m}^2/\text{s}$ was estimated by monitoring the diffusion of labeled primers in the conduit in the absence of amplification reaction (See below - Estimating the Diffusion Coefficient (D)). The

$$\text{boundary and interfacial conditions are: } \frac{\partial \hat{c}(-\hat{d}, \hat{t})}{\partial \hat{x}} = 0 \quad ;$$

$$\hat{c}(0^-, \hat{t}) - \hat{c}(0^+, \hat{t}) = \frac{\partial \hat{c}(0^-, \hat{t})}{\partial \hat{x}} - \frac{\partial \hat{c}(0^+, \hat{t})}{\partial \hat{x}} = 0 \quad (\text{the interface between the well and the conduit});$$

and $\hat{c}(\infty, \hat{t}) = 0$. The initial conditions are $\hat{c}(\hat{x}, 0) = \hat{c}_0$ when $-\hat{d} < \hat{x} < 0$ (sample well) and $\hat{c}(\hat{x}, 0) = 0$ when $0 < \hat{x} < \infty$ (conduit).

[0172] The predictions of equation (5) (solid lines in FIGS. 2A and B) closely resemble the experimental data. FIG. 2H depicts the position of the predicted reaction front as a function of time for different initial concentrations $\hat{c}_0$. Although at short times, the front velocity varies as

a function of $\hat{c}$, soon enough all the curves asymptote to straight lines with a slope independent of time and the initial target concentration. The *dimensionless* predicted reaction front velocity is

2. The *dimensional* predicted reaction front velocity $v_0^{theory} = 2\sqrt{kD} \sim 1.8 \mu m/s$ is very close to the experimentally measured one. Moreover, consistent with experiments, the theory predicts a constant reaction front velocity independent of target concentration. When $\hat{t} > \hat{t}_1 > \hat{t}_0$, the front location can be estimated with equation (1), where $\hat{t}$ depends on the initial concentration through equation (2). FIG. 2I depicts the predicted t_0 as a function of the number of target molecules using an estimated value of c_{max} . FIG. 2I is in qualitative agreement with the experimental data of FIG. 2F.

Estimation of the reaction rate constant

[0173] To estimate the reaction rate constant, LAMP amplification was carried out in the nucleometer's sample well (FIG. 10). The amplification process was approximated with

$$\frac{dc}{dt} = kc \left(1 - \frac{c}{c_{max}}\right) \quad (6)$$

where c is the concentration (mol/m^3), k (s^{-1}) is the reproductive parameter, and c_{max} (mol/m^3) is the saturation concentration. The fluorescence emission intensity was assumed to be proportional to the concentration. This assumption is not critical and equation (6) could have been formulated in terms of the emission intensity instead of the concentration. It is convenient to introduce the

normalized concentration $\hat{c} = \frac{c}{c_{max}}$. Accordingly, equation (6) reduces to

$$\frac{d\hat{c}}{dt} = k\hat{c}(1 - \hat{c}) \quad (7)$$

with the initial condition:

$$\hat{c}(0) = \hat{c}_0 \quad (8)$$

Equation (7) with initial condition (8) admits the solution:

$$\hat{c}(t) = \frac{\hat{c}_0}{(1 - \hat{c}_0)\exp(-kt) + \hat{c}_0} \quad (9)$$

By minimizing the discrepancy between the predictions of equation (9) and the experimental data, the reaction rate constant k and $\hat{c}$ were estimated. FIG. 10D depicts the predictions of

equation (9) with the optimal estimate $k = 0.008 \text{ s}^{-1}$ and $\hat{c} = 0.006$ (solid lines) along with the experimental data (symbols). The number of target molecules is 10^3 copies.

Estimating the Diffusion Coefficient (D)

[0174] To estimate the diffusion coefficient of nucleic acids in the HPMC polymer solution, oligonucleotides tagged with a fluorescent dye (HEX-GGTGTCTCATTGTTTATACTA) were introduced into the sample well and monitored the nucleic acid diffusion in the microconduit as a function of time (FIG. 11A). This experiment was carried out at the LAMP incubation temperature of 62.5°C , but in the absence of enzymes so that no amplification took place. The normalized signal intensity (normalized concentration) is depicted (dashed lines) in FIG. 10B as a function of position along the conduit at various times.

[0175] The concentration distribution was modeled with a diffusion equation in a semi-infinite medium:

$$\frac{\partial \hat{c}}{\partial t} = D \frac{\partial^2 \hat{c}}{\partial x^2} \quad 0 \leq t \leq \tau, \quad x \geq 0 \quad (10)$$

with the initial and boundary conditions

$$\hat{c}(0, t) - 1 = \hat{c}(\infty, t) = \hat{c}(x, 0) = 0 \quad (11)$$

In the above, $\hat{c} = \frac{c}{c(0, t)}$ where $c(0, t)$ is assumed to be time-independent. Since the volume of the sample chamber far exceeds the volume of the conduit, the error introduced by assuming that $c(0, t)$ remains constant throughout the process is quite small, as has been verified both by scaling analysis and by obtaining a more accurate solution for equation (10) that allows for time-dependence of $c(0, t)$ as mandated by mass conservation.

[0176] Equations (10) and (11) admit the classical solution

$$\hat{c} = \text{erfc}\left(\frac{x}{2\sqrt{Dt}}\right) \quad (12)$$

[0177] Using the MatLab Optimization toolbox (The MathWorks, Inc., Natick, MA), it was observed that $D \sim 10^{-10} \text{ m}^2\text{s}^{-1}$ minimized the discrepancy between the predictions of equation (12) and the experimental data. The predictions of equation (12) with the optimal D are depicted (dashed lines) along with the experimental data (symbols) in FIG. 11B.

[0178] Disclosed herein are devices and endpoint methods for the quantification of nucleic acids undergoing enzymatic amplification. Unlike traditional quantitative fluorescence enzymatic amplification detection, the disclosed devices and methods do not require continuous monitoring of the amplicons and is based on inferring the number of target nucleic acid molecules from the position of the amplification reaction front. The nucleometer requires only a single image to quantify the nucleic acid at a prescribed time or, in the presence of a calibration column, at any time. The nucleometer is compatible with multiplexing, allowing low-cost, high throughput nucleic acid screening, and it may be readily combined with modules for nucleic acid isolation, concentration, purification, and self-heating to facilitate inexpensive non-instrumented, quantitative, on-site molecular detection.

[0179] Those skilled in the art will appreciate that numerous changes and modifications can be made to the preferred embodiments of the invention and that such changes and modifications can be made without departing from the spirit of the invention. It is, therefore, intended that the appended claims cover all such equivalent variations as fall within the true spirit and scope of the invention.

[0180] The disclosures of each patent, patent application, and publication cited or described in this document are hereby incorporated herein by reference, in its entirety.

EMBODIMENTS

The following list of embodiments is intended to complement, rather than displace or supersede, the previous descriptions.

Embodiment 1. A device for monitoring nucleic acid amplification comprising a nucleometer comprising a sample chamber and at least one reaction-diffusion conduit in fluid communication with the sample chamber, each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.

Embodiment 2. The device of embodiment 1, wherein the sample chamber has a size and shape suitable for containing a nucleic acid amplification reaction.

Embodiment 3. The device of embodiment 1 or 2, wherein the sample chamber and the conduit are separated with a barrier.

Embodiment 4. The device of any one of the previous embodiments, wherein the barrier is removable, is comprised of a material that degrades, dissolves, or melts, is movable, or any combination thereof.

Embodiment 5. The device of any one of the previous embodiments, having a plurality of reaction-diffusion conduits in fluid communication with the sample chamber.

Embodiment 6. The device of any one of the previous embodiments, having a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith.

Embodiment 7. The device of embodiment 6, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.

Embodiment 8. The device of embodiment 7, at least one sample chamber having a known quantity of a preselected, target nucleic acid molecule.

Embodiment 9. The device of any one of the previous embodiments, the sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification.

Embodiment 10. The device of any one of the previous embodiments, further comprising a waste conduit, a light source, an optical imager, a heat source, or any combination thereof.

Embodiment 11. The device of embodiment 10, wherein the optical imager is a fluorescent microscope or a camera.

Embodiment 12. The device of embodiment 10 or 11, wherein the heat source is a thermoelectric unit, an electrical heater, an exothermic reactor, or a laser.

Embodiment 13. The device of embodiment 12, wherein the heat source provides a spatially varying temperature distribution along the length of the reaction-diffusion conduit.

Embodiment 14. The device of embodiment 12 or 13, wherein the temperature is regulated with a phase change material.

Embodiment 15. The device of any one of embodiments 12-14, wherein the temperature in the sample chamber is different from the temperature in the conduit.

Embodiment 16. The device of any one of the previous embodiments, said device being operatively connected to a controller or computer.

Embodiment 17. The device of any one of the previous embodiments, further comprising a ruler along the length of the reaction-diffusion conduit.

Embodiment 18. The device of any one of the previous embodiments, further comprising a blend of reactants.

Embodiment 19. The device of embodiment 18, wherein the blend of reactants comprises a plurality of primers and enzymes.

Embodiment 20. The device of embodiment 19, having at least two nucleometers, each containing different primers.

Embodiment 21. The device of embodiment 19, wherein the nucleometer has multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers.

Embodiment 22. The device of any one of the previous embodiments, wherein the reaction diffusion conduit contains at least one polymer.

Embodiment 23. The device of any one of the previous embodiments, wherein the sample chamber contains at least one polymer.

Embodiment 24. The device of embodiment 23, wherein the at least one polymer is a gel.

Embodiment 25. The device of embodiment 24, wherein the primers, enzymes, or both are immobilized to the gel.

Embodiment 26. The device of any one of embodiments 23-25, wherein the polymer is a solid at an ambient temperature and a liquid at an amplification temperature.

Embodiment 27. The device of any one of the previous embodiments, wherein the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.

Embodiment 28. A nucleic acid amplification monitor comprising a substrate and, on or forming a part of the substrate, a plurality of nucleometers, each nucleometer comprising an sample chamber and at least one reaction-diffusion conduit in fluid communication with the chamber, each of the chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.

Embodiment 29. The nucleic acid amplification monitor of embodiment 28, wherein the sample chamber has a size and shape suitable for containing a nucleic acid amplification reaction.

Embodiment 30. The nucleic acid amplification monitor of embodiment 28 or 29, wherein the sample chamber and the conduit are separated with a barrier.

Embodiment 31. The nucleic acid amplification monitor of embodiment 30, wherein the barrier is removable, is comprised of a material that degrades, dissolves, or melts, is movable, or any combination thereof.

Embodiment 32. The nucleic acid amplification monitor of any one of embodiments 28-31, having a plurality of reaction-diffusion conduits in fluid communication with the sample chamber.

Embodiment 33. The nucleic acid amplification monitor of any one of embodiments 28-32, having a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith.

Embodiment 34. The nucleic acid amplification monitor of embodiment 33, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.

Embodiment 35. The nucleic acid amplification monitor of embodiment 34, at least one sample chamber having a known quantity of a preselected, target nucleic acid molecule.

Embodiment 36. The nucleic acid amplification monitor of any one of embodiments 28-35, the sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification.

Embodiment 37. The nucleic acid amplification monitor of any one of embodiments 28-36, further comprising a waste conduit, a light source, an optical imager, a heat source, or any combination thereof.

Embodiment 38. The nucleic acid amplification monitor of embodiment 37, wherein the optical imager is a fluorescent microscope or a camera.

Embodiment 39. The nucleic acid amplification monitor of embodiment 37 or 38, wherein the heat source is a thermoelectric unit, an electrical heater, an exothermic reactor, or a laser.

Embodiment 40. The nucleic acid amplification monitor of embodiment 39, wherein the heat source provides a spatially varying temperature distribution along the length of the reaction-diffusion conduit.

Embodiment 41. The nucleic acid amplification monitor of embodiment 39 or 40, wherein the temperature is regulated with a phase change material.

Embodiment 42. The nucleic acid amplification monitor of any one of embodiments 39-41, wherein the temperature in the sample chamber is different from the temperature in the conduit.

Embodiment 43. The nucleic acid amplification monitor of any one of embodiments 28-42, said device being operatively connected to a controller or computer.

Embodiment 44. The nucleic acid amplification monitor of any one of embodiments 28-43, further comprising a ruler along the length of the reaction-diffusion conduit.

Embodiment 45. The nucleic acid amplification monitor of any one of embodiments 28-44, further comprising a blend of reactants.

Embodiment 46. The nucleic acid amplification monitor of embodiment 45, wherein the blend of reactants comprises a plurality of primers and enzymes.

Embodiment 47. The nucleic acid amplification monitor of embodiment 46, wherein the nucleometer has multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers.

Embodiment 48. The nucleic acid amplification monitor of any one of embodiments 28-47, wherein the reaction diffusion conduit contains at least one polymer.

Embodiment 49. The nucleic acid amplification monitor of embodiment 48, wherein the at least one polymer is a gel.

Embodiment 50. The nucleic acid amplification monitor of embodiment 49, wherein the primers, enzymes, or both are immobilized to the gel.

Embodiment 51. The nucleic acid amplification monitor of any one of embodiments 48-50, wherein the polymer is a solid at an ambient temperature and a liquid at an amplification temperature.

Embodiment 52. The nucleic acid amplification monitor of any one of embodiments 28-51, wherein the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.

Embodiment 53. The nucleic acid amplification monitor of any one of embodiments 28-52, wherein the substrate is a polymeric chip.

Embodiment 54. The nucleic acid amplification monitor of embodiment 53, wherein the polymeric chip comprises polymethyl methacrylate (PMMA).

Embodiment 55. A method of quantifying nucleic acid amplification comprising amplifying a sample comprising a nucleic acid molecule in the device of any one of embodiments 1-27, or the nucleic acid amplification monitor of any one of embodiments 28-54 to generate an amplified nucleic acid molecule and measuring a reaction-diffusion length of said amplified nucleic acid molecule through the one or more conduits, wherein reaction-diffusion length is proportional to the number of nucleic acid copies in the sample.

Embodiment 56. The method of embodiment 55, wherein amplifying the nucleic acid molecule comprises thermo-cycling or isothermal nucleic acid amplification.

Embodiment 57. The method of embodiment 55 or 56, further comprising comparing the reaction-diffusion length of the one or more reaction-diffusion conduits.

Embodiment 58. The method of embodiment 55-57, further comprising comparing the reaction-diffusion length of the one or more reaction-diffusion conduits to a control

Embodiment 59. The method of any one of embodiments 55-58, wherein the nucleic acid molecule is RNA.

Embodiment 60. The method of embodiment 59, further comprising, prior to the amplifying, reverse transcribing the RNA to generate cDNA.

Embodiment 61. A method of identifying an unknown nucleic acid molecule, comprising: amplifying a sample comprising an unknown nucleic acid molecule in the device of any one of embodiments 1-27, or the nucleic acid amplification monitor of any one of embodiments 28-54, wherein each of the at least one conduits contain a plurality of primers specific for a different known nucleic acid molecule; and measuring a reaction-diffusion length within the reaction-diffusion conduit, wherein the presence of the reaction-diffusion length

indicates that the sample having an unknown nucleic acid molecule comprises the known nucleic acid molecule to which the primers in the at least one conduit are specific and the extent of the reaction-diffusion length within the conduit indicates the number of the unknown nucleic acid molecules in the sample.

Embodiment 62. The method of embodiment 61, comprising determining the difference in length of reaction-diffusion from a nucleometer having a sample having a known quantity of a nucleic acid and a nucleometer having a sample having an unknown nucleic acid molecule, wherein the difference in the length of the reaction-diffusion between the known quantity of a nucleic acid molecule and the sample having an unknown nucleic acid molecule indicates a number of nucleic acid molecules in the sample having the unknown nucleic acid molecule.

Embodiment 63. A system for monitoring nucleic acid amplification comprising: a nucleometer comprising a sample chamber, at least one reaction-diffusion conduit in fluid communication with the sample chamber, each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification; an optical imager; and a heat source.

Embodiment 64. The system of embodiment 63, further comprising a controller.

Embodiment 65. The system of any one of embodiments 63 or 64, further comprising an output device.

Embodiment 66. The system of embodiment 65, each of the optical imager, heat source and output device being operably controlled by the controller.

What is claimed:

1. A device for monitoring nucleic acid amplification comprising:
 - a nucleometer comprising:
 - a sample chamber; and
 - at least one reaction-diffusion conduit in fluid communication with the sample chamber; each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.
2. The device of claim 1, wherein the sample chamber has a size and shape suitable for containing a nucleic acid amplification reaction
3. The device of claim 1 or 2, wherein the sample chamber and the conduit are separated with a barrier.
4. The device of any one of the previous claims, wherein the barrier is removable, is comprised of a material that degrades, dissolves, or melts, is movable, or any combination thereof.
5. The device of any one of the previous claims, having a plurality of reaction-diffusion conduits in fluid communication with the sample chamber.
6. The device of any one of the previous claims, having a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith.
7. The device of claim 6, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.
8. The device of claim 7, at least one sample chamber having a known quantity of a preselected, target nucleic acid molecule.

9. The device of any one of the previous claims, the sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification.
10. The device of any one of the previous claims, further comprising a waste conduit, a light source, an optical imager, a heat source, or any combination thereof.
11. The device of claim 10, wherein the optical imager is a fluorescent microscope or a camera.
12. The device of claim 10 or 11, wherein the heat source is a thermoelectric unit, an electrical heater, an exothermic reactor, or a laser.
13. The device of claim 12, wherein the heat source provides a spatially varying temperature distribution along the length of the reaction-diffusion conduit.
14. The device of claim 12 or 13, wherein the temperature is regulated with a phase change material.
15. The device of any one of claims 12-14, wherein the temperature in the sample chamber is different from the temperature in the conduit.
16. The device of any one of the previous claims, said device being operatively connected to a controller or computer.
17. The device of any one of the previous claims, further comprising a ruler along the length of the reaction-diffusion conduit.
18. The device of any one of the previous claims, further comprising a blend of reactants.
19. The device of claim 18, wherein the blend of reactants comprises a plurality of primers and enzymes.
20. The device of claim 19, having at least two nucleometers, each containing different primers.

21. The device of claim 19, wherein the nucleometer has multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers.
22. The device of any one of the previous claims, wherein the reaction diffusion conduit contains at least one polymer.
23. The device of any one of the previous claims, wherein the sample chamber contains at least one polymer.
24. The device of claim 23, wherein the at least one polymer is a gel.
25. The device of claim 24, wherein the primers, enzymes, or both are immobilized to the gel.
26. The device of any one of claims 23-25, wherein the polymer is a solid at an ambient temperature and a liquid at an amplification temperature.
27. The device of any one of the previous claims, wherein the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.
28. A nucleic acid amplification monitor comprising:
 - a substrate and, on or forming a part of the substrate, a plurality of nucleometers, each nucleometer comprising:
 - an sample chamber; and
 - at least one reaction-diffusion conduit in fluid communication with the chamber;
 - each of the chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification.
29. The nucleic acid amplification monitor of claim 28, wherein the sample chamber has a size and shape suitable for containing a nucleic acid amplification reaction.
30. The nucleic acid amplification monitor of claim 28 or 29, wherein the sample chamber and the conduit are separated with a barrier.

31. The nucleic acid amplification monitor of claim 30, wherein the barrier is removable, is comprised of a material that degrades, dissolves, or melts, is movable, or any combination thereof.
32. The nucleic acid amplification monitor of any one of claims 28-31, having a plurality of reaction-diffusion conduits in fluid communication with the sample chamber.
33. The nucleic acid amplification monitor of any one of claims 28-32, having a plurality of sample chambers, each sample chamber having at least one reaction-diffusion conduit in fluid communication therewith.
34. The nucleic acid amplification monitor of claim 33, wherein at least some of the sample chambers have a plurality of reaction-diffusion conduits in fluid communication therewith.
35. The nucleic acid amplification monitor of claim 34, at least one sample chamber having a known quantity of a preselected, target nucleic acid molecule.
36. The nucleic acid amplification monitor of any one of claims 28-35, the sample chamber having a porous membrane for nucleic acids extraction, concentration, and purification.
37. The nucleic acid amplification monitor of any one of claims 28-36, further comprising a waste conduit, a light source, an optical imager, a heat source, or any combination thereof.
38. The nucleic acid amplification monitor of claim 37, wherein the optical imager is a fluorescent microscope or a camera.
39. The nucleic acid amplification monitor of claim 37 or 38, wherein the heat source is a thermoelectric unit, an electrical heater, an exothermic reactor, or a laser.
40. The nucleic acid amplification monitor of claim 39, wherein the heat source provides a spatially varying temperature distribution along the length of the reaction-diffusion conduit.

41. The nucleic acid amplification monitor of claim 39 or 40, wherein the temperature is regulated with a phase change material.
42. The nucleic acid amplification monitor of any one of claims 39-41, wherein the temperature in the sample chamber is different from the temperature in the conduit.
43. The nucleic acid amplification monitor of any one of claims 28-42, said device being operatively connected to a controller or computer.
44. The nucleic acid amplification monitor of any one of claims 28-43, further comprising a ruler along the length of the reaction-diffusion conduit.
45. The nucleic acid amplification monitor of any one of claims 28-44, further comprising a blend of reactants.
46. The nucleic acid amplification monitor of claim 45, wherein the blend of reactants comprises a plurality of primers and enzymes.
47. The nucleic acid amplification monitor of claim 46, wherein the nucleometer has multiple reaction-diffusion conduits, at least two of said reaction-diffusion conduits containing different primers.
48. The nucleic acid amplification monitor of any one of claims 28-47, wherein the reaction diffusion conduit contains at least one polymer.
49. The nucleic acid amplification monitor of claim 48, wherein the at least one polymer is a gel.
50. The nucleic acid amplification monitor of claim 49, wherein the primers, enzymes, or both are immobilized to the gel.
51. The nucleic acid amplification monitor of any one of claims 48-50, wherein the polymer is a solid at an ambient temperature and a liquid at an amplification temperature.

52. The nucleic acid amplification monitor of any one of claims 28-51, wherein the sample chamber, reaction-diffusion conduit, or both are suitable for reverse transcription.
53. The nucleic acid amplification monitor of any one of claims 28-52, wherein the substrate is a polymeric chip.
54. The nucleic acid amplification monitor of claim 53, wherein the polymeric chip comprises polymethyl methacrylate (PMMA).
55. A method of quantifying nucleic acid amplification comprising:
 amplifying a sample comprising a nucleic acid molecule in the device of any one of claims 1-27, or the nucleic acid amplification monitor of any one of claims 28-54 to generate an amplified nucleic acid molecule; and
 measuring a reaction-diffusion length of said amplified nucleic acid molecule through the one or more conduits, wherein the reaction-diffusion length is proportional to the number of nucleic acid copies in the sample.
56. The method of claim 55, wherein amplifying the nucleic acid molecule comprises thermo-cycling or isothermal nucleic acid amplification.
57. The method of claim 55 or 56, further comprising comparing the reaction-diffusion length of the one or more reaction-diffusion conduits.
58. The method of claim 55-57, further comprising comparing the reaction-diffusion length of the one or more reaction-diffusion conduits to a control.
59. The method of any one of claims 55-58, wherein the nucleic acid molecule is RNA.
60. The method of any one of claims 59, further comprising, prior to the amplifying, reverse transcribing the RNA to generate cDNA.
61. A method of identifying an unknown nucleic acid molecule, comprising:

amplifying a sample comprising an unknown nucleic acid molecule in the device of any one of claims 1-27, or the nucleic acid amplification monitor of any one of claims 28-54, wherein each of the at least one conduits contain a plurality of primers specific for a different known nucleic acid molecule; and measuring a reaction-diffusion length within the reaction-diffusion conduit, wherein the presence of the reaction-diffusion length indicates that the sample having an unknown nucleic acid molecule comprises the known nucleic acid molecule to which the primers in the at least one conduit are specific and the extent of the reaction-diffusion length within the conduit indicates the number of the unknown nucleic acid molecules in the sample.

62. The method of claim 61, comprising:

determining the difference in length of reaction-diffusion from a nucleometer having a sample having a known quantity of a nucleic acid and a nucleometer having a sample having an unknown nucleic acid molecule, wherein the difference in the length of the reaction-diffusion between the known quantity of a nucleic acid molecule and the sample having an unknown nucleic acid molecule indicates a number of nucleic acid molecules in the sample having the unknown nucleic acid molecule.

63. A system for monitoring nucleic acid amplification comprising a nucleometer comprising:

a sample chamber;
at least one reaction-diffusion conduit in fluid communication with the sample chamber; each of the sample chamber and the conduit being capable of holding a blend of reactants for nucleic acid amplification;
an optical imager; and
a heat source.

64. The system of claim 63, further comprising a controller.

65. The system of any one of claims 63 or 64, further comprising an output device.

66. The system of claim 65, each of the optical imager, heat source and output device being operably controlled by the controller.

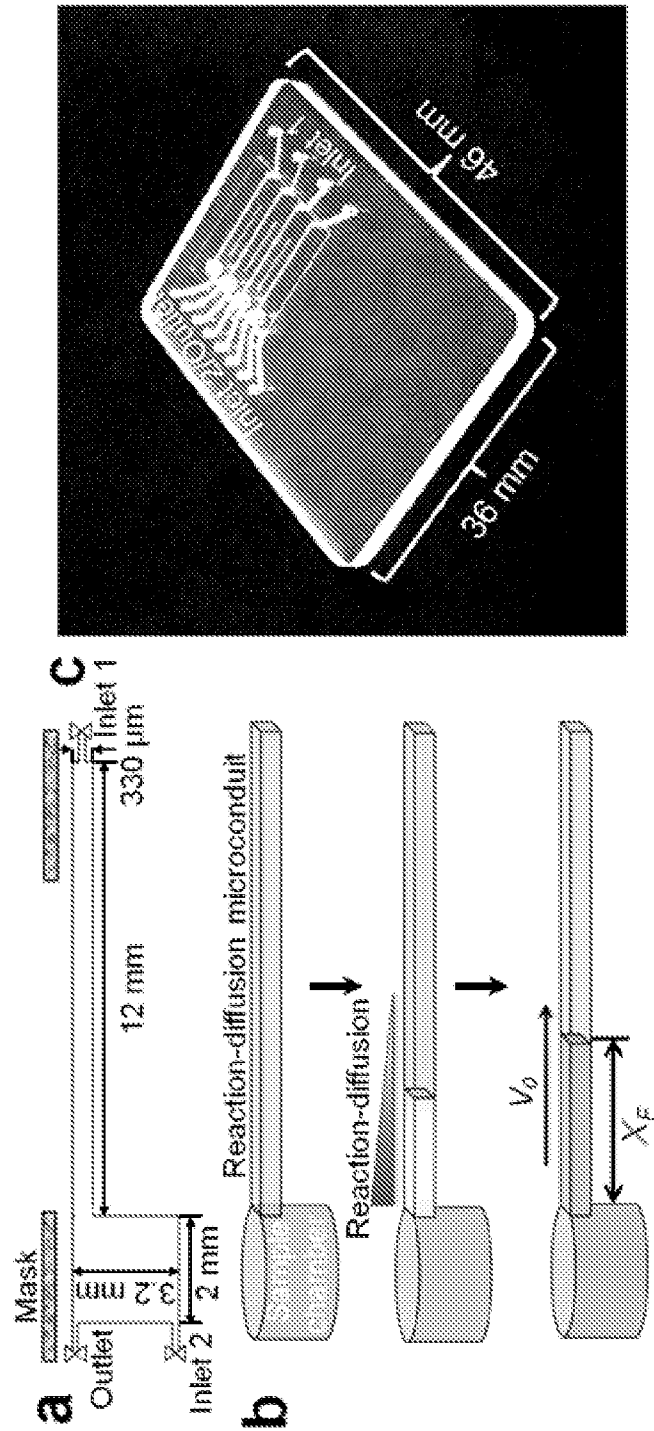


FIGURE 1

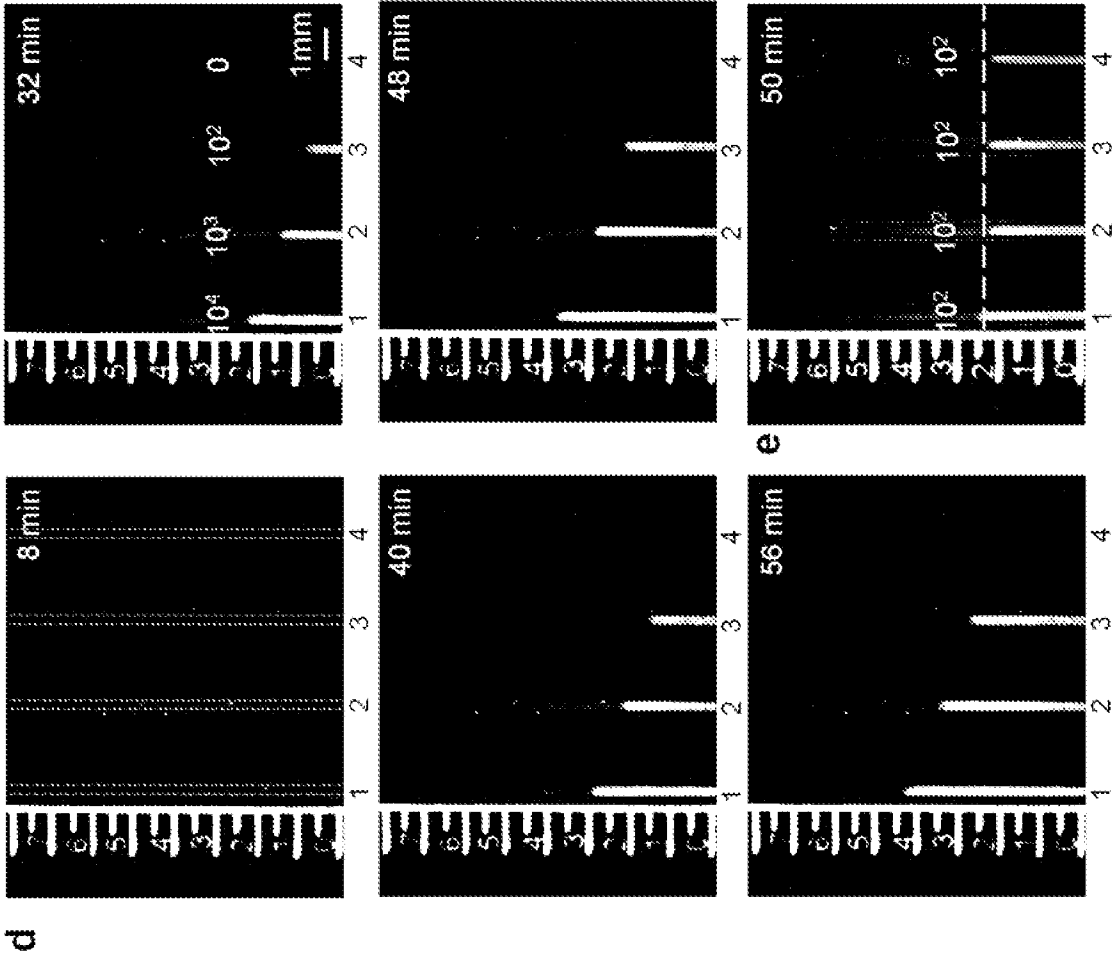


FIGURE 1 cont.

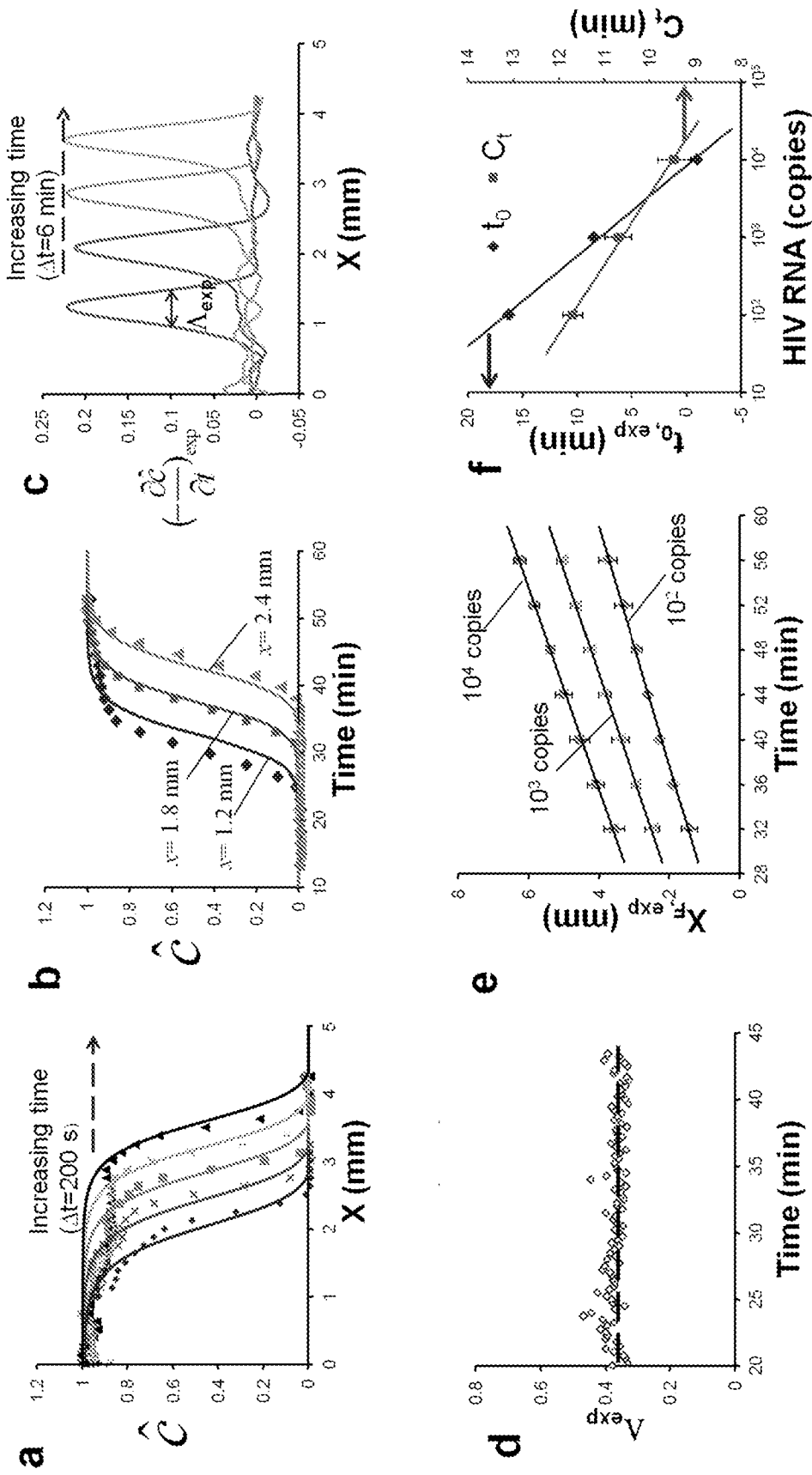


FIGURE 2

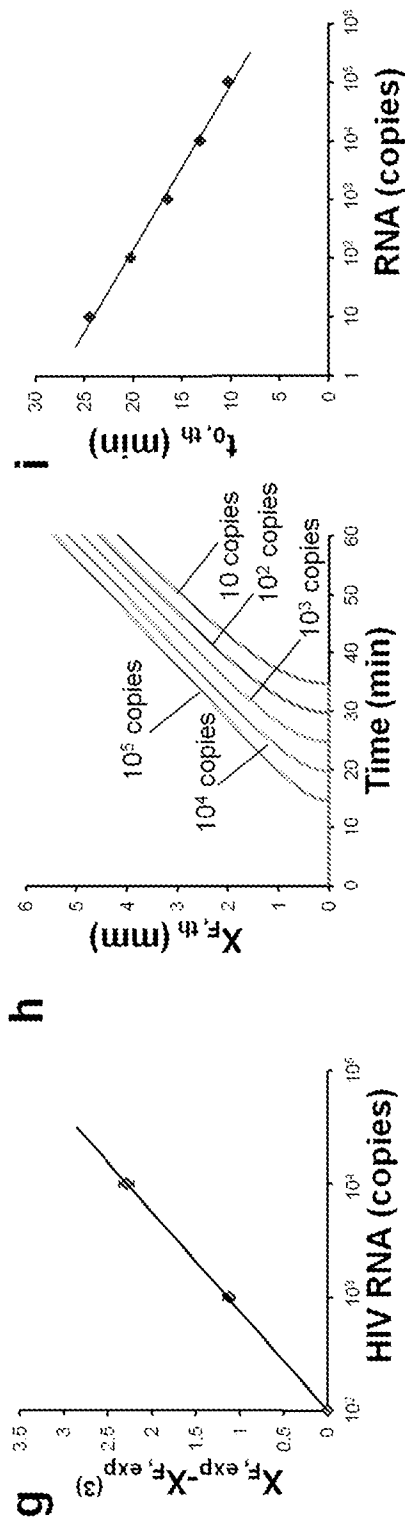


FIGURE 2 cont.

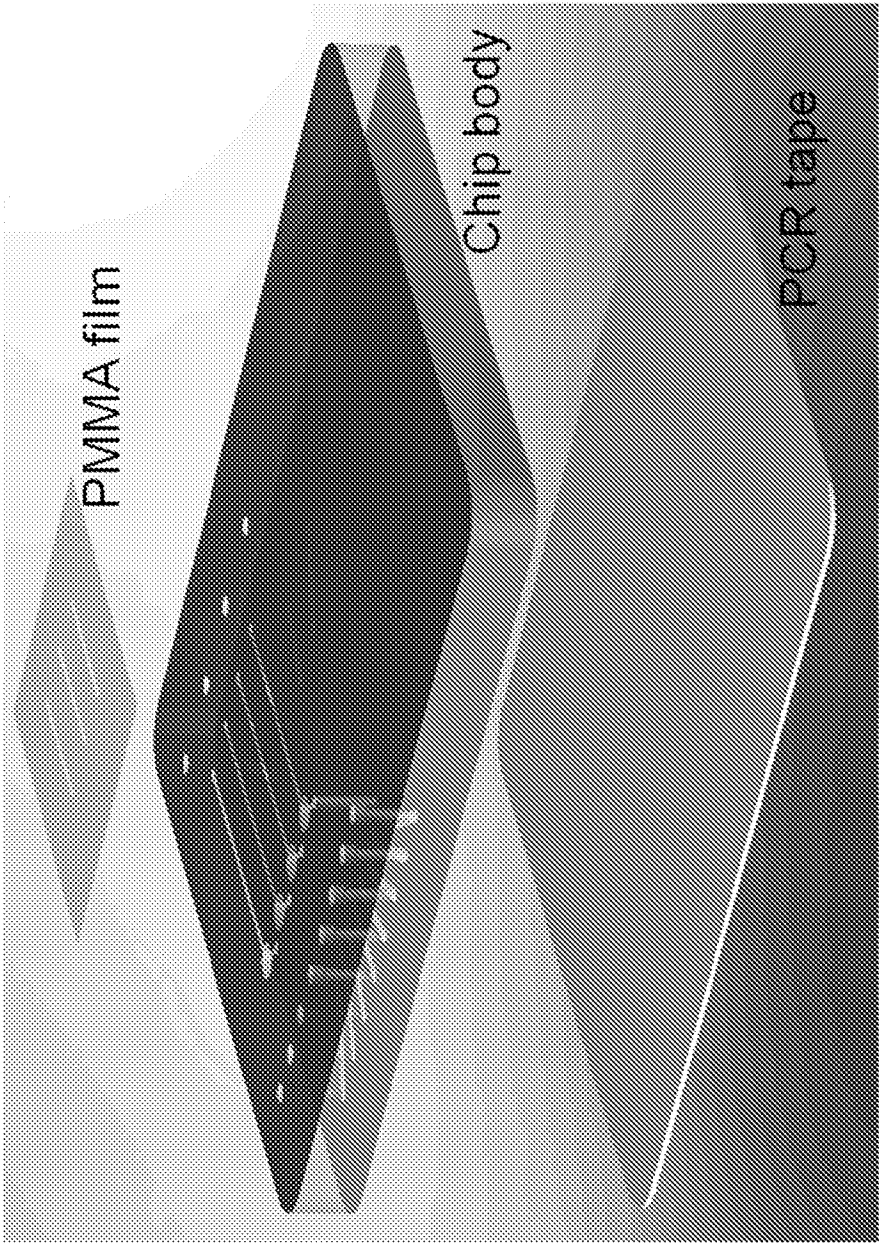


FIGURE 3

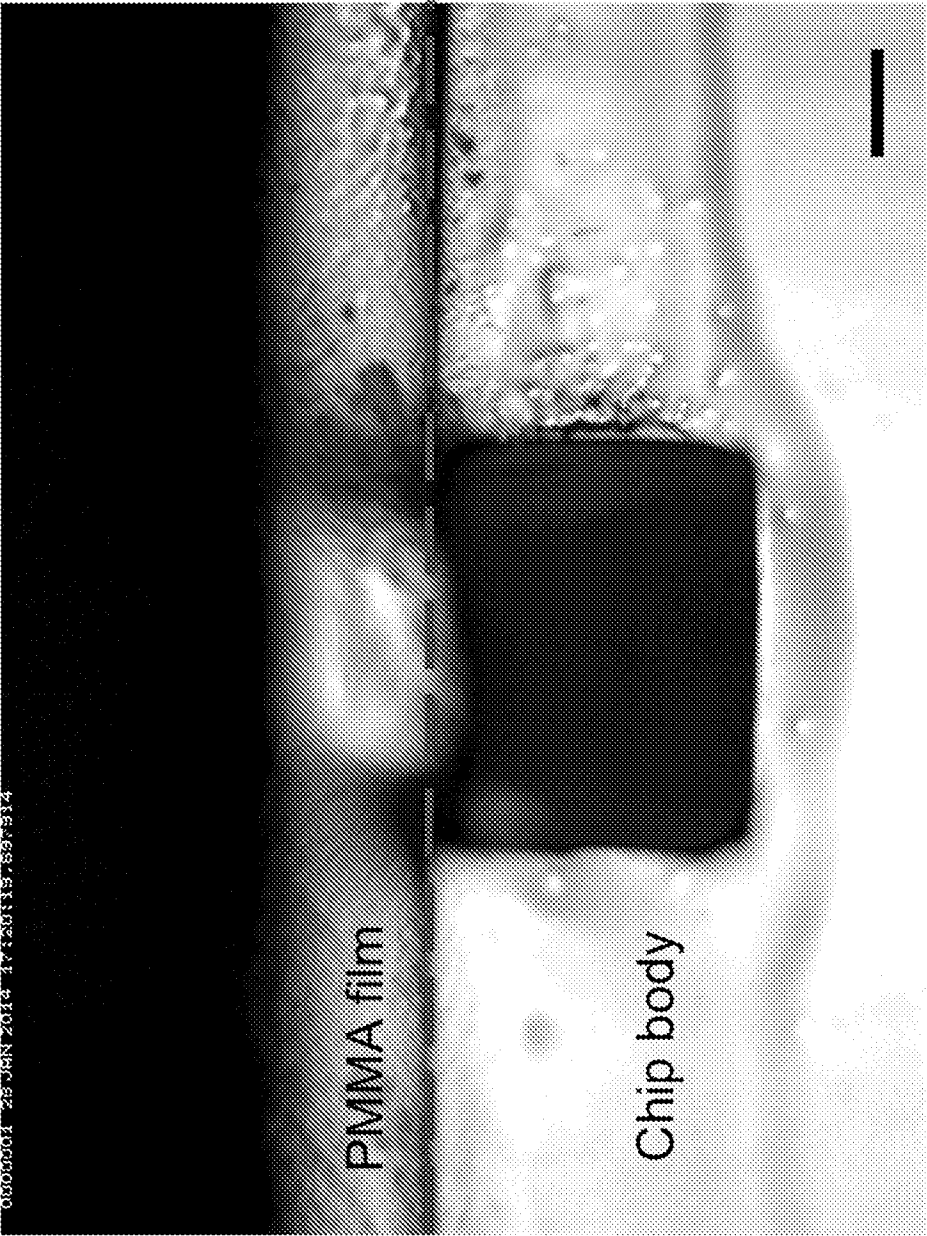


FIGURE 4

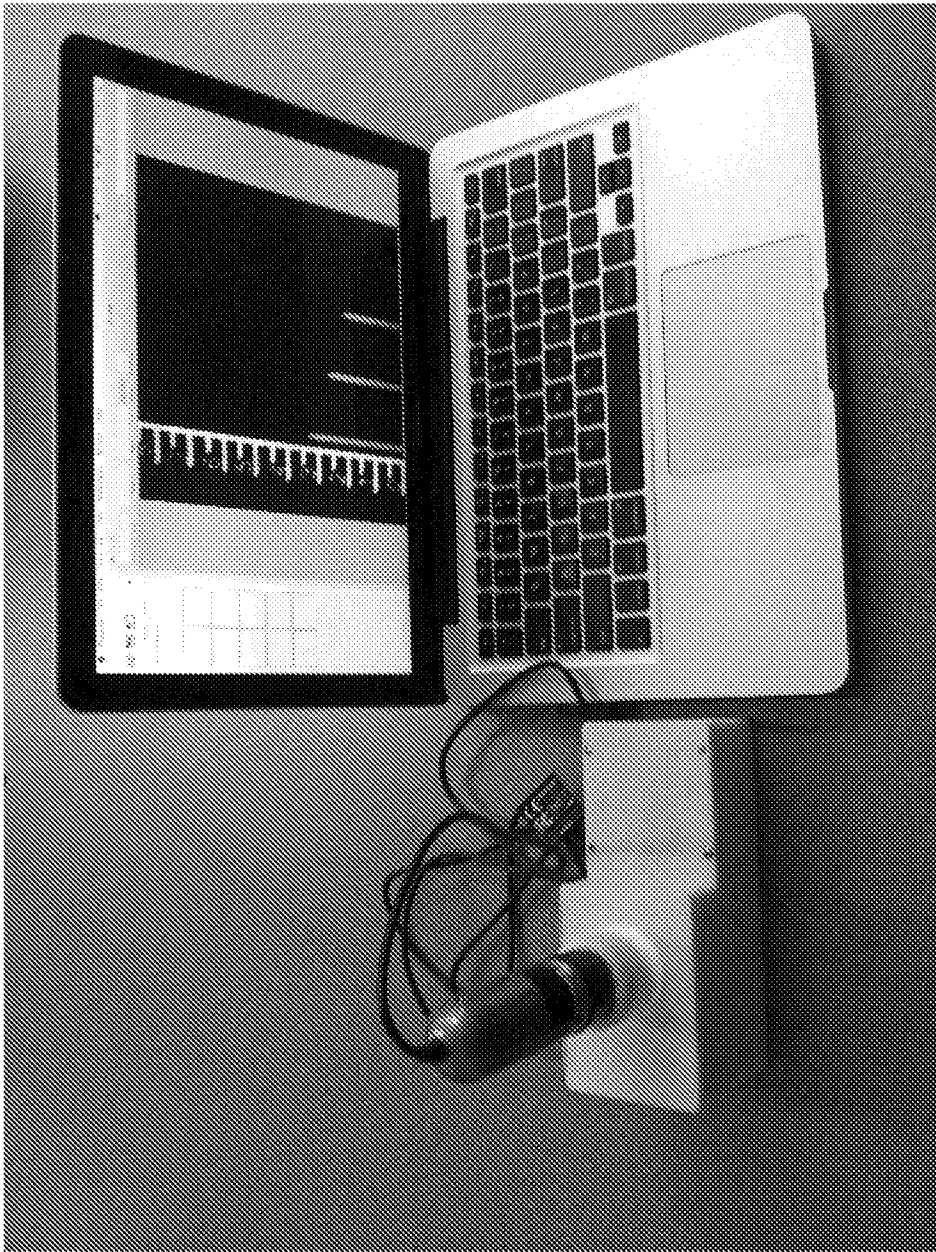


FIGURE 5

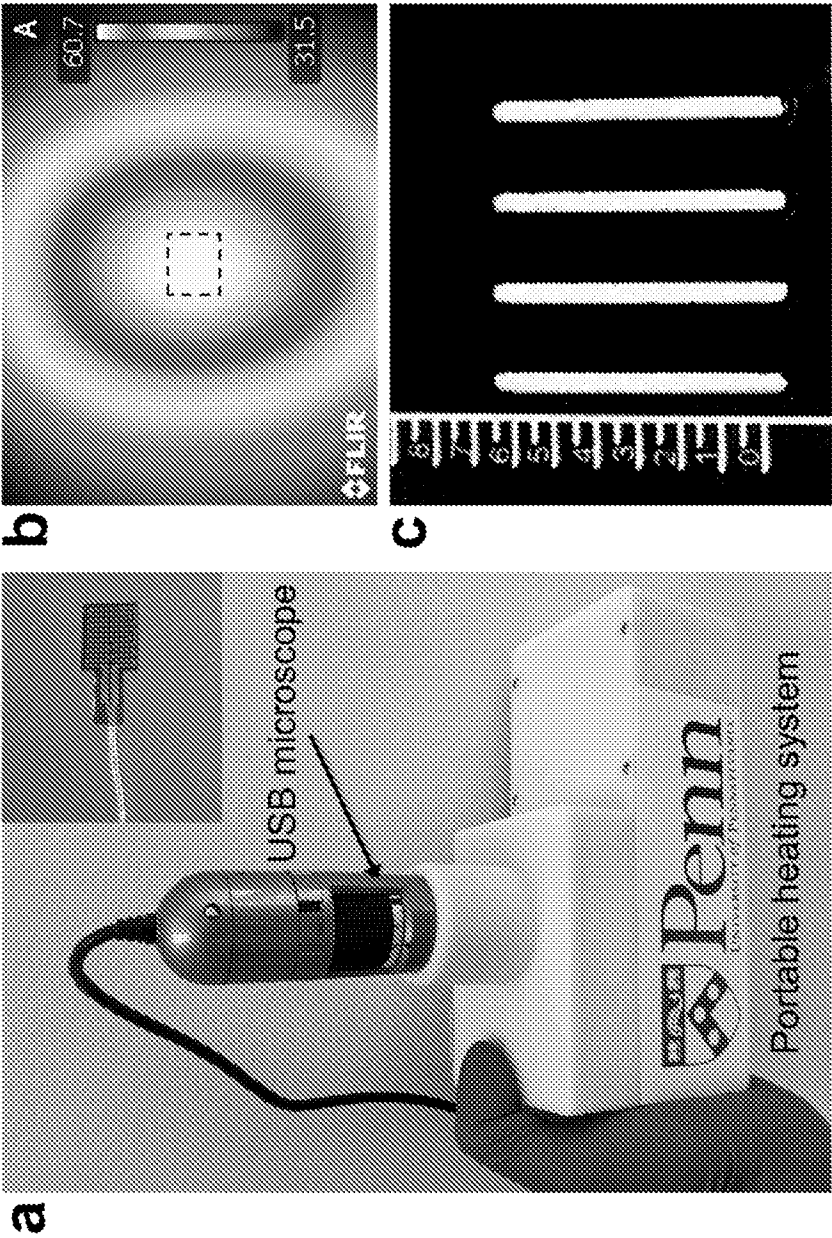


FIGURE 6

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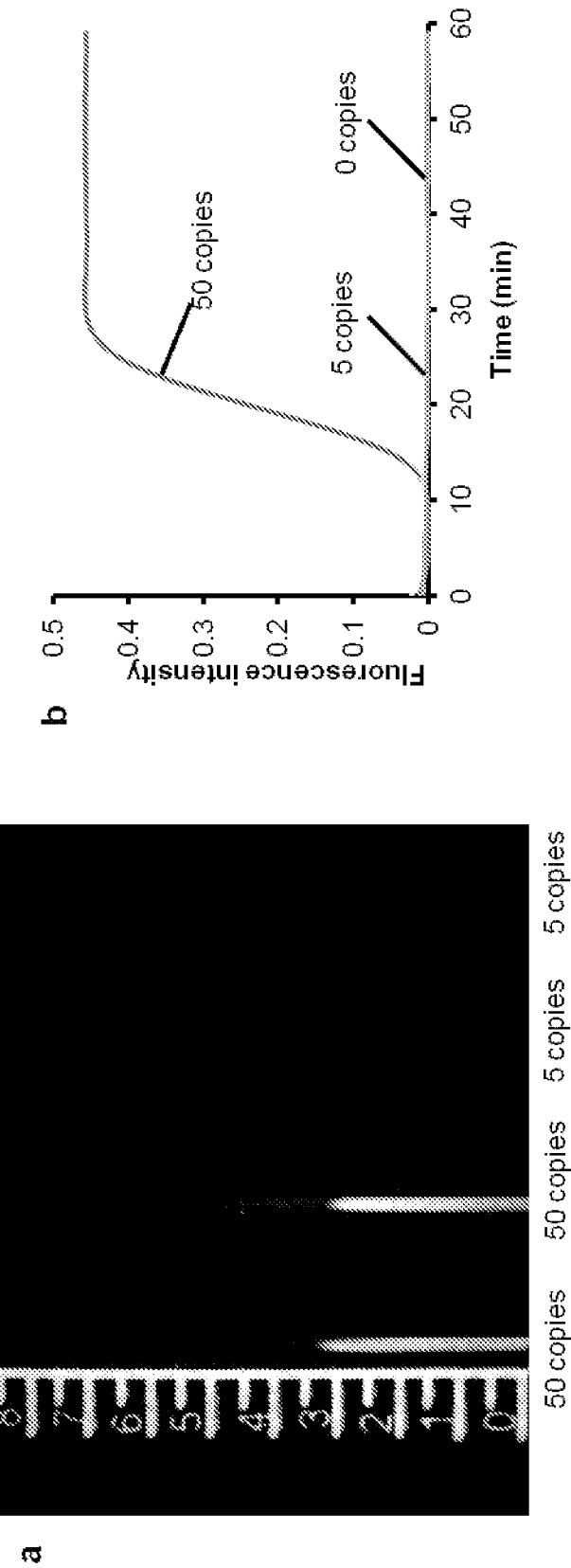


FIGURE 7

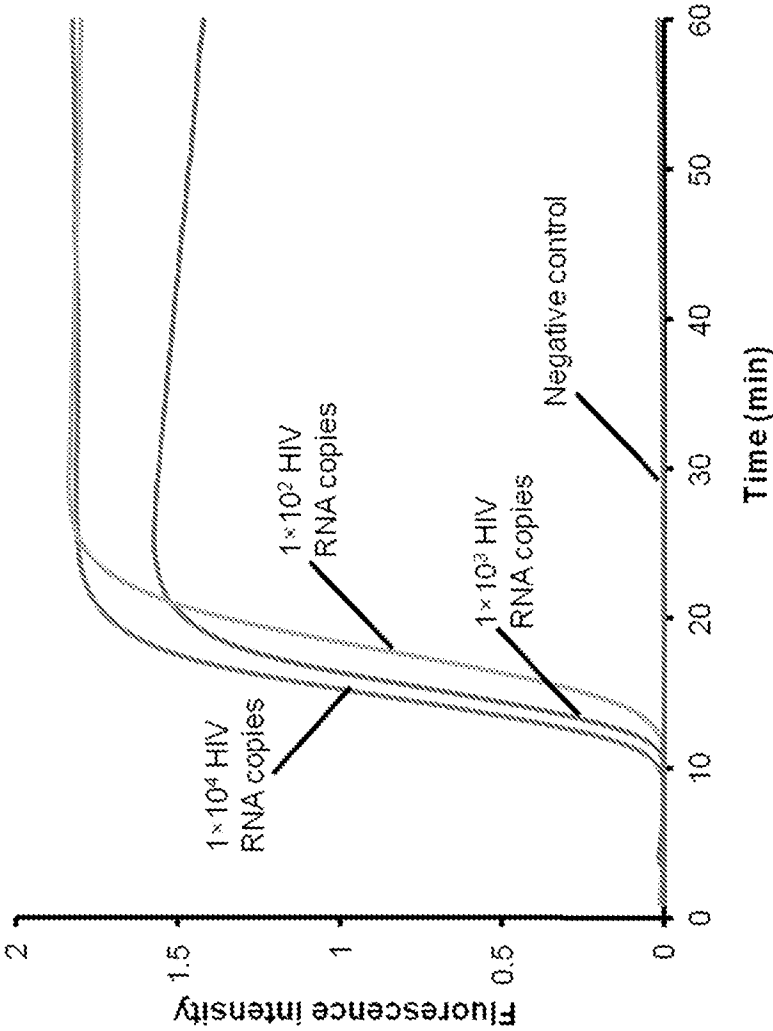


FIGURE 8

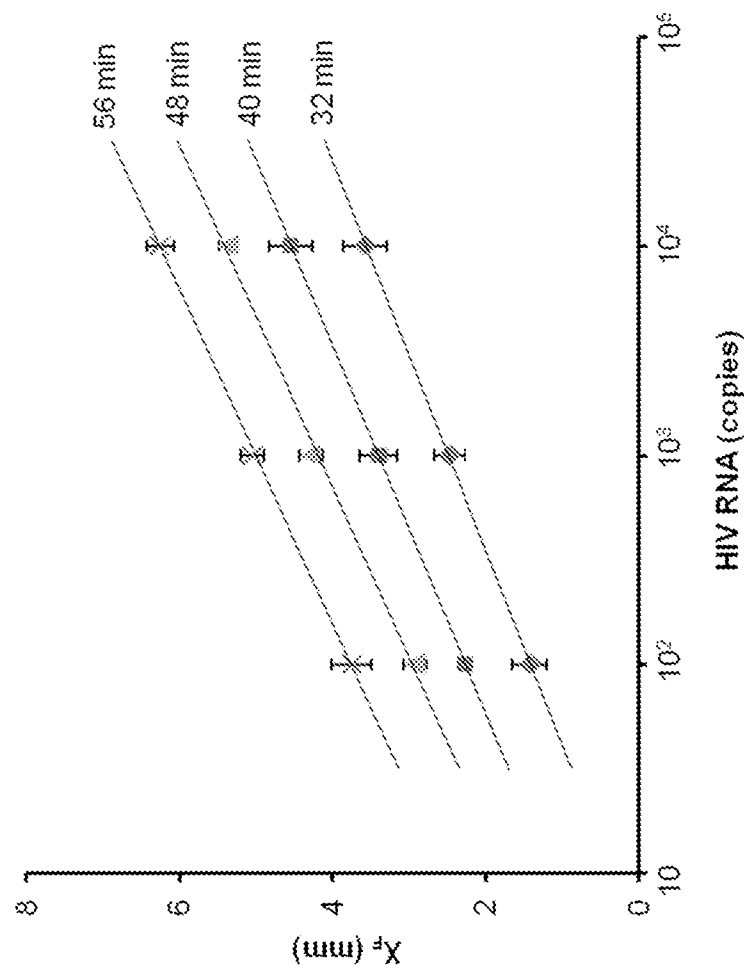


FIGURE 9

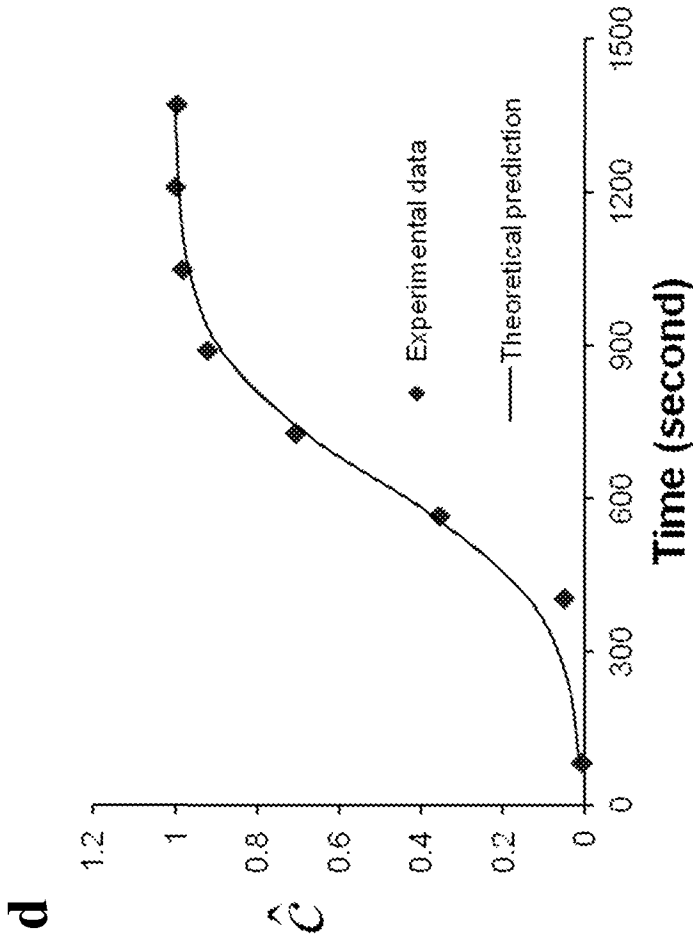
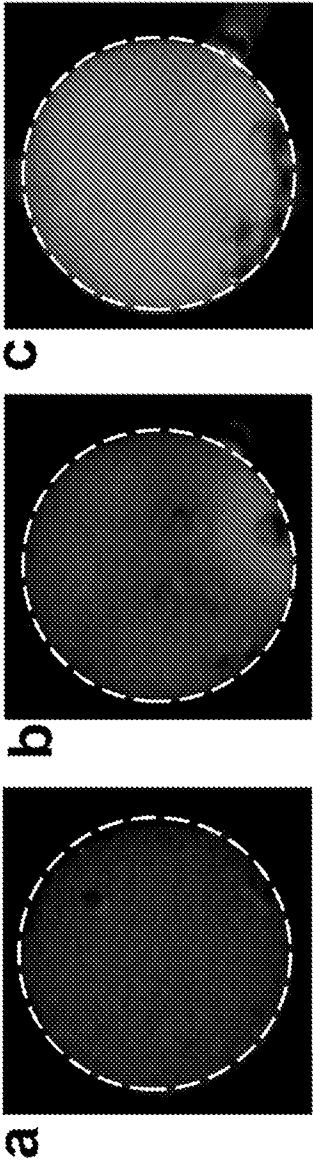


FIGURE 10

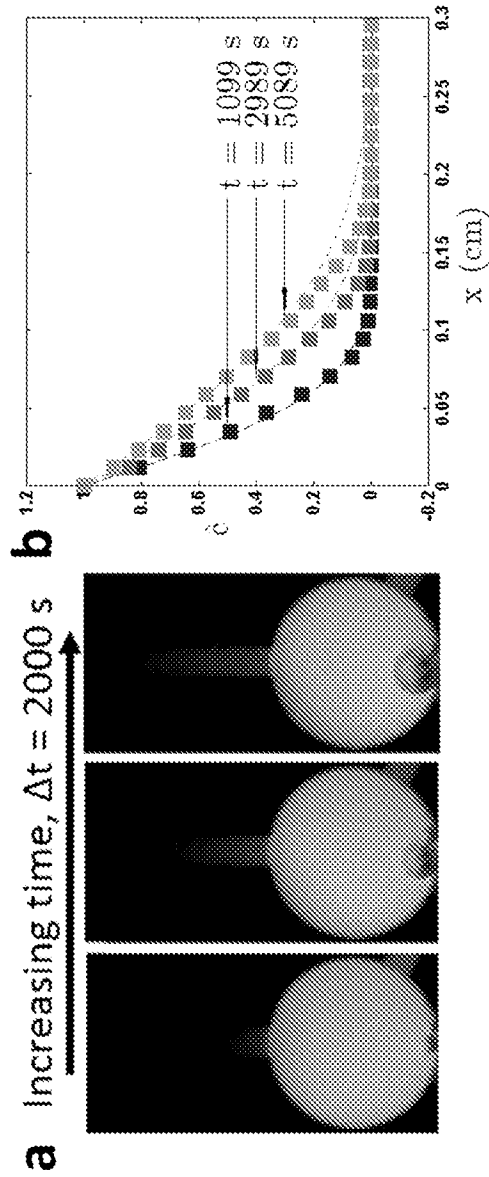


FIGURE 11

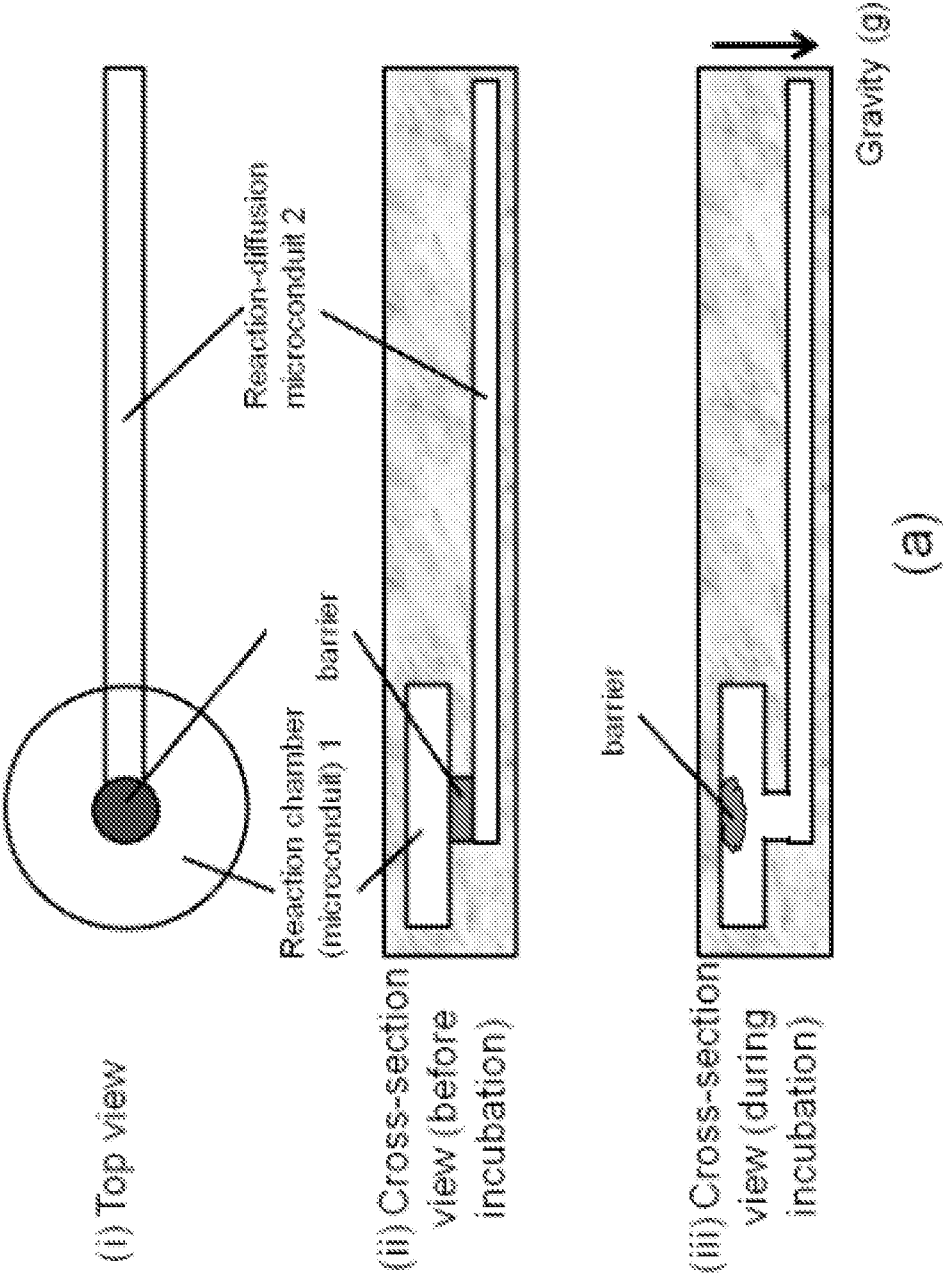


FIGURE 12

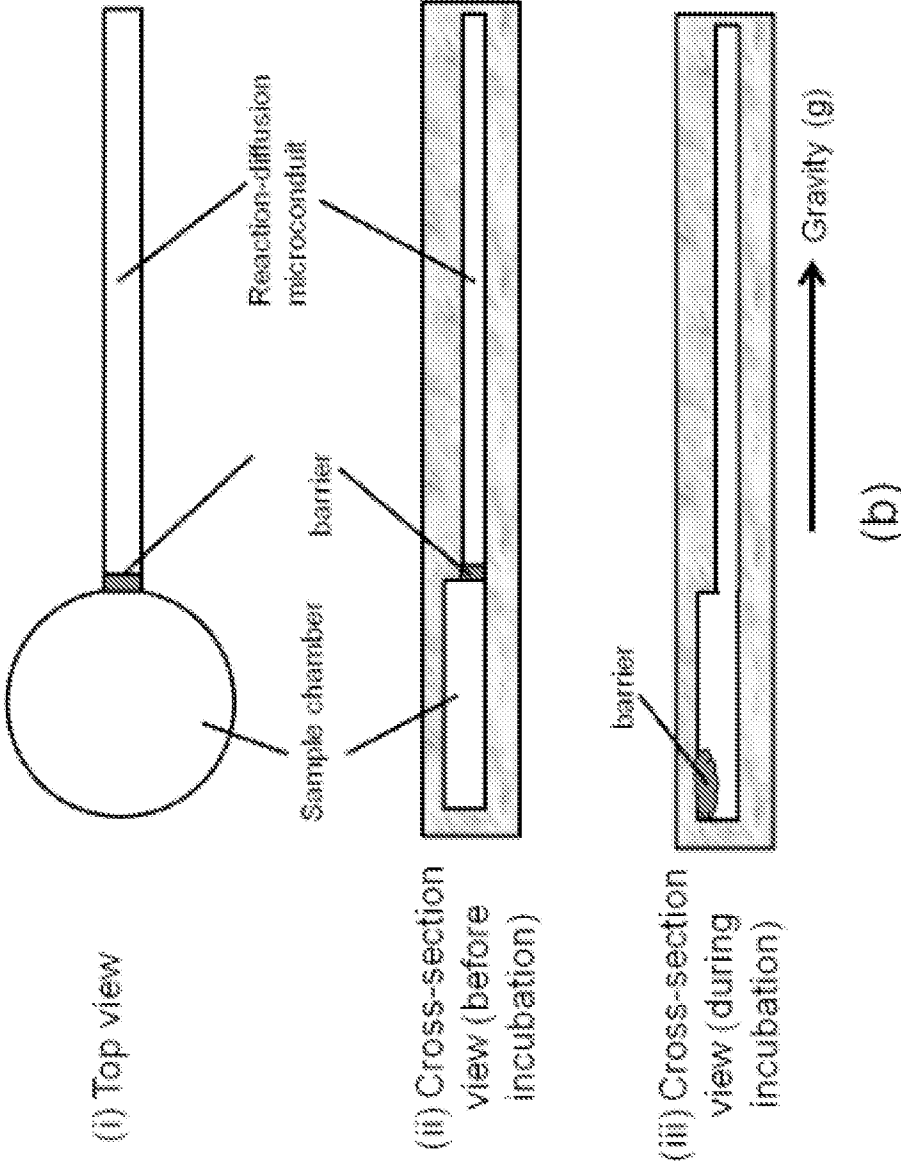


FIGURE 12 cont.

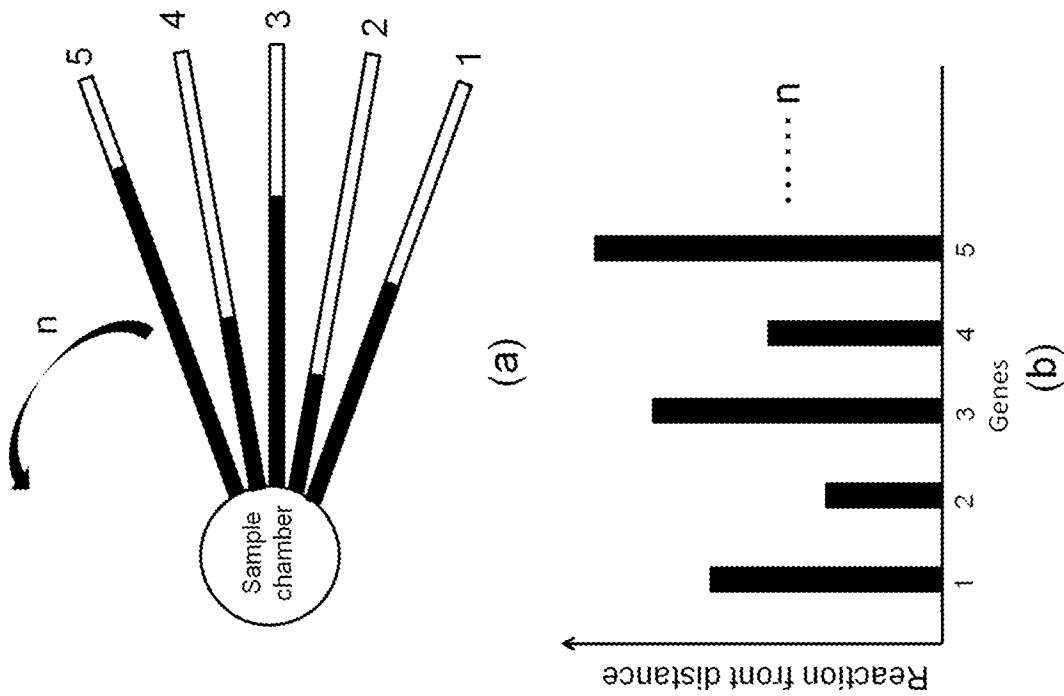


FIGURE 13

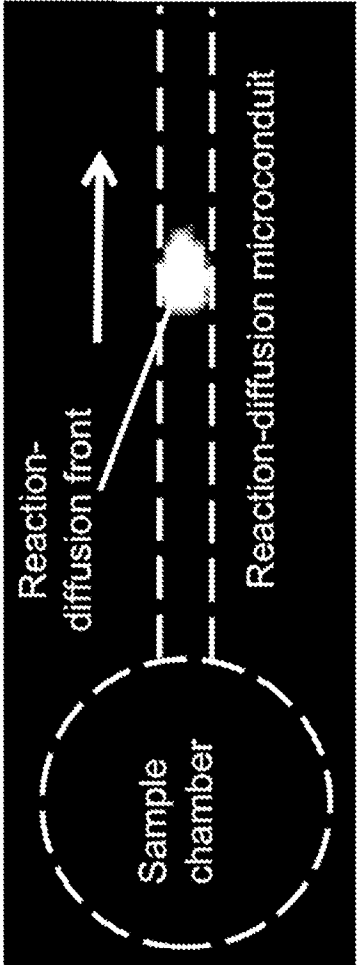


FIGURE 14

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US15/38739

A. CLASSIFICATION OF SUBJECT MATTER

IPC(8) - C12Q 1/68 (2015.01)

CPC - C12Q 1/6844

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC(8): C12Q 1/68 (2015.01)

CPC: C12Q 1/6844

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

PatSeer (US, EP, WO, JP, DE, GB, CN, FR, KR, ES, AU, IN, CA, RU, AT, CH, TH, BR, PH, Other Countries (INPADOC)); PubMed; Google; Google Scholar; Google Patents; 'PCR,' nucleic, amplification, chip, microfluidic, 'reaction-diffusion'

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	CHEN, D et al. An integrated, self-contained microfluidic cassette for isolation, amplification, and detection of nucleic acids. Biomed Microdevices, August 2010, Vol. 12, No. 4, pages 705-719. doi:10.1007/s10544-010-9423-4.	1-3, 28-31, 63-66
A	BAROUD, CN et al. Reaction-Diffusion Dynamics: Confrontation Between Theory And Experiment In A Microfluidic Reactor. arXiv:physics/0208091v3 [physics.flu-dyn] 07 Jul 2003, 5 pages.	1-3, 28-31, 63-66
A	ZHANG, C et al. Survey And Summary: Miniaturized PCR Chips For Nucleic Acid Amplification And Analysis: Latest Advances And Future Trends. Nucleic Acids Research, 2007, Vol. 35, No. 13, pages 4223-4237. DOI:10.1093/nar/gkm389.	1-3, 28-31, 63-66
A	US 2008/0280285 A1 (CHEN, Z et al.) November 13, 2008; abstract	1-3, 28-31, 63-66

☐ Further documents are listed in the continuation of Box C.☐ See patent family annex.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&" document member of the same patent family

Date of the actual completion of the international search

15 September 2015 (15.09.2015)

Date of mailing of the international search report

06 OCT 2015

Name and mailing address of the ISA/

Mail Stop PCT, Attn: ISA/US, Commissioner for Patents
P.O. Box 1450, Alexandria, Virginia 22313-1450
Facsimile No. 571-273-8300

Authorized officer

Shane Thomas

PCT Helpdesk: 571-272-4300
PCT OSP: 571-272-7774

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US15/38739

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☒ Claims Nos.: 4-27, 32-62
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

- ☐ The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.
- ☐ The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.
- ☐ No protest accompanied the payment of additional search fees.