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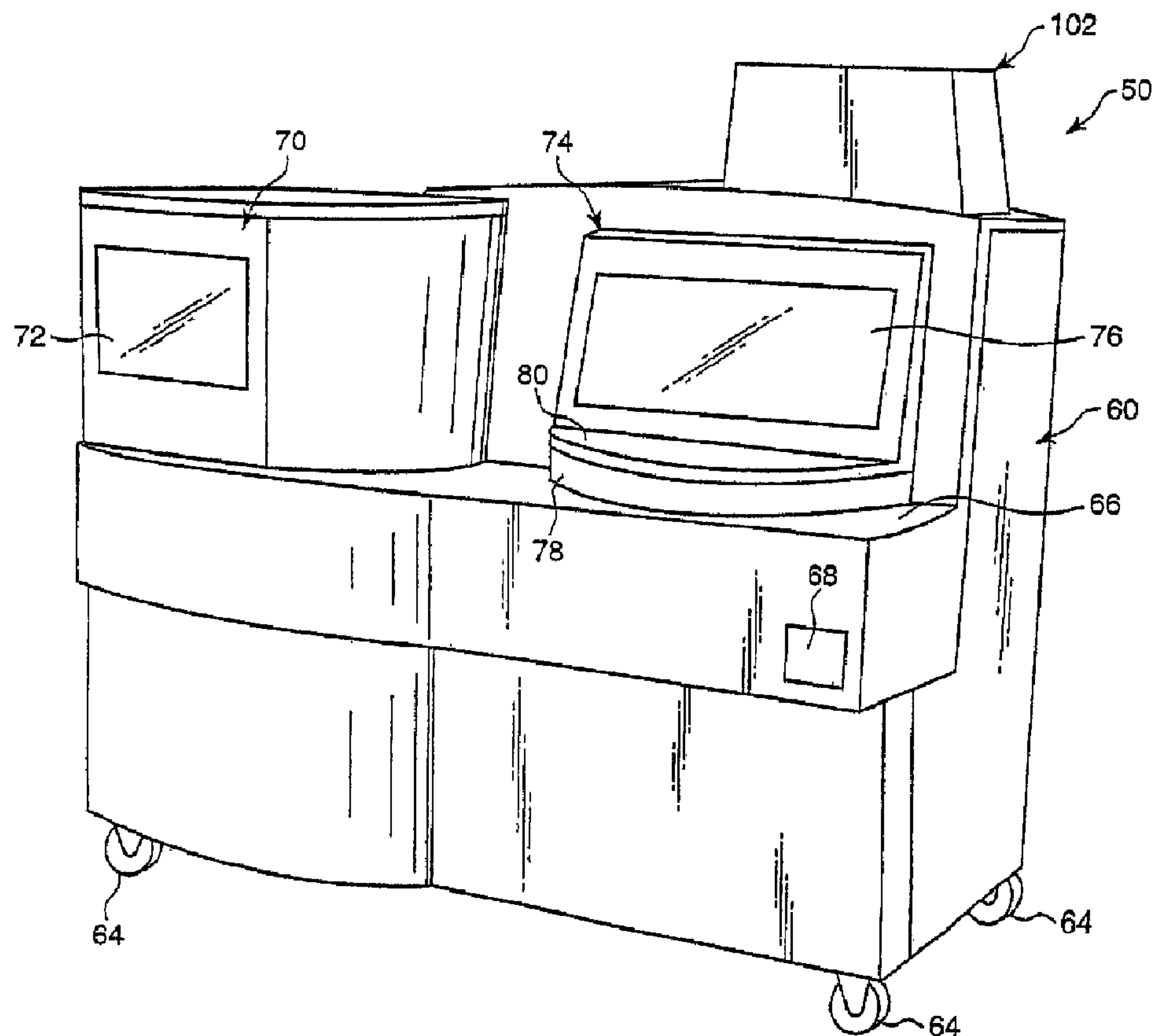
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(54) Titre : APPAREIL ET METHODES DE DETECTION DE SIGNAUX OPTIQUES MULTIPLES

(54) Title: SYSTEM AND METHODS FOR DETECTING MULTIPLE OPTICAL SIGNALS



(57) Abrégé/Abstract:

An automated analyzer for performing multiple diagnostic assays simultaneously includes multiple stations in which discrete aspects of the assay are performed on fluid samples contained in sample vessels. The analyzer includes stations for automatically

(57) **Abrégé(suite)/Abstract(continued):**

preparing a sample, incubating the sample, performing an analyte isolation procedure, ascertaining the presence of a target analyte, and analyzing the amount of a target analyte. An automated receptacle transporting system moves the sample vessels from one station to the next. A method for performing an automated diagnostic assay includes an automated process for isolating and amplifying a target analyte, and, in one embodiment, a method for real-time monitoring of the amplification process.

ABSTRACT

An automated analyzer for performing multiple diagnostic assays simultaneously includes multiple stations in which discrete aspects of the assay are performed on fluid samples contained in sample vessels. The analyzer includes stations for automatically preparing a sample, incubating the sample, performing an analyte isolation procedure, ascertaining the presence of a target analyte, and analyzing the amount of a target analyte. An automated receptacle transporting system moves the sample vessels from one station to the next. A method for performing an automated diagnostic assay includes an automated process for isolating and amplifying a target analyte, and, in one embodiment, a method for real-time monitoring of the amplification process.

SYSTEM AND METHODS FOR DETECTING MULTIPLE OPTICAL SIGNALS

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FIELD OF THE INVENTION

0002 The present invention relates generally to an automated analyzer for simultaneously performing multiple nucleic acid-based assays, and more specifically to a system and method for performing multiple nucleic acid amplification assays, including both real-time and end-point amplifications assays. The present invention also relates to an apparatus and method for continuously processing the contents of a plurality of reaction receptacles following a real-time amplification procedure. The present invention further relates to a method for reducing the presence of amplification inhibitors in reaction receptacles prior to performing nucleic acid amplification reactions.

BACKGROUND OF THE INVENTION

0003 Nucleic acid-based assays can enable highly specific and sensitive detection of nucleic acid analytes from a variety of sources, including clinical, industrial, environmental, and food sources. These assays can be used to determine or monitor for the presence or amount of biological antigens (e.g., prions), cell abnormalities, disease states, and disease-associated pathogens, including parasites, fungi, bacteria and viruses present in a host organism or sample. Nucleic acid-based assays may be qualitative or quantitative, with the quantitative assays providing useful information to practitioners for evaluating the extent of infection or disease or to determine the state of a disease over time. Quantitative assays can also be used, for example, to assess the effectiveness of a therapeutic treatment program or, alternatively, to determine the extent of an infection or contamination by a particular organism or virus.

0004 All nucleic acid-based assay formats involve a number of process steps leading to the

identification, detection or quantification of one or multiple target nucleic acids in a sample. When necessary, the specifically targeted nucleic acid sequences of a nucleic acid-based assay may be unique to an identifiable group of organisms (as used herein, the term "organisms" is inclusive of viruses), where the group is defined by at least one shared nucleic acid sequence that is common to all members of the group and that is specific to that group. (A "group" of organisms is generally a phylogenetic grouping of organisms, such as a strain, species or genus of organisms and may be limited to a single organism.) Generally, the uniqueness of the targeted nucleic acid sequence or sequences need only be limited to the particular sample type being assayed (e.g., a human sample versus an industrial or environmental sample). Nucleic acid-based methods and means for detecting individual and groups of organisms are disclosed by Kohne, "Method for Detection, Identification and Quantitation of Non-Viral Organisms," U.S. Patent No. 4,851,330, and Hogan et al., "Nucleic Acid Probes for Detection and/or Quantitation of Non-Viral Organisms," U.S. Patent No. 5,541,308.

0005 After determining what organisms are to be targeted by an assay, the first step is to select or design a probe which exhibits specificity for a nucleic acid sequence belonging to those organisms which define the group. Nucleic acid-based assays can be designed to detect either deoxyribonucleic acid (DNA) or ribonucleic acid (RNA), including ribosomal RNA (rRNA), transfer RNA (tRNA) or messenger RNA (mRNA). For prokaryotic and eukaryotic organisms, rRNA or the encoding DNA (rDNA) is generally a preferred target for detection. Ribosomal RNA sequences are particularly preferred targets for non-amplified, nucleic acid-based assays because of their relative abundance in cells, and because rRNA contains regions of sequence variability that can be exploited to design probes capable of distinguishing between even closely related organisms. Viruses, which do not contain ribosomal nucleic acid, and cellular changes are often best detected by targeting DNA, RNA, or a messenger RNA (mRNA) sequence. See, e.g., McDonough et al., "Detection of Human Immunodeficiency Virus Type 1, U.S. Patent No. 6,649,749; and Fradet et al., "Methods to Detect Prostate Cancer in a Sample," U.S. Patent Application Publication No. US 2005-0282170 A1. Such viruses may include positive-strand RNA viruses (e.g., hepatitis C virus), where the RNA genome is mRNA, negative-strand RNA viruses (e.g., influenza viruses), retroviruses (e.g., human immunodeficiency virus), single-stranded DNA viruses (e.g., parvoviruses), and double-stranded DNA viruses (e.g.,

adenoviruses), which would require a melting step to render the double-stranded target region sufficiently single-stranded for amplification or detection. When the focus of a nucleic acid-based assay is the detection of a genetic abnormality, then the probes are usually designed to detect identifiable changes in the genetic code, an example of which is the abnormal Philadelphia chromosome associated with chronic myelocytic leukemia. See, e.g., Stephenson et al., "Deoxynucleic Acid Molecules Useful as Probes for Detecting Oncogenes Incorporated Into Chromosomal DNA," U.S. Patent No. 4,681,840.

0006 When performing a nucleic acid-based assay, preparation of the sample is necessary to release and stabilize target nucleic acids which may be present in the sample. Sample preparation can also serve to eliminate nuclease activity and remove or inactivate potential inhibitors of nucleic acid amplification (discussed below) or detection of the target nucleic acids. See, e.g., Ryder et al., "Amplification of Nucleic Acids From Mononuclear Cells Using Iron Complexing and Other Agents," U.S. Patent No. 5,639,599, which discloses methods for preparing nucleic acid for amplification, including the use of complexing agents able to complex with ferric ions released by lysed red blood cells. The method of sample preparation can vary and will depend in part on the nature of the sample being processed (e.g., blood, urine, stool, pus or sputum). When target nucleic acids are being extracted from a white blood cell population present in a diluted or undiluted whole blood sample, a differential lysis procedure is generally followed. See, e.g., Ryder et al., "Preparation of Nucleic Acid From Blood," European Patent Application No. 0 547 267 A2. Differential lysis procedures are well known in the art and are designed to specifically isolate nucleic acids from white blood cells, while limiting or eliminating the presence or activity of red blood cell products, such as heme, which can interfere with nucleic acid amplification or detection. Other lytic methods are disclosed by, for example, Cummins et al., "Methods of Extracting Nucleic Acids and PCR Amplification Without Using a Proteolytic Enzyme," U.S. Patent No. 5,231,015, and Clark et al., "Methods for Extracting Nucleic Acids From a Wide Range of Organisms by Nonlytic Permeabilization," U.S. Patent No. 5,837,452; and Cunningham et al., "Compositions, Methods and Kits for Determining the Presence of Cryptosporidium Organisms in a Test Sample," U.S. Patent Application Publication No. US 2002-0055116 A1.

0007 To purify the sample and remove nucleases and other materials capable of interfering with

amplification or detection, the targeted nucleic acid can be isolated by target-capture means using a "capture probe" which binds the target nucleic acid and is or becomes either directly or indirectly bound to a solid substrate, such as a magnetic or silica particle. See, e.g., Ranki et al., "Detection of Microbial Nucleic Acids by a One-Step Sandwich Hybridization Test," U.S. Patent No. 4,486,539; Stabinsky, "Methods and Kits for Performing Nucleic Acid Hybridization Assays," U.S. Patent No. 4,751,177; Boom et al., "Process for Isolating Nucleic Acid," U.S. Patent No. 5,234,809; Englehardt et al., "Capture Sandwich Hybridization Method and Composition," U.S. Patent No. 5,288,609; Collins, "Target and Background Capture Methods and Apparatus for Affinity Assays," U.S. Patent No. 5,780,224; and Weisburg et al., "Two-Step Hybridization and Capture of a Polynucleotide," U.S. Patent No. 6,534,273. When the solid support is a magnetic particle, magnets in close proximity to the reaction receptacle are used to draw and hold the magnetic particles to the side of the receptacle, thereby isolating any bound nucleic acid within the reaction receptacle. Other methods for isolating bound nucleic acid in a reaction receptacle include centrifugation and immobilizing the capture probe on the reaction receptacle. See, e.g., Boom et al., *supra*, and Urdea, "Polynucleotide Capture Assay Employing in Vitro Amplification," U.S. Patent No. 5,200,314. Once the bound nucleic acid is thus isolated, the bound nucleic acid can be separated from unbound nucleic acid and other cellular and sample material by aspirating the fluid contents of the reaction receptacle and optionally performing one or more wash steps with a wash solution.

0008 In most cases, it is desirable to amplify the target sequence. Nucleic acid amplification involves the use of nucleic acid polymerases to enzymatically synthesize nucleic acid amplification products (copies) containing a sequence that is either complementary or homologous to the template nucleic acid sequence being amplified. The amplification products may be either extension products or transcripts generated in a transcription-based amplification procedure. Examples of nucleic acid amplification procedures practiced in the art include the polymerase chain reaction (PCR), strand displacement amplification (SDA), loop-mediated isothermal amplification (LAMP) ligase chain reaction (LCR), immuno-amplification, and a variety of transcription-based amplification procedures, including transcription-mediated amplification (TMA), nucleic acid sequence based amplification (NASBA), and self-sustained sequence replication (3SR). See, e.g., Mullis, "Process for Amplifying, Detecting, and/or

Cloning Nucleic Acid Sequences," U.S. Patent No. 4,683,195; Walker, "Strand Displacement Amplification," U.S. Patent No. 5,455,166; Notomi et al., "Process for Synthesizing Nucleic Acid," U.S. Patent No. 6,410,278; Birkenmeyer, "Amplification of Target Nucleic Acids Using Gap Filling Ligase Chain Reaction," U.S. Patent No. 5,427,930; Cashman, "Blocked-Polymerase Polynucleotide Immunoassay Method and Kit," U.S. Patent No. 5,849,478; Kacian et al., "Nucleic Acid Sequence Amplification Methods," U.S. Patent No. 5,399,491; Malek et al., "Enhanced Nucleic Acid Amplification Process," U.S. Patent No. 5,130,238; and Lizardi et al., *BioTechnology*, 6:1197 (1988). Nucleic acid amplification is especially beneficial when the amount of target sequence present in a sample is very low. By amplifying the target sequences and detecting the synthesized amplification product, the sensitivity of an assay can be vastly improved, since fewer target sequences are needed at the beginning of the assay to ensure detection of the targeted nucleic acid sequences.

0009 Detection of a target nucleic acid requires the use of a probe having a nucleotide base sequence which binds to a target sequence contained within the target nucleic acid or, alternatively, amplification product containing the target sequence or its complement. Probes useful for distinguishing between sources of nucleic acid are selected or designed such that they do not detectably bind to nucleic acid from non-target organisms which may be present in the sample under the selected assay conditions. While probes may include non-nucleotide components, the target binding portion of a probe will include DNA, RNA and/or analogs thereof in order to effect hybridization to the target sequence or its complement. See, e.g., Becker et al., "Modified Oligonucleotides for Determining the Presence of a Nucleic Acid Analyte in a Sample," U.S. Patent Application No. US 2003-0036058 A1 (discloses the use of 2'-O-methyl modified probes); and Nielsen et al., "Peptide Nucleic Acids," U.S. Patent No. 5,539,082 (discloses the use of probes having a 2-aminoethylglycine backbone which couples the nucleobase subunits by means of a carboxymethyl linker to the central secondary amine). For detection purposes, probes may include a detectable label, such as a radiolabel, fluorescent dye, biotin, enzyme or chemiluminescent compound, where the label may be provided either before, during or after hybridization to the probe to the target sequence or its complement. See, e.g., Higuchi, "Homogenous Methods for Nucleic Amplifications and Detection," U.S. Patent No. 5,994,056 (discloses the use of intercalating agents such as ethidium bromide); and Urdea et al.,

“Solution Phase Nucleic Acid Sandwich Assays Having Reduced Background Noise,” U.S. Patent No. 5,635,352 (discloses use of label probes for binding to cruciform structures containing a target nucleic acid).

0010 Nucleic acid-based assays may be based on a homogenous or a heterogenous format. One form of a heterogenous assay involves preferentially binding a probe:target complex to a solid support, such as glass, minerals or polymeric materials, and removing any unbound probe prior to detection. In an alternative approach, it is the unbound probe which is associated with the solid support while probe complexed with the target sequence remains free in solution and can be separated for detection. Homogenous assays generally take place in solution without a solid phase separation step and commonly exploit chemical differences between a probe free in solution and a probe which has formed part of a target:probe complex. An example of a homogenous assay is the Hybridization Protection Assay (HPA), the particulars of which are disclosed by Arnold et al., “Homogenous Protection Assay,” U.S. Patent No. 5,639,604. Detection in HPA is based on differential hydrolysis which permits specific detection of an acridinium ester-labeled probe hybridized to the target sequence or its complement. See, e.g., Arnold et al., “Protected Chemiluminescent Labels,” U.S. Patent No. 4,950,613; Campbell et al., “Chemiluminescent Acridium Labelling Compounds,” U.S. Patent No. 4,946,958; Arnold et al., “Acridinium Ester Labelling and Purification of Nucleotide Probes,” U.S. Patent No. 5,185,439; and Arnold et al., “Linking Reagents for Nucleotide Probes,” U.S. Patent No. 5,585,481. This detection format includes both a hybridization step and a selection step. In the hybridization step, an excess of acridinium ester-labeled probe is added to the reaction receptacle and permitted to anneal to the target sequence or its complement. Following the hybridization step, label associated with unhybridized probe is rendered non-chemiluminescent in the selection step by the addition of an alkaline reagent. The alkaline reagent specifically hydrolyzes only that acridinium ester label associated with unhybridized probe, leaving the acridinium ester of the probe:target hybrid intact and detectable. Chemiluminescence from the acridinium ester of the hybridized probe can then be measured using a luminometer and signal is expressed in relative light units or RLU.

0011 Other homogenous assays include those disclosed by the following: Gelfand et al., “Reaction Mixtures for Detection of Target Nucleic Acids,” U.S. Patent No. 5,804,375; Nadeau

et al., "Detection of Nucleic Acids by Fluorescence Quenching," U.S. Patent No. 5,958,700; Tyagi et al., "Detectably Labeled Dual Conformation Oligonucleotide Probes, Assays and Kits," U.S. Patent No. 5,925,517; Morrison, "Competitive Homogenous Assay," U.S. Patent No. 5,928,862; and Becker et al., "Molecular Torches," U.S. Patent No. 6,849,412. These patents each describe unimolecular or bimolecular probes which may be used to determine the amount of a target nucleic acid in an amplification procedure in real-time, where signal changes associated with the formation of probe:target complexes are detected during amplification and used to calculate an estimated amount of a target nucleic acid present in a sample. Algorithms for calculating the quantity of target nucleic acid originally present in a sample based on signal information collected during an amplification procedure include that disclosed by Wittwer et al., "PCR Method for Nucleic Acid Quantification Utilizing Second or Third Order Rate Constants," U.S. Patent No. 6,232,079; Sagner et al., "Method for the Efficiency-Corrected Real-Time Quantification of Nucleic Acids," U.S. Patent No. 6,691,041; McMillan et al., "Methods for Quantitative Analysis of a Nucleic Acid Amplification Reaction," U.S. Patent No. 6,911,327; and Chismar et al., "Method and Algorithm for Quantifying Polynucleotides," U.S. Provisional Application No. 60/693,455, which enjoys common ownership herewith.

0012 After the nucleic acid-based assay is run, and to avoid possible contamination of subsequent amplification reactions, the reaction mixture can be treated with a deactivating reagent which destroys nucleic acids and related amplification products in the reaction receptacle. Such reagents can include oxidants, reductants and reactive chemicals which modify the primary chemical structure of a nucleic acid. These reagents operate by rendering nucleic acids inert towards an amplification reaction, whether the nucleic acid is RNA or DNA. Examples of such chemical agents include solutions of sodium hypochlorite (bleach), solutions of potassium permanganate, formic acid, hydrazine, dimethyl sulfate and similar compounds. More details of deactivation protocols can be found in Dattagupta et al., "Method and Kit for Destroying the Ability of Nucleic Acid to Be Amplified," U.S. Patent No. 5,612,200, and Nelson et al., "Reagents, Methods and Kits for Use in Deactivating Nucleic Acids," U.S. Patent Application Publication No. US 2005-0202491 A1.

0013 Given the large number of complex steps associated with nucleic acid-based amplification assays, and the different processing and equipment requirements of each type of amplification

assay, a need exists for an automated system capable of processing the contents of a plurality of reaction receptacles according to different amplification assay protocols, and most especially for performing both real-time and end-point amplification assays on the same platform and/or within a self-contained housing. Real-time amplification assays involve periodically determining the amount of targeted amplification products as the amplification reaction is taking place, thereby making it easier to provide quantitative information about target nucleic acids present in a sample, whereas end-point amplifications determine the amount of targeted amplification products after the amplification reaction has occurred, generally making them more useful for providing qualitative information about target nucleic acids. To improve flow through and, thereby, to reduce the amount of time needed to process large volumes of samples, there is also a need for a system capable of continuously processing the contents of multiple reaction receptacles according to a real-time amplification protocol without having to interrupt the system to manually or automatically load a new batch of reaction receptacles for processing. Additionally, there is a need for a reagent and method for reducing the amount of amplification inhibitors in reaction receptacles that could affect qualitative or quantitative determinations.

SUMMARY

0014 The above-described needs are addressed by an automated analyzer constructed and operated as described herein. In general, the automated analyzer integrates and coordinates the operation of various automated stations, or modules, involved in performing one or more assays on a plurality of reaction mixtures contained in reaction receptacles. The analyzer is preferably a self-contained, stand alone unit. Assay sample materials and reaction receptacles, as well as the various solutions, reagents, and other materials used in performing the assays are preferably stored within the analyzer, as are the waste products generated when assays are performed. The analyzer is an integrated nucleic acid testing system that fully automates all assay steps from sample processing through amplification and multi-format detection. In a preferred embodiment, the instrument is capable of running both real-time and end-point amplification assays. After daily set up is completed, the operator may choose to run an end-point amplification assay, a real-time amplification assay, or both. For real-time amplification assays, the analyzer of the present invention is capable of processing the contents

of multiple reaction receptacles in a continuous as opposed to a batch mode, thereby greatly increasing the speed at which results can be calculated and reported. In operation, the analyzer can also be used to reduce the presence of amplification inhibitors by providing a surface treating agent that is used to coat the inner surfaces of reaction receptacles prior to or during an isolation and purification step to remove sample material and/or reagents from reaction receptacles.

0015 The analyzer includes a computer controller which runs analyzer-controlling and assay-scheduling software to coordinate operation of the stations of the analyzer and movement of each reaction receptacle through the analyzer.

0016 Reaction receptacles can be loaded in an input queue which sequentially presents each receptacle at a pick-up position to be retrieved by a transport mechanism, which automatically transports the reaction receptacles between the stations of the analyzer.

0017 Sample containers are carried on a first ring assembly, and disposable pipette tips are carried on a second ring assembly. Containers of target capture reagent, including a suspension of solid support material, are carried on an inner rotatable assembly constructed and arranged to selectively agitate the containers or present the containers for access by the probe of an automatic robotic pipette system. Reaction mixtures, including fluid sample material and target capture reagent, are prepared by the pipette system within each reaction receptacle.

0018 The analyzer further includes receptacle mixers for mixing the contents of a receptacle placed therein. The mixer may be in fluid communication with fluid containers and may include dispensers for dispensing one or more fluids into the receptacle. One or more incubators carry multiple receptacles in a temperature-controlled chamber and permit individual receptacles to be automatically placed into and removed from the chamber. Magnetic separation stations automatically perform a magnetic separation and wash procedure on the contents of a receptacle placed in the station.

0019 In the preferred method of operation, assay results may be ascertained by the amount of light emitted from a receptacle at the conclusion of the appropriate preparation steps. Accordingly, the analyzer includes a luminometer (a type of signal detecting device) for detecting and/or quantifying the amount of light emitted by the contents of the reaction receptacle. A deactivation queue may be provided to deactivate the contents of a reaction receptacle placed

therein at the conclusion of the assay.

0020 Reaction receptacles can be independently transported between stations by the transport mechanism, and the stations can be operated in parallel to perform different assay procedures simultaneously on different reaction receptacles, thereby facilitating efficient, high through-put operation of the analyzer. Moreover, the present disclosure facilitates arranging the various stations associated with a nucleic acid-based assay onto a single, contained platform, thereby achieving efficient space utilization.

0021 Various embodiments of the claimed invention relate to a method for detecting two or more different optical emission signals emitted from each of a plurality of reaction receptacles, said method comprising, for each optical emission signal to be detected, the steps of: (a) modulating an optical excitation signal at a known excitation signal frequency, the excitation signal having a known excitation wavelength that excites an associated optical emission signal having a known emission wavelength; (b) directing the optical excitation signal at each receptacle; (c) filtering an optical signal, including any optical emission signal emitted by the contents of each receptacle, to transmit a portion of the optical signal that is at the emission wavelength; (d) detecting the portion of the optical signal transmitted in step (c); (e) determining an emission signal frequency of the optical signal detected in step (d); and (f) discarding any portion of the optical signal detected in step (d) having an emission signal frequency determined in step (e) that is inconsistent with the known excitation signal frequency to determine the portion of the optical signal detected in step (d) that corresponds to the associated optical emission signal.

0021A Various embodiments of the claimed invention relate to a system for detecting two or more different optical emission signals emitted from each of a plurality of reaction receptacles, wherein the reaction receptacles are disposed in a fixed arrangement with respect to each other, said system comprising: (a) a plurality of optical sensors, each of said optical sensors being configured to detect one of the two or more optical emission signals and being associated with one of the reaction receptacles so that each reaction receptacle has associated therewith one optical sensor for each optical emission signal to be detected, whereby said optical sensors associated with each of the plurality of reaction receptacles are arranged in a configuration that enables each optical sensor to detect the associated optical emission signal emitted from the associated reaction receptacle; (b) a transport mechanism constructed and arranged to effect relative movement between the plurality of reaction receptacles and the optical sensors to sequentially place each reaction receptacle into a signal-detecting position with respect to one of its associated optical sensors, wherein (1) each

optical sensor is configured to direct an optical excitation signal at an associated reaction receptacle that is in a signal-detecting position with respect to the optical sensor and to detect an optical emission signal emitted from the reaction receptacle that is associated with the excitation signal, and (2) each optical sensor is further configured to direct an excitation signal of a excitation frequency at an associated receptacle, and wherein optical sensors configured to detect at least two different optical emission signals are configured to direct excitation signals of two different excitation frequencies; and (c) circuitry configured to discard any portion of an optical signal detected by an optical sensor having an emission frequency that is inconsistent with the excitation frequency of the associated excitation signal.

0021B Other objects, features, and characteristics of the present invention, including the methods of operation and the function and interrelation of the elements of structure, will become more apparent upon consideration of the following description and the appended claims, with reference to the accompanying drawings, all of which form a part of this disclosure, wherein like reference numerals designate corresponding parts in the various figures.

DESCRIPTION OF THE DRAWINGS

0022 FIGURE 1 is a perspective view of an automated nucleic acid-based diagnostic analyzer according to the present invention;

0023 FIGURE 2 is a perspective view of the structural frame of the analyzer of the present invention;

0024 FIGURE 3 is a plan view of a portion of the assay processing deck of the analyzer of the present invention;

0025 FIGURE 4 is an exploded perspective view of the assay processing deck;

0026 FIGURE 5 is a plan view of a sample ring and a pipette tip wheel of the assay processing deck of the analyzer of the present invention;

0027 FIGURE 6 is a perspective view showing the sample ring and the pipette tip wheel;

0028 FIGURE 6 A is a partial cross-sectional view along the line 6A-6A in FIGURE 5;

0029 FIGURE 7 is a perspective view of a multi-axis mixer of the processing deck of the analyzer of the present invention;

0030 FIGURE 8 is a plan view of the multi-axis mixer;

0031 FIGURE 9 is a side elevation of the multi-axis mixer;

0032 FIGURE 10 is a plan view of the multi-axis mixer with container holders and a turntable cover removed therefrom;

0033 FIGURE 11 is a cross-sectional view of the multi-axis mixer taken in the direction 11-11 in FIGURE 10;

0034 FIGURE 12 is a perspective view of a drive assembly of the multi-axis mixer;

0035 FIGURE 13 is a perspective view of a transport mechanism of the processing deck of the analyzer of the present invention;

0036 FIGURE 14 is a perspective view of a manipulating hook mounting plate and a manipulating hook actuating mechanism of the transport mechanism, with the manipulating hook member engaged with a reaction receptacle and in a retracted position;

0037 FIGURE 15 is the same as FIGURE 14, except with the manipulating hook member in the extended position;

0038 FIGURE 16 is an exploded perspective view of the transport mechanism;

0039 FIGURE 17 is a side-elevation of a temperature ramping station of the processing deck of the analyzer of the present invention;

0040 FIGURE 18 is a front-elevation of the temperature ramping station;

0041 FIGURE 19 is a perspective view of a rotary incubator of the processing deck of the analyzer of the present invention;

0042 FIGURE 20 is an exploded view of a portion of a housing and access opening closure mechanisms according to a first embodiment of the rotary incubator;

0043 FIGURE 21 is a partial view of a skewed disk linear mixer of the rotary incubator, shown engaged with a reaction receptacle employed in a preferred mode of operation of the analyzer of the present invention;

0044 FIGURE 22 is an exploded perspective view of the first embodiment of the rotary incubator;

0045 FIGURE 23 is a perspective view of the rotary incubator according to a second embodiment thereof;

0046 FIGURE 23A is an exploded perspective view of the second embodiment of the rotary incubator;

- 0047 FIGURE 23B is a partial exploded perspective view of an access opening closure mechanism of the second embodiment of the rotary incubator;
- 0048 FIGURE 23C is an exploded view of a receptacle carrier carousel of the second embodiment of the rotary incubator;
- 0049 FIGURE 24 is a perspective view of a particles of the processing deck of the present invention with a side plate thereof removed;
- 0050 FIGURE 25 is a partial transverse cross-section of the particles;
- 0051 FIGURE 25A is a partial transverse cross-section of a tip of an aspirating tube of the particles with a contamination-limiting triplet carried on the end thereof;
- 0052 FIGURE 26 is an exploded perspective view of a receptacle carrier unit, an orbital mixer assembly, and a divider plate of the particles;
- 0053 FIGURE 27 is a partial cross-sectional view of a wash solution dispenser nozzle, an aspirator tube with a contamination-limiting triplet engaged with an end thereof, and a receptacle carrier unit of the particles, showing a multi-tube unit reaction receptacle employed in a preferred mode of operation of the analyzer carried in the receptacle carrier unit and the aspirator tube and contamination-limiting triplet inserted into a reaction tube of the multi-tube unit;
- 0054 FIGURE 28 is a partial cross-sectional view of the wash solution dispenser nozzle, the aspirator tube, and the receptacle carrier unit of the particles, showing the multi-tube unit carried in the receptacle carrier unit and the aspirator tube engaging the contamination-limiting triplet held in a contamination-limiting element holding structure of the multi-tube unit;
- 0055 FIGURES 29A-29D show a partial cross-section of a first embodiment of a triplet stripping hole of a triplet stripping plate of the particles and a triplet stripping operation using the triplet stripping hole;
- 0056 FIGURES 30A-30D show a partial cross-section of a second embodiment of a triplet stripping hole and a triplet stripping operation using the triplet stripping hole;
- 0057 FIGURE 31A is a plan view of a third embodiment of a triplet stripping hole of a triplet stripping plate of the particles;
- 0058 FIGURES 31B-31C show a partial cross-section of the third embodiment of the triplet stripping hole and a triplet stripping operation using the triplet;

- 0059 FIGURE 32 is a perspective view of an orbital mixer with a front plate thereof removed;
- 0060 FIGURE 33 is an exploded view of the orbital mixer of the processing deck of the analyzer of the present invention;
- 0061 FIGURE 34 is a top-plan view of the orbital mixer;
- 0062 FIGURE 35 is a top perspective view of a reagent cooling bay of the processing deck of the analyzer of the present invention;
- 0063 FIGURE 36 is a top perspective view of a reagent cooling bay with the container tray removed therefrom;
- 0064 FIGURE 37 is a bottom plan view of the reagent cooling bay;
- 0065 FIGURE 38 is an exploded view of the reagent cooling bay;
- 0066 FIGURE 39 is a top perspective view of a modular container tray of the reagent cooling bay;
- 0067 FIGURE 40 is a perspective view of a first embodiment of a luminometer of the processing deck of the analyzer of the present invention;
- 0068 FIGURE 41 is a partial exploded perspective view of the luminometer of the first embodiment;
- 0069 FIGURE 42A is a partial perspective view of a receptacle transport mechanism of the first embodiment of the luminometer;
- 0070 FIGURE 42B is an end view of the receptacle transport mechanism of the first embodiment of the luminometer;
- 0071 FIGURE 42C is a top view of the receptacle transport mechanism of the first embodiment of the luminometer;
- 0072 FIGURE 43 is a break away perspective view of a second embodiment of the luminometer of the present invention;
- 0073 FIGURE 44 is an exploded perspective view of a multi-tube unit door assembly for the luminometer of the second embodiment;
- 0074 FIGURE 45 is an exploded perspective view of a shutter assembly for a photosensor aperture for the luminometer of the second embodiment;
- 0075 FIGURE 45A is a perspective view of an aperture plate of the shutter assembly of the

luminometer of the second embodiment;

0076 FIGURE 46 is a perspective view of a reaction tube positioner assembly of the luminometer of the second embodiment, including a reaction tube positioner disposed within a reaction tube positioner frame;

0077 FIGURE 47 is a perspective view of the reaction tube positioner;

0078 FIGURE 48 is a side elevation of the reaction tube positioner assembly;

0079 FIGURE 49 is a perspective view showing the reaction tube positioner of the reaction tube positioner assembly operatively engaging a multi-tube unit employed in a preferred mode of operation of the analyzer;

0080 FIGURE 50 is a perspective view of a multi-tube unit transport mechanism of the luminometer of the second embodiment;

0081 FIGURE 51 is a partial perspective view showing a multi-tube unit transport and drive screw of the multi-tube unit transport mechanism of the luminometer;

0082 FIGURE 52 is a perspective view of a lower chassis of the analyzer of the present invention;

0083 FIGURE 53 is a perspective view of a right-side drawer of the lower chassis;

0084 FIGURE 54 is a perspective view of a left-side drawer of the lower chassis;

0085 FIGURE 55 is a perspective view of a sample tube tray employed in a preferred mode of operation of the analyzer of the present invention;

0086 FIGURE 56 is a top plan view of the sample tube tray;

0087 FIGURE 57 is a partial cross-section of the sample tube tray through line "57-57" in FIGURE 55;

0088 FIGURE 58 is a perspective view of a multi-tube unit employed in a preferred mode of operation of the analyzer of the present invention;

0089 FIGURE 59 is a side elevation of a contact-limiting pipette triplet employed in a preferred mode of operation of the analyzer of the present invention and carried on the multi-tube unit shown in FIGURE 58; and

0090 FIGURE 60 is an enlarged bottom view of a portion of the multi-tube unit, viewed in the direction of arrow "60" in FIGURE 58.

- 0091 FIGURE 61 is a side elevation in cross-section showing an optical detection module and portions of a real-time fluorometer and a multi-tube unit;
- 0092 FIGURE 62 is an exploded perspective view of the housing of the optical detection module;
- 0093 FIGURE 63 is an exploded perspective view of the optical detection module;
- 0094 FIGURE 64 is a top plan view of a real-time fluorometer showing preferred positions of the optical detection modules;
- 0095 FIGURE 65 is a schematic view of a real-time fluorometer showing preferred positions of the optical detection modules;
- 0096 FIGURE 66 is a graph showing excitation spectra of preferred amplification detection dyes;
- 0097 FIGURE 67 is a graph showing emission spectra of preferred amplification detection dyes;
- 0098 FIGURES 68A-68F show a diagram of a circuit for the optical detection module;
- 0099 FIGURE 69 is a perspective view showing the optical detector scanning assembly of a scanning real-time fluorometer;
- 0100 FIGURE 70 is a side elevation of the detector scanning assembly;
- 0101 FIGURE 71 is a top plan view of the detector scanning assembly;
- 0102 FIGURE 72 is a bottom plan view of the detector scanning assembly.
- 0103 FIGURE 73 is a perspective view of a portion of a carousel of the real-time fluorometer;
- 0104 FIGURE 74 is an exploded perspective view showing the carousel of the real-time fluorometer and a magnetic divider;
- 0105 FIGURE 75 is a bottom plan view of the carousel of the real-time fluorometer showing a single magnetic divider attached thereto;
- 0106 FIGURE 75A is a partial cross-sectional view taken along the line A-A in FIGURE 75;
- 0107 FIGURE 76A is a flow chart showing the protocols of a preferred real-time amplification assay and a portion of a preferred end-point amplification assay which stops after exposure to amplification conditions, both assays being in accordance with the present invention;
- 0108 FIGURE 76B is a flow chart showing the remainder of the protocol for the preferred end-

point amplification assay of FIGURE 76A following exposure to amplification conditions;

0109 FIGURE 77 is a flow chart showing an analyte quantification process;

0110 FIGURE 78 is a time plot of real-time fluorometer data; and

0111 FIGURE 79 is a plot showing a method for fitting a curve to real-time fluorometer data and using the fit to determine a threshold time.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENT

0112 While the present invention may be embodied in a variety of forms, the following descriptions and accompanying drawings are merely intended to disclose some of those forms as specific examples of the present invention. Accordingly, the present invention is not intended to be limited to the forms or embodiments so described and illustrated. Instead, the full scope of the present invention is set forth in the appended claims.

ANALYZER OVERVIEW

0113 An automated diagnostic analyzer according to the present invention is designated generally by reference number 50 in FIGURES 1 and 2. Analyzer 50 includes a housing 60 built over an internal frame structure 62, preferably made of steel. The analyzer 50 is preferably supported on caster wheels 64 structurally mounted to the frame structure 62 so as to make the analyzer movable.

0114 The various stations involved in performing an automated assay and the assay samples are housed within housing 60. In addition, the various solutions, reagents, and other materials used in performing the assays are preferably stored within the housing 60, as are the waste products generated when assays are performed with the analyzer 50.

0115 Housing 60 includes a test receptacle loading opening 68, which is shown in FIGURE 1 to be disposed in a forwardly facing panel of the housing 60, but could as well be located in other panels of the housing 60. A pipette door 70 having a view window 72 and a carousel door 74 having a view window 76 are disposed above a generally horizontal work surface 66. A forwardly protruding arcuate panel 78 accommodates a sample carousel, which will be described

