Geno-Sen's

Chlamydia Pneumonia (Rotor Gene)

Real Time PCR Kit

Quantitative

For use with the

Rotor Gene™2000/3000/6000

(Corbett Research Australia)

For In-Vitro Diagnostic Use

PACK INSERT

Revised 2017

GenomeDiagnosticsPvt.Ltd.

(AnISO13485:2012, 9001:2008 Certified Company)



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CHLAMYDIA PNEUMONIA Geno-Sen's Real Time PCR Kit for use with the *Rotor Gene*[™] 2000/3000/6000^{*} (Corbett Research).

| Color Code | Contents | Cat. No. 9111307 100 rxns | Cat. No. 9111308 50 rxns | Cat. No. 9111309 25 rxns |
|---------------------|---------------------------|------------------------------|-----------------------------|-----------------------------|
| R1 | CHLAMYDIA PNEUMONIA | 25 rxns x 4 Vials | 25 rxns x 2 Vials | 25 rxns x 1 Vials |
| Blue | Super mix. | | | |
| R2 | Mg Sol RT. | 1 Vial | 1 Vial | 1 Vial |
| <mark>Yellow</mark> | | | | |
| C.pneumo- | CHLAMYDIA PNEUMONIA | 1 Vial of 300µl | 1 Vial of 300µl | 1 Vial of 300µl |
| S1 | Standard 1 | | | |
| Red | 1 Χ 10 ⁵ IU/μl | | | |
| C.pneumo - | CHLAMYDIA PNEUMONIA | 1 Vial of 300µl | 1 Vial of 300µl | 1 Vial of 300µl |
| S2 | Standard 2 | | | |
| Red | $1 X 10^4$ IU/ μ I | | | |
| C.pneumo - | CHLAMYDIA PNEUMONIA | 1 Vial of 300µl | 1 Vial of 300µl | 1 Vial of 300µl |
| S3 | Standard 3 | | | |
| Red | 1 Χ 10 ³ IU/μl | | | |
| C.pneumo - | CHLAMYDIA PNEUMONIA | 1 Vial of 300µl | 1 Vial of 300µl | 1 Vial of 300µl |
| S4 | Standard 4 | | | |
| Red | 1 Χ 10² IU/μl | | | |
| C.pneumo - | CHLAMYDIA PNEUMONIA | 1 Vial of 300µl | 1 Vial of 300µl | 1 Vial of 300µl |
| S5 | Standard 5 | | | |
| Red | 1 Χ 10 ¹ ΙU/μΙ | | | |
| W | Molecular Grade Water. | 1 Vials of 1 ml | 1 Vial of 1 ml | 1 Vial of 1 ml |
| White | | | | |
| IC-1 (R3) | IC-1 RG (R3) | 1 Vial of 1 ml | 1 Vial of 1 ml | 1 Vial of 1 ml |
| Green | | | | |

1. Contents of the Kit

R= Reagents

S= Quantitation Standards

W= Molecular Grade Water.

All Vials have Color Coder tops to distinguish between different reagents.

2. Storage of the Kit.

All the reagents of the kit should be stored at -20°C and is stable till expiry at this temperature. Repeated thawing and freezing (>3x) should be avoided, as this may reduce the sensitivity of the assay. If the kit is to be used only occasionally, then the reagents should be frozen in aliquots. Storage at +4°C is not recommended & should not exceed a period of 2 hours in any case.

3. CHLAMYDIA PNEUMONIA Information & Application

Chlamydia pneumoniae is a type of bacteria that causes lung infections, such as pneumonia. The bacteria cause illness by damaging the lining of the respiratory tract (throat, windpipe, and lungs).

How It Spreads

A person who is sick with *C. pneumoniae* infection has the bacteria in their nose, throat, windpipe, and lungs. *C. pneumoniae* is spread from person to person when people who are sick cough or sneeze while in close contact with others, who then breathe in the bacteria. If someone who is infected coughs into their hands, they can spread the bacteria to others, by shaking hands for example, who can also become infected if they touch their nose or mouth while the bacteria are on their hands.

Most people who are exposed for a short amount of time to someone with *C. pneumoniae* infection do not get sick. However, it is common for this illness to spread between family members who live together. *C. pneumoniae* infections are known to have long incubation periods (the time between first catching the bacteria from an ill person and development of symptoms), with symptoms beginning 3 to 4 weeks after exposure.

People at Increased Risk

People of all ages can get sick from *C. pneumoniae*. It most commonly infects people for the first time when they are school-aged children or young adults; however, reinfection is most common in older adults.

People at increased risk include those who live or work in crowded settings where outbreaks most commonly occur, such as:

- Schools
- College residence halls (dormitories)
- Military barracks
- Nursing homes
- Hospitals
- Prisons

Older adults are at increased risk for severe disease caused by *C. pneumoniae* infection, including pneumonia.

Diagnosis

Clinicians may test to see if a patient has Chlamydia pneumoniae infection by:

- Laboratory test that involves taking a sample of sputum (phlegm) or swab from the nose or throat, PCR test.
- Blood test

<u>Treatment</u>

C. pneumoniae infection is routinely treated with antibiotics (medicines that kill bacteria in the body), although most people will recover from the illness on their own without medicine.

There are several types of antibiotics that can treat *C. pneumoniae* infections. If you or your child is diagnosed with a *C. pneumoniae* infection, your doctor will decide how to best treat it.

Complications

While *C. pneumoniae* infection is most often a mild disease, serious complications can occur that result in needing care or treatment in a hospital. Complications that have been reported include:

- Pneumonia (lung infection)
- Swelling in the brain (encephalitis)
- Swelling of the heart (myocarditis)
- Chronic infection

Prevention

While there is no vaccine to prevent *Chlamydia pneumoniae* infection, there are things you can do to protect yourself and others.

Hygiene

Like many respiratory diseases, *C. pneumoniae* infection is spread by coughing and sneezing. The best way to keep from getting or spreading the bacteria is to wash your hands often, especially after coughing or sneezing. To practice good hygiene, you should:

- Cover your mouth and nose with a tissue when you cough or sneeze.
- Put your used tissue in the waste basket.
- Cough or sneeze into your upper sleeve or elbow, not your hands, if you don't have a tissue.
- Wash your hands often with soap and water for at least 20 seconds.
- Use an alcohol-based hand rub if soap and water are not available.

Previous Infections

If you get sick from *C. pneumonia*, you may still get sick from it again in the future. Lifelong immunity to *C. pneumonia* likely does not occur.

Preventive Antibiotics for Close Contacts

Antibiotics (medicines that kill bacteria in the body) to prevent *C. pneumoniae* infection are not usually recommended for people who are close contacts, including household members, of someone who is sick with *C. pneumoniae* infection. Preventive antibiotics might be recommended if an exposed person is at increased risk for developing serious complications if they get sick (this is best determined by your doctor).

Antibiotics should not be used unless they are absolutely necessary. Not only can they cause harmful side effects, but every time a person takes antibiotics, sensitive bacteria (bacteria that antibiotics can still attack) are killed, but resistant bacteria are left to grow and multiply. Repeated use of antibiotics can increase the number of drug-resistant bacteria.

Samples which can be used for Extraction: C.S.F, Bronchial Swabs, Bronchial lavage, etc.

4. Precautions for PCR

The following aspects should always be taken care of:

- Store positive material (Specimens, Standards or amplicons) separately from all other reagents and add it to the reaction mix in a separate facility.
- Thaw all components thoroughly at room temperature before starting the assay.
- When thawed, mix the components and centrifuge briefly.
- Work quickly on ice or in the Cooling Block.
- All the reagents including the NTC (except for standards & specimens) should be mixed & dispensed in pre-mix area.
- All the standards & specimens should be mixed & dispensed in extraction area.
- Use pipette tips with filters only.
- Always use disposable powder-free gloves.

5. Additionally Required Materials and Devices

- DNA isolation kit (see 8.a. DNA extraction)
- 0.2 ml PCR tubes for use with 36-well rotor (Corbett Research, Cat.-Nr.: SE-1003F) alternatively 0.1 ml PCR tubes for use with 72-well rotor (Corbett Research, Cat.-Nr.: ST-1001)
- Micro Pipettes Variable Volume 2-20µl, 10-100µl, 100-1000µl,
- Sterile pipette tips with aerosol barrier 2-20µl, 10-100µl, 100-1000µl,

- Disposable powder-free gloves
- Vortex mixer
- Centrifuge Desktop with rotor for 1.7 ml reaction tubes
- Rotor Gene[™] 2000,3000 or Rotor Gene[™] 6000, Corbett Research (The Real time PCR Instrument)

6. Principle of Real-Time PCR

The robust assay exploits the so-called Taqman principle. During PCR, forward and reverse primers hybridize to a specific sequence product. A TaqMan probe, which is contained in the same reaction mixture and which consists of an oligonucleotide labeled with a 5'-reporter dye and a downstream, 3'-quencher dye, hybridizes to a target sequence within the PCR product. A Taq polymerase which possesses 5' - 3' exonuclease activity cleaves the probe. The reporter dye and quencher dye are separated upon cleavage, resulting in an increase in fluorescence for the reporter. Thus, the increase in fluorescence is directly proportional to the target amplification during PCR.

7. Description of the Product.

The Geno-Sen's Chlamydia pneumonia PCR Reagents constitute a ready to use system for detection and quantification of Chlamydia pneumonia using Polymerase chain reaction (PCR) in the Rotor Gene 2000/3000/6000 (Corbett Research). The Specific Master mix contains reagents and enzymes for the specific amplification of Chlamydia pneumonia and for the direct detection of the specific amplification in fluorescence channels Cycling A.FAM of the Rotor Gene 2000/3000/6000 & the Reference gene on Cycling A. Joe. External positive Standards (C.pneumo S 1-5) are supplied which allow the determination of the gene load. For further information, please refer to section 8.c Quantitation.

8. Procedure

8.a DNA Extraction

DNA Extraction kits are available from various manufacturers. Sample volumes for the DNA Extraction procedure depend on the protocol used. Please carry out the DNA Extraction according to the manufacturer's instructions. The recommended Extraction kits are the following:

| Sample Material | Nucleic Acid Isolation Kit | Cat. Num. | •••• |
|---|--|-----------|---|
| Bronchial Swab, Bronchial Lavage,etc | Geno Sen's® Genomic DNA Extraction Mini Kit — for DNA purification OR | | Genome Diagnostics Pvt. Ltd. India. |

| Pure Link Viral R Kit. OR | NA/DNA | Mini | 12280-050 | InvitroGen USA. | Inc. |
|-------------------------------------|--------|------|-----------|--------------------|------|
| QIAamp Viral extraction Kit (50) | DNA | Mini | 51104 | QIAGEN | |

Note:

All the Above Kits were evaluated in conjuction with the *Geno Sen's*[®] *Chlamydia pneumonia Real Time PCR RG Kit on the Rotor Gene 3000/6000. The performance of* Pure Link Viral RNA/DNA Mini Kit manufactured by InvitroGen Inc. USA was found to be almost equivalent to the *Geno Sen's*[®] Genomic DNA Extraction Mini Kit — for DNA purification. However the sample size was limited for this study hence for all practical purposes all the validations were conducted with the *Geno Sen's*[®] Genomic DNA Extraction Kit consistently had a Ct value difference of 1.0 or more. Which clearly means that the yield by Qiagen Kit was low, hence preferably *Geno Sen's*[®] Genomic DNA Extraction Mini Kit or the Pure Link Viral RNA/DNA Mini Kit by InvitroGen Inc. USA is strongly recommended.

However the customer can use their own extraction systems depending on how good the yield is. Always use an extraction kit with a higher DNA yield.

Blood collection tubes coated with anticoagulants may inhibit the PCR; however these inhibitors will be eliminated by the use of the isolation kits given above. It is recommended to avoid the usage of heparin blood.

When using Extraction protocols with ethanol-containing washing buffers, please carry out an additional centrifugation step before the elution to remove any remaining ethanol. This prevents possible inhibition of PCR.

The *Chlamydia pneumonia* Rotor Gene PCR Reagents should not be used with phenol based isolation methods.

8.b Inhibition Control:

Inhibition Control Gene allows the user to determine & control possible PCR inhibition. The Inhibition Control gene reagents are in built in the premix provided and need not be run separately. There is need to add internal control Gene in the reaction mix which has been provided as IC-1 RG (R3). To add the IC Gene 0.5 μ I/rxn of R3 is added to the premix as shown in the Figure 4 & 5. The results can be visualized in the Joe channel. However if there is a need for the Internal control to be used as an extraction efficiency control, then 5 μ I of R3 i.e. Inhibition control should be added to each sample before adding the Buffer MBD in the extraction process.

However in such a case, the IC-1 (R3) should not be mixed with the pre-mix as described above & instead 0.5 μ l/rxn Molecular Grade water should be added to the pre-mix.

Also Note that in such a case there will be no amplification of Internal control in the Joe Channel in the NTC, & Standards. There will only be amplification in the Joe Channel in the Samples to which the IC-1 (R3) has been added before extraction.

Whereas if the IC-1 (R3) is added to the premix then there will be amplification in Joe Channel in all the samples, standards & NTC.

Quantitation

The quantitation standards provided in the kit (**Chlamydia pneumonia** S 1-5) are treated in the same way as extracted samples and the same volume is used i.e. (10µl) instead of the sample. To generate a standard curve in the *Rotor* GeneTM 2000/3000/6000, all 5 Standards should be used as defined in the menu window *Edit Samples* of the *Rotor* GeneTM software. The same should also be defined as standards with the specified concentrations (see *Rotor* GeneTM Manual). The standard curve generated as above can also be used for quantitation in subsequent runs, provided that at least one standard curve needs to be imported (see *Rotor* GeneTM 2000/3000/6000 Manual). However, this quantitation method may lead to deviations in the results due to variability between different PCR runs & due to varying Reaction efficiencies.

Attention:

The standards are defined as Copies/µl. The following formula has to be applied to convert the values determined using the standard curve into copies/ml of sample material:

Result (Copies/µl) x Elution Volume (µl)

Result (Copies/ml) =

Sample Volume (ml)

Or else direct conversion can be done keeping in mind the starting volume of the sample & the final eluted Volume e.g.

If the starting volume of the sample while using the **Geno Sen's**[®] Genomic DNA Extraction Mini Kit is 200µl & the final Eluted Volume is 60µl then to obtain the direct values i.e. Copies/ml for the patient samples the following values for the standards can be fed directly into the operating software which is again based on the above formula and will yield direct conversion of per ml of patient sample data.

| S1: 10^5 Copies /µl = | 3000000 Copies/ml |
|-------------------------|-------------------|
| S2: 10^4 Copies /µl = | 3000000 Copies/ml |
| S3: 10^3 Copies /µl = | 300000 Copies/ml |

| S4: 10^2 Copies /µl = | 30000 Copies/ml |
|-------------------------|-----------------|
| S5: 10^1 Copies /µl = | 3000 Copies/ml |

If the starting volume of the sample while using the **Pure Link Viral RNA/DNA Mini Kit.**

Or Qiagen QIAamp DNA Mini extraction kit is 200µl & the final Eluted Volume is 50µl then to obtain the direct values i.e. copies/ml for the patient samples the following values for the standards can be fed directly into the operating software which is again based on the above formula and will yield direct conversion of per ml of patient sample data.

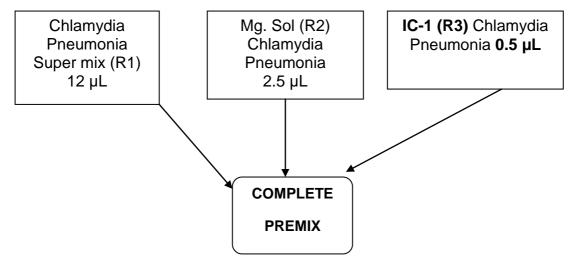
| | 10 ⁵ Copies /μl = | 25000000 Copies/ml |
|-----|------------------------------|--------------------|
| | 10^4 Copies /µl = | 2500000 Copies/ml |
| | 10^3 Copies /µl = | 250000 Copies/ml |
| | 10^2 Copies /µl = | 25000 Copies/ml |
| S5: | 10^1 Copies /µl = | 2500 Copies/ml |

8.d Preparation for PCR

8. e. Preparation for PCR amplification:

First make sure that the Cooling Block (accessory of the *Rotor Gene*TM, Corbett Research) is pre-cooled to +4°C in a Refrigerator or Deep Freezer. Place the desired number of PCR tubes into the Cooling Block. Make sure that the tubes for standards & at least one negative control (*Water, PCR grade*) are included per PCR run. To generate a standard curve, use all supplied Standards (*C.pneumo S 1-5*) for each PCR run. Before each use, all reagents need to be thawed completely and mixed (by pipetting or by brief vortexing).

Please follow the pipeting scheme mentioned below for each sample depending upon the number of samples a mix can be prepared as follows. Each sample



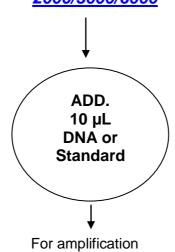


Fig. 4.

Depending upon the number of samples the following pipetting scheme can be followed. e.g. for 10 rxns.

| Chlamydia pneumonia MASTER MIX | 1 rxns. | 10 rxns. |
|------------------------------------|---------|----------|
| Chlamydia Pneumonia Super Mix (R1) | 12 µL | 120 µL |
| Chlamydia Pneumonia Mg Sol. (R2) | 2.5 µL | 25 µL |
| IC-1 (R3) | 0.5 µL | 5 µL |
| Total | 15µL | 150µL |

Fig. 5.

Pipette 15 µl of the Master Mix into each labelled PCR tube. Then add 10 µl of the earlier extracted DNA to each sample tube and mix well by pipeting up and down. Correspondingly, 10 µl of the Standards (C.pneumo *S1-5*) must be used as a positive control and 10 µl of water (*Water, PCR grade*) as a negative control. Close the PCR tubes and transfer the C.pneumonia tubes into the rotor of the *Rotor Gene*TM instrument. The *Rotor Gene*TM software versions 5.0.53 and higher require a Locking Ring (accessory of the *Rotor Gene*TM, Corbett Research) to be placed on top of the rotor to prevent accidental opening of the tubes during the run.

8.f. Programming the *Rotor Gene*[™] 2000/3000

The *Rotor Gene*[™] 2000/3000 PCR program for the detection of C.pneumonia can be divided into following steps:

- A. Setting of general assay parameters & reaction volume
- B. Thermal Profile & Calibration

- C. Cycling profile/ & Initial activation of the Hot Start enzyme
- D. Cycling for Amplification of DNA
- E. Adjustment of the sensitivity of the fluorescence channels
- F. Starting of the Rotor Gene™ run

Program the *Rotor Gene*TM 2000/3000 for these 5 steps according to the parameters shown in Figures 6-20 below All specifications refer to the *Rotor Gene*TM software version 6.0.33. Please find further information on programming the *Rotor Gene*TM in the *Rotor Gene*TM 2000/3000 Operator's Manual.In the illustrations these settings are shown by arrows.

a) Setting of general assay parameters & Reaction volume.

First, confirm whether the PCR tubes used are No Domed PCR tubes by clicking in the box as shown in the figure. Please note that tubes with domes cannot be used in Rotor gene hence click in the box as shown below

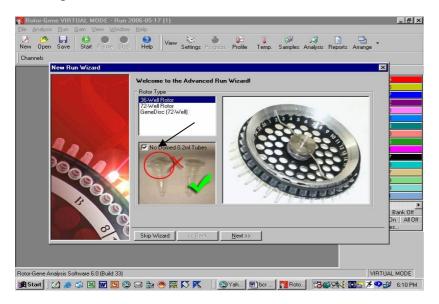


Fig.6.

Confirmation of Reaction Volume as follows.

| 💦 Rotor-Gene VIRTUAL MODE - Run 2008-08-14 (1) | _ = × |
|---|--|
| File Analysis Run Gain View Window Help | |
| New Open Save Start Pause Stop Help | |
| Channels New Run Wizard | |
| This screen displays miscellaneous options for the run. Complete the fields, circling Next when you are ready to move to the next page. Operator: FETER Notes: CHLAMYDIA PNEUMONIA QUANTITATION Fescellan Reaction Sample Layout: 1.2.3 Skip Wicard << Back Next> 2 2 3 3 4 4 5 5 5 5 1 | over here been been been been been been been b |
| | |
| | |
| Rotor-Gene Analysis Software 6.0 (Build 33) | VIRTUAL MODE |
| | 🤟 🧟 J🚵 7:10 PM |

Fig.7.Setting of general assay parameters.

Please click on the volume buttons to make sure that 25 ul is reflected in the window as shown above.

Then click next and a new window will open as shown below.

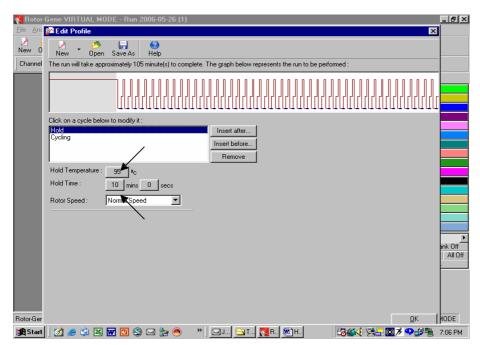
THERMAL PROFILE & CALIBRATION:

Here the thermal profile for the assay will be defined.

| 💦 Rotor-Gene VIRTUAL MODE - Run 2 | | _ 8 × |
|--|---|---|
| <u>File Analysis Bun G</u> ain <u>V</u> iew <u>W</u> indow | Help | |
| New Open Save Start Pause Stop | | Pa . rrange |
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| | the vizar on an iter your mound term for h | lements in d. For help n, hover se over the se over the physical sectors the sectors and the sectors |
| e (| Skip Wizard << Back Next >> | es |
| | | |
| Rotor-Gene Analysis Software 6.0 (Build 33) | | VIRTUAL MODE |
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Fig.8.

Programming the temperature profile is done by activating the button *Edit Temperature Profile* in the next *New Experiment Wizard* menu window as shown above.



CYCLINGPROFILE: Hold 95°C for 10 minutes as below

Fig.9.Initial activation of the Hot Start enzyme.

Setting up of denaturation step in the cycling profile as depicted below i.e.

95°C for 15 seconds.

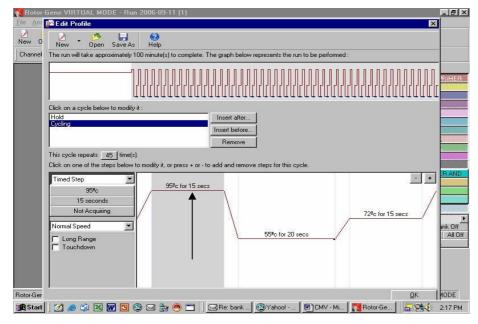


Fig.10.

Setting up of Annealing step in the cycling profile as depicted below i.e. 55°C for 20 Seconds and defining the Data acquiring channel i.e. FAM & JOE.

| 💦 Rotor- | | _ 8 × |
|-----------|---|--------------------|
| | 🗱 Edit Profile 🛛 🗙 | |
| New 0 | New Open SaveAs Help | |
| Channel | The run will take approximately 100 minute(s) to complete. The graph below represents the run to be performed : | |
| | | |
| | Click on a cycle below to modify it : | |
| | Hold Insert after | |
| | Remove | |
| | This cycle repeats 45 [time(s). | |
| | Click on one of the steps below to modify it, or press + or - to add and remove steps for this cycle. | |
| | Timed Step • + 55%c | RAND |
| | 20 seconds | |
| | Acquiring to Cycling A 72tc for 15 secs 72tc for 15 secs | Þ |
| | Namal Speed 55% for 20 secs | ank Off All Off |
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| Rotor-Ger | <u>O</u> K | IODE |
| Start | 📝 🥔 🏟 📓 🖬 📴 🕲 🖂 🎥 🖲 🗂 🗍 🛶 Ret bank 🕲 Yahool 🖻 CMV - Mi 🞇 Rotor-Ge 🕴 层 🧏 🤅 | 2:18 PM |

Fig.11.

Setting up of Extension step in the cycling profile as depicted below i.e. 72°C for 15 Seconds.

| 💦 Rotor-I | Gene VIRTUAL MODE - Run 2007-09-22 (1) | - 8 × |
|-------------|---|------------------------|
| | 😰 Edit Profile 🛛 🔀 | |
| New 0 | New Open SaveAs Help | |
| Channel | The run will take approximately 100 minute(s) to complete. The graph below represents the run to be performed : | |
| | | |
| |) Click on a cycle below to modify it : | |
| | Inset ater Inset ater Remove | |
| | · · · · · · · · · · · · · · · · · · · | |
| | This cycle repeats 45 time(s). Click on one of the steps below to modify it, or press + or - to add and remove steps for this cycle. | |
| | Timed Step | |
| | Normal Speed T Long Range 55% for 20 secs | ▶ nk Off All Off |
| Data Ca | | 1005 |
| Rotor-Ger | - 11 | IODE |
| Start Start | 🛛 💋 🥔 🖾 🖬 🔣 🙆 🕘 🚇 📕 📋 🔂 REAL TIME P 🖉 HBV - Microsof 🔣 Rotor-Gene VI 🛛 🐯 🖼 🗉 | 5:50 PM |

Fig.12.

Setting up of Number of Cycles to 45 cycles in the cycling profile as depicted below.

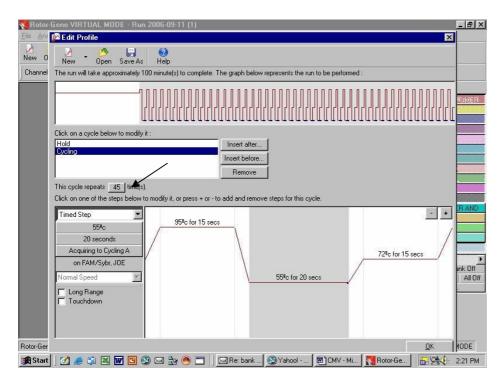


Fig.13.

Final Confirmation of the Thermal profile by pressing OK button as shown below.

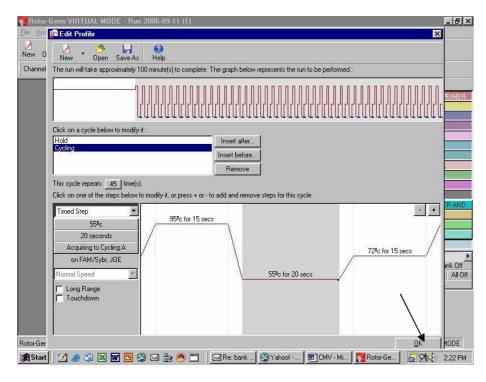


Fig.14.

e) Setting the gains for the acquiring channel by clicking at the Calibrate button as shown below.

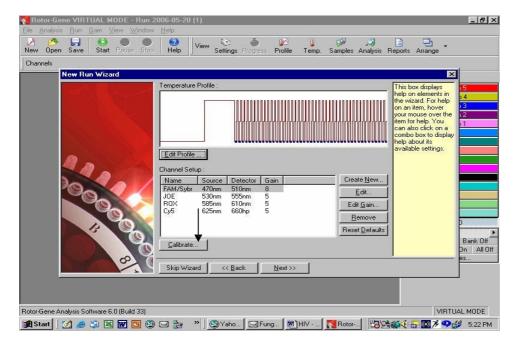


Fig.15.

The detection range of the fluorescence channels has to be determined according to the fluorescence intensities in the PCR tubes. This adjustment is done in the menu window *Auto Gain Calibration Setup* (activation in menu window *New Experiment Wizard* under *Calibrate*). Please set the calibration temperature to the annealing temperature of the amplification program (compare Fig. 16).

Adjustment of the fluorescence channel sensitivity as shown below.

| 😽 Rotor-Gene VIRTUAL MODE - Run 2006-05-26 (1) | _ 8 × |
|--|--|
| Eile Analysis Bun Gain Yiew Window Help | |
| New Open Save Start Pause Stop Help View Settings Progress Profile Term Samples Analysis Be | 🗈 🔁 🗸 ports Arrange |
| Channels Auto Gain Calibration Setup | |
| Calibration : | × |
| Auto-taan Labitation will read the fluoresence on the mosted sample at different gain levels until finds one at which the fluorescence levels are acceptable. The range of fluorescence you are looking for depends on the chemistry you are performing. Set temperature to 55 degrees. Calibrate All Calibrate Acquiring Perform Calibration Before 1st Acquirition | This box displays help on elements in he wizard. For help no an item, hover roour mouse over the tem for help. You can also click on a combo box to display help about its available settings. Bank Off n All Off es |
| Rotor-Gene Analysis Software 6.0 (Build 33) | VIRTUAL MODE |
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Fig.16.

| Rotor-Gene VIRTUAL MODE - Run 2006-05-26 (1) | |
|---|-----------------|
| Rotor-Gene VIRTUAL MODE - Run 2006-05-25 [1] File Analysis Bun Bain View Window Help New Open Save Stat Pouce 500 Help View Settinns Pouces Profile Tenn Samples Analysis Channels New Run Wizard New Run Wizard Oralibration Setup Calibration will read the fluorescence on the inserted sample at comparison on the inserted sample at comparison on the chemistry you are performing. Set temperature to 55 degrees. Calibration At 55 Degrees At Beginning Of Runi Channel Setting: Name Tube Position Min Reading Max Reading Edit RAM/Sybr 1 5FI JOE 1 Set Tube Position Min Reading Max Reading Edit Remove All | Peports Arrange |
| Start Manual Close Help Rotor-Gene Analysis Software 6.0 (Build 33) | VIRTUAL MODE |
| | See 2.16 PM |



Please do not forget to click on the box against "Perform calibration at 55°C at beginning of the run." After that press Close and a new window will open as shown below.

The gain values determined by the channel calibration are saved automatically and are listed in the last menu window of the programming procedure.

PRESS NEXT

| Rotor-Gene VIRTUAL MODE - Run 2006-05-18 (1) Ile Analysis Run Gain View Window Help | × |
|--|---|
| New Open Save Start Pouse Stop Help View Settings Progress Profile Temp. Samples Analysis Reports Anange | |
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| New Run Wizard | |
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| Rotor-Gene Analysis Software 6.0 (Build 33) IRTUAL MODE IR Start / / / / / / / / / / / / / / / / / / / | |

Fig.18.

Starting of the *Rotor Gene*™run.

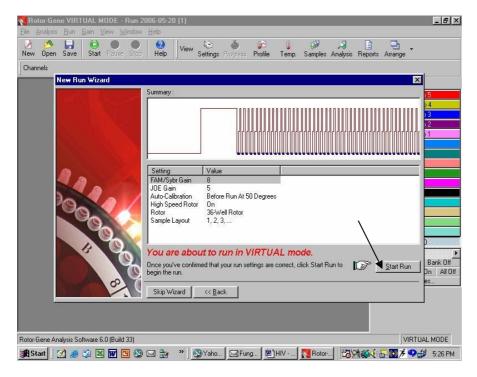


Fig.19:

Press Start Run Button.

| 💦 Rotor-Gene VIRTUAL MODE - Run 2006-10-19 (1) | _ 8 × |
|---|-----------------------------------|
| <u>Eile Analysis Bun Gain View Window H</u> elp | |
| New Open Save Start Parse Stop Help View Settings Progress Profile Temp. Samples Analysis Reports Arrange | |
| Channels | |
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| Save in: Rotor-Gene 6000 Software Changes Demonstrations Export Cod Archive | STD. PREP |
| Reports Templates | 0 |
| File name: C. FNEUMONIA 181105 Save Save as (ype: Run File (*:rex) Cancel | |
| | ▶ Bank Off Dn All Off es |
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| Rotor-Gene Analysis Software 6.0 (Build 33) VIRTUA | L MODE |
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Fig.20.

SAVING THE RUN FILE AS ABOVE

Store the run file either in my documents or a designated folder as shown above. The moment save button is clicked after the file name, machine will start.

8. Programming the Rotor Gene[™] 6000

The *Rotor Gene*[™] 6000 PCR program for the detection of **Chlamydia pneumonia** can be divided into following steps:

- A. Setting of general assay parameters & reaction volume
- B. Thermal Profile & Calibration
- C. Cycling profile/ & Initial activation of the Hot Start enzyme
- D. Cycling for Amplification of DNA
- E. Adjustment of the sensitivity of the fluorescence channels
- F. Starting of the Rotor Gene™ run

Program the *Rotor Gene*TM 6000 for these 5 steps according to the parameters shown in Figures 21-37 below All specifications refer to the *Rotor Gene*TM 6000 software version 1.7 Please find further information on programming the *Rotor Gene*TM in the *Rotor Gene*TM 6000 *Operator's Manual*,. In the illustrations these settings are shown by arrows.

g)

Setting of general assay parameters & Reaction volume.

Please see to it that you in advanced mode and then click Hydrolysis Probes.

On a double click the software will automatically go to the next function. If there is just a single click then click new after highlighting the Hydrolysis Probes.

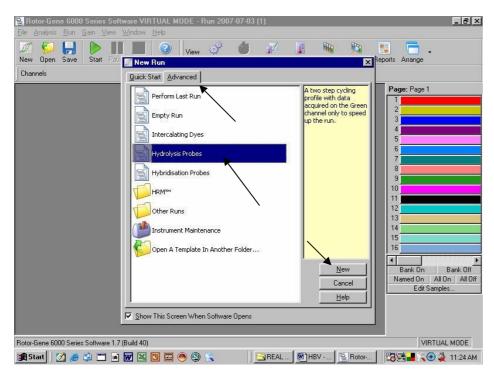


Fig.21.

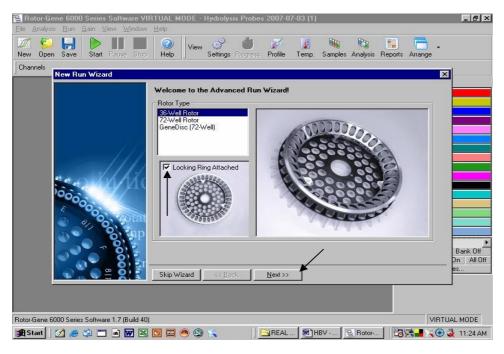


Fig.22.

First, confirm by clicking in the box as shown in the figure above whether a locking ring has been attached. Then click next to proceed to the next step.

Confirmation of reaction Volume as follows.

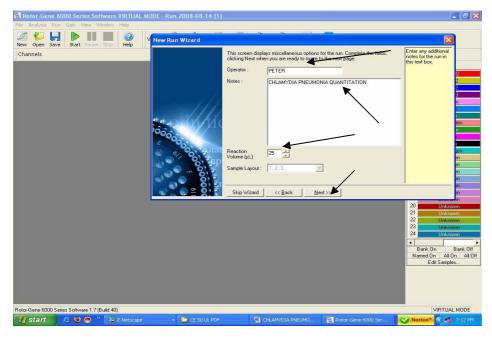


Fig.23.

- Please click on the volume buttons to make sure that 25µl is reflected in the window as shown above.
- In case required Operators Name can be Fed into the system.
- Any Notes which need to be either printed into the Report can be typed out in the Box of Notes.
- Then click next and a new window will open as shown below.

h) THERMAL PROFILE & CALIBRATION:

Here the thermal profile for the assay will be defined.

| Rotor-Gene 6000 Series Software V Ele Analysis Bun Gain View Window | RTUAL MODE - Hydrolysis Probes 2007-07-03 (1) Help | _ <u> </u> |
|--|---|--|
| New Open Save Start Pause Stop | Help View Settings Progress Profile Temp. Samples Analysis Reports Arran | je - |
| Channels New Run Wizard | | × |
| | Temperature Profile : This box display to the wead P Edk Profile Finite Souther Sou | ents in or help over You k on a o display |
| | Gain Optimisation | Bank Off Dn All Off |
| | Skip Wizard << Back Next >> | es |
| | | |
| Rotor-Gene 6000 Series Software 1.7 (Build 40 | 2 | VIRTUAL MODE |
| | | 11.20 AM |

Fig.24.

Programming the temperature profile is done by activating the button *Edit Profile* in the next *New RUN Wizard* menu window as shown above.

CYCLING PROFILE: First hold 95°C for 10 minutes as below

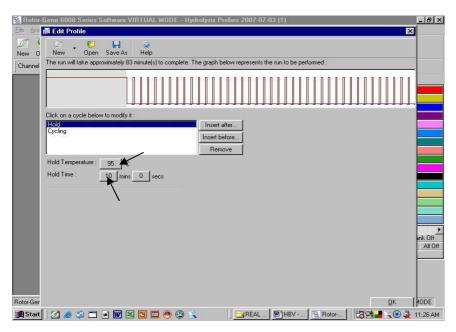


Fig.25. Initial activation of the Hot Start enzyme. Generally the window will open with the Hold Temp as95°Cand the Hold Time as10minutes. In case it is different then set the same to as shown above.

j) Setting the Cycling and acquisition.

When clicked on Cycling the window will open as below.

| Image: Control of the control of th | |
|--|---------|
| New 0 New Open Save As Help | |
| Channel The run will take approximately 83 minute(s) to complete. The graph below represents the run to be performed : | |
| | |
| | |
| Click on a cycle below to modify it : | |
| Hold Insett after | |
| Insert before | |
| Remove | |
| This cycle repeats 40 time(s). | |
| Click on one of the steps below to modify it, or press + or - to add and remove steps for this cycle. | |
| Timed Step | |
| 95*C 95*C 95*C for 10 secs | |
| 10 seconds | |
| Not Acquiring | |
| Long Range 60°C for 45 secs | ink Off |
| | All Off |
| | |
| | |
| | |
| Rotor-Ger OK | IODE |
| 1100 com (1100 | |

Fig.26.

Click on the Plus sign at the right hand as shown in the figure to make the Cycling a three step process. A new window as shown below will be there.

Setting up of denaturation step in the cycling profile as depicted below i.e. 95°C for 15 seconds.

| 😫 Rotor-G | iene 6000 Series Software VIRTUAL MODE - Hydrolysis Probes 2007-07-03 (1) | - 8 × |
|-----------|---|---------|
| Eile Ana | 🛛 Edit Profile 🛛 🔀 | |
| Mew 0 | New Open Save As Help | |
| Channel | The run will take approximately 100 minute(s) to complete. The graph below represents the run to be performed : | |
| | | |
| | Click on a cycle below to modify it : | |
| | Hold Inset after | |
| | Remove | |
| , | This cycle repeats 40 [time(s). | |
| | Click on one of the steps below to modify it, or press + or - to add and remove steps for this cycle. | |
| | Timed Step 95°C 95°C 15 seconds Not Acquiring Cong Range Touchdown 60°C for 45 secs | nk Off |
| Rotor-Ger | <u>o</u> k (| 10DE |
| 🚮 Start | 🗹 🥔 🛱 🖬 🖬 🖬 🖪 🧮 🖲 🕲 🐒 👘 🔰 📴 🖓 👔 👘 🖉 👘 👘 👘 🖉 | 1:27 AM |

Fig.27.

Setting up of Annealing step in the cycling profile as depicted below i.e.55°C for 20 Seconds.

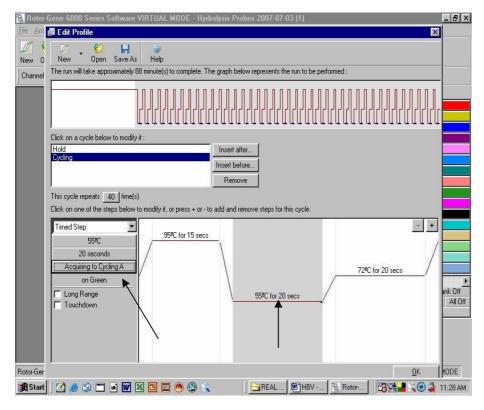


Fig.28.

After setting the Annealing temperature and the time for annealing click on the "Acquiring to Cycling A" as shown by arrow. A New window will open as shown below.

Defining the Data acquiring channel i.e. Green (FAM) & Yellow (JOE)

| | | | | E - Hydrolysis Probes 2007-07-03 (1) | |
|--|--|--|--|--|------|
| 📶 Edit Profile | | | | | > |
| | 📁 🛃 pen Save, | |) slp | | |
| The run will take | Acquisitio | n | | | |
| | Same as P | revious : | (New Acqui | isition) | |
| | | n Configu | | | |
| | Available | Channels | 1 | Acquiring Channels : | |
| | Name Crimson | | | Name Name | |
| Click on a cycle t Hold | Orange | | | Green | |
| Cycling | Red Yellow | | | << | |
| | reliow | | | | |
| | , To acqui | re from a c | hannel sele | ect it from the list in the left and click >. To stop acquiring from a | |
| | | | | | |
| This cycle repeat | channel, | | | and list and click <. To remove all acquisitions, click <<. | |
| This cycle repeat Click on one of th | channel, | | | | |
| | channel, | select it in | | and list and click <. To remove all acquisitions, click <<. | |
| Click on one of th | Dve Char | select it in | | | |
| Click on one of th Timed Step | channel, Dye Char Dye Char | select it in t>> | the right-ha | and list and click <. To remove all acquisitions; click <<. <table> </table> | |
| Click on one of th Timed Step 55ª | Dye Char Dye Char Dye Char Channel | select it in t>> nnel Sele Source | the right-ha | and list and click <. To remove all acquisitions; click <<. <u> QK</u> Don't Acquire <u>Help</u> art Dyes | |
| Click on one of th Timed Step 55ª 20 sec | Dye Char Dye Char Dye Char Channel Green | select it in t>> nnel Sele Source 470nm | the right-ha ection Cha Detector 510nm | and list and click <. To remove all acquisitions; click <<. <table> DK Don't Acquire Help art FAM^D, SYBR Green 1^D, Fluorescein, EvaGreen^D, Alexa Fluor 488^D</table> | secs |
| Click on one of th Timed Step 55° 20 sec Acquiring to | Dye Char Dye Char Dye Char Green Yellow | select it in t>> nnel Sele Source 470nm 530nm | the right-ha ection Cha Detector 510nm 555nm | Ist and click <. To remove all acquisitions; click <<. | |
| Click on one of th Timed Step 55° 20 sec Acquiring to on Gr | Dye Char Dye Char Channel Green Yellow Drange | select it in t>> Source 470nm 530nm 585nm | ction Cha Detector 510nm 555nm 610nm | Ist and click <. To remove all acquisitions; click <<. | |
| Click on one of the Step Step Step Step Step Step Step Ste | Dye Char Dye Char Channel Green Yellow Drange | select it in t >> select it in t >> source 470nm 530nm 585nm 625nm | ection Cha Detector 510nm 555nm 610nm 660nm | and list and click <. To remove all acquisitions; click <<. <table> DK Don't Acquire Help art Image: Control of the state o</table> | |
| Click on one of the Step Step Step Step Step Step Step Ste | Dye Char Dye Char Channel Green Yellow Drange | select it in t>> Source 470nm 530nm 585nm | ction Cha Detector 510nm 555nm 610nm | Ist and click <. To remove all acquisitions; click <<. | secs |
| Click on one of the Step Step Step Step Step Step Step Ste | Dye Char Dye Char Channel Green Yellow Drange | select it in t >> select it in t >> source 470nm 530nm 585nm 625nm | ection Cha Detector 510nm 555nm 610nm 660nm | and list and click <. To remove all acquisitions; click <<. <table> DK Don't Acquire Help art Image: Control of the state o</table> | |
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Fig.29.

Highlight the Yellow and then press the right arrow. Just see before shifting the yellow to right that there is no other channel in the right except for Green. In case any other Channel appears besides Green on the right then the same be shifted to the left.

Confirmation of Channels as shown below.

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| The run will take 🛛 | Acquisitio | on | | | - |
| | Same as P | revious : | (New Acqu | isition) | |
| | Acquisitio | on Configu | ration : | | |
| | Available | Channels | 1 | Acquiring Channels : | |
| | Name | | | > Name | haaaaaa |
| Click on a cycle t | Crimson | | | < Green Yellow | |
| Hold Cycling | Red | | | <<1 (i) | |
| | | | | | |
| | 1 | | | | |
| This cycle repeat | | | | act it from the list in the left and click >. To stop acquiring from a and list and click <. To remove all acquisitions, click <<. | |
| Click on one of th | errarinten, | | i dhe ngrit ne | | |
| Timed Step | | | | | . + |
| | Dye Char | t>> | | <u>O</u> K Don't Acquire <u>H</u> elp | |
| 20 sec | Dye Chai | nnel Sele | ection Cha | nt | |
| Acquiring to | Channel | Source | Detector | Dyes | |
| on Gr | Green | 470nm | 510nm | FAM ⁽¹⁾ , SYBR Green 1 ⁽¹⁾ , Fluorescein, EvaGreen ⁽¹⁾ , Alexa Fluor 488 ⁽¹⁾ | secs/ |
| Long Range | Yellow | 530nm | 555nm | JOE ⁽¹⁾ , VIC ⁽¹⁾ , HEX, TET ⁽¹⁾ , CAL Fluor Gold 540 ⁽¹⁾ , Yakima Yellow ⁽¹⁾ | |
| Touchdown | Orange | 585nm | 610nm | R0X ⁽¹⁾ , CAL Fluor Red 610 ⁽¹⁾ , Cy3.5 ⁽¹⁾ , Texas Red ⁽¹⁾ , Alexa Fluor 568 ⁽¹⁾ | |
| | Red | 625nm | 660nm | Cy5 ⁽¹⁾ , Quasar 670 ⁽¹⁾ , LightCycler Red640 ⁽¹⁾ , Alexa Fluor 633 ⁽¹⁾ | |
| | Crimson | 680nm | 710hp | Quasar705 ⁽¹⁾ , LightCycler Red705 ⁽¹⁾ , Alexa Fluor 680 ⁽¹⁾ | |
| | | | | | |
| | | | | | |
| | | | | | |
| | | | | | <u>0</u> K |

Fig.30.

Once the Yellow and Green Channels are on the Right side then press OK as shown by the arrow.

Setting up of Extension step in the cycling profile as depicted below i.e.72°C for 15Seconds

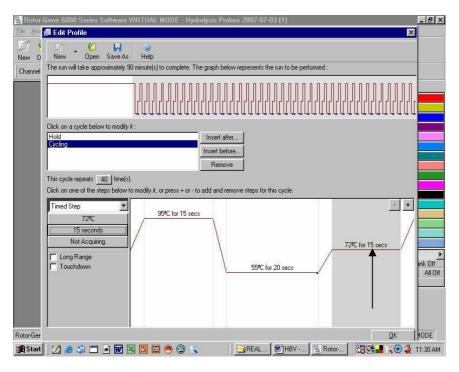


Fig.31.

Setting up of Number of Cycles to 45cycles in the cycling profile as depicted below.

| otor-Gene 6000 Series Softwa | re VIRTUAL MODE - H | ydrolysis Probes 2 | 007-07-03 (1) | | | |
|--------------------------------------|--------------------------------|---|------------------------|---------------|--------------------------------|-----|
| Ane 📶 Edit Profile | | | | | > | < |
| New Open Save | | | | | | |
| nnel The run will take approximat | ely 100 minute(s) to complete | e. The graph below re | presents the run to be | e performed : | | |
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| | and and an analysis and | | | | | - |
| | | | ┍╎┍╎┍╎┍╎┍╎┍╵┍ | | | ſ |
| | | | | | | |
| Click on a cycle below to mo Hold | dify it : | Insert after | | | | |
| Cycling | | Insert before | | | | |
| | | | | | | |
| | | Remove | | | | |
| This cycle repeats 45 tip | | | | | | |
| Click on one of the steps be | ow to modify it, or press + or | to add and remove : | steps for this cycle. | | | - |
| Timed Step | 95°C for 15 s | 808 | | | <u> </u> | ᆘ |
| 72ºC | | | | | | /[|
| 15 seconds | - 7 | | | | / | |
| Not Acquiring | _/ | 1 | | 7 | ⁷ 2ºC for 15 secs / | |
| Long Range | | | 022233 - 1033 | / | | an |
| I ouchdown | | | 55ºC for 20 secs | | | Ĩ |
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| | | | | | | |
| Ger | | | | | ▼ <u>_</u> K | 140 |

Fig.32.

After setting the number of Cycles Press OK.

Setting the gains for the acquiring channel by clicking at the Gain Optimization button as shown below.

| and the second se | | RTUAL MODE - Hydrolysis Probes 2007-07-03 (1) | _ & × |
|---|--|--|--------------------------------|
| Eile Analysi | s <u>R</u> un <u>G</u> ain ⊻iew <u>W</u> indow | Help | |
| M Kew Oper | Save Start Pause Stop | Help View 🔗 🍏 💋 📕 Karange | |
| Channels | New Run Wizard | × | r I |
| | New Hun Wizard | Temperature Profile : This box displays help on elements in the ward. For help on an item, hove more than hove more and the ward. For help on an item, hove more than hove more help on an item hove more help on an item hove more help on an item hove more than hove more help on an item hove more help on an an andite setting on an item hove more help on an an anditem h | Bank Off Pn All Off es |
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Fig.33

The detection range of the fluorescence channels has to be determined according to the fluorescence intensities in the PCR tubes. This adjustment is done in the menu window *Auto Gain optimization Setup* (activation in menu window *New Experiment Wizard* under Gain Optimization). Please set the temperature to the annealing temperature of the amplification program (compare Fig.34.)

| 😫 Rotor-Gene 6000 Serie | | | | | | | _ 8 × |
|--------------------------------|--|--|--|--|-------------------------------------|-------|------------------------------|
| <u>Eile Analysis Run G</u> ain | ⊻iew <u>W</u> indow <u>H</u> elp | | | | | | |
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| Channels | | r setup | | | | | |
| Channels New Run W | different gai chemistry yo Set temperal Optimise All € Perform Optimisation Perform Optimisation Channel Settings : Name Tube Position Green 1 Yellow 1 | u are performing. ure to 55 ± constant optimise Acquiring Before 1st Acquisitio At 55 Degrees At Brit 5FI 1 5FI 1 5FI 1 | ne at which the fluor. sence you are lookin prees. n sginning <u>DI Run</u> <u>Max Reading Mini</u> <u>Max Reading Mini</u> <u>10</u> 10FI -10 | escence levels are g for depends on the | Add Edit Bemove Remove All | - | Bark Off Dn All Off es |
| | <u>Start</u> Ma | rual Clo: | se <u>T</u> <u>H</u> elp | | | | |
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| | | | | | | | THUS AM |

Fig.34.

The following needs to be done.

- Click on the Optimise Acquiring
- Click on Set temperature To Adjust the Temperature to 55°C.
- Click on the Box Perform Optimisation At 55 degree At beginning of Run.
- Just see that below the channel settings there appear only two channels i.e. Green and Yellow. In case it is different than the channels have not been set properly. Close the window and reset the channels.
- Then Press Close.

L) PRESS Start RUN

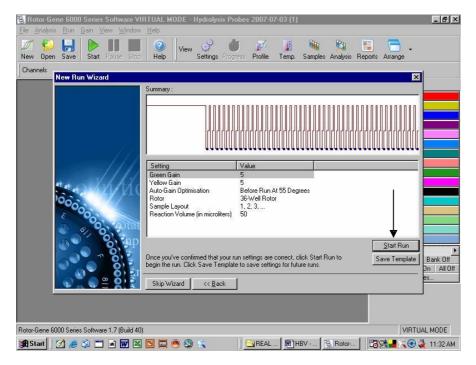


Fig.35.

Saving the RUN File.

| le | 000 Series Soft | ware VIRTUAL MODE | Run 2008-08-05 (1) | | | 2 |
|---|--|--|--|---|---|--|
| Save As A Je Save | e in: 🞯 Desktop | | | 2 🛛 | | |
| Ct Wy Recent Documents Desktop My Document My Document | Rotor-Gene Agreement ARTICLES A checklists_f corbett documents esurgicalsC GAURAV GAURAV Genome | er Dord (Places kapp Camonstrations later Commonstrations later Commonstrations later Commonstrations later Commonstrations later (Places later Commonstrations) Commonstrations Common | an II 50.55 t docs t docs so t docs so t docs ports t files ht files at yung genes | us softwares iortcut to data iortcut to New Folde | Value 4 4 8 9 Before Run At 58 Degrees 36:Well Floor 36:Well Floor 30:Well Floor microfilere) 50 | |
| My Network Places | < | Chlamydia pneumonia | | Save Cancel | d that your run settings are correct, click ave Template to save settings for future n | Start Run Save Template |
| | Lave di (jper | j nurrae (16x) | | Cancer | (Book | 19 20 21 22 23 24 * Bark On Bark On Named On All Da All Da All Da E dit Samples |
| otor-Gene 6000 Se | ries Software 1.7 (Bu | ild 40) | | | | VIRTUAL MODE |
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Fig.36.

Store the run file either in my documents or a designated folder as shown above. The moment save button is clicked after the file name, machine will start.

9. General data interpretation and Analysis

a) For Rotor Gene 2000/3000:

Data analysis is performed with the *Rotor Gene*[™] software according to the manufacturer's instructions (*Rotor Gene*[™] 3000 Operator's Manual).

The following results are possible:

1. A signal is detected in fluorescence channel Cycling A.FAM.

The result of the analysis is positive: The sample contains Chlamydia pneumonia DNA.

2. In fluorescence channel Cycling A.FAM no signal is detected.

No Chlamydia pneumonia DNA detectable. It can be considered negative if there is amplification in the Joe Channel. Please refer for details on PCR inhibition in the section PCR inhibition.

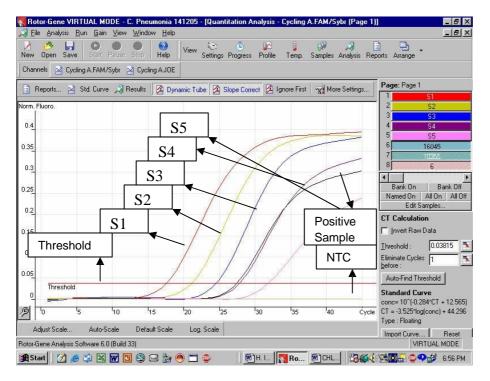


Fig. 37: Detection of the quantitation standards *(C.Pneumo. S 1-5)* in fluorescence channel Cycling A.FAM. NTC: non-template control.

Examples of positive and negative PCR reactions are given in the above figure.

| | 1 1 | <u>G</u> ain <u>V</u> iew <u>W</u> i | | 147 2525 | 100 C 100 | | | |
|-----------------------------|------------------|--------------------------------------|-----------------|---------------|--------------------|---|--------------------------|-------------------------------|
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| 0 | uantitation An | alysis - Cycling | A.FAM/Sybr (| Page 1) | | | _ 🗆 × | Page: Page 1 |
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| 10 | uant. Results | - Cycling A.FAM | 1/Sybr (💻 | 🗆 🗙 📝 Star | ndard Curve - Cyc | cling A.FAM/ | Sybr 💶 🗖 🗙 | Threshold: 0.03815 |
| No. | Name | Type Ct | Given | | | Cuoling & E | AM/Sybr (Page 1): | 1 |
| | S1 | Standard | | 35 30 | | R=0.99947 | Amoyor (ruge r). | Eliminate Cycles 1 |
| | S2 | Standard | 21.25 | 3! | | | 34 | Auto-Find Threshold |
| 2 | \$3 | Standard | 24.69 | : 525 | | M=-3.525 | | Auto-Find Threshold |
| 1 2 3 | | Standard | 27.96 | | | B=44.296 | | Standard Curve |
| 2 | S4 | | 31.98 | 20. | | Efficiency= | 0.82 | conc=10^(-0.284*CT + 12.565 |
| 2 | S4 S5 | Standard | | | 1 | + + - + - + + + + - + | | CT = -3.525*log(conc) + 44.29 |
| 1 2 3 4 5 11 | S4 S5 p00L | Unknown | 27.56 | - the second | | | 6 10^07 | T EL C |
| 2 | S4 S5 | | 27.56 | _ | 10^04 10 | 1977 - 1971 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 - 1977 | 0 10 0r | Type : Floating |
| 2 3 1 5 | S4 S5 p00L | Unknown | 27.56 | × | | Concentration | 0 10 07 | Import Curve Reset |

Fig.38.

Example of analysed data for C.pneumonia where all the amplification curves can be seen as well as the standard curve, reaction efficiency, M value etc. can be seen and the quantitative data for the samples can be seen.

Inhibition Control gene amplification.

Analyze the Joe Channel

PCR Inhibition

A signal is detected in fluorescence channel Cycling A. Joe: **No PCR inhibition**

In fluorescence channel Cycling A. Joe no signal is detected.

But signal detected in FAM: The sample is positive for Chlamydia pneumonia DNA.

In fluorescence channel Cycling A. Joe no signal is detected.

No signal detected in FAM as well possible PCR inhibition has occurred.

9b)

Generated Data Interpretation & Analysis for Rotor Gene 6000

Data analysis is performed with the *Rotor Gene*[™] software according to the manufacturer's instructions (*Rotor Gene*[™] 6000 Operator's Manual).

The following results are possible:

A signal is detected in fluorescence channel Cycling A. Green.

The result of the analysis is positive: The sample contains Chlamydia pneumonia DNA.

In fluorescence channel Cycling A.Green no signal is detected.

No Chlamydia pneumonia DNA detectable. It can be considered negative if there is amplification in the Yellow Channel. Please refer for details on PCR inhibition in the section PCR inhibition.

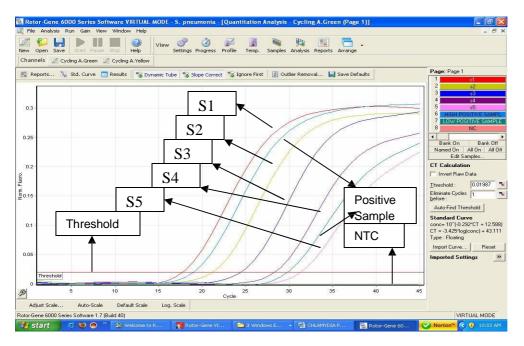


Fig. 39: Detection of the quantitation standards *(Chlamydia pneumonia S 1-5)* in fluorescence channel Cycling A. Green. NTC: non-template control.

Examples of positive and negative PCR reactions are given in the above figure.

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| 0 Threshold 5 | | | | 24 | Cycle ²⁵ | 30 | 35 | 40 | 45 | ☐ Invert Raw Data <u>Threshold</u> : 0.01987 Eliminate Cycles 1 <u>b</u> efore: 1 |
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| Adjust Scale Quant. Results - 2 0 Name 5 0 Name 5 63 84 85 | Auto-Scale Cycling A.G Type Ct Standard Standard Standard Standard Standard | Default Scale reen (Page 1) Giver 17.76 21.22 24.47 28.35 31.32 | Log. Scale n Conc (Cop Ca 2500000 2500000 250000 | - Conc (Cop 2521077 247060 27697 2040 2777 | Cycle Standard (30 28 | | | 1) | | Invert Raw Data [number Raw [number Raw Data [number Raw [numb |
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| Adjust Scale Quant. Results - 5 Name 91 52 93 54 95 HildH POSITIVE S LOW POSITIVE S | Auto-Scale Cycling A.G Type Ct Standard Standard Standard Standard Unknown Unknown | Default Scale reen (Page 1) Given 17.76 21.22 24.47 28.35 31.32 18.98 29.78 | Log. Scale n Conc (Cop Cd 2500000 250000 250000 250000 25000 | - Conc (Cop 2521077 247060 27697 2040 2777 | Cycle Standard (30 28 | | | 1) Cycling A.Green R=0.99941 R^2=0.99882 M=-3.425 B=43.111 | | Invert Raw Data Inveshold : 0.01987 Eliminate Cycles 1 Eliminate Cycles 1 Standard Enves conce 101/0.282*CT + 12.9 CT = 3.425*log(conc) + 43: Type : Floating Import Curve Rese |
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| o Threshold 5 Adjust Scale Quant. Results c. Name c1 c2 c3 c3 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c5 c4 c4 c5 c5 c4 c4 c4 c5 c5 c4 c4 c4 c5 c5 c4 c4 c5 c5 c5 c5 c5 c5 c5 c5 c5 c5 c5 c5 c5 | Auto-Scale Cycling A.G Type Ct Standard Standard Standard Standard Standard Unknown Unknown NTC I | Default Scale reen (Page 1) Given 17.76 21.22 24.47 28.35 31.32 18.99 29.78 NEG (NTC) | Log. Scale n Conc (Cop Cd 2500000 250000 250000 250000 25000 | - Conc (Cop 2521077 247060 27697 2040 2779 1114704 779 | Cycle Standard (30 28 5 24 22 20 | Curve - Cycling J | A.Green (Page | 1) Cycling A.Green R=0.99941 R^2=0.99882 M=-3.425 B=43.111 | (Page 1): | Invert Raw Data [number Raw [number Raw Data [number Raw [numb |
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Fig.40.

Example of analysed data for C.pneumonia where all the amplification curves can be seen as well as the standard curve, reaction efficiency, M value etc. can be seen and the quantitative data for the samples can be seen.

Inhibition Control gene amplification.

Analyze the Yellow Channel

PCR Inhibition

A signal is detected in fluorescence channel Cycling A. Yellow: No PCR inhibition.

In fluorescence channel Cycling A. Yellow no signal is detected. But signal detected in Green: The sample is positive for Chlamydia pneumonia DNA.

In fluorescence channel Cycling A. Yellow no signal is detected. No signal detected in Green as well: A possible PCR inhibition has occurred.

10. Troubleshooting

10. A) *a) For Rotor Gene 2000/3000:*

1. No signal with positive Standards (C.pneumonia S *1-5*) in fluorescence channel Cycling A.FAM.

- Incorrect programming of the *Rotor-Gene*[™] 2000/3000.
 - → Repeat the PCR with corrected settings.

2. Weak or no signal in fluorescence channel Cycling A. FAM:

- The PCR conditions do not comply with the protocol.
 - → Repeat the PCR with corrected settings.
 - →
- The **C.pneumonia** Super Mix *R1* has been thawed and frozen too often.
- The **C.pneumonia** Super Mix R1 has been kept at +4°C for longer than 5 hours.
 - → Please mind the storage conditions given in the **Storage**.
 - → Repeat the assay using a new HBV super mix (R1).
- The PCR was inhibited.
 - ➔ Make sure that you use a recommended extraction method (see 8.a. DNA extraction) and stick closely to the manufacturer's instructions.

b) For Rotor Gene 6000:

1. No signal with positive Standards (C.pneumonia *S 1-5*) in fluorescence channel Cycling A. Green.

- Incorrect programming of the *Rotor-Gene*[™] 6000.
 - ➔ Repeat the PCR with corrected settings.

2. Weak or no signal in fluorescence channel Cycling A. Green:

- The PCR conditions do not comply with the protocol.
 - → Repeat the PCR with corrected settings.
- The **C.pneumonia** Super Mix *R1* has been thawed and frozen too often.
- The **C.pneumonia** Super Mix R1 has been kept at +4°C for longer than 5 hours.
 - → Please mind the storage conditions given in the **Storage**.
 - → Repeat the assay using a new **C.pneumonia** super mix (R1).

• The PCR was inhibited.

Make sure that you use a recommended extraction method (see **8.a. DNA extraction**) and stick closely to the manufacturer's instructions.

11.Specifications

a. Sensitivity and Reproducibility.

In order to determine the sensitivity of the *Geno Sen's*[®] Chlamydia pneumonia *Real Time PCR RG Kit*, a dilution series has been set up from 10⁷ copies/ul down to 10⁻¹ Copies/µl of Chlamydia pneumonia DNA In-Vitro Transcription and analyzed with the *Geno Sen's*[®] Chlamydia pneumonia Real Time PCR RG kit *using RotorGene 3000/6000. The assays were carried out in triplicate except for the last* dilution run in 5 tubes. The results were determined by a probit analysis. The detection limit as determined for *Geno Sen's*[®] Chlamydia pneumonia Real Time *PCR RG Kit* is consistently 90 Copies/ml. This means that there is 95% probability that 90 copies/ml will be detected

Analytical Sensitivity

Analytical Sensitivity in Conjunction with the Geno Sen's[®] Genomic DNA Extraction Mini Kit — for DNA purification (Cat . No. 98021) of the Geno Sen's[®] Chlamydia pneumonia Real Time PCR RG Kit on Rotor Gene 3000/6000 was determined by Spiking a known negative Serum to a nominal 95 copies/ml. This was subjected to extraction using the Geno Sen's[®] Genomic DNA Extraction Mini Kit — for DNA purification (Cat. No. 98021) eight times with starting volume of 200µl & elution volume of 60µl. All the Eight extractions were then analyzed with the GenoSen's[®] Chlamydia pneumonia Real Time PCR RG Kit along with the standards & the NTC.

All the eight extractions were amplified correctly & the mean value obtained for the eight replicates was 93 copies/ml.

Hence Analytical Sensitivity in Conjunction with the Geno Sen's.[®] Genomic DNA Extraction Mini Kit — for DNA purification (Cat . No. 98021) of the Geno Sen's[®] Chlamydia pneumonia Real *Time PCR RG Kit on Rotor Gene 3000/6000 was determined to be 95 copies/ml.*

11.b Specificity

The specificity of the **Geno Sen's**[®] **Chlamydia pneumonia** Real Time PCR RG Kit is ensured by selection of very specific Primers & probes. This is also ensured by stringent reaction conditions. The primers & probes were blasted for possible homologies to all sequences published in the GeneBank. It was found that the primers & probes are very specific to the **Chlamydia pneumonia** DNA sequences & picks up all the known strains of **Chlamydia pneumonia** DNA.

To further Validate the stringent data In order to check the specificity of the **Geno Sen's**[®] **Chlamydia pneumonia** Real Time PCR RG Kit, different RNA & DNA listed below were analyzed with **Geno Sen's**[®] **Chlamydia pneumonia** Real Time PCR RG Kit. None of these led to a positive signal with the **Geno Sen's**[®] **Chlamydia pneumonia** Real Time PCR RG Kit.

| Varicella Zoster Virus | Hepatitis B Virus | N. Meningitis |
|------------------------|------------------------|----------------------------|
| S. Pneumonia | Hepatitis C Virus | ChikunGunya Virus |
| Epstein barr Virus | Hepatitis E Virus | JEV |
| Cytomagalovirus | HIV-1 | Mycobacterium tuberculosis |
| S.pneumonia | HIV-2 | Hepatitis A |
| Parvovirus B 19 | Leptospira nterrogans. | Staphylococcus aureus |
| Dengue Virus 1-4 | H. Influenza | Salmonella enteritidis |
| Leprosy | Malaria | Scrub typhus |
| B.pseudomallie | West Nile Virus | Leptospira interrogans. |

In order to validate the Specificity further with wet lab testing the following Known clinical samples were analyzed using the *Geno Sen's.*[®] Chlamydia pneumonia Real Time PCR RG kit on RotorGene 3000/6000. The extraction was carried out with the Geno Sen's[®] Genomic DNA Extraction Mini Kit — for DNA purification (Cat. No. 98021)

The run was carried out with the known set of standards in order to quantiate the **Chlamydia pneumonia** DNA Gene.

| Sample Type | B.swab | B.lavage |
|---------------------|--------|----------|
| High +ve's | 1 | |
| Medium +ve's | 4 | 2 |
| Low +ve's | 5 | 2 |
| Extremely low +ve's | 1 | 2 |
| Negative samples. | 3 | 3 |
| | 14 | 9 |

All the above samples were correctly identified by the *Geno Sen's*[®] Chlamydia pneumonia Real Time PCR RG kit & all the 3 extremely low samples were accurately detected by the *Geno Sen's*[®] Chlamydia pneumonia Real Time PCR RG kit & exhibited copies around 80 copies /ml or less than 80 copies/ml.

12. Warranty::

Products are guaranteed to confirm to the quality and content indicated on each vial and external labels during their shelf life. Genome Diagnostics Pvt. Ltd. obligation and purchaser's rights under the warranty are limited to the replacement by Genome Diagnostics Pvt. Ltd of any product that is shown defective in fabrication, and that must be returned to Genome Diagnostics Pvt. Ltd Freight prepaid, or at Genome Diagnostics Pvt. Ltd. option, replacement of the purchasing price. Any complaint on damaged goods during transport must be directed to the handling or transport agent.

In Vitro Diagnostic Medical device, as per the Directive 98/79/EC specifications. This product must be used by qualified professionals only. It is the responsibility of the user to ascertain that a given product is adequate for a given application. Any product not fulfilling the specifications included in the product sheet will be replaced. This warranty limits our responsibility to the replacement of the product. No other warranties, of any kind, express or implied, including without limitation, implicit warranties of commercialization ability or adequacy for a given purpose, are provided by Genome Diagnostics Pvt. Ltd. Genome Diagnostics Pvt. Ltd will not be responsible for any direct, indirect, consequential or incidental damage resulting of the use, misuses, results of the use or inability to use any product.

13. Limitations of product use:

- a.) All reagents may exclusively be used for *in vitro* diagnostics.
- b.) The product is to be used by personnel specially instructed and trained in for the *in-vitro* diagnostics procedures only.
- c.) It is important to pipet the indicated quantities, and mix well after each reagent addition. Check pipettes regularly.
- d.) Instructions must be followed Correctly in order to obtain correct results. If the

user has any questions, please contact our Technical Dept. (<u>dharam@vsnl.com</u> or at pbpl@vsnl.net).

- e.) This test has been validated for use with the reagents provided in the kit. The use of other Reagents or methods, or the use of equipment not fulfilling the specifications, may render equivocal results. User is responsible for any modifications done in any of the indicated parameters. Compliance with the kit protocol is required.
- f.) Detection of Viral DNA depends on the number of Viruses present in the sample, and can be affected by sample collection methods, patient-related factors (e.g. age, symptoms), or for infection stage and sample size.
- g.) False negative results may be obtained due to polymerase inhibition. It is recommended to perform control reactions to distinguish between inhibition and true negatives. This is achieved by the inhibition Control included in the kit.
- h.) Cross contamination between samples and exogenous DNA can only be avoided by following good laboratory practice. Instructions in this document must be strictly followed.
- i.) Attention should be paid to expiration dates printed on the kit box and labels of all components. Do not use expired components.

If you have any further questions or problems, please contact our technical support at <u>dharam@vsnl.com</u> OR <u>pbpl@vsnl.net</u>.

14. Publication and Reference

15. LIST OF GENO-SEN'S RANGE OF REAL TIME PCR KITS

| S.NO. | PRODUCT |
|-------|--|
| | |
| 1 | HIV-1 RG quantitative Real time PCR kit. |
| | |
| 2 | HBV RG quantitative Real time PCR kit. |
| - | |
| 3 | HCV RG quantitative Real time PCR kit. |
| 4 | HCV Genotyping 1/2/3/4 RG qualitative Real time PCR kit. |
| 5 | HEV RG quantitative Real time PCR kit. |
| 6 | HAV RG quantitative Real time PCR kit. |
| | |
| 7 | JEV RG quantitative Real time PCR kit. |
| 8 | ENTEROVIRUS RG quantitative Real time PCR kit. |
| | |
| 9 | DENGUE RG quantitative Real time PCR KIT |
| | |

| 10 | HSV 1 & 2 RG quantitative Real time PCR kit. |
|----|---|
| 11 | CMV RG quantitative Real time PCR kit. |
| 12 | Hanta Virus RG quantitative Real time PCR kit. |
| 13 | Measles Virus RG quantitative Real time PCR kit. |
| 14 | West Nile Virus RG quantitative Real time PCR kit. |
| 15 | H5 N1 (Bird Flu) RG quantitative Real time PCR kit. |
| 16 | Chikungunya RG quantitative Real time PCR kit. |
| 17 | TTV RG quantitative Real time PCR kit. |
| 18 | SARS RG quantitative Real time PCR kit. |
| 19 | JC/BK Virus RG quantitative Real time PCR kit. |
| 20 | MTb Complex RG quantitative Real time PCR kit. |
| 21 | MTb Complex /MOTT RG quanlitative Real time PCR kit. |
| 22 | Chlamydia pneumonia RG quantitative Real time PCR kit. |
| 23 | Streptococcous pneumonia RG quantitative Real time PCR kit. |
| 24 | N. Meningitis RG quantitative Real time PCR kit. |
| 25 | H. Influenza RG quantitative Real time PCR kit. |
| 26 | Leprosy RG quantitative Real time PCR kit. |
| 27 | Helicobacter Pylori RG quantitative Real time PCR kit. |
| 28 | Scrub Typhus RG quantitative Real time PCR kit. |
| 29 | B. Pseudomalie RG quantitative Real time PCR kit. |
| 30 | Filaria RG quantitative Real time PCR kit. |
| 31 | Leptospira(pathogenic) RG quantitative Real time PCR kit. |
| 32 | CCL3-L1 RG quantitative Real time PCR kit. |

| 33 | Malaria (P. Vivax) RG quantitative Real time PCR kit. |
|----|---|
| | |
| 34 | Bcr/abl Major RG quantitative Real time PCR kit. |
| 35 | Bcr/abl Minor RG quantitative Real time PCR kit. |
| | |
| 36 | PML/RARA RG quantitative Real time PCR kit. |
| | |
| 37 | RARA/PML RG quantitative Real time PCR kit. |
| | |
| 38 | GAPDH RG quantitative Real time PCR kit. |
| | |
| 39 | β-Actin RG quantitative Real time PCR kit. |
| | |
| 40 | β-Globin RG quantitative Real time PCR kit. |
| | |
| 41 | Abl gene RG quantitative Real time PCR kit. |
| | |
| 42 | Rabies RG quantitative Real time PCR kit. |
| | |
| 43 | Factor V Leiden detection RG Real time PCR kit. |

GENOME DIAGNOSTICS PVT. LTD. (AN ISO 13485 :2012,9001 : 2008 CERTIFIED COMPANY.) Up Mohal Naryal, KHASRA NO : 427,Opp. Divya Packers, Old Timber Depot Road, Near Sector 4,Parwanoo Distt. Solan, Himachal Pradesh-173220 ,INDIA. Tel No : 00-91-1792-234285, +91 9810037653 Fax : 00-91-1792-234286 E-mail: genome24@rediffmail.com dharam@vsnl.com pbpl@vsnl.net Version : 005 Websites : www.genomediagnostics.in www.genomediagnostics.com www.genomediagnostics.co.in www.diagnostics.genome.com