Geno-Sen's

PML/RARA (APML) (Rotor Gene)

Real Time PCR Kit

Quantitative

for use with the

Rotor Gene™ 2000/3000/6000

(Corbett Research Australia)



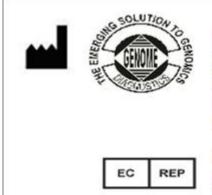
PACK INSERT

Revised July 2016



Genome Diagnostics Pvt. Ltd.

(An ISO 13485:2012, 9001:2008 Certified Company)



Genome Diagnostics Pvt. Ltd Up Mohal Naryal, Khasra No. 427, Old Timber Depot Road, Parwanoo, Dist Solan Himachal Pradesh 173220 Ph: +91 9810037653 Fax: 01792-234286 Email: genome24@rediffmail.com

EMERGO EUROPE Prinsessegracht 20,2514 AP, The Hague The Netherlands

Table of Contents

1.	Contents of Kit.	Page 3				
2.	Storage of the kit.	Page 4				
3.	PML/RARa information	Page 4				
4.	Precautions for PCR					
5.	Additionally required Materials & Devices					
6.	Principle of Real Time PCR	Page 6				
7.	Description of the Product.	Page 6				
8.	Procedure					
	 8.a RNA Extraction 8.b Reference Gene 8.c Quantitation 8.d Preparation for PCR 8.e Preparation for PCR amplification 8.f Programming of the Rotor Gene 2000/3000 8.g Programming of the Rotor Gene 6000 	Page 7 Page 7 Page 7 Page 8 Page 8 Page 10 Page 18				
9.	a) Generated Data Interpretation & Analysis 2000/3000 b) Generated Data Interpretation & Analysis 6000	Page 28 Page 29				
10.	a) Trouble shooting 2000/3000 b) Trouble shooting 6000					
11.	Specifications 11.a Sensitivity & Reproducibility 11.b Specificity					
12.	Warranty	Page 34				
13.	Limitations of product use.	Page 35				
14.	List of GENO-SEN'S range of Real Time PCR kits	Page 35				

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

1. Contents of the Kit:

Color	Contents	Cat. No.	Cat. No.	Cat. No.
Code		91116010	91116011	91116012
oouc				
		100 rxns	50 rxns	25 rxns
R1	PML-Bcr-1 Super Mix	25 rxns x 4 Vials	25 rxns x 2 Vials	25 rxns x 1 Vials
Blue	RG.			
R2 Blue	PML Bcr-3 Super Mix RG .	25 rxns x 4 Vials	25 rxns x 2 Vials	25 rxns x 1 Vials
R3	Abl Super Mix . RG	25 rxns x 4 Vials	25 rxns x 2 Vials	25 rxns x 1 Vials
Blue R4 Yellow	PML/RARa Mg Sol RG	1 vials	1 vial	1 vial
PML Bcr-1-	PML Bcr-1 Standard 1	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S1 Red	1 X 10 ⁶ copies/µl			
PML-Bcr-1-	PML Bcr-1 Standard 2	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S2 Red	1 X 10⁵ copies/µl			
PML-Bcr-1-	PML Bcr-1 Standard 3	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S 3	1 X 10⁴ copies/µl	-		-
Red PML -Bcr-	PML Bcr-1 Standard 4	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
1-S4	1 X 10 ³ copies/µl			
Red				
PML-Bcr-1-	PML Bcr-1 Standard 5	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S5	1 X 10 ² copies/µl	-	-	
Red Abl-S1	abl Standard 1	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Red	1 X 10 ⁶ copies/µl	ι νιαι οι 200μι		
Abl-S2	abl Standard 2	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Red	1 X 10⁵ copies/µl			
AbI-S3 Red	abl Standard 3 1 X 10⁴ copies/µl	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Abl-S4	abl Standard 4	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Red	1 X 10 ³ copies/µl			
Abl-S5	abl Standard 5	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Red	1 X 10 ² copies/µl PML Bcr-3 Standard 1			
PML Bcr-3- S1	1 X 10 ⁶ copies/µl	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
Red				
PML-Bcr-3-	PML Bcr-3 Standard 2	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S2	1 X 10⁵ copies/µl		•	-
Red				
PML-Bcr-3-	PML Bcr-3 Standard 3	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S3 Red	1 X 10 ⁴ copies/µI			
PML -Bcr-	PML Bcr-3 Standard 4	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
3-S4	1 X 10 ³ copies/µl			
Red				
PML-Bcr-3-	PML Bcr-3Standard 5	1 Vial of 200µl	1 Vial of 200µl	1 Vial of 200µl
S5	1 X 10 ² copies/µl	-	-	-
Red	Malaanka Oosta	A Mala of A sol	4 \/:=1	4 \(-1 \) (4 \) \
W	Molecular Grade	1 Vials of 1 ml	1 Vial of 1 ml	1 Vial of 1 ml
White	Water.			

R = Reagents

S = Quantitation Standards

W = Molecular Grade Water.

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

^{*} The Rotor Gene™ 2000/3000/6000 is a registered trademark of Corbett Research, Australia.

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, 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.

PML/RARa Information

3. Application

Acute promyelocytic leukemia (APL) known as acute myeloid leukemia-3, AML3, or M3 according to the French-American-British (FAB) classification is characterized by a predominance of malignant promyelocytes that carry a reciprocal translocation between the long arms of chromosomes 15 and 17: t(15;17)(q22;q11.2-q12). This translocation is diagnostic for APL as it is present in almost 100% of cases. De The et al. showed that the (15;17) translocation fuses the retinoic acid receptor alpha (*RARA*) locus to a gene, initially called myl and then renamed *PML*. Retinoic acid, a metabolite of vitamin A, can overcome the block of maturation at the promyelocytic stage and induce the malignant cells to terminally mature into granulocytes resulting in complete albeit transient disease remission.Comparison of the cDNA structures obtained by multiple groups showed variation in the amount of PML sequences included in the fusion protein. The *RARA* portion was invariant, containing the DNA-binding and ligand-binding motifs. The *PML* sequence variation seen among patients was generated by heterogeneous breakpoint cluster regions as well as by alternative splicing.

Bcr1 is more 3' within the PML gene and includes sequences from PML exons 5 and 6. Breakage in bcr2 involves sites in and around exon 6 of PML and leads to an intermediate length of PML sequence. The most 5' breakpoint, bcr3, fuses PML exons 1 through 3 of PML to RARA exon 3. This breakpoint yields short PML-RARA fusion proteins. In general, about 50% of patients exhibit long PML-RARA fusion proteins associated with bcr1.

Many cases of leukemia are caused by translocations that result in the formation and expression of chimeric fusion genes. The majority of APL patients harbor a translocation between chromosomes 15 and 17, with resultant fusion of the PML gene, at 15q12, with the retinoic acid receptor α (RAR α) gene, at 17q22. This gene fusion results in production of a leukemia-specific chimeric mRNA, PML-RAR α . Acute promyelocytic leukemia is defined by a characteristic morphology (AML FAB M3/M3v), by the specific translocation t(15;17) and its molecular correlates (PML/RARa and RARa/PML). Thereby it can be separated from all other forms of acute leukemia. Now it is routinely possible to detect extremely small numbers of transcripts from these leukemia-associated fusion genes in the blood or bone marrow of affected patients with the help of Real Time PCR. Geno Sen's PML/RARa Real

Time PCR kit is a sensitive and quantitative assay that can expeditiously analyze large numbers of samples from the patients samples.

Fusion transcripts containing the RARa/PML can be measured with a distinct Geno-Sen's Kit.

The kit is optimally suited for quantification of residual amounts of leukemia-specific Ph⁺ cells and for the follow up of leukemia patients undergoing therapy. The increase or decrease of PML/RARa transcripts normalized to abl transcripts may be an additional tool for early treatment decisions.

The Geno-Sen's PML/RARa cDNA Quantification assay is developed for laboratory scale or high-throughput quantitative transcript analysis by real time quantitative fluorescence PCR.

Geno-Sen's standardized ready-to-use Control and Reaction mix allow fast processing of the samples.

Module specificity

In vitro quantification of PML/RARa fusion transcripts Bcr1 isoform (type A) & Bcr 3 isoform (Type B) in whole RNA/mRNA samples which are reverse transcribed into cDNA using random hexanucleotides, optimized for transcripts prepared from crude or purified human white blood cells (WBC).

Samples which can be used for Extraction: Whole Blood or cultured cells.

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

- RNA isolation kit (see 8.a. RNA 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 PML/RARa** PCR Reagents constitute a ready to use system for detection and quantification of **PML/RARa** 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 either the Bcr 1 isoform or Bcr 3 isoform of PML/RARa and for the direct detection of the specific amplicon in fluorescence channels Cycling A.FAM/GREEN of the *Rotor Gene 2000/3000/6000 & the Reference gene on* Cycling A.FAM. In a separate reaction to determine ratio of PML Vs ABL gene. External positive Standards (PML Bcr-1 S 1-5), (PML Bcr-3 S 1-5) & (ABL S 1-5) are supplied which allow the determination of the gene load. For further information, please refer to section 8.3 Quantitation.

8. Procedure

8.a RNA Extraction

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

Sample Material	Nucleic Acid Isolation Kit	REF Cat. Num.	
	QIAamp RNA Blood Mini extraction Kit (50)	52304	QIAGEN
cells.	GenoSen's whole blood RNA Extraction kit (50)	98007	Genome Diagnostics Pvt.Ltd.

However the customer can use their own extraction systems depending on how good the yield is. Always use an extraction kit with a higher RNA 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 *PML/RARa Rotor Gene PCR Reagents* should not be used with phenol based isolation methods.

8.b *Reference gene:*

Reference Gene i.e. **abl** allows the user to determine the PML/RARa Vs abl ratio & simultaneously control possible PCR inhibition. The Reference gene control reagents are provided separately and have to be run separately on FAM channel itself. There is no need to add separate internal control in the reaction mix.

8.c Quantitation

Geno-Sen's PML/RARa Real Time PCR Kit for Rotor Gene 2000/3000/6000

The quantitation standards provided in the kit (*PML Bcr-1 S 1-5*), (*PML Bcr-3 S 1-5*) & (*abl S 1-5*) are treated in the same way as extracted samples and the same volume is used i.e. (2µl) instead of the sample. To generate a standard curve in the *RotorGene*TM 2000/3000/6000, all 5 Standards should be used as defined in the menu window *Edit Samples* of the *RotorGene*TM software. The same should also be defined as standards with the specified concentrations (see *RotorGene*TM Manual). The standard curve generated as above can also be used for quantitation in subsequent runs, provided that at least one standard is used in the current run. For this purpose, the previously generated standard curve needs to be imported (see *Rotor Gene*TM 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.

<u>Attention</u>: 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/ml) =

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

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 Qiagen QIAamp RNA Blood Mini extraction kit is 1.5 ml & the final Eluted Volume is 40µl 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 ⁶ Copies/μl =	26660000 Copies/ml
S2: 10 ⁵ Copies/µl =	2666000 Copies/ml
S3: 10 ⁴ Copies/μl =	266600 Copies/ml
S4: 10 ³ Copies/μl =	26660 Copies/ml
S5: 10 ² Copies/μl =	2666 Copies/ml

8.d Preparation for PCR

8. e. Preparation for PCR amplification:

Geno-Sen's PML/RARa Real Time PCR Kit for Rotor Gene 2000/3000/6000

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 X 3 (in triplicate). 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 (*PML Bcr-1 S 1-5*) (*PML Bcr-3 S 1-5*) & (*abl S1-5*) 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

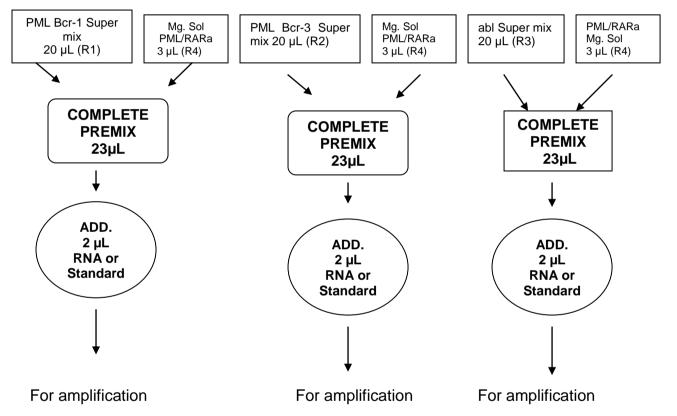


Fig. 4.

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

For Bcr 1 isoform

Bcr	1 rxns.	10 rxns.
Bcr Super Mix (R1)	20 µL	200 µL
Bcr Mg Sol. (R4)	3 µL	30 µL
Total	23µL	230µL

For Bcr 3 isoform

Bcr 1 rxns. 10 rxns.

Bcr Super Mix (R2)	20 µL	200 µL
Bcr Mg Sol. (R4)	3 µL	30 µL
Total	23µL	230µL

For abl

abl	1 rxns.	10 rxns.
abl Super Mix (R3)	20 µL	200 µL
PML/RARa Mg Sol. (R4)	3 µL	30 µL
Total	23µL	230µL

Fig. 5.

Pipette 23 µl of the Master Mix into each PCR tube labelled separately for Bcr 1 isoform, Bcr 3 isoform & *abl* respectively. Then add 2 µl of RNA to each sample tube in both sets and mix well by pipeting up and down. Correspondingly, 2 µl of the Standards (PML Bcr-1 *S1-5*), (PML Bcr-3 *S1-5*) & (abl S 1-5) must be used as a positive control and 2 µl of water (*Water, PCR grade*) as a negative control. Close the PCR tubes and transfer the PML tubes into the rotor of the *RotorGene*TM instrument whereas that of abl into the deep freezer at –20°C so that the same can be run later if required or else if the sample batch is small then the same can be run simultaneously in the same batch. The *RotorGene*TM software versions 5.0.53 and higher require a Locking Ring (accessory of the *RotorGene*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

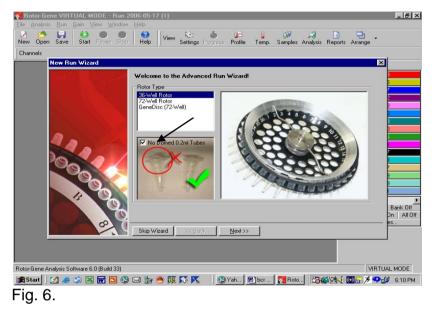
The *RotorGene*[™] 2000/3000 PCR program for the detection of PML/RARa can be divided into following steps:

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

Program the *RotorGene*TM 2000/3000 for these 5 steps according to the parameters shown in Fig. 6-19. All specifications refer to the *RotorGene*TM software version 6.0.33. Please find further information on programming the *RotorGene*TM in the *RotorGene*TM 2000/3000 Operator's Manual,. In the illustrations these settings are shown by arrows

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.



Confirmation of Reaction Volume as follows.

			_	dODE - Ru			(1)										_ & ×
	ynalysis	<u>B</u> un	<u>G</u> ain	⊻iew <u>W</u> ind													
Mew New	🤌 Open	↓ Save	Start	Pause S		🕜 Help	View	Settings	Progress	Profile	🌡 Temp.	💞 Samples	Analysis	E Reports	Arrange .	•	
Chan	nels																
-	_	New Ru	ın Wiz	ard												×	
					Thi	is screen :king Ne:	i displaj xt wher	ys miscell n you are	aneous or ready to n	tions for th iove to the	e run. Cor Next pag	mplete the f e.	ields,	Enter a notes l this te:	any addition for the run i «t box.	ial n	i ABL
					Op	erator :		PETER	-								
					No	tes :		PML/RA	ARa 🖌	K							BL
																	DL
				11													Ni¢
		20		0						/							1A
		-	30														NA NA
			X	9		action lume (μL)):	25									NA
			8					1, 2, 3		-	/					Ē	Þ
					0					_	/						Bank Off On All Off
				8	(Skip Wiz	ard 1	<< Ba	ek 1	Next>>	1					-	98
	1									<u>11</u> 0/07/77		_	_	_			
	40.5	nalysis So		6.0 (Build 33)				1.1					1.1		· · ·		L MODE
🚮 Sta	art	🥖 🥭	🗯 🛯	g 👿 🖸 I	🔊 🖂) 눩 🥘) 🗖	(M)	PML RAR/	A - Mic	Rotor-	Gene VIR	. 23	24 4	🚽 💽 考 <	<mark>> 4</mark>	8:35 AM

Fig. 7. Setting of general assay parameters.

- Please click on the volume buttons to make sure that 25µl 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 2006-05-17 (1)	_ & ×
<u>Fi</u> le <u>A</u> nalysis <u>B</u> un <u>G</u> ain <u>View</u> <u>W</u> indow <u>H</u> elp	
New Open Save Statt Pause Stop Help View Settings Progress Profile Temp. Samples Analysis Reports Arrange	•
Channels	
New Run Wizard	×
Image: Second state of the second s	s in 5 elp 7 rr 7 u 3 splay 3
	IRTUAL MODE
🏽 🕅 Start 🛛 🧭 🗯 🖾 🚾 📴 🕲 🖙 🛬 🔿 🎆 🚿 📉 🔤 💁 🖓 Vah 🖻 bor 💦 Roto	🤗 💕 6:03 PM

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.

CYCLING PROFILE: First hold 50°C for 15 minutes i.e. cDNA synthesis step as below.

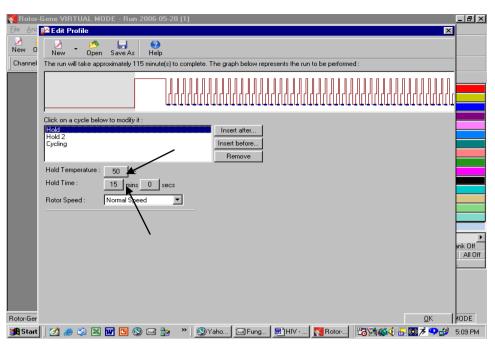


Fig. 9.

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

💦 Rotor-Gene VIRTUAL MODE - Run 2009-0	01-29 (1)	_ 7 🗙
File Analysis Run Gain View Window Help		
New Open Save Start Pause Stop Help	🏴 Edit Profile	
Channels	New Open Save As Help	
	The run will take approximately 109 minute(s) to complete. The graph below represents the run to be performed :	
New Run Wize	Cick on a cycle below to modify it :	
	Hold Cycling Control Holdy 1. Inset alter Cycling Remove	CDNA NA IEW OL
	Hold Time : 95 % Hold Time : 10, mins 0 secs Rotor Speed : Nome Speed •	CDNA CDNA NA
		L.6MM
		nk Off
Rotor-Gene Analysis Software 6.1 (Build 93)	with EU Rep word	VIRTUAL MODE
🛃 start 📄 🧔 🕲 🎕 🐣 🤤 Inbo	ox for dharam@d 🔁 with EU Rep word 🛛 🐨 4 Microsoft Office 👻 🌄 Rotor-Gene VIRTUAL	🔹 🛒 🛃 12:04 PM

Fig. 10. 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.

Geno-Sen's PML/RARa Real Time PCR Kit for Rotor Gene 2000/3000/6000

💦 Rotor-Gene VIRTUAL MODE - Run 2009-0	1-29 (1)			🗖 🗗 🗖
File Analysis Run Gain View Window Help				
New Open Save Start Pause Stop Help	🔑 Edit Profile			\mathbf{X}
Channels	New Open Save As Help			
	The run will take approximately 109 minute(s) to	complete. The graph below represents	the run to be performed :	
New Run Wiza				
	Click on a cycle below to modify it : Hold Hold 2 Cycling	Insert after		CDNA CDNA NA
	This cycle repeats 45 time(s). Click on one of the steps below to modify it, or p		this cycle.	
	Timed Step 95% 95% 15 seconds Not Acquiring Normal Speed Cong Range Touchdown	tor 15 secs	0/ 20 secs	or 15 secs
				<u>0</u> K
Rotor-Gene Analysis Software 6.1 (Build 93)				VIRTUAL MODE
🛃 start 💋 😂 🎕 👋 🤹 Inbo	< for dharam@d 🔁 with EU Rep word	😿 4 Microsoft Office 👻	Rotor-Gene VIRTUAL	🔇 💕 💕 12:05 PM

Fig. 11.

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

💦 Rotor-Gene VIRTUAL MODE - Run 2009-0	I-29 (1)	ĸ
File Analysis Run Gain View Window Help		
New Open Save Start Pause Stop Help	😰 Edit Profile 🛛 🔀	
Channels	New Open Save As Help	
	The run will take approximately 109 minute(s) to complete. The graph below represents the run to be performed :	_
New Run Wiza		
	Click on a cycle below to modify it : 1 CDNA Hold : Insert after 2 DDNA Hold 2 Insert after NA Cycling Remove If KW 00	
	This cycle repeats _45_ line(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 for 15 secs 20 seconds CDNA Acquing to Cycling A 72% (or 15 secs)	
	Acquiring to Lyceing A on FAM Normal Speed	
	Cong Range Touchdown	•
		-
Rotor-Gene Analysis Software 6.1 (Build 93)	VIRTUAL MODE	
	for dharan@d 🎓 with EU Rep word 🛛 🛛 4 Microsoft Office 🔹 🏹 Rotor-Gene VIRTUAL 🌒 🌾 🐉 12:06 PM	

Fig. 12.

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

💦 Rotor-Gene VIRTUAL MODE - Run 2009-0	-29 (1)			_ 7 🗙
File Analysis Run Gain View Window Help				
New Open Save Start Pause Stop Help	P Edit Profile			
Channels	New Open Save As Help			
	The run will take approximately 109 minute(s) to co	mplete. The graph below represents th	e run to be performed :	
New Run Wiza				
	Click on a cycle below to modify it : Hold Hold 2 Cycling	Insert after Insert before Remove		CDNA CDNA NA
	This cycle repeats 45 time(s). Click on one of the steps below to modify it, or pres		is cycle.	
	1/2°C 15 seconds Not Acquiring Normal Speed	r 15 secs	72% for 15:	- + CDNA CDNA NA L6MM
	Long Range Touchdown			CL:NA rk Off
				<u>o</u> k
Rotor-Gene Analysis Software 6.1 (Build 93)				VIRTUAL MODE
🛃 start 🛛 🔅 🥹 🍣 👋 🏹 Inbos	for dharam@d 🔁 with EU Rep word	4 Microsoft Office 👻	💦 Rotor-Gene VIRTUAL	C 🖉 🖉 12:09 PM

Fig. 13.

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

be	low.

Rotor-Gene VIRTUAL MODE - Run 2009-01	l-29 (1) 🖉 🖉 🗙
File Analysis Run Gain View Window Help	
New Open Save Start Pause Stop Help	📭 Edit Profile 🛛 🔀
Channels	New Open Save As Help
	The run will take approximately 109 minute(s) to complete. The graph below represents the run to be performed :
New Run Wizar	Click on a cycle below to modify it : Hold Insert after Cycling Insert before
	This cycle repeats 45 mole). Elik on one of the steps below to modify it, or press + or - to add and remove steps for this cycle. Image: Comparison of the steps below to modify it. or press + or - to add and remove steps for this cycle.
	72% 95% for 15 secs CDNA 15 seconds 72% for 15 secs CDNA Normal Speed 72% for 15 secs CDNA I Long Range 55% for 20 secs If referse If referse Touchdown 55% for 20 secs If referse If referse
	<u>K</u>
Rotor-Gene Analysis Software 6.1 (Build 93)	VIRTUAL MODE
🛃 start 👘 😂 🥹 🔌 🕌 🤤 Inbox	for dharam@d 🍃 with EU Rep word 🛛 👿 4 Microsoft Office 🔹 🍢 Rotor-Gene VIRTUAL



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

🛜 Rotor-Gene VIRTUAL MODE - Run 2009-0	I-29 (1) 📃 🗗 🗙
File Analysis Run Gain View Window Help	
New Open Save Start Pause Stop Help	😰 Edit Profile 🛛 🔀
Channels	New Open Save As Help
	The run will take approximately 109 minute(s) to complete. The graph below represents the run to be performed :
New Run Wiza	
	Click on a cycle below to modify it: ICDNA Hold Insert after 50156 Hold 2 Insert after NA Gysters NA NA Bysters Remove IEW OIL
	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 95% for 15 secs CDNA 15 seconds Nat Acquiring CDNA Normal Speed 72% for 15 secs LBMM Long Range 55% for 20 secs CDNA Touchdown DK CDNA
Rotor-Gene Analysis Software 6.1 (Build 93)	VIRTUAL MODE
🛃 start 🔰 😂 🍣 🦉 🤤 Inbo	: for dharam@d 🎽 with EU Rep word 🛛 🛛 4 Microsoft Office 🔹 🌄 Rotor-Gene VIRTUAL 🔍 🔊 😰 12:11 PM

Fig. 15.

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

	ene VIRTUAL MODE - Run 2006-05-18 (1)		
<u>File</u> <u>A</u> nalysis	sis <u>R</u> un <u>G</u> ain <u>V</u> iew <u>W</u> indow <u>H</u> elp		
New Open		ples Analysis Reports Ar	Tange
Channels			
	New Run Wizard		×
	EAM/Sybz 470nm 510nm 8 PROX 505nm 570nm 570nm 5 PROX 585nm 510nm 5 5 Cy5 525nm 650hp 5 5	Create New Edt Edt.gain Reset Defaults	the 0 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
40.5	Analysis Software 6.0 (Build 33)	Hereinsteine Autom	VIRTUAL MODE
Start	🗹 🥭 🕸 🖬 🖬 🕲 🕲 😂 🤌 🔹 🧕 🕲 Ya 🔤 Re 💦 Ro 👼 bor	1797 - 19	🔘 差 ᡐ 🚅 🛛 12:54 PM

Fig. 16.

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. 17).

Teb Analysis Run Gain Vere Worder Hell Wer Open Sove Stat Fraze 200 Hell Vere Run U Channels Person Sove Stat Fraze 200 Hell Vere Run U Free Run	💦 Rotor-Gene VIRTUAL MODE - Run 2009-01-29 (1)	. . .
New Open Save Start Pasie Start Pasie Start Pasie Start Pasie Page: Page 1 Channels Intro Gain Calibration Setup Intro Calibrat		
Channels		
Auto Gain Calibration Sciup Image: Calibration Sciup Image: Calibration Windows New Run W Calibration: Image: Calibration Windows Image: Calib		
Auto Gain Calibration Setup Image: Calibration New Run W Image: Calibration Image: Calibration Calibration I		Page: Page 1
Altro Calin Calindration Setup 3 New Run W Calination W do Can Calination Auto Can Calination W do Can Calination W dipleter Set temperature to Set temperature to Calination Calination V do Can Calination Actors Calination Set temperature to Set temperature to Calination Calination Calination Min Gain Manne Calination Manne Calination Manne Calination Manne Calination Manne Calination Manu Set <td< td=""><td></td><td>1</td></td<>		1
New Run VI Calkration Ado Gain Calkration will read the fluorescence on the inserted sample at different gain levels util it fluors one at which the fluorescence levels are chemistry on use tooking for depends on the chemistry ou are performing. Image: Calkration of fluorescence on the inserted sample at different gain levels util it fluors one at which the fluorescence levels are chemistry ou are performing. Image: Calkration of fluorescence on the inserted sample at different gain levels util it fluors one at which the fluorescence levels are chemistry ou are performing. Image: Calkration of the inserted sample at different gain levels util it fluorescence levels are chemistry ou are performing. Image: Calkration of the inserted sample at different gain levels util it fluorescence levels are chemistry ou are performed at the chemistry ou are performed at a calkration of the inserted sample at different gain levels util it fluorescence levels are chemistry ou are performed at the chemistry ou are performed at the chemistry ou are performed at a calkration of the chemistry ou are performed at the chemistry ou are performed at the chemistry out and performed at the chemistry out and themistry out and th	Auto Gain Calibration Setup	
Deter Care Andrés Coloure (1,0,210)	Auto Gain Calibration will read the fluorescence on the instreted sample at acceptable. The range of knorescence levels are acceptable. The range of knorescence levels are ac	
Data Cara Arabin Calmar C1 (0.3400)		
Nucl-dene Analysis Sociwale 6.1 (poliu 35)	Rotor-Gene Analysis Software 6.1 (Build 93)	VIRTUAL MODE
🔰 start 🔰 S 🤒 🌯 🛸 🕼 Inbox for dharan@d 🔪 with EU Rep word 🛛 🖾 4 Microsoft Office 🔹 💽 Rober-Gene VIRTUAL 🔹 📀 🛃 12:13 PM	🤁 start 💫 😂 🍣 🧉 🍯 Inbox for dharam@d 📄 with EU Rep word 🛛 👿 4 Microsoft Office 🔹 🌄 Rotor-G	ene VIRTUAL 🔿 🗊 🐉 12:13 PM

Adjustment of the fluorescence channel sensitivity as shown below.

Fig. 17.

Please do not forget to click on the box against "Perform calibration before 1st, acquisition." 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 (Fig. 8).

PRESS NEXT

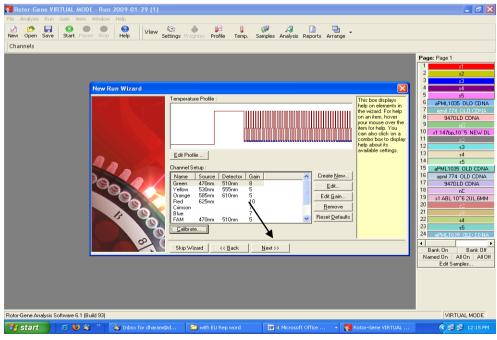


Fig. 18.

Starting of the *Rotor Gene*[™] run. **FIG: AMPLIFICATION OF THE cDNA**

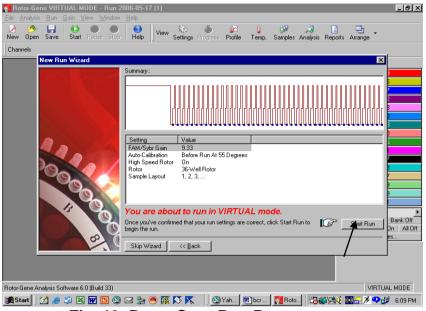


Fig. 19: Press Start Run Button.

🗙 Rotor-Gene VIRTUAL MODE - Run 2006-05-20 (1)
File Analysis Bun Gain View Window Help
New Open Save Start Parse Stop Help View Settings Progress Profile Temp. Samples Analysis Reports Arrange
Channels
New Run Wizard
Save As
Save jn: RotorGene 6 Save jn: RotorGene 6 Changes Changes Changes Personstations File name: PML-RARa 200506 Save Persons Save Persons Save
Save as type: Run File (*:rex) Cancel Cancel Concel Conce
Rotor-Gene Analysis Software 6.0 (Build 33) VIRTUAL MODE
🏽 🗿 Start 🛛 🖉 🤌 🖾 🐨 📴 🕲 🖙 谢 🕲 quan 🖂 appl 🕲 PML 💦 Roto 🛛 🐯 🥵 🌾 🌄 💭 🌮 💋 🖄 10:14 Ab
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.g. Programming the *RotorGene*[™] 6000

The *RotorGene*[™] 6000 PCR program for the detection of PML/RARa can be divided into following steps:

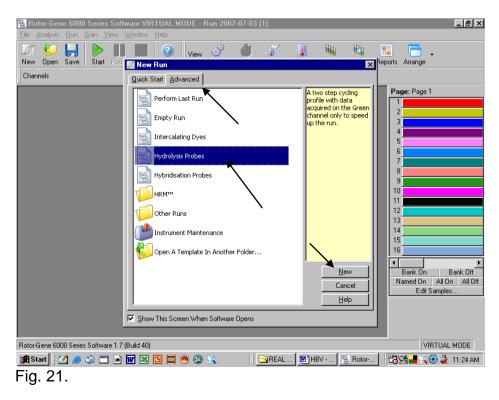
- H. Setting of general assay parameters & reaction volume
- I. Thermal Profile & Calibration
- J. Synthesis of cDNA.
- K. Cycling profile/ Initial activation of the Hot Start enzyme
- L. Cycling for Amplification of cDNA
- M. Adjustment of the sensitivity of the fluorescence channels
- N. Starting of the Rotor Gene™ run

Program the *RotorGene*TM 6000 for these 5 steps according to the parameters shown in Figures 21-38 below All specifications refer to the *RotorGene*TM 6000 software version 1.7 Please find further information on programming the *RotorGene*TM in the *RotorGene*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.



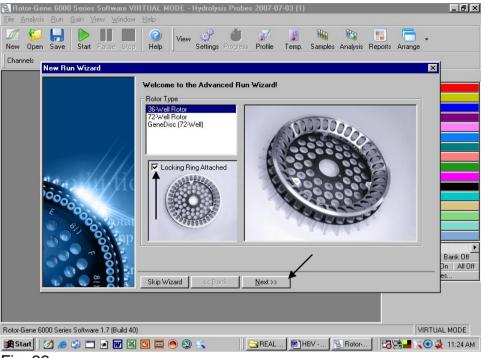


Fig. 22.

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.

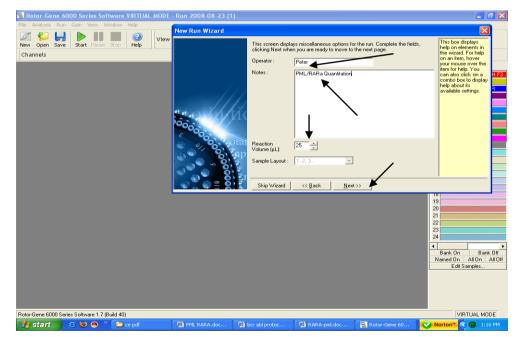


Fig. 23.

- 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.

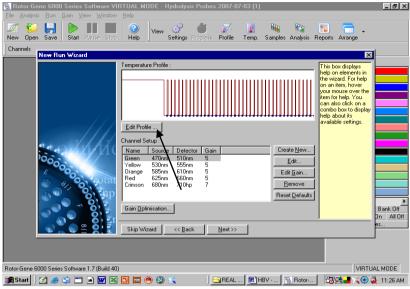
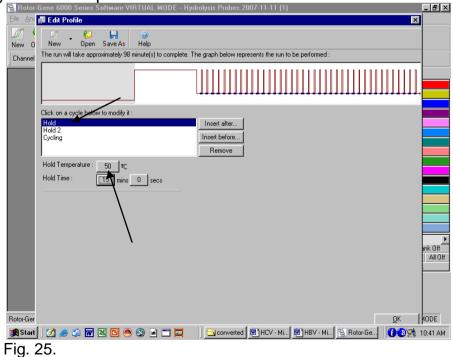


Fig. 24.

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

i) CYCLING PROFILE: First hold 50°C for 15 minutes i.e. cDNA synthesis step as below



Second hold 95°C for 10 minutes as below

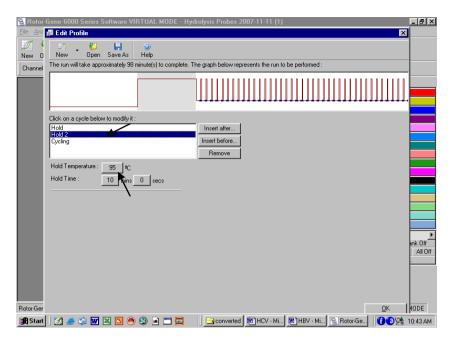


Fig. 26. Initial activation of the Hot Start enzyme. Generally the window will open with the Hold Temp as 95°C and the Hold Time as 10 minutes. In case it is different then set the same to as shown above.

j) Setting the Cycling and acquisition.

When clicked on Cycling the window will open as below.

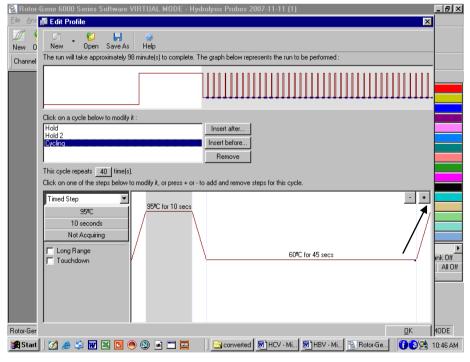


Fig. 27.

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.

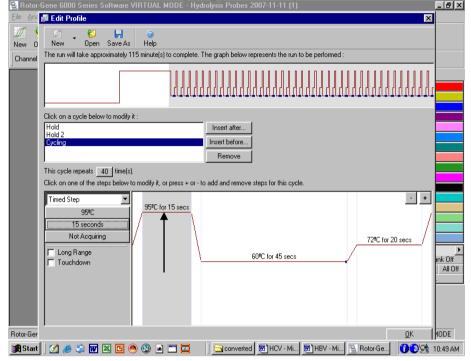


Fig. 28.

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

Seconds

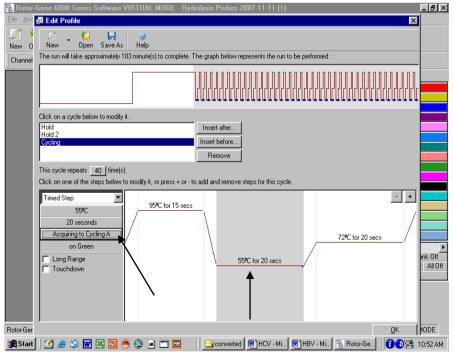


Fig. 29.

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

	Def	ining	the	Data	acquiring channel i.e Green (FA	AM)	
😫 Rotor-	Gene 6000 Serie	es Softwa	re VIRTL	JAL MODE	- Hydrolysis Probes 2007-11-11 (1)		_ B ×
<u>File Ana</u>	🛃 Edit Profile						×
Mew D	New Op	5 🛃 Den Save.	As He				
Channel	The run will take	Acquisitio	n				
		Same as P	revious :	(New Acqui	sition)		
		- Acquisitio	on Configu	ration :			
		Available	Channels	:	Acquiring Channels :		
	<u> </u>	Name			> Name		
	Click on a cycle t	Crimson Orange			Green		
	Hold Hold 2	Red			~~		
	Cycling	Yellow					
	This cycle repeat				ct it from the list in the left and click >. To stop acquiring from a nd list and click <. To remove all acquisitions, click <<.		
	Click on one of th				· · · · · · · · · · · · · · · · · · ·		
	Timed Step						
	55º	Dye Char	t>>		<u> </u>		
	20 sec	Dye Char	nnel Sele	ection Cha	rt		
	Acquiring to	Channel	Source	Detector	Dyes	/	
	on Gr	Green	470nm	510nm	FAM [®] , SYBR Green 1 [®] , Fluorescein, EvaGreen [®] , Alexa Fluor 488 [®]	secs /	
	Long Range	Yellow	530nm	555nm	JOE ⁽¹⁾ , VIC ¹⁾ , HEX, TET ⁽¹⁾ , CAL Fluor Gold 540 ⁽¹⁾ , Yakima Yellow ⁽¹⁾		ank Off
	Touchdown	Orange	585nm	610nm	ROX ¹ , CAL Fluor Red 610 ¹ , Cy3.5 ¹ , Texas Red ¹ , Alexa Fluor 568 ¹		All Off
		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⁾		
						1	
Rotor-Ger						<u>0</u> K	HODE
🄀 Start] 🧭 🧶 🚺	V 🛛 🖸	0	🖻 🗖 🕻	🔤 🔢 🔄 converted 📓 HCV - Mi 👼 HBV - Mi 😒 Rotor-Ge		👌 10:54 AM

Fig. 30.

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.

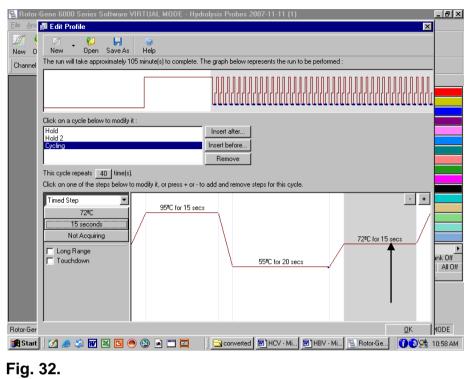
强 Rotor-Gene 6000 Series Software VIR	UAL MODE - Run 2009-01-27 (1)	- 8 🗙			
File Analysis Run Gain View Window Help					
Image: Start Palse Stop Heb View Image: Start Palse Stop Heb View Image: Start Palse Stop Heb Settings Progress					
Channels		X			
		Page: Page 1			
	en Save As Heln	ST RCR ARL WITH OLD			
The run will take	Acquisition	2 1845 II RNA 2UL 3 1040 PNA			
	Same as Previous : (New Acquisition)	4 aSHA HIX BNA			
	- Acquisition Configuration :	5 hEMENT MIX			
	Available Channels : Acquiring Channels :	6 mOHINDER 7 vijki			
	Name > Name -	8 BUENDER			
Click on a cycle	Crimson Green	9 oLD RNA			
Hold 2	Red <<	10 nC 11 st set with OLE FITM			
Cycling	Tellow	12 1845 II RNA 2UL			
	To acquire from a channel, select it from the list in the left and click >. To stop acquiring from a	13 1840 RNA			
This cycle repea	channel, select it in the right-hand list and click <. To remove all acquisitions, click <<.	14 aSHA MIX RNA 15 bEMENT MIX			
Click on one of t		16 MOHINDER			
Timed Step	Dye Chart >> Don't Acquire Help	- + 17 VUAY			
55		18 BJENDER 19 OLD RNA			
20 sec	Channel Source Detector Dues	20			
Acquiring to on G	Green 470nm 510nm FAM ¹⁰ , SYBB Green 1 ¹⁰ , Fluorescein, EvaGreen ¹⁰ , Alexa Fluor 488 ¹⁰	ecs 21			
Long Bang		22			
Touchdowr		24			
	Red 625nm 660nm Cy5 ⁽¹⁾ , Quasar 670 ⁽¹⁾ , LightCycler Red640 ⁽¹⁾ , Alexa Fluor 633 ⁽¹⁾	•			
	Crimson 680nm 710hp Quasar705 ^{,0} , LightCycler Red705 ^{,0} , Alexa Fluor 680 ^{,0}	Bank On Bank Off Named On All On All Off			
		Edit Samples			
1					
	_	<u>0</u> K			
Rotor-Gene 6000 Series Software 1.7 (Build 40)		VIRTUAL MODE			
🎒 start 👘 🧉 😂 🍣 🦈 🕅 🛙	Microsoft Office 🔹 🍓 Inbox for dharam@d 🚞 NOT TO BE DONE 🛛 👹 Rotor-Gene	6000 Seri 🔇 🖄 💭 🦻 5:56 PM			

Fig. 31.

Once the 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 15

Seconds



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

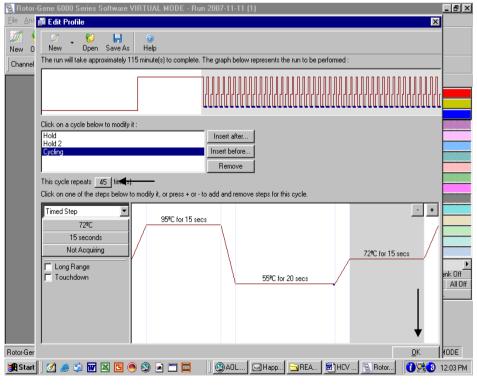


Fig. 33.

After setting the number of Cycles Press OK.

M) Setting the gains for the acquiring channel by clicking at the Gain Optimisation

S Rotor-Ge File Analysi		RTUAL MODE - Run 2007-11-11 (1) Help	_ # ×
Mew Oper		Image: Constraint of the second sec	
Channels	New Run Wizard	×	1
-	New Hun Wizard	Temperature Profile : This box displays	
		Edit Profile Periodic State Edit Profile Edit Profile Channel Setup : Create New Read \$25rm Yellow \$30rm Source Detector Green 470rm State Edit Palow \$30rm Yellow \$30rm State Edit Gain Red \$25rm Crimson \$60rm Yellow \$30rm Yellow \$30rm State \$25rm Times \$25rm Born \$10rm State \$25rm Born \$25rm State \$25rm State \$25rm State \$25rm State \$25rm S	Bank Off Dr. All Off
		Skip Wizard << Back Next >>	es
Rotor-Gene 6	000 Series Software 1.7 (Build 40)		AL MODE
🚮 Start	🧭 🖻 🔛 🔛 🍮	🕲 🖬 🚍 🔢 🔯 AOL 🖾 Happ 🔄 REA 🕲 HCV 😒 Rotor 🚺 😯) 12:05 PM

button as shown below.

Fig. 34

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.35.

<complex-block> Reverse for interverse wirder Wirder interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse for interverse wirder Reverse f</complex-block>	🙀 Rotor-Gene 6000 Series Software VIRTUAL MODE - Run 2009-01-27 (1)	- B X
New Open Save Start Pacie Start Pacie Progress Progre	File Analysis Run Gain View Window Help	
Image: State Stat		
New Runt Optimisation Setup Image: Set Works (200) Image: Set Works (200) Added to the function of the set of of t	Channels	
Auto-Gan Optimization Sotup Image: Control optimization will read the fluoresence on the instead sample at control optimization will read the fluoresence on two is an acceptable. The areas of fluorese		Page: Page 1
New Run W Optimization With Run W Optimization	Lutz Cain Ontinination Satur	
	Optimisation: Auto Gan Optimisation will need the fluxescence on the insented sample at Auto Gan Optimisation will need the fluxescence scenes looking for depends on the chemistry you are performing. Set temperature to Compare the fluxescence on the insented sample at a Optimise Automatic Automa	A daglages default as d
	Rotor-Gene 6000 Series Software 1.7 (Build 40)	VIRTUAL MODE
		Gene 6000 Seri 🔇 🖄 🛱 🦻 💈 5:57 PM

Fig. 35.

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 before 1st Acquisition.
- Just see that below the channel settings there appear only one channels i.e. Green. In case it is different then the channels have not been set properly. Close the window and reset the channels.
 Then Press Close.

The press Next as shown below.	
Rotor-Gene 6000 Series Software VIRTUAL MODE - Run 2007-11-11 (1)	_ @ ×
File Analysis Bun Gain View Window Help	
New Open Save Start Parse Stop Help View 3 to 12 Temp. Samples Ar	halysis Reports Arrange
Channels New Run Wizard	x
Temperature Profile :	This box displays
Edit Profile Edit Profile Channel Setup : Image Source Detector Gain Green 470tm 510tm 5 Yellow 550tm 55 Drange 585m 610tm 5 Bed 625m 660tm 5 Chinson 680tm 710tp 7 Reset D	cin
Gain Optimisation]	Bank Off
Skip Wizard <u>Keack</u>	
Rotor-Gene 6000 Series Software 1.7 (Build 40)	VIRTUAL MODE
🏽 🕅 Start 🛛 💋 🍮 🗐 🔣 🖸 🕘 🕲 🖬 💳 🧮 🕼 🕲 AOL 🖾 Happ 🔄 REA 👼 Hi	CV 🖹 Rotor 👔 🐏 🕄 12:09 PM
Fig. 36.	

N) PRESS Start RUN

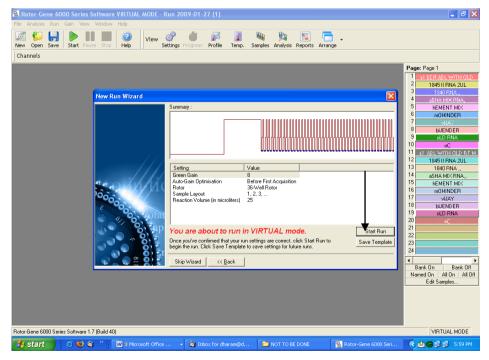


Fig. 37.

2 🗙 Save in: 🔯 Desktop • 🗕 🖆 📰 • My Documents 🚞 genome 1 Crecent run files Ò Genome Export S My Computer My Recent genomediagnostics 👝 s.o.pii Rotor-Gene Demonstrations HBV GENOTYPING safdar iung genes importnat files ARTICLES AND PAPERS -kana aq auns for literature 600 ce.co.in. for loading latest docs i ce... cordett documents etc esurgicalsCD GaluraV GAURAV GalurAV 2 Genome latest docs norway olfert photos poland presentations professional qc reports recent files Shortcut to New F 10 Before Run At 72 Degrees 36-Well Rotor 1, 2, 3, ... 25 My Documents olitore) My Corr to run in VIRTUAL mode. Start Run ed that your run settings are correct, click Start Run to Save Template to save settings for future runs. Save Template PML/RARa quantitation File name: • Save < Back Save as type Run File (".rex) • Cancel 24 Bank Off All On All Off Bank On Named On All Un Edit Samples. VIRTUAL MODE tor-Gene 6000 Series Software 1.7 (Build 40) 🦺 start 🔰 🧉 🙂 🔍 M PML RARA bcr abl prof **1** Rotor-Gene 60. 1:22 PM 📄 ce pd

Saving the RUN File.

Fig. 38.

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. a) Generated Data Interpretation & Analysis for RotorGene 2000/3000

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

The following results are possible:

Important Note:

Please turn off rest of the tubes while reading the results of Bcr-1 including other standards as well. Only the Standards & the samples run in the Bcr-1 tubes should be read. Similar procedure should be followed for BCR-3 & Abl while reading the results.

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

The result of the analysis is positive: The sample contains PML Bcr-1/RARa cDNA or PML Bcr-3/RARa cDNA depending upon the tubes it has amplified. Please look for the results in respective Bcr-1 or Bcr-3 tubes.

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

No PML/RARa cDNA detectable. It can be considered negative if there is amplification in the abl run. Please refer for details on PCR inhibition in the section PCR inhibition.

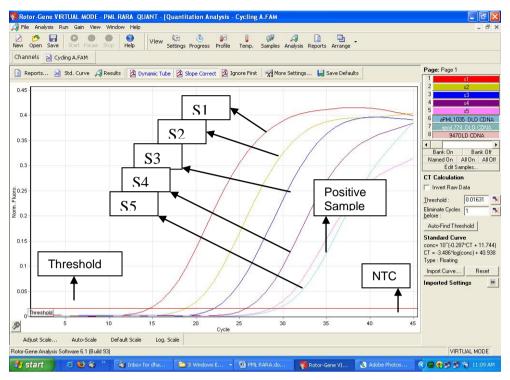


Fig. 39: Detection of the quantitation standards in fluorescence channel Cycling A.FAM. NTC: non-template control.

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

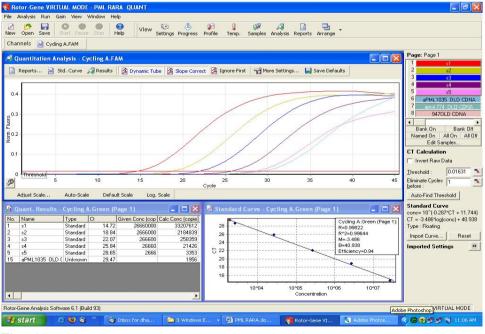


Fig. 40.

Example of analysed data for PML/RARa 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.

Abl reference gene amplification.

Run the diluted samples for abl stored at -20 °C in exactly the similar fashion as described for PML/RARa so far. After the run the quantitative results for the abl gene can be obtained & then the ratio of PMLRARa/ABL can be determined from the Values obtained. These can also be run simultaneously in the same run as Bcr in case there is additional space in the Rotor to do so.

PCR Inhibition

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

In fluorescence channel Cycling A. FAM in **abl** no signal is detected.

A possible PCR inhibition has occurred.

9. b) Generated Data Interpretation & Analysis for RotorGene 6000

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

The following results are possible:

Important Note:

Please turn off rest of the tubes while reading the results of Bcr-1 including other standards as well. Only the Standards & the samples run in the Bcr-1 tubes should be read. Similar procedure should be followed for BCR-3 & Abl while reading the results.

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

The result of the analysis is positive: The sample contains PML Bcr-1/RARa cDNA or PML Bcr-3/RARa cDNA depending upon the tubes it has amplified. Please look for the results in respective Bcr-1 or Bcr-3 tubes.

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

No PML/RARa cDNA detectable. It can be considered negative if there is amplification in the abl run. Please refer for details on PCR inhibition in the section PCR inhibition.

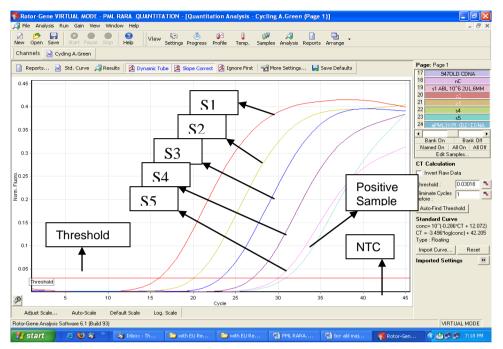


Fig. 41: Detection of the quantitation standards in fluorescence channel Cycling A.Green. NTC: non-template control.

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

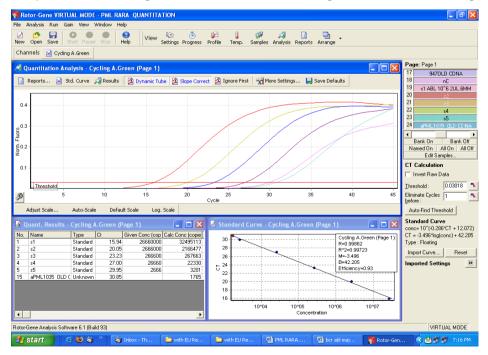


Fig. 42.

Example of analysed data for PML/RARa 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.

Abl reference gene amplification.

Run the diluted samples for abl stored at -20 °C in exactly the similar fashion as described for PML/RARa so far. After the run the quantitative results for the abl gene can be obtained & then the ratio of PMLRARa/ABL can be determined from the Values obtained. These can also be run simultaneously in the same run as Bcr in case there is additional space in the Rotor to do so.

PCR Inhibition

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

In fluorescence channel Cycling A. Green in **abl** no signal is detected.

A possible PCR inhibition has occurred.

10. a) Troubleshooting for RotorGene 2000/3000

1. No signal with positive Standards (PML Bcr-1/, PML Bcr-3/, abl 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 Super Mix *R1*, *R2*, *R3* has been thawed and frozen too often.
- The Super Mix R1, R2, R3 has been kept at +4°C for longer than 5 hours.

→ Please mind the storage conditions given in the **Storage**.

→ Repeat The steps with using a new PML/RARa Super mix R1, R2 & abl R3 super Master Mix.

The PCR was inhibited.

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

10. b) Troubleshooting for RotorGene 6000

1. No signal with positive Standards (PML Bcr-1/, PML Bcr-3/, abl 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 Super Mix *R1*, *R2*, *R3* has been thawed and frozen too often.
- The Super Mix R1, R2, R3 has been kept at +4°C for longer than 5 hours.
 - → Please mind the storage conditions given in the **Storage**.
 - → Repeat The steps with using a new PML/RARa Super mix R1, R2 & abl R3 super Master Mix.

• The PCR was inhibited.

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

11. Specifications

11.a Sensitivity and Reproducibility

In order to determine the sensitivity of the **Geno-Sen's** *PML/RARa Real Time PCR Kit*, a dilution series has been set up from 10^7 down to 10^0 Copies/µl of PML/RARa cDNA and analyzed with the **Geno-Sen's** *PML/RARa Real Time PCR Kit*. The assays were carried out on three different days in the form of 8-fold determinations. The results were determined by a probit analysis. The detection limit as determined for **Geno-Sen's** *PML/RARa Real Time PCR Kit* is consistently 140 copies/ml. This means that there is 95% probability that 140 copies/ml will be detected.

Analytical Sensitivity

Analytical Sensitivity in Conjunction with the *Geno Sen's*[®] Whole Blood RNA Extraction Mini Kit — for RNA purification (Cat. No. 98007, 98008) of the *Geno Sen's*[®] *Geno-Sen's PML RARA Real Time PCR RG Kit on ROTOR GENE 3000/6000 was determined by Spiking a known negative blood to a nominal 100 copies/ml. This was subjected to extraction using the Geno-Sen's[®] Whole Blood RNA Extraction Mini Kit — for RNA purification (Cat. No. 98007, 98008) eight times with starting volume of 1.5 ml & elution volume of 40µl.*

All the Eight extractions were then analyzed with the Geno-Sen's[®] PML RARA 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 98 copies/ml.

Hence Analytical Sensitivity in Conjunction with the *Geno Sen's*[®] Whole Blood RNA Extraction Mini Kit — for RNA purification (Cat. No. 98007, 98008) of the *Geno Sen's*[®] *PML RARAReal Time PCR RG was determined to be 100 copies/ml.*

11.b Specificity

In order to check the specificity of the *Geno-Sen's* PML/RARa Real Time PCR kits, different cDNA and plasmids listed below were analyzed with *Geno-Sen's* PML/RARa Real Time PCR Kit. None of these led to a positive signal with the *Geno-Sen's* PML/RARa Real Time PCR kit. Gene sequence analysis of the amplified region of PML/RARa shows a pronounced homology among the various PML/RARa cDNA, and no homology with other cDNA. Besides which utmost care has been taken in selection of the primers & probes being used in the kit.

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

TNF α
IFN γ
Bcr Major
Bcr Minor

h-tert

In order to validate the Specificity further with wet lab testing the following Known clinical samples were analyzed using the *Geno Sen's*[®] PML/RARa Real Time PCR RG kit on ROTOR GENE 3000/6000 machine. The extraction was carried out with the QIAamp RNA Blood Mini extraction Kit (Cat. No. 52304)

The run was carried out with the known set of standards in order to quantitate the PML/RARa RNA Gene.

Sample Type	WHOLE BLOOD	CELL CULTURE
High +ve's	6	7
Medium +ve's	6	5
Low +ve's	9	4
Extremely low +ve's	4	3
Negative samples.	4	4
	29	23

All the above samples were correctly identified by the **Geno Sen's**[®] PML/RARa Real Time PCR RG kit & all the 7 extremely low samples were accurately detected by the **Geno Sen's**[®] PML/RARa Real Time PCR RG kit & exhibited copies around 140 copies /ml or less than 140 copies/ml.

Further studies are underway on this aspect.

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 RNA depends on the number of RNA 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 dharam@vsnl.com OR pbpl@vsnl.net.

14. 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



EMERGO EUROPE Prinsessegracht 20, 2514 AP , The Hague The Netherlands