RealMOD[™] Green W² 2x qPCR mix

For Real-time quantitative PCR

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RUO
       Research Use Only
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INTRODUCTION

Real-time PCR (gPCR) is the preferred method for DNA and cDNA guantification because of i ts high sensitivity, reproducibility and wide dynamic range. Real-time detection of PCR produc ts makes it possible to include the reaction of fluorescent molecule that reports an increase i n the amount of DNA with a proportional increase in fluorescent signal. The fluorescent chem istries employed for this purpose include DNA- binding dyes and fluorescently labeled sequen ce-specific primers or probes. RealMOD[™] Green W² 2x qPCR mix is an optimized ready-to-use solution for real-time quantitative PCR assays, incorporating SYBR Green I dye. It comprises all the components necessary to perform qPCR: Taq DNA Polymerase, ultrapure dNTPs, MgCl and SYBR Green I dve. The user simply needs to add water, template and primers. Hot start DNA Polymerase is activated by a 5 minutes incubation step at 95°C. This prevents extensio n of nonspecifically annealed primers and primer dimers formed at low temperatures during q PCR setup. The kit includes the components necessary for performing PCR amplification, and has been successfully used to amplify and detect a variety of DNA targets such as genomic DNA, cDNA and plasmidDNA.

KIT CONTENTS	
Label	Volume
RealMOD [™] Green W ² 2x qPCR mix	1 ml

t Spin down before use

- STORAGE AND STABILITY
- Storage condition : Store the product at -20 °C
- RealMOD[™] Green W² 2x qPCR mix are light-sensitive; avoid prolonged exposure to intense I iaht.
- Expiration date : The solution is stable for 1 year from the date of shipping when stored an d handled properly.

WIDE INSTRUMENT COMPATIBILITY

RealMOD[™] Green W² 2x qPCR mix is designed for use with standard cycling mode on standar d and fast qPCR platforms regardless of requirements in ROX. Our product is compatible with

- Applied BioSystems : Quant Studio[™] 12K Flex, ViiA[™] 7, 7900HT, 7500, 7700, StepOne[™] & StepOnePlus™
- Stratagene : MX3000P™, MX3005™
- Bio-Rad : CFX96™ & CFX384™, iQ™5 & MyiQ™, Chromo4™, Opticon® 2 & MiniOpticon®
- Qiagen : Rotor-Gene® Q, Rotor-Gene® 6000
- Eppendorf: Mastercycler®: ep realplex2 & ep realplex4
- Illumina: The Eco™
- Roche : LightCycler® 480

PRODUCT WARRANTY AND SATISFACTIONGUARANTEE

At iNtRON we pride ourselves on the quality and availability of our technical support. Our CRT center is staffed by experienced scientists with extensive practical and theoretical expertise i n molecular biology and the use of iNtRON products. If a iNtRON product does not meet your expectations, simply call your local distributor. If you have questions about product specifica tions or performance, please call iNtRON Technical Services or your local distributor.

NOTICE BEFORE USE

The RealMOD[™] Green W² 2x qPCR mix is intended for research use only. This product is not i ntended for the diagnosis, prevention, or treatment of disease. All due care and attention sho uld be exercised in the handling of the products. Do not use internally or externally in humans or animals. Please observe general labolatory precaution and utilize safety while using this kit

APPLICATIONS

REF

- Real-time PCR : Detection and quantification of DNA and cDNA targets •
- Gene expression profiling : Gene knockdownverification
- · Microbial detection : Viral load determination
- · Array validation : SNP genotyping

PROTOCOL

- 1. Thaw the RealMOD[™] Green W² 2x qPCR mix, template DNA, primers and DNase/RNase fr ee Water on ice. Mix each solutionwell.
- 2. Mix the reaction mix thoroughly, and centrifuge briefly to collect solutions at the bottom of PCR tubes or plates, and then store on ice protected from light.

Reagent	20 µl Reaction	50 µl Reaction	Final Concentration
RealMOD [™] Green W ² 2x qPCRmix	10 µl	25 µl	1X
Forward Primer (10 µM)	0.2 - 2.0 µl	0.5 - 5.0 µl	0.1 - 1.0 µM
Reverse Primer (10 µM)	0.2 - 2.0 µl	0.5 – 5.0 µl	0.1 - 1.0 µM
Template DNA	Variable*	Variable*	Variable*
DNase/RNase free Water	Up to 20 µl	Up to 50 µl	-

* Concentration of cDNA : 0.1 pg/µl - 10 ng/µl / Plasmid DNA < 50 ng / gDNA : 500 - 1000 ng/µl

3. Perform qPCR reactions using the following cycling program :

qPCR Steps	Temp.	Time	Cycle(s)
Initial activation*	95℃	10min*	1
Denaturation	95℃	20 sec	
Annealing	50℃ - 65℃	40 sec	25 - 40
Elongation	72 ℃	60 sec./kb	
Final Extension	72 ℃	5min.	1
Melting curve	Refer to specific guidelines for instrument used		

* To activate the polymerase, include an incubation step at 95 °C for 10 minutes at the beginning of the qPCR cvcle

4. Place the PCR tubes or plates in the Real-time cycler, and start the cycling program.

5. After the reaction is completed, perform analysis.

GENERAL CONSIDERATION

1. Primer design guidelines

The specific amplification, yield and overall efficiency of any Real-time PCR can be criticall y affected by the sequence and concentration of the primers, as well as by the amplicon le ngth. We strongly recommend taking the following points into consideration when designi ng and running your Real-time PCR.

- Use primer-design software, such as Primer3 (http://frodo.wi.mit.edu/primer3/) or vis 1) ual OMPTM (http://dnasoftware.com/).
- GC contents should be between 30% and 80% (ideally 40-60%). 2)
- 3) Avoid runs of identical nucleotides, especially of 3 or more Gs or Cs at the 3' end.
- The Tm should be between 58 $^\circ\!\!\!C$ and 60 $^\circ\!\!\!C$. 4)
- 5) Keep the GC contents in the 30-80% range.
- Avoid runs of identical nucleotides. If repeats are present, there must be fewer than fo 6) ur consecutive G residues.
- 7) Make sure the five nucleotides at the 3' end contain no more than two G and/or C bas es.
- 2. Primer design guidelines
 - It is important to detect the presence of contaminating DNA that may affect the reliability of the data. Always include a no-template control (NTC), replacing the template with PCR grade water.



TERMS USED IN REAL-TIMEPCR	
Term	Definition
Baseline	The initial cycles of Real-time PCR in which there is little or no change in fluorescence signal.
Threshold	A level of Δ Rn - automatically determined (or manually set) by the Real-time PCR system software – used for Ct determination in real time assays. The level is set to be above the baseline and sufficiently low to be within the exponential growth region of the amplification curve. The threshold is the line whose intersection with the amplification plot defines the Ct.
Threshold cycle (Ct)	The fractional cycle number at which the fluorescence passes the threshold value.
Passivereference	A dye that provides an internal fluorescence reference to which the reporter dye signal can be normalized during data analysis. Normalization is necessary to correct for fluorescence fluctuations caused by changes in concentration or in volume. Passive reference can be optionally detected on certain real- time instruments (CCD detector type).
Normalized reporter (Rn)	The ratio of the fluorescence emission intensity of the reporter dye to the fluorescence emission intennsity of the passive reference dye.
Delta Rn (ΔRn)	The magnitude of the signal generated by the specified set of PCR conditions ($\Delta Rn = Rn$ - baseline).

TROUBLE SHOOTING GUIDE

This troubleshooting guide may be helpful in solving problems that may frequently arise. The sci entists at iNtRON are always happy to answer any questions you may have about the information or protocol in this manual or other molecular biology applications.

Problem / Possibl	e cause	Recommendation
No Product, or weak pr	oductsignal in qPCR	
1) Pipetting error or agent 2) No detection activated	including primers, to	tions and storage conditions of the reagents, missing r emplate DNA. Repeat the PCR. ce detection was activated in the cycling
3)Problems with starting t emplate	Check the concentra emplate. If necessar	tion, storage conditions, and quality of the starting t y, make new serial dilutions of template DNA from t tepeat the PCR using the new dilutions. of cycles.
 Insufficient number of c ycles 	Decrease annealing	emperature in steps of 2° C.
5) Annealing temperature toohigh	Increase annealing te	mperature in steps of 2 $^\circ \mathbb{C}$.
6) Annealing temperatur e too low	Reposition the sample	e tubes.
 7) Incorrect setting for sample position. 8) Incorrect setting for data collection 	Confirm the data col	ection setting.
Variation in detection		
 Inappropriate concentrat n of primers 	io • Optimize primer o	concentration according to the instructions.
2) Failure or malfunction of e	Check the device. device	
3) Variation of dispensed • Ir	crease the reaction volu	me. volume
 Inappropriate cycle nditions 	Confirm Tm of the p	imers. co
Poor dynamic range of	CTvalue	
1) Template amount too h igh	Do not exceed maxim more than 500ng ten	num recommended amount of template. Do not use nplate.
2) Template amount too • In	crease template amount,	if possible. low
Signals in blank reaction	ons	
 Contamination of mplicons or sample DNAs 	 Use fresh PCR gra mix. 	de water. Re-make primer solution and master a
2) Detection of a non- specific amplification	Optimize the prim	er and cycle conditions.
Primer-dimmers and/o	r nonspecific PCRPr	oducts
1) Annealing too low	Temperature increDecrease the amount	ase annealing temperature in increments of $2{}^\circ\!\mathbb{C}.$ Int of primer.
2) To much amount of p rimer		

ORDERING INFORMATION

Product Name	Amount	Cat. No.
ConinTM Tatal DNA Extraction Mini Vit	50 col.	17045
G-spin™ Total DNA Extraction Mini Kit	200 col.	17046
HiSenScript™RH(-) cDNA Synthesis Kit	50 rxn.	25014
G-spin™Genomic DNA Extraction Kit (for Bacteria)	50 col.	17121
G-DEX [™] IIc Genomic DNA Extraction Kit (Cell/Tissue)	300 T	17231
G-DEX [™] IIb Genomic DNA Extraction Kit (For blood)	200 T	17241

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