

FastFire qPCR PreMix (Probe)

For fast, quantitative, specific real-time
PCR using sequence-specific probe

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Cat. no. 4992220/4992221

Kit Contents

Contents	4992220 20 μ l \times 125 rxn	4992221 20 μ l \times 500 rxn
2 \times FastFire qPCR PreMix (Probe)	1.25 ml	4 \times 1.25 ml
50 \times ROX Reference Dye	250 μ l	1 ml
RNase-Free ddH ₂ O	2 \times 1 ml	5 \times 1 ml
Handbook	1	1

Storage

FastFire qPCR PreMix (Probe) can be stored at -30~-15°C for one year. It should be stored immediately upon receipt at -30~-15°C. 2 \times FastFire qPCR PreMix (Probe) and 50 \times ROX Reference Dye should be thawed and then mixed upside down gently to be homogenous before using. If the Reagents have been thawed but not used, it is important to thoroughly mix prior to re-freezing. (The layering of salts during the thawing process and subsequent crystallization during freezing will damage the enzyme and decrease product performance.) The reagents could be stored for up to 3 months at 2-8°C if frequently used. Please avoid refreezing and thawing repeatedly.

Introduction

FastFire qPCR PreMix (Probe) is designed for probe based quantitative PCR assays, enables fast and specific quantitative results with total running time 60% reduced compared with regular real-time PCR.

FastFire qPCR PreMix is ready to use with premix including hot-start DNA Polymerase and unique PCR buffer. It could ensure a sensitive PCR detection on any Real-Time PCR thermal cyclers.

Important Notes

1. The hot start DNA polymerase is recommended to be activated at the start of reaction: for cDNA template by a 30-60 sec, 95°C incubation step; for gDNA template by 98°C, 1-2 min.
2. The performance will decrease if the reagents are not mixed thoroughly. Gently mix the reagents by inverting the tubes and centrifuge briefly prior to use. DO NOT vortex and avoid producing bubble.
3. Fluorescence probe is not included in this kit.
4. 0.2 μM concentration of probe, 0.3 μM concentration of primer could achieve excellent amplification result for most systems. If need further optimization, primer concentration could be adjusted between 0.05-0.9 μM , and probe concentration between 0.1-0.5 μM .
5. In a 20 μl reaction volume, the amounts of genome DNA or cDNA template is usually less than 100 ng. The reverse transcription product, if used as template, should not comprise more than 20% of the total PCR reaction volume.

Protocol

<1> Set up the Real-Time reaction

1. Thaw FastFire qPCR PreMix (if stored at -30~-15°C), ROX Reference Dye, template, primers, probe and RNase-free ddH₂O. Completely mix and equilibrate all the reagents to room temperature before use.
2. Prepare a reaction solution according to the following table. All the steps should be operated on ice.

Component	50 μ l volume	25 μ l volume	20 μ l volume	Final concentration
2 \times FastFire qPCR PreMix	25 μ l	12.5 μ l	10 μ l	1 \times
Forward Primer (10 μ M)	1.5 μ l	0.75 μ l	0.6 μ l	0.3 μ M ^{*1}
Reverse Primer (10 μ M)	1.5 μ l	0.75 μ l	0.6 μ l	0.3 μ M ^{*1}
fluorescence probe (10 μ M)	1.0 μ l	0.5 μ l	0.4 μ l	0.2 μ M ^{*2}
DNA template	-	-	-	\leq 200 ng
50 \times ROX Reference Dye ^{*3}	-	-	-	-
RNase-Free ddH ₂ O	Up to 50 μ l	Up to 25 μ l	Up to 20 μ l	-

^{*1} A final primer concentration of 0.3 μ M is optimal for most applications. Higher concentration can be used when the amplification efficiency is not favorable. If non-specific amplification is observed, the primer concentration should be decreased. For further optimization, a primer titration from 0.05 μ M to 0.9 μ M can be performed.

^{*2} The optimal concentration of probe is related with Real-time PCR instruments, probe form and fluorescence dyes according to the information provided in handbook with the instruments or fluorescence probe. A final probe concentration of 0.2 μ M is optimal for most applications. For further optimization, a probe titration from 0.1 μ M to 0.5 μ M can be performed.

^{*3} The optimal concentration of ROX Reference Dye for commonly used Real-Time PCR instruments is as below:

Instrument	Final Concentration
ABI PRISM 7000/7300/7700/7900HT/ Step One etc.	5 \times (e.g. 5 μ l ROX/ 50 μ l volume)
ABI 7500, 7500 Fast; Stratagene Mx3000P, Mx3005P and Mx4000 etc.	1 \times (e.g. 1 μ l ROX/ 50 μ l volume)
Instruments of Roche, Bio-Rad and Eppendorf etc.	No need

<2> Real-Time Amplification

Typically, optimal results are obtained using a two-step PCR. Three-step PCR could be adopted because low copies of template, etc. cause the low amplification efficiency.

Two-step PCR procedure

Stage	Cycle	Temperature	Time	Step	Signal Collection
Initial denaturation	1×	95°C ^{△ 1}	1 min	Initial denaturation	N
PCR	40×	95°C ^{△ 1}	5 sec	Denaturation	N
		60°C ^{△ 2}	15 sec ^{△ 3}	Annealing/ Extension	Y

Three-step PCR procedure

Stage	Cycle	Temperature	Time	Step	Signal Collection
Initial denaturation	1×	95°C ^{△ 1}	1 min	Initial denaturation	N
PCR	40×	95°C ^{△ 1}	5 sec	denaturation	N
		50-60°C ^{△ 4}	10 sec	Annealing	N
		72°C	15 sec ^{△ 3}	Extension	Y

^{△ 1} the initial denaturation temperature is 95°C for cDNA template and 98°C for gDNA.

^{△ 2} 60°C, 15 sec is optimal for most application. If for further optimization, please try 56-66°C.

^{△ 3} Set time should be referred to the manual of specific instrument. The optimal annealing and extended time for commonly used Real-Time PCR instruments is as below:

Roche/ABI 7500 Fast/BioRad/Agilent: 15 sec.
ABI 7900HT/7900HT Fast/ViiA 7/StepOne/StepOnePlus: 20 sec.
ABI 7000/7300: 31 sec.
ABI 7500: 32 sec.

^{△ 4} Annealing temperature of primers is usually 5°C lower than Melting temperature (T_m). The annealing temperature could be increased properly if the base number is low, which could increase the specification. On the contrary, the annealing temperature could be decreased if the base number is high.

3. Close the tubes and mix samples gently. Briefly centrifugation can be performed to collect residual liquid from the walls of the tubes.
4. Place the PCR tubes in the thermal cycler and then start the PCR cycle.

Trouble shooting

1. Chaos of amplification curve and weak fluorescence signal due to low concentration of template.

Comments	Suggestions
Insufficient amount of starting template	Increase the amount of template, if possible. Ensure that sufficient copies of the target nucleic acids are present in your sample.
Competition with primer dimer	Amplification of target DNA will be weakened by amplification of primer dimer. Optimize the PCR thermal cycling or redesign the primers to prevent the formation of primer dimers.
Template loss by adsorption of DNA to the reaction tube	Long-term storage of low concentration of template or diluted template will cause template loss by adsorption to the tube. Please raise the template concentration. PCR reaction should be performed immediately after the sample is diluted.

2. Positive signal in no-template control (NTC)

Comments	Suggestions
Cross contamination	Discard all the components of the assay (e.g., master mix, primers, and probes). Repeat the assay using new components.
PCR instruments configuration error	When using several fluorescence probes together, ensure that the correct detection channel is activated or the correct filter set is chosen for the reporter dye.

3. Poor repeatability of CT value

Comments	Suggestions
Instrument malfunction	Thermo cycling is not suitable or performs in improper temperature or has poor repeatability. Operate and checkup the Real-Time PCR instrument according to the instruction manual.
Contamination of templates	Contamination of templates may lead to the poor linearity.
Long-term stored of diluted template	DNA concentration will decrease due to adsorption to the tube for long term storage. Make new serial dilutions of template from the stock solutions. Repeat the PCR using the newly diluted template immediately.
Poor quality of primers and probes	The batch difference of primers should be avoided. Good quality of primers previously synthesized could be used as the control.
Unsuitable PCR programs, primer concentration or primer sequence	Poor amplification efficiency will cause poor repeatability. Please use optimal primer and probe concentrations and modify the PCR thermal cycling according to the information provided in this handbook. Generally, lower annealing temperature, higher concentration of primers, prolongation of the extension time will help. For high GC content template, please prolong the denature time. If still no improvement, redesign the primers and probes.
Metering inaccuracies	Too small volume of reaction will decrease the detection accuracy. Use the recommended volume in manual and repeat the PCR amplification.

4. Amplification efficiency less than 90% (slope<-3.6)

Comments	Suggestions
Poor quality of primers	Poor quality of primers will decrease the amplification efficiency largely. Make new dilution of primer from the stock solution or re-synthesize the primers.
Non-exclusion of deviation CT	The deviation CT will increase the calculation error. Please Exclude the deviating CT value when calculating the amplification efficiency
Incorrect reaction setup	Optimize the concentration of primers, probes and PCR reaction condition.

5. Amplification efficiency over 110% (slope>-3.1)

Comments	Suggestions
Non-exclusion of deviation CT	The deviation CT will increase the calculation error. Please Exclude the deviating CT value when calculating the amplification efficiency
Impurities in sample	Impurities in sample inhibit the amplification of high concentration of template largely and therefore increase the amplification efficiency. Please lower the concentration of sample or purify the template further.

6. Poor fluorescence signal or zigzag amplification curve

Comments	Suggestions
Incorrect detection filter/channel	The detection spectroscopy differs with different qPCR instruments. Please select the luminophore and quencher according to different instrument models. Please refer to the instruction manual and reset the parameters.
Low fluorescence probe purity	Please select the probe with purity above HPLC grade, otherwise the remaining unbinded fluorescence dye will cause high background signal and affect the sensitivity of detection.
Poor quality of fluorescence probe	Degradation of probe will raise the baseline and decrease the fluorescence signals of amplification products at the same time. Besides, some fluorescence dyes are not suitable for storage in the buffer contains EDTA. Please use the condition recommended by probes synthesizer.
The fluorescence signals collection time is too short	For some instruments, collection time should be prolonged to collect enough fluorescence signals. Please set 45-60 sec for extension time to improve the zigzag amplification curve.