SUPRA SYBR MASTER MIX

Master mix (2X)





SUPRA Sybr Master mix (2x)

Cat. no. EC05

Kit Contents

| Contents | EC05-01 25 μl × 100 rxns | EC05-02 25 μl × 400 rxns |
|---|-----------------------------|-----------------------------|
| 2× Supra Sybr Master Mix (with SYBR Green I) | 1.25 ml | 4 × 1.25 ml |
| 50× ROX Reference Dye | 250 μΙ | 1 ml |
| RNase-free ddH ₂ O | 1 ml | 5 × 1 ml |
| Handbook | 1 | 1 |

Storage

The Supra Sybr Master Mix (SYBR Green) Kit should be stored immediately upon receipt at -20°C, protected from light. Thaw the 2× Supra Sybr Master Mix and 50× ROX Reference Dye and mix thoroughly before use. If the 2× Supra Sybr Master Mix and 50× ROX Reference Dye are thawed but not used, it is important to thoroughly mix them 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. For frequent use, Supra Sybr Master Mix can be stored at 2-8°C for 3 months. Repeated freeze-thaw cycles should be avoided.

Introduction

Supra Sybr Master Mix (SYBR Green) Kit is specially designed to perform Real-time PCR in SYBR Green I fluorescent-based detection assays. The Real-Time PCR reaction buffer, a 2× pre-mixed solution included in this kit, provides an optimum concentration of SYBR Green I solution, which greatly facilitates the preparation of qPCR



reaction mixture. Supra Sybr Master Mix adopts a unique dual hotstart enzymes system (chemically modified HotStar *Taq* DNA polymerase and antibody modified Anti *Taq* DNA Polyerase), which, plus the pre-optimized buffer solution, provides a convenient format for highly sensitive and specific qPCR amplification.

Important Notes

- 1. The initial denaturation conditions must be 95°C 15 min, in order to activate the hot start enzymes.
- 2. Supra Sybr Master Mix includes the SYBR Green I. Store the reagent in dark and avoid direct exposition to strong light during the preparation of PCR reaction mixtures.
- 3. Gently mix the reagents by inverting the tubes and centrifuge briefly prior to use. Do NOT vortex and avoid producing bubble.
- The purity of primers is important for the specificity of PCR.
 Primers purified by PAGE or more superior methods are recommended.
- 5. Typically, best amplification results can be obtained using a primer concentration of 0.3 μ M. However, for individual determination of optimal primer concentration, a primer titration from 0.2 μ M to 0.5 μ M can be performed.
- 6. In a 20 μ l reaction volume, the amounts of genome DNA or cDNA template is usually less than 100 ng. The reverse transcription products, if used as template, should not comprise more than 20% of the total PCR reaction volume.

Protocol

<1> Set up the Real-Time reaction system

Note: 2× Supra Sybr Master Mix and 50× ROX Reference Dye should be stored protected from light.

 Thaw 2× Supra Sybr Master Mix (if stored at -20°C), 50× ROX Reference Dye, template, primers and RNase-free ddH₂O. Completely mix and equilibrate 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 μl volume | 25 μl volume | 20 μl volume | Final concentration |
|--------------------------|-----------------|-----------------|-----------------|----------------------|
| 2× Supra Sybr Master Mix | 25 μΙ | 12.5 μΙ | 10 μΙ | 1× |
| Forward Primer (10 μM) | 1.5 µl | 0.75 μΙ | 0.6 µl | 0.3 μM* ¹ |
| Reverse Primer (10 μM) | 1.5 µl | 0.75 μΙ | 0.6 µl | 0.3 μM* ¹ |
| cDNA template | - | - | - | -ng-pg |
| 50× ROX Reference Dye*2 | - | - | - | - |
| RNase-free ddH₂O | Up to 50 μl | Up to 25 μl | Up to 20 μl | - |

 $^{^{*1}}$ A final primer concentration of 0.3 μM is optimal for most applications. Higher concentration can be used when the amplification efficiency is not favorable. If non-specific amplification is observed, however, the primer concentration should be reduced. For further optimization, a primer titration from 0.2 μM to 0.5 μM can be performed.

^{*&}lt;sup>2</sup> The optimal concentration of ROX Reference Dye for commonly used Real-Time PCR instruments:

| Instrument | Final Concentration |
|--|-------------------------------------|
| ABI PRISM 7000/7300/7700/7900HT | 5× (e.g. 5 μl ROX/ 50 μl volume) |
| ABI 7500, 7500 Fast; Stratagene Mx3000P, Mx3005P and Mx4000 | 1× (e.g. 1 μl ROX/ 50 μl volume) |
| Instruments of Roche, Bio-Rad and Eppendorf | No need to add |

<2> Real-Time Amplification

Typically, best results are obtained using a two-step PCR. However, if two-step PCR does not yield favorable results (e.g. non-specific amplification caused by low template concentration or reduced amplification efficiency induced by low Tm value) the three-step PCR is recommended.



Two-step PCR

| Stage | Cycle | Temperature | Time | Step | Signal Collection |
|----------------------------------|-----------------------|----------------------------|-------------------------|-------------------------|----------------------|
| Initial denaturation | 1× | 95°C | 15min | Initial denaturation | N |
| DCD | 40% | 95°C | 10s | Denaturation | N |
| PCR 40× | 60-66°C* ¹ | 20-32 sec* ³ | Annealing/ Extension | Υ | |
| Melting/Dissociation Curve Stage | | | | | |

Three-step PCR

| Stage | Cycle | Temperature | Time | Step | Signal Collection |
|-------------------------|-------|-------------|--------|-------------------------|----------------------|
| Initial denaturation | 1× | 95°C | 15min | Initial denaturation | N |
| PCR | 40× | 95°C | 10 sec | Denaturation | N |



| | | 50-60°C* ² | 20 sec | Annealing | N |
|----------------------------------|--|-----------------------|----------------------------|-----------|---|
| | | 72°C | 20-32 sec* ³ | Extension | Y |
| Melting/Dissociation Curve Stage | | | | | |

^{*1} An Annealing/Extension temperature of 60°C is optimal for most applications. However, if further optimization is required, temperature from 60°C to 66°C can be performed.

- *2 Normally, annealing temperature would be 5°C lower than primer's Tm value. If primers are relatively short, the annealing temperature can be increased to improve the specificity. Otherwise, the opposite treatment should be taken.
- *3 For a certain Real-Time instrument, the extension time should be set according to its instruction manual. For the guidelines for commonly used instruments please see the list below.

| Roche LightCycler | 20 sec |
|---------------------------|--------|
| ABI 7700/7900HT/7500 Fast | 30 sec |
| ABI 7000/7300 | 31 sec |
| ABI 7500 | 32 sec |

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.

Take ABI 7500 Real Time PCR Instrument as an example. Optimization strategies to improve amplification efficiency in this instrument:

| | Basic program | | Optimized program 1 (extending the elongation time) | Optimized pro (using three-sto | _ |
|-------|---------------|--------|--|-----------------------------------|--------|
| Cycle | Temperature | Time | Time | Temperature | Time |
| 1× | 95°C | 15min | 15min | 95°C | 15min |
| | 95°C | 10 sec | 10 sec | 95°C | 10 sec |
| 40× | 60°C | 32 sec | 32-60 sec | 55°C | 30 sec |
| NA | | | A | 72°C | 32 sec |

Optimization strategy to improve specificity in ABI 7500 Real Time PCR Instrument:

| Basic program | | Optimized p | | |
|---------------|-------------|-------------|-------------|--------|
| Cycle | Temperature | Time | Temperature | Time |
| 1× | 95°C | 15min | 95°C | 15min |
| 40× | 95°C | 10 sec | 95°C | 10 sec |
| 40× | 60°C | 32 sec | 60-64°C | 32 sec |

Troubleshooting Guide

1. No signal, or signal detected late in PCR, or only primer–dimers detected.

| Comments | Suggestions |
|---|--|
| HotStarTaq DNA Polymerase not activated | Ensure that the cycling program includes the initial denaturation step (15 min at 95°C) to activate the hot-start enzymes. |



| PCR programs or primer concentration not optimal | Use optimal primer concentrations and check for possible degradation of primers. Modify the PCR thermal cycling according to the information provided in this handbook. If necessary, redesign the primers. |
|--|--|
| Problems with starting template | Check the concentration, storage conditions, and quality of the starting template. If necessary, make new serial dilutions of template nucleic acids from the stock solutions. Repeat the PCR using the new dilutions. |

2. High fluorescence in "No Template" control

| Comments | Suggestions |
|-------------------------------------|---|
| Contamination of reagents | Discard reaction components and repeat PCR with new reagents. |
| Contamination during reaction setup | Take appropriate safety precautions (e.g., use filter tips). |
| Primer degradation | Check for possible degradation of primers on a denaturing polyacrylamide gel. |

3. Primer–dimers and/or non-specific PCR products

| Comments | Suggestions |
|--|--|
| Mg ²⁺ concentration not optimal | The Mg ²⁺ concentration provided in 2x Supra Sybr Master Mix is 2 mM. For few targets, an increase up to 5 mM Mg2+ may be helpful. Perform the titration in 0.5 mM steps. |
| Annealing temperature too low | Increase annealing temperature in increments of 2°C. |
| Primer design not optimal | Review primer design. |
| PCR product too long | For optimal results, PCR products should be between 100 and 150 bp. PCR products should not exceed 500 bp. |



| Primers degraded | Check for possible degradation of primers on a denaturing polyacrylamide gel. |
|-----------------------|--|
| Metering inaccuracies | Too small reaction volume may reduce the accuracy of detection. Use the volume recommended in instruction manual and repeat the PCR. |

4. No linearity in ratio of CT value/crossing point to log of the template amount

| Comments | Suggestions |
|--|---|
| Instrument malfunction | Operate the Real-Time PCR instrument according to the instruction manual. |
| Contamination of templates | Contamination of templates may lead to the poor linearity. |
| Long stored dilutions of template | Make new serial dilutions of template nucleic acids from the stock solutions. Repeat the PCR using the new dilutions. |
| PCR programs or primer concentration not optimal | Use optimal primer concentrations and check for possible degradation of primers. Modify the PCR thermal cycling according to the information provided in this handbook. If necessary, redesign the primers. |
| Metering inaccuracies | Too small reaction volume can reduce the accuracy of detection. Use the volume recommended in instruction manuals and repeat the PCR. |