Associated Products

| Product | Description | Pack Size | Cat No. |
|----------------------------------|--|-----------------------------------|-------------------------------------|
| ISOLATE II Genomic DNA Kit | Rapid isolation of high-quality genomic DNA from many different starting material | 10 Preps 50 Preps 250 Preps | BIO-52065 BIO-52066 BIO-52067 |
| ISOLATE II Plant DNA Kit | Rapid isolation of high-quality genomic DNA from a wide variety of plant species | 10 Preps 50 Preps 250 Preps | BIO-52068 BIO-52069 BIO-52070 |
| ISOLATE II RNA Mini Kit | Isolation of high-yield and extremely pure total RNA from a variety of samples | 10 Preps 50 Preps 250 Preps | BIO-52071 BIO-52072 BIO-52073 |
| ISOLATE II RNA Plant Kit | Isolation of high-yield and extremely pure total RNA from a wide variety of plant species | 10 Preps 50 Preps | BIO-52076 BIO-52077 |
| TRIsure™ | Quick isolation of high-quality RNA from a variety of sources for subsequent use in cDNA synthesis | 100 mL 200 mL | BIO-38032 BIO-38033 |
| SensiFAST™ cDNA Synthesis Kit | Fully optimized to generate maximum yields of full-length and low abundance cDNA from RNA | 50 Reactions 250 Reactions | BIO-65053 BIO-65054 |
| Agarose | arose Molecular biology grade agarose | | BIO-41026 BIO-41025 |

TRADEMARK AND LICENSING INFORMATION

1) Trademarks: SensiMix™ (Bioline Reagents Ltd), SYBR[®] (Molecular Probes), ROX™, LightCycler™ (Roche), StepOne™ (ABI), RotorGene™ (Qiagen), LightCycler[®] (Roche). iCycler™ MyiQ™, IQ™ (Bio-Rad).

2) Purchase of this product conveys a licence from Life Technologies to use this SYBR® containing reagent in an end-user RUO assay. Parties wishing to incorporate this SYBR® containing reagent into a downstream kit, should contact Life Technologies for SYBR® Licencing information

Storage and Stability:

The SensiMix SYBR® & Fluorescein Kit is shipped on dry/blue ice. All kit components should be stored at -20°C upon receipt. Excessive freeze/thawing is not recommended

Expiry:

When stored under the recommended conditions and handled correctly, full activity of the kit is retained until the expiry date on the outer box label.

Quality control:

The SensiMix SYBR[®] & Fluorescein and its components are extensively tested for activity, processivity, efficiency, heat activation, sensitivity, absence of nuclease contamination and absence of nucleic acid contamination prior to release

Safety precautions:

Please refer to the material safety data sheet for further information.

Notes:

Research Use Only

Description

The SensiMix[™] SYBR[®] & Fluorescein Kit is a high-performance product designed for superior sensitivity and specificity on real-time instruments, in which a fluorescein passive reference signal optionally used. The SensiMix SYBR[®] & Fluorescein Kit employs a hot-start DNA polymerase, for high PCR specificity and sensitivity. SensiMix SYBR[®] & Fluorescein is inactivated and possesses no polymerase activity during the reaction set-up, preventing non-specific amplification including primer-dimer formation.

For ease-of-use and added convenience SensiMix SYBR[®] & Fluorescein Kit is provided as a 2x mastermix containing all the components necessary for real-time PCR including the SYBR[®] Green I dye, dNTPs, stabilisers and enhancers. As a ready to use premix, only primers and template need to be added.

Kit Components

| Reagent | 250 x 50 μL reactions | 500 x 50 μL reactions | 2000 x 50 µL Rreactions |
|-------------------------------|--------------------------|--------------------------|----------------------------|
| SensiMix™ SYBR [®] & | 5 x 1.25 mL | 10 x 1.25 mL | 40 x 1.25 mL |
| Fluorescein (2X) | (6.25 mL) | (12.5 mL) | (50 mL) |

Kit Compatibility

The 2X SensiMix SYBR® & Fluorescein Kit has been optimized for use in SYBR Green-based real-time PCR on the real-time instruments listed in the following compatibility table, each of these instruments having the capacity to analyze the real-time PCR data with the passive reference signal either on or off. The kit is also compatible with several instruments that do not require the use of fluorescein, such as the BMS Mic, Qiagen (Corbett) Rotor-Gene[™] 6000, Bio-Rad CFX96 or Roche LightCycler[®] 480.

| Manufacturer | Model |
|--------------|------------------------------------|
| Bio-Rad | ICycler [®] , MyiQ™, IQ™5 |

General Considerations

To help prevent any carry-over DNA contamination we recommend that separate areas are maintained for PCR set-up, PCR amplification and any post-PCR gel analysis. It is essential that any amplified PCR product should not be opened in PCR set-up area.

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Primers: the sequence and concentration of primer as well as the amplicon length can be critical for specific amplification, yield and overall efficiency of any real-time PCR. We strongly recommend taking the following into consideration when designing and running your PCR reaction:

- use primer-design software, such as Primer3 or visual OMP[™] (http://frodo.wi.mit.edu/primer3/ and DNA Software, Inc; http:// dnasoftware.com/ respectively). Primers should have a melting temperature (Tm) of approximately 60 °C
- optimal amplicon length should be 50-150 bp
- · a final primer concentration of 250 nM is suitable for most PCR conditions, however to determine the optimal concentration we recommend a primer titration in the range of 0.1–1 μM
- use equimolar primer concentrations
- when amplifying from cDNA use gene-specific primers. If possible use intron-spanning primers to avoid amplification from genomic DNA

Template: it is important that the DNA template is suitable for use in PCR in terms of purity and concentration. Also, the template needs to be devoid of any contaminating PCR inhibitors (e.g. EDTA). The recommended amount of template for PCR is dependent upon the type of DNA used. The following should be considered when using genomic DNA and cDNA templates:

- Genomic DNA: use up to 1 µg of complex (e.g. eukaryotic) genomic DNA in a single PCR. We recommend using the Bioline ISOLATE II Genomic DNA Mini Kit (BIO-52066) for high yield and purity from both prokaryotic and eukaryotic sources
- **cDNA:** the optimal amount of cDNA to use in a single PCR is dependent upon the copy number of the target gene. We suggest using 100 ng cDNA per reaction, however it may be necessary to vary this amount. To perform a two-step RT-PCR, we recommend using the SensiFAST cDNA Synthesis Kit (BIO-65053) for reverse transcription of the purified RNA. For high yield and purity of RNA, use the Bioline ISOLATE II RNA Mini Kit (BIO-52072)

MgCl₂: The MgCl₂ concentration in the 1x reaction mix is 3 mM. In the majority of qPCR conditions this is optimal for both the reverse transcriptase and the hot-start DNA polymerase. If necessary, we suggest titrating the MgCl₂ to a maximum of 5mM.

PCR controls: 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. When performing a two-step RT-PCR, set-up a no RT control as the NTC for the PCR.

Optional Fluorescein well-factor correction: SYBR Fluorescein Kit is premixed with fluorescein, so that fluorescence emitted by fluorescein can be optionally detected on certain real-time instruments. If your real-time instrument has the capability of using fluorescein and you wish to use this option, then this option must be selected by the user in the software.

Procedure

Troubleshooting Guide

Reaction mix composition: Prepare a PCR master mix. The volumes given below are based on a standard 50 µL final reaction mix and can be scaled accordingly.

| Reagent | Volume | Final concentration |
|---|-------------|---------------------|
| 2x SensiMix™ SYBR [®] & Fluorescein | 25 μL | 1x |
| 25 μM Forward Primer | 0.5 μL | 250 nM |
| 25 μM Reverse Primer | 0.5 μL | 250 nM |
| H ₂ 0 | Up to 45 µL | - |
| Template | 5 μL | |
| 50 μL Final volume | | |

Suggested Thermo-cycling conditions

The following PCR conditions are suitable for SensiMix SYBR[®] & Fluorescein Kit with a majority of amplicons and real-time PCR instruments. However, the cycling conditions can be varied to suit customer or machine-specific protocols. The critical step of the PCR is the 10 minute initial activation at 95 °C. The detection channel on the real-time instrument should be set to (SYBR®) Green or FAM.

| Cycles | Temperature | Time | Notes |
|--------|-------------------------|----------------------|---|
| 1 | *95 °C | *10 min | Polymerase activation |
| 40 | 95 ℃ 55-60 ℃ 72 ℃ | 15 s 15 s 15 s | Temp. depends on the Tm of primers Acquire at end of step |

*Non-variable parameter

Optional analysis:

After the reaction has reached completion refer to the instrument instructions for the option of melt-profile analysis.

| Problem | Possible Cause | Recommendation | |
|---------------------------|---|--|--|
| | Activation time too short | Make sure SensiMix is activated for 10 min at 95 °C before cycling | |
| | Error in protocol setup | Verify that correct reagent concentrations, volumes, dilutions and storage conditions have been used | |
| | Suboptimal primer design | Use primer design software or validated primers. Test primers on a control template | |
| No amplification | Incorrect concentration of primers | Use primer concentration between 100 nM and 1 μM | |
| trace AND | Template degraded | Re-isolate your template from the sample material or use freshly prepared template dilution | |
| No product on agarose gel | Primers degraded | Use newly synthesized primers | |
| | Template contaminated with PCR inhibitors | Further dilute template before PCR or purify template and resuspend it in PCR-grade H_2O | |
| | Template concentration too low | Increase concentration used | |
| | Cycling conditions not optimal | Increase extension/annealing times, increase cycle number, reduce annealing temperature | |
| No amplification trace | | | |
| AND | Error in instrument setup | Check that the acquisition settings are correct during cycling | |
| Product on agarose gel | | | |

Troubleshooting Guide (Continued)

| | č | |
|---|--|---|
| Problem | Possible Cause | Recommendatio |
| | Suboptimal primer design | Redesign primers |
| | Primer concentration too high | Test dilution serie products disappea |
| Non-specific amplification product AND | Primer concentration too low | Titrate primers in |
| | Primer annealing temperature too low | Increase PCR and amplification prod |
| Primer-dimers | Template concentration too low | Increase template |
| | Template concentration too high | Reduce template |
| | Extension time too long | Reduce extension |
| | Activation time too short | Ensure the reaction |
| | Annealing temperature too high | Decrease anneali |
| | Extension time too short | Double extension |
| Late | Template concentration too low | Increase concentr |
| amplification trace | Template with high secondary structure | Increase reverse |
| | Template is degraded | Re-isolate templa |
| | Suboptimal design of primers | Redesign primers |
| | Primer concentration too low | Increase concentr |
| DOD | Extension time is too short | Increase extensio |
| PCR efficiency below 90% | Primer concentration too low | Increase concentr |
| 5CIOW 5070 | Suboptimal design of primers | Redesign primers |
| PCR | Template is degraded or contains PCR inhibitors | Re-isolate template template and result |
| efficiency above 110% | Non specific amplification and/or primer dimers | Use melt analysis amplification prod |

Technical Support

If the troubleshooting guide does not solve the difficulty you are experiencing, please contact technical support with details of reaction setup, cycling conditions and relevant data

Email: tech@bioline.com

s using appropriate software or use validated primers

es of primer concentrations until primer dimer/non-specific amplification ear

the concentration range of 100 nM - 1 μ M

nnealing temperature in increments of 2 °C until primer dimer/non-specific ducts disappear

e concentration

concentration until non-specific products disappear

n time to determine whether non-specific products are reduced

ion is activated for 10 min at 95 °C before cycling

ling temperature in steps of 2 °C

n time to determine whether the cycle threshold (C_T) is affected

tration if possible

transcription reaction time up to 30 min

transcription reaction temperature up to 45 °C

ate from sample material or use freshly prepared template dilution

s using appropriate software or use validated primers

tration of primer in 100 nM increments

on time

tration of primer in 100 nM increments

s using appropriate software or use validated primers

ate from sample material or use freshly prepared template dilution or purify uspend it in H₂O

s and 4% agarose gel electrophoresis to confirm presence of non-specific ducts. See above for preventing/removing non-specific products