### **Associated products**

| Product                  | Description  | Pack Size                         | Cat No.                             |
|--------------------------|--|-----------------------------------|-------------------------------------|
| ISOLATE II RNA Mini Kit  | Fast and efficient isolation of extremely pure total RNA from a variety of samples                 | 10 Preps<br>50 Preps<br>250 Preps | BIO-52071<br>BIO-52072<br>BIO-52073 |
| ISOLATE II Plant RNA Kit | Fast and efficient isolation of extremely pure total RNA from a variety of plant samples           | 10 Preps<br>50 Preps              | BIO-52076<br>BIO-52077              |
| TRIsure™                 | Quick isolation of high-quality RNA from a variety of sources for subsequent use in cDNA synthesis | 100 mL<br>200 mL                  | BIO-38032<br>BIO-38033              |
| Agarose                  | Molecular biology grade agarose  | 100 g<br>500 g                    | BIO-41026<br>BIO-41025              |

### 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: mbi.tech@meridianlifescience.com

Bioline Reagents Ltd UNITED KINGDOM Tel: +44 (0)20 8830 5300 Fax: +44 (0)20 8452 2822

Meridian Life Science Inc. USA

Tel: +1 901.382.8716 Tel: +49 (0)3371 60222 03 Fax: +1 901.382.0027 Fax: +49 (0)3371 60222 01

Bioline GmbH GERMANY

> Tel: +61 (0)2 9209 4180 Fax: +61 (0)2 9209 4763

Bioline (Aust) Pty. Ltd

AUSTRALIA

Website: www.bioline.com/sensifast email: info@meridianlifescience.com

### Storage and stability:

MyTaq<sup>™</sup> One-Step RT-PCR Kit is shipped on dry/blue ice. All kit components should be stored at -20 °C upon receipt. Excessive freeze/thawing is not recommended. Thaw, mix, and briefly centrifuge each component before use

### 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:

MyTaq One-Step RT-PCR Kit and its components are extensively tested for activity, processivity, efficiency, heat activation, sensitivity, absence of nuclease contamination and absence of nucleic acid contamination.

### Safety precautions:

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

### Notes:

For research or further manufacturing use only.

### Trademarks:

MyTaq and TRIsure are trademarks of Bioline Reagents Ltd.

# MyTaq™ One-Step RT-PCR Kit

Shipping: On dry/blue ice Catalog numbers

Batch No.: See vial BIO-65048: 25 Reactions

Concentration: See vial BIO-65049: 100 Reactions

Store at -20 °C



## Description

MyTaq™ One-Step RT-PCR Kit has been formulated for highly reproducible first-strand cDNA synthesis and subsequent PCR in a single tube. A combination of the latest advances in buffer chemistry together with a reverse transcriptase and hot-start DNA polymerase system ensures that MyTaq One-Step RT-PCR Kit produces fast, highly-specific and ultrasensitive one-step RT-PCR, from either total RNA or mRNA using gene-specific primers.

MyTaq One-Step Kit consists of a 2x MyTaq One-Step mix, as well as separate reverse transcriptase and RiboSafe RNase Inhibitor.

### Kit components

| Reagent                            | 25 Reactions | 100 Reactions |
|------------------------------------|--------------|---------------|
| MyTaq One-Step mix (2x)            | 1 x 625 μL   | 2 x 1.25 mL   |
| RiboSafe RNase Inhibitor (10 u/μL) | 1 x 25 μL    | 1 x 100 μL    |
| Reverse transcriptase              | 1 x 12.5 μL  | 1 x 50 μL     |
| DEPC-treated water                 | 1 x 1.8 mL   | 1 x 1.8 mL    |

## General considerations

When handling RNA, it is important to use RNase-free plasticware and reagents. We also recommend performing RNA work in an RNase-free area. To help prevent any carry-over DNA contamination, we recommend that separate areas are maintained for reaction set-up, PCR amplification and any post-PCR gel analysis. It is essential that any tubes containing amplified DNA product are not opened in the reaction set-up area.

**Primers:** The sequence and concentration of the primers, as well as amplicon length, can be critical for specific amplification, yield and overall efficiency of any RT-PCR.

We strongly recommend taking the following points into consideration when designing and running your RT-PCR:

- The use of gene-specific primers is recommended for use with the MyTaq One-Step RT-PCR Kit. The use of oligo dT or random hexamers is not recommended with a One-Step RT-PCR set-up since this can result in the generation of non-specific products.
- use primer-design software, such as Primer3 (<a href="http://frodo.wi.mit.edu/primer3/">http://frodo.wi.mit.edu/primer3/</a>) or visual OMP<sup>TM</sup> (<a href="http://dnasoftware.com/">http://dnasoftware.com/</a>). Primers should have a melting temperature (Tm) of approximately 60 °C
- final primer concentration of 400 nM is suitable for most RT-PCR, however, to determine the optimal concentration we recommend titrating in the range 0.1-1 μM
- · use an equimolar primer concentration

**Template:** It is important that the RNA template is intact and devoid of both DNA and contaminating inhibitors which can effect both reverse transcription and PCR. For high purity RNA, we recommend using the Bioline ISOLATE RNA Mini Kit (BIO-52043). RNA stocks and dilutions should be made in DEPC-treated water to avoid any RNase-mediated degradation.

The recommended amount of template for one-step RT-PCR is dependent upon the type of RNA used. The following should be considered when using total RNA and isolated mRNA:

- **total RNA:** purified total RNA can be used in the range from 1 pg to 1 μg per 50 μL reaction
- mRNA: purified mRNA can be used from 0.01 pg per 50 μL reaction

**MgCl<sub>2</sub>:** The MgCl<sub>2</sub> concentration in the 1x reaction mix is 3 mM. In the majority of RT-PCR conditions, this is optimal for both the reverse transcriptase and the hot-start DNA polymerase. If necessary, we suggest titrating the MgCl<sub>2</sub> to a maximum of 5 mM.

**RT-PCR controls:** It is important to detect the presence of contaminating DNA that may affect the reliability of the data. Always include a no-RT control, by omitting the reverse transcriptase from the reaction.

## Procedure

The following RT-PCR instructions are intended as a guideline for setting-up and running RT-PCR.

**Reaction mix preparation:** Before preparing an RT-PCR master mix, ensure all reagents are mixed thoroughly. The volumes given below are based on a standard 50  $\mu$ L final reaction mix and can be scaled accordingly.

# Kit Components:

| Reagent                  | Volume      | Final concentration |
|--------------------------|-------------|---------------------|
| 2x MyTaq One-Step Mix    | 25.0 μL     | 1x                  |
| Forward Primer (10 µM)   | 2.0 μL      | 400 nM              |
| Reverse Primer (10 µM)   | 2.0 μL      | 400 nM              |
| Reverse transcriptase    | 0.5 μL      | -                   |
| RiboSafe RNase Inhibitor | 1.0 μL      | -                   |
| DEPC-H <sub>2</sub> O    | Up to 45 μL | -                   |
| Template                 | 5 μL        | -                   |
| Final volume             | <b>50</b> μ | L                   |

**Suggested RT-PCR conditions:** The following RT-PCR conditions are suitable for the MyTaq One-Step RT-PCR Kit with the majority of amplicons.

| Cycles | Temperature             | Time                 | Notes                                  |
|--------|-------------------------|----------------------|--|
| 1      | 45 °C                   | 20 min               | Reverse transcription                  |
| 1      | 95 °C                   | 1 min                | Polymerase activation                  |
| 40     | 95 °C<br>60 °C<br>72 °C | 10 s<br>10 s<br>30 s | Denaturation<br>Annealing<br>Extension |

**RT-PCR optimization:** The following optimization may be necessary to improve the efficiency of some reactions:

- The reverse transcription reaction temperature can be increased from 45 °C to 50 °C for high GC-rich templates
- The reverse transcription time can be increased from 20 to 40 minutes for amplicons over 1 kb
- Increase the extension time of the PCR for target sequences longer than 1kb or if running multiplex RT-PCR

# Troubleshooting guide

| Problem                   | Possible Cause                               | Recommendation  |
|---------------------------|--|---|
| No product on agarose gel | Activation time too short                    | Ensure MyTaq One-Step mix is activated for a minimum of 1 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   |
|                           | Incorrect concentration of primers           | Use primer concentrations between 100 nM and 1 μM   |
|                           | Template degraded                            | Re-isolate your template from the sample material or use freshly prepared template dilution. We recommend using the ISOLATE RNA kits for template preparation and DEPC-treated water for resuspension or dilution of the template |
|                           | Template degraded                            | Verify the integrity of RNA using agarose gel electrophoresis   |
|                           |  | Ensure RNase inhibitor is added before addition of template   |
|                           | Primers degraded                             | Do not use old diluted primer stock, use newly synthesized primers and store in TE (pH 8.5) rather than water. Avoid repeated freeze/thawing of stock solution  |
|                           | Template contaminated with RT-PCR inhibitors | Further dilute template before PCR or purify template and resuspend it in DEPC treated water  |
|                           | Template concentration too low               | Increase concentration used   |
|                           | Cycling conditions not optimal               | Increase extension/annealing time, increase cycle number, reduce annealing temperature  |

# Troubleshooting Guide (Continued)

| Problem                            | Possible Cause                       | Recommendation   |
|------------------------------------|--------------------------------------|--|
|                                    | Non-specific priming                 | Use gene-specific primers rather than Oligo dT or random hexamers.   |
|                                    | Inefficient reverse transcription    | Increase the reverse transcription temperature up to 48 °C   |
|                                    | Suboptimal primer design             | Redesign primers using appropriate software, or use validated primers  |
| Non-specific amplification product | Primer concentration too high        | Test dilution series of primer concentrations until primer-dimer/non-specific amplification products disappear         |
| AND / OR                           | Primer concentration too low         | Use primer concentration between 100 nM and 1 μM   |
| Primer-dimers                      | Primer annealing temperature too low | Increase PCR annealing temperature up to 65 °C or until primer-dimer/<br>non-specific amplification products disappear |
|                                    | Template concentration too low       | Increase template concentration  |
|                                    | Template concentration too high      | Reduce template concentration until non-specific products disappear  |
|                                    | Extension time too long              | Reduce extension time to determine whether non-specific products are reduced   |
| Variability<br>between             | Error in reaction set-up             | Prepare large volume master mix  |
| replicates                         | Air bubbles in reaction mix          | Centrifuge reaction samples/plate prior to running RT-PCR  |
|                                    | Inefficient reverse transcription    | Increase the reverse transcription temperature up to 48 °C, and increase the RT incubation time                        |
|                                    | Activation time too short            | Ensure MyTaq One-Step mix is activated for a minimum of 1min at 95 °C before cycling                                   |
|                                    | Annealing temperature too high       | Decrease annealing temperature in steps of 2 °C  |
| Low yield of PCR products          | Extension time too short             | Double extension time to determine whether the yield is affected   |
|                                    | Template concentration too low       | Increase concentration if possible   |
|                                    | Template is degraded                 | Re-isolate template from sample material or use freshly prepared template dilution                                     |
|                                    | Suboptimal primer design             | Redesign primers using appropriate software, or use validated primers  |
|                                    | Primer concentration too low         | Increase concentration of primers in 100 nM increments   |

Website: www.bioline.com/ email: info@meridianlifescience.com