

## SuperScript® III Platinum® One-Step Quantitative RT-PCR System

Cat. no. 11732-020

Cat. no. 11732-088

Size: 100 reactions

Size: 500 reactions

Store at -20°C

### Description

The SuperScript® III Platinum® One-Step Quantitative RT-PCR System is a one-step, quantitative RT-PCR (qRT-PCR) kit for the detection and quantification of RNA using real-time detection instruments. This system combines SuperScript® III Reverse Transcriptase (RT) and Platinum® *Taq* DNA Polymerase in a single enzyme mix. Both cDNA synthesis and PCR are performed in a single tube using gene-specific primers and either total RNA or mRNA. Reagents are provided for 100 or 500 amplification reactions of 50 µL each.

The system enables highly sensitive detection from as few as 10 copies of RNA template, with a broad dynamic range that supports accurate quantification of high-copy mRNA at up to 1 µg of total RNA.

- **SuperScript® III Reverse Transcriptase** is a version of M-MLV RT that has been engineered to reduce RNase H activity and provide increased thermal stability<sup>1</sup>. The enzyme can synthesize cDNA at a temperature range of 42°C –60°C. Because SuperScript® III RT is not significantly inhibited by ribosomal and transfer RNA, it can be used to synthesize cDNA from total RNA.
- **Platinum® *Taq* DNA Polymerase** is recombinant *Taq* DNA polymerase complexed with a proprietary antibody that blocks polymerase activity at ambient temperatures<sup>2,3</sup>. Activity is restored after the denaturation step in PCR cycling, providing an automatic “hot start” in PCR for increased sensitivity, specificity, and yield.
- **2X Reaction Mix** consists of a proprietary buffer system, MgSO<sub>4</sub>, dNTPs, and stabilizers. A tube of 50 mM MgSO<sub>4</sub> is included in the kit for further optimization of the Mg<sup>2+</sup> concentration.

This one-step qRT-PCR kit has been formulated for use with fluorogenic primers (e.g., LUX™ Primers) or fluorogenic probe-based technology (e.g., TaqMan® probes)<sup>4-7</sup>. For one-step qRT-PCR using SYBR® Green I dye, we recommend the SuperScript® III Platinum® SYBR® Green One-Step qRT-PCR System (see **Additional products**, below).

**Note:** This kit has been optimized for real-time qRT-PCR. For end-point RT-PCR, we recommend the SuperScript® III One-Step RT-PCR System with Platinum® *Taq* DNA Polymerase (Catalog nos. 12574-018 and 12574-026).

### Component

	<b>100 reaction kit</b>	<b>500 reaction kit</b>
SuperScript® III RT/Platinum® <i>Taq</i> Mix	100 µL	500 µL
2X Reaction Mix (a buffer containing 0.4 mM of each dNTP and 6 mM MgSO <sub>4</sub> )	2 × 1.25 mL	12.5 mL
50 mM Magnesium Sulfate (MgSO <sub>4</sub> )	1 mL	2 × 1 mL
ROX™ Reference Dye (25 µM)	100 µL	500 µL

### Storage

Store components at -20°C. Stability can be extended by storing at -80°C. ROX™ Reference Dye must be stored in the dark.

### Additional products

The following products are also available. Visit [www.lifetechnologies.com](http://www.lifetechnologies.com)

Product	Amount	Catalog no.
SuperScript® III Platinum® One-Step Quantitative RT-PCR System with ROX™ Reference Dye	100 reactions	11745-100
	500 reactions	11745-500
SuperScript® III Platinum® SYBR® Green One-Step qPCR System	100 reactions	11736-051
	500 reactions	11736-059
PureLink® Micro-to-Midi™ Total RNA Purification System	50 reactions	12183-018
TRIzol® Reagent	100 mL	15596-026
	200 mL	15596-018
RNaseOUT™ Recombinant Ribonuclease Inhibitor	5000 units	10777-019
DNase I, Amplification Grade	100 units	18068 015
LUX™ Fluorogenic Primers		visit <a href="http://www.lifetechnologies.com">www.lifetechnologies.com</a>

## Recommendations and guidelines for One-Step qRT-PCR

### Instrument compatibility

This kit can be used with a variety of real-time instruments, including but not limited to the Applied Biosystems 7000, 7300, 7500, 7700, and 7900HT Real-Time PCR Systems; the Applied Biosystems GeneAmp® 5700; the Bio-Rad iCycler®; the Stratagene Mx3000P®, Mx3005P™, and Mx4000; the Corbett Research Rotor-Gene®, the MJ Research DNA Engine Opticon™, Opticon® 2, and Chromo 4™ Real-Time Detector; and Cepheid Smart Cycler® thermal cyclers. For instrument-specific protocols, go to [www.lifetechnologies.com/qpcr](http://www.lifetechnologies.com/qpcr). Optimal cycling conditions will vary with different instruments.

### Template

Starting material can range from 1 pg to 1 µg of purified total RNA. If you are using purified mRNA, the amount of template may be reduced to as low as 0.5 pg.

RNA should be free of RNase contamination and aseptic conditions should be maintained. RNA may be treated with DNase I (Catalog no. 18068-015) to remove any contaminating genomic DNA.

**Optional:** An RNase inhibitor such as RNaseOUT™ (Cat. no. 10777-019) may be added to the reaction after the 2X Reaction Mix to safeguard against degradation of target RNA due to ribonuclease contamination.

### Isolate total RNA

To isolate total RNA, we recommend the PureLink™ Micro-to-Midi™ Total RNA Purification System (Cat. no. 12183-018), TRIzol® Reagent (Cat. nos. 15596-026 and 15596-018), or the PureLink™ 96 Total RNA Purification Kit for high-throughput applications (Cat. no. 12173-011). Isolation of mRNA from total RNA is typically not necessary, although incorporating this step may improve the yield of specific cDNAs.

### Magnesium concentration

The 2X Reaction Mix includes magnesium at a final concentration of 3 mM. This works well for most targets; however, the optimal concentration may range from 3 mM to 6 mM. If necessary, use the separate tube of 50 mM magnesium sulfate to increase the magnesium concentration. Use the following table to determine the amount of MgSO<sub>4</sub> to add to achieve the specified concentration (in a 50 µL PCR with 25 µL of 2X Reaction Mix):

<u>Volume of 50 mM MgSO<sub>4</sub> (per 50 µL rxn)</u>	<u>Final MgSO<sub>4</sub> conc.</u>
1 µL	4.0 mM
2 µL	5.0 mM
3 µL	6.0 mM

Decrease the amount of water in the reaction accordingly.

### ROX™ Reference Dye

ROX™ Reference Dye can be included in the reaction to normalize the fluorescent reporter signal, for instruments that are compatible with that option. ROX™ Reference Dye is supplied at a 25 µM concentration, and is composed of a glycine conjugate of 5-carboxy-X-rhodamine, succinimidyl ester in 10 mM Tris-HCl (pH 8.4), 0.1 mM EDTA, and 0.01% Tween® 20. Use the following table to determine the amount of ROX™ Reference Dye to use with a particular instrument:

Instrument	ROX™ dye per 50-µL reaction	Final ROX™ dye concentration
Applied Biosystems 7000, 7300, 7700, 7900HT, and 7900HT Fast	1.0 µL	500 nM
Applied Biosystems 7500; Stratagene Mx3000™, Mx3005P™, and Mx4000	0.1 µL*	50 nM

\*To accurately pipet 0.1 µL per reaction, we recommend that you dilute ROX™ Reference Dye 1:10 immediately before use and use 1 µL of the dilution.

Note that the SuperScript® III Platinum® One-Step Quantitative RT-PCR System with ROX™ Reference Dye includes ROX™ Reference Dye in the 2X Reaction Mix at a final concentration of 500 nM (see **Additional products**, page 1).

### Primers

Gene-specific primers are required. LUX™ Fluorogenic Primers ([www.lifetechnologies.com](http://www.lifetechnologies.com)) are available separately.

A final concentration of 200 nM per primer is effective for most reactions. Doubling the amount of reverse primer (to 400 nM) may improve the performance of certain reactions. Optimal results may require a primer titration between 100 nM and 500 nM.

### Dual-labeled probes

A final probe concentration of 100 nM is effective for most reactions. The optimal concentration may vary between 50 nM and 500 nM.

### Reaction setup and conditions

Keep all components, reaction mixes and samples on ice. For most templates, efficient cDNA synthesis can be accomplished in a 15 minute incubation at 50°C. For problematic templates, or to increase the specificity of cDNA priming, increase the cDNA synthesis temperature up to 60°C. The cDNA synthesis temperature can range from 42°C to 60°C.

## General protocol for ABI instruments

Follow the protocol below for one-step qRT-PCR using either LUX™ primers or TaqMan® probes on ABI real-time instruments. Note the separate cycling conditions for the ABI 7500 in Fast Mode, and the lower amount of ROX™ Reference Dye required for the ABI 7500 and 7500 Fast systems. This generic protocol may also be used for other real-time instruments.

For more instrument-specific protocols, go to [www.lifetechnologies.com/qpcr](http://www.lifetechnologies.com/qpcr). A standard 50 µL reaction size is provided; component volumes can be scaled as desired (e.g., scaled down to a 20 µL reaction volume for 384-well plates).

1. Program your real-time instrument to perform cDNA synthesis immediately followed by PCR amplification, as shown below. Optimal temperatures and incubation times may vary for different target sequences.

Standard cycling program	Fast cycling program (for the ABI 7500 in Fast Mode)
50°C for 15 minutes hold (cDNA synthesis temperature may range from 42°C–60°C)	Select <i>Fast Mode</i> on the Thermal Profile tab
95°C for 2 minutes hold	50°C for 5 minutes hold
40 cycles of:	95°C for 2 minutes hold
95°C, 15 seconds	40 cycles of:
60°C, 30 seconds (60 seconds for the 7900HT)	95°C, 3 seconds
	60°C, 30 seconds
Melting curve analysis (LUX™ primers only): Refer to instrument documentation.	Melting curve analysis (LUX™ primers only): Refer to instrument documentation.

2. Set up reactions **on ice**. Volumes for a single 50 µL reaction are listed below. For multiple reactions, prepare a master mix of common components, add the appropriate volume to each tube or plate well on ice, and then add the unique reaction components (e.g., template). **Note:** Preparation of a master mix is *crucial* in qRT-PCR to reduce pipetting errors.

LUX™ primers reaction mix		TaqMan® probes reaction mix	
Component	Single rxn	Component	Single rxn
SuperScript® III RT/Platinum® Taq Mix	1 µL*	SuperScript® III RT/Platinum® Taq Mix	1 µL*
2X Reaction Mix	25 µL	2X Reaction Mix	25 µL
LUX™ labeled primer, 10 µM	1 µL	Forward primer, 10 µM	1 µL
Unlabeled primer, 10 µM	1 µL	Reverse primer, 10 µM	1 µL
ROX™ Reference Dye (optional)	1 µL/0.1 µL**	Fluorogenic probe, 10 µM	0.5 µL
RNaseOUT™ (optional)	1 µL	ROX™ Reference Dye (optional)	1 µL/0.1 µL**
Template (1 pg to 1 µg total RNA)	≤ 10 µL	RNaseOUT™ (optional)	1 µL
DEPC-treated water	to 50 µL	Template (1 pg to 1 µg total RNA)	≤ 10 µL
		DEPC-treated water	to 50 µL

\*To test for genomic DNA contamination of the RNA template, prepare a control reaction containing 2 units of Platinum® Taq DNA Polymerase (Catalog no. 10966-018) instead of the SuperScript® III RT/Platinum® Taq Mix.

\*\*See the table on page 2 for the amount/concentration of ROX™ Reference Dye to use for your specific instrument.

3. Cap or seal the reaction tube/PCR plate, and gently mix. Make sure that all components are at the bottom of the tube/plate; centrifuge briefly if needed.
4. Place reactions in a preheated real-time instrument programmed as described above. Collect data and analyze results.

## References

1. Kotewicz, M.L., D'Alessio, J.M., Driftmier, K.M., Blodgett, K.P., and Gerard, G.F. (1985) Cloning and overexpression of Moloney murine leukemia virus reverse transcriptase in *Escherichia coli*. *Gene* 35, 249.
2. Chou, Q., Russel, M., Birch, D., Raymond, J., and Bloch, W. (1992) Prevention of pre-PCR mispriming and primer dimerization improves low-copy-number amplifications. *Nucl. Acids Res.* 20, 1717.
3. Sharkey, D.J., Scalice, E.R., Christy, K.G., Atwood, S.M., and Daiss, J.L. (1994) Antibodies as thermolabile switches: high temperature triggering for the polymerase chain reaction. *BioTechnology* 12, 506.
4. Lowe, B., Avila, H. A., Bloom, F., Gleeson, M., and Kusser, W. (2003) Quantitation of gene expression in neural precursors by reverse-transcription polymerase chain reaction using self-quenched, fluorogenic primers. *Anal. Biochem.* 315, 95.
5. Tyagi, S. and Kramer, F.R. (1996) Molecular beacons: probes that fluoresce upon hybridization. *Nature Biotechnology* 14, 303.
6. Kostrikis, L.G., Tyagi, S., Mhlanga, M.M., Ho, D.D., and Kramer, F.R. (1998) Spectral genotyping of human alleles. *Science* 279, 1228.
7. Holland, P.M., Abramson, R.D., Watson, R., and Gelfand, D. H. (1991) Detection of specific polymerase chain reaction product by utilizing the 5'–3' exonuclease activity of *Thermus aquaticus* DNA polymerase. *Proc. Natl. Acad. Sci. USA* 88, 7276.

## Troubleshooting guide

Problem	Possible cause	Solution
No amplification product; Relative fluorescent signal $\leq$ background or no-template control	cDNA synthesis temperature too high, low priming efficiency	Lower incubation temperature.
	RT or cDNA primer blocked by secondary structure	Raise incubation temperature. Redesign primer(s).
	RNA has been damaged/degraded	Replace RNA if necessary.
	RNase contamination	Maintain aseptic conditions; add RNase inhibitor.
	Fluorescent probe not functional	Validate probe design and presence of fluorophore and quencher: Treat TaqMan <sup>®</sup> probe with DNase, and check for increase in fluorescence. Redesign and/or resynthesize probe if necessary.
Poor sensitivity	Not enough template RNA	Increase concentration of template RNA; use 10 ng–1 $\mu$ g total RNA.
Product detected at higher than expected cycle number	RNA has been damaged/degraded	Replace RNA if necessary.
	RNase contamination	Maintain aseptic conditions; add RNase inhibitor.
	RT inhibitors are present in RNA	Remove inhibitors in the RNA preparation by an additional 70% ethanol wash. Inhibitors of RT include SDS, EDTA, guanidium salts, formamide, sodium phosphate and spermidine.
	Inefficient cDNA synthesis	Adjust cDNA synthesis temperature and/or primer design. Double the amount of reverse primer ( <i>e.g.</i> , to 400 nM).
	Inefficient PCR amplification	Optimize PCR conditions: <ul style="list-style-type: none"> <li>• Adjust annealing temperature as necessary.</li> <li>• Increase magnesium concentration.</li> <li>• Redesign primers.</li> </ul>
Higher than expected signal	Too much sample added to reactions	Decrease the concentration of template RNA.
Product detected at lower-than-expected cycle number, and/or positive signal from no-template controls	Template or PCR carry-over contamination	<ul style="list-style-type: none"> <li>• Isolate source of contamination and replace reagent(s).</li> <li>• Use separate dedicated pipettors for reaction assembly and post-PCR analysis.</li> <li>• Assemble reactions (except for target addition) in a DNA-free area.</li> <li>• Use aerosol-resistant pipet tips or positive displacement pipettors.</li> </ul>
Unexpected bands after electrophoresis	Genomic DNA contamination	Pre-treat RNA with DNase I.
	Oligo(dT) or random primers used for cDNA synthesis	Use only gene-specific primers.
	Low specificity in PCR	Optimize PCR conditions as described above.

### Certificate of Analysis

The Certificate of Analysis provides detailed quality control and product qualification information for each product. Certificates of Analysis are available from [www.lifetechnologies.com](http://www.lifetechnologies.com).

### Important Licensing Information

These products may be covered by one or more Limited Use Label Licenses. By use of these products, you accept the terms and conditions of all applicable Limited Use Label Licenses.

### Limited Product Warranty

Life Technologies Corporation and/or its affiliate(s) warrant their products as set forth in the Life Technologies' General Terms and Conditions of Sale found on Life Technologies' website at [www.lifetechnologies.com/termsandconditions](http://www.lifetechnologies.com/termsandconditions). If you have any questions, please contact Life Technologies at [www.lifetechnologies.com/support](http://www.lifetechnologies.com/support).

**DISCLAIMER:** LIFE TECHNOLOGIES CORPORATION AND/OR ITS AFFILIATE(S) DISCLAIM ALL WARRANTIES WITH RESPECT TO THIS DOCUMENT, EXPRESSED OR IMPLIED, INCLUDING BUT NOT LIMITED TO THOSE OF MERCHANTABILITY, FITNESS FOR A PARTICULAR PURPOSE, OR NON-INFRINGEMENT. TO THE EXTENT ALLOWED BY LAW, IN NO EVENT SHALL LIFE TECHNOLOGIES AND/OR ITS AFFILIATE(S) BE LIABLE, WHETHER IN CONTRACT, TORT, WARRANTY, OR UNDER ANY STATUTE OR ON ANY OTHER BASIS FOR SPECIAL, INCIDENTAL, INDIRECT, PUNITIVE, MULTIPLE OR CONSEQUENTIAL DAMAGES IN CONNECTION WITH OR ARISING FROM THIS DOCUMENT, INCLUDING BUT NOT LIMITED TO THE USE THEREOF.

©2015 Thermo Fisher Scientific Inc. All rights reserved. All trademarks are the property of Thermo Fisher Scientific and its subsidiaries unless otherwise specified. Rotor-Gene is a trademark of Qiagen GmbH. Mx3000P, Mx3005P are trademarks of Agilent Technologies, Inc. iCycler, DNA Engine Opticon, Opticon, and Chrome 4 are trademarks of Bio-Rad Laboratories, Inc. SmartCycler1 is a trademark of Cepheid Corporation. TaqMan is a registered trademark of Roche Molecular Systems, Inc.