



SG OneStep RT-qPCR kit

SG OneStep RT-qPCR kit is one-step RT-qPCR kit that provides accurate real-time quantification of RNA targets. Kit is composed of unique reverse transcriptase and highly processive hot start on Taq DNA Polymerase in easy to use format.

SG OneStep RT-qPCR kit

Kit Components

Component	Cat. No. E0810-01 25 reactions of 25 μl	Cat. No. E0810-02 100 reactions of 25 μl
2 x RT-qPCR SG Buffer	1 x 350 μl	2 x 0.7 ml
SG Enzyme Mix	25 μΙ	100 μΙ
Thermolabile UNG (uracil-N-glycosylase) 1 U/μl	10 μΙ	30 μΙ
Water, nuclease free	1 x 0.5 ml	2 x 1 ml

SG OneStep RT-qPCR kit, plus ROX Solution

Kit Components

Component	Cat. No. E0811-01 25 reactions of 25 μl	Cat. No. E0811-02 100 reactions of 25 μl
2 x RT-qPCR SG Buffer	1 x 350 μl	2 x 0.7 ml
ROX Solution, 25 μM	15 μΙ	60 μΙ
SG Enzyme Mix	25 μΙ	100 μΙ
Thermolabile UNG (uracil-N-glycosylase) 1 U/μl	10 μΙ	30 μΙ
Water, nuclease free	1 x 0.5 ml	2 x 1 ml

Storage

Store at -20°C in the dark.

The kit provides:

- 2 x RT-qPCR SG Buffer that is a universal reaction buffer with dNTPs (dTTP is partially replaced with dUTP) that can be used on most real-time PCR cyclers available. Allows as option to use thermolabile uracil-N-glycosylase (UNG).
- The SG Enzyme Mix contains unique highly sensitive reverse transcriptase, hot start on Taq DNA Polymerase, RNase Inhibitor and SYBR Green I dye.
- Reverse transcriptase that works in high range of temperatures from 35-55°C without loss of specificity and sensitivity.
- Both cDNA synthesis and PCR are performed in a single tube using gene-specific primers and either total RNA or mRNA.
- onTaq DNA Polymerase is a modified "hot start" enzymes which provides very tight inhibition of the polymerase activity at moderate temperatures which is restored during the initial denaturation step at 95°C for at least ten minutes.

- SYBR Green I is a fluorescent dye which binds all doublestranded DNA molecules and emits a fluorescent signal on binding. The excitation and emission maxima of SYBR Green I are at 494 nm and 521 nm, respectively, which are compatible with any real-time cycler.
- Kit contains thermolabile uracil-N-glycosylase (UNG) that is optimized for RT-qPCR reactions.
- If cyclers from Applied Biosystems are used ROX passive reference dye is necessary. SG OneStep RT-qPCR kit is provided in two variants: without ROX and with ROX Solution provided separately. The table below shows recommended amount of ROX (25 μ M) required for a specific PCR cycler.

Recommended amounts of ROX for a specific real-time PCR cycler

Instrument	Amount of ROX per 25 µl reaction	Final ROX concentration
Applied Biosystems: 7300, 7900HT, StepOne, StepOnePlus, ABI PRISM 7000 and 7700	0.5 யி	500 nM
Applied Biosystems: 7500, ViiA 7, Stratagene: Mx3000P, Mx3005P, Mx4000	0.5 μl 10 x diluted (in water)	50 nM
PCR machines from other manufacturers: Bio-Rad, Roche, Corbett, Eppendorf, Cepheid, etc.	Not required	-

Protocol

Preparation of PCR reaction:

Component	Volume/reaction	Final concentration
2 x RT-qPCR SG Buffer	12.5 μΙ	1 x
Forward Primer	Variable	0.4 μΜ
Reverse Primer	Variable	0.4 μΜ
Template RNA	Variable	≤500 ng
SG Enzyme Mix	1 μΙ	1 μl/ reaction
Optional: ROX Solution, 25 μΜ	0.5 μl or 0.5 μl 10 x diluted	500 nM 50 nM
Optional: Thermolabile UNG (uracil-N-glycosylase) 1 U/μl	0.25 µl	0.25 U/reaction
Water, nuclease free	To 25 μl	-
Total volume	25 μΙ	-

Notes:

- Minimize thaw-freeze cycles of 2 x RT-qPCR SG Buffer, keep SG Enzyme Mix and ROX solution on ice and minimize light exposure during handling to avoid loss of fluorescent signal intensity.
- 2. Thaw and gently vortex before use 2 x RT-qPCR SG Buffer.
- 3. A reaction volume of 25 μ l should be used with most real-time cyclers. Other reaction volumes may be used if recommended for a specific instrument.
- 4. Optimal amplicon length in real-time RT-PCR using SYBR Green I is 70-150 bp.
- 5. To avoid amplification from genomic DNA design exon-exon primers.
- 6. Set up RT-PCR reactions on ice to minimize RNA template degradation.
- 7. The RNA template (≤500 ng/reaction) should be added to the individual PCR tubes or wells containing the whole reaction mix. Centrifuge briefly before placing into a cycler. Check if there are no bubbles left, if yes, spin again.
- 8. Place the samples in the cycler and start the program.

- Reverse transcriptase works in a wide range of temperatures 35-55°C. The recommended starting temperature for reverse transcription is 50°C. For individual experiment temperature might be changed.
- 10. Standard concentration of $MgCl_2$ in real-time RT-PCR reaction is 3 mM (as provided with the 1 x RT-qPCR SG Buffer) in most cases this concentration will produce optimal results. However, if a higher $MgCl_2$ concentration is required, prepare a 25 mM $MgCl_2$ stock solution and add to reaction.
- 11. A final primer concentration of 0.3-0.5 μ M is usually optimal, but can be individually optimized in the range of 0.1 μ M to 1 μ M. The recommended starting concentration is 0.4 μ M. Raising primer concentration may increase PCR efficiency, but negatively affect RT-PCR specificity. Optimal primer concentration depends on the individual reaction and the real-time PCR cycler used.
- 12. Readjust the threshold value for analysis of every run.
- 13. If using Bio-Rad iCycler iQ or MyiQ instruments collect well factors at the beginning of each experiment. Use an external well factor plate according to the manufacturer's recommendations. Well factors are used to compensate for any excitation or pipetting variations.

Thermal Cycling Conditions:

Step	Temperature	Time	Number of Cycles
Reverse Transcription	50°C	20 min	1
Initial Denaturation	95°C	15 min	1
Denaturation Annealing/Extension/Data acquisition	94°C 60°C	15 s 60 s	40-45
Cooling	4°C	Indefinite	1

Notes:

- 1. During reverse transcription step of 50°C also thermolabile uracil-N-glycosylase might be used. Do not use UNG from *E.coli*, UNG will degrade all newly synthetized cDNA.
- 2. Thermolabile UNG is inactivated at 50°C during RT.
- Melting curve analysis should be performed to verify the specificity and identity of PCR products. Melting curve analysis is an analysis step built into the software of real-time cyclers. Melting curve data between 65°C and 95°C should be acquired.
- 4. Data acquisition should be performed during the extension step. To suppress fluorescence readings caused by the generation of primer-dimers an additional data acquisition step can be added to the protocol. It is possible when T_m of primer-dimers differs from T_m of the specific product (T_m are generated during melting curve analysis). The temperature of the data acquisition step should be above T_m of primer-dimers but approximately 3°C below the T_m of the specific product.
- 5. Always check the RT-PCR product specificity by gel electrophoresis when designing a new assay. Melting temperatures of the specific product and primer-dimers may overlap.