For life science research only. Not for use in diagnostic procedures.



Titan One Tube RT-PCR Kit

Content Version: February 2021

Cat. No. 11 939 823 001 1 kit 50 rea

1 kit 50 reactions including 10 control reactions

Store the kit at −15 to −25°C.

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1. General Information

1.1. Contents

| Vial / bottle | Label | Function / description | Content |
|---------------|--|---|-----------------------|
| 1 | Titan One Tube RT-PCR Kit, Enzyme mix | Expand High Fidelity enzyme mix and AMV reverse transcriptase in storage buffer. 1 reaction/µl | 1 vial, 50 µl |
| 2 | Titan One Tube RT-PCR Kit, dNTP mix | 10 mM (total), 2.5 mM each dNTP. | 1 vial, 200 µl |
| 3 | Titan One Tube RT-PCR Kit, RNase inhibitor | 5 U/μl | 1 vial, 50 µl |
| 4 | Titan One Tube RT-PCR Kit, Human control RNA | 2 pg/µl K-562 total RNA with MS2 carrier RNA. | 1 vial, 50 µl |
| 5 | Titan One Tube RT-PCR Kit, Control primer mix | 20 μM (total) Human β-actin upstream and downstream primer | 1 vial, 20 µl |
| 6 | Titan One Tube RT-PCR Kit, Water, PCR Grade | To adjust the final reaction volume. | 2 vials, 1 ml each |
| 7 | RT-PCR buffer, 5x conc. | Optimized RT-PCR buffer with 7.5 mM MgCl ₂ and DMSO. | 1 vial, 1 ml |
| 8 | Dithiothreitol (DTT) solution | 100 mM solution | 1 vial, 1 ml |
| 9 | MgCl ₂ 25 mM Stock Solution | To adjust final Mg ² + concentration. | 1 vial, 1 ml |

1.2. Storage and Stability

Storage Conditions (Product)

When stored at -15 to -25° C, the kit is stable through the expiry date printed on the label.

| Vial / bottle | Label | Storage | |
|---------------|--|------------------------------------|--|
| 1 | Enzyme mix | Store at -15 to -25° C. | |
| 2 | dNTP mix | | |
| 3 | RNase inhibitor | | |
| 4 | Human control RNA | Store at −60°C or below. | |
| 5 | Control primer mix | Store at -15 to -25° C. | |
| 6 | Water, PCR Grade | | |
| 7 | RT-PCR buffer, 5x conc. | | |
| 8 | Dithiothreitol (DTT) solution | | |
| 9 | MgCl ₂ 25 mM Stock Solution | | |

1.3. Additional Equipment and Reagent required

Standard laboratory equipment

- Nuclease-free, aerosol-resistant pipette tips
- · Pipettes with disposable, positive-displacement tips
- Sterile reaction tubes for preparing PCR mixes and dilutions
- PCR reaction vessels, such as 0.2 ml thin-walled PCR tubes or plates
- Standard benchtop microcentrifuge
- Thermal block cycler

For RT-PCR

- Template RNA
- Mineral oil (optional)

1.4. Application

The Titan One Tube RT-PCR Kit is used to analyze RNA (total, poly (A)+, or viral) from a variety of sources, such as animal, plant, human, viral, and bacterial. The kit can be used for:

- Qualitative, semi quantitative, or quantitative analysis of RNA transcription levels even in single cells in combination with gel-based or ELISA-based detection methods.
- Cloning of RNA up to 6 kb.
- Mutation analysis at RNA level in combination with sequencing or other mutation scanning techniques.

1.5. Preparation Time

Assay Time

3 to 6 hours, depending on fragment length.

2. How to Use this Product

2.1. Before you Begin

Sample Materials

Isolated total or poly(A)+ RNA up to 1 µg or single cells.

Control Reactions

DNA contamination

To exclude artifacts from DNA targets, such as residual genomic DNA contaminations from RNA preparations, and contaminating DNA from previous amplifications, appropriate positive and negative control reactions should be included in the experimental design. Consider setting up the following 3 controls:

To control contaminating DNA amplicons from previous PCR, always set up a control reaction without RNA template.

- No PCR product should be visible.

2 Set up a control reaction where the RT reaction is not performed.

– Instead of the Titan Enzyme mix in the kit, use the Expand Enzyme mix from the Expand High Fidelity or Expand Long Template Systems*, or use 2.5 U Taq DNA Polymerase*.

- Alternatively, heat the Master Mix 2 for 2 minutes at +94°C to inactivate reverse transcriptase AMV, then follow Step 1 in the RT-PCR protocol and proceed with Step 3 (do not use Step 2).

i Consider that even Taq DNA polymerase has a low reverse transcriptase activity.

3 Treat your RNA preparation with RNase A, then proceed with the RT-PCR protocol.

RT-PCR control reaction

Amplification of a 587 bp fragment out of human β-actin mRNA target.

Primers

Primer design

Sequence-specific primers must be used for one tube RT-PCR.

- To differentiate between amplification of cDNA and amplification of contaminating genomic DNA, primers can be designed that anneal to sequences in exons on both sides of an intron. PCR products from genomic DNA will be much longer compared to the intronless mRNA-derived products.
- Alternatively, a primer designed on an exon/exon boundary of the mRNA should not amplify genomic DNA.
- The final concentration of primers in the reaction should range from 0.2 to 1 $\mu M.$

Primer sequences

Human β -actin forward primer: 5'-CCA-AGG-CCA-ACC-GCG-AGA-AGA-TGA-C Human β -actin reverse primer: 5'-AGG-GTA-CAT-GGT-GGT-GCC-GCC-AGA-C

Mg2+ Concentration

1.5 mM

The optimum magnesium concentration in RT-PCR depends on the final concentration of dNTPs, primers, and template. For most applications, 1.5 mM magnesium will allow amplification of your target RNA using the standard protocol. Depending on the respective target/primer combination, a titration of the magnesium concentration may significantly improve sensitivity and specificity.

General Considerations

The optimal conditions, including incubation times and temperatures, concentration of enzyme, template DNA, Mg²+ vary from system to system and must be determined for each individual experimental system. At the very least, titrate the Mg²+ concentration and the amount of enzyme mix used per assay to ensure optimal efficiency of DNA synthesis. As a starting point, use the following guidelines:

- Optimal Mg²+: In most cases, a Mg²+ concentration of 1.5 mM will produce satisfactory results.
- dNTP concentration: For the amplification of RT-PCR products >2 kb, use the sodium salts of dNTPs. Always use equal concentrations of all four dNTPs. The most commonly used concentration is 200 µM. RT-PCR performance can be improved by increasing the concentration to 500 µM dNTPs, each. Since nucleotides strongly bind magnesium ions, the concentration of magnesium will also have to be increased accordingly. A final concentration of 500 µM dNTP typically requires a concentration of 3 mM magnesium chloride.

RNA preparation

Any contamination with RNases from other potential sources, such as glassware, plasticware, and reagent solutions must be avoided.

- Integrity of mRNA is particularly important for the generation of long RT-PCR products. The size of the mRNA can be determined by gel electrophoresis and ethidium bromide staining. The mRNA should appear as a smear between approximately 500 bp and 8 kb. The bulk of the mRNA should be between 1.5 and 2 kb.
- For high quality eukaryotic mRNA preparations, minimize the activity of RNases released during cell lysis by using inhibitors of RNases or methods that disrupt cells and simultaneously inactivate RNases.

RNase inhibitor

For best results, add 5 to 10 U of RNase inhibitor (Vial 3) to minimize RNase activity. This is particularly important when RT-PCR is started with low amounts of RNA (<10 ng).

Prevention of Carryover Contamination

If dTTP is completely substituted by dUTP, as generally done for carryover prevention using Uracil DNA-Glycosylase, heat-labile*, the yield of PCR products incorporating modified nucleotides can decrease significantly.

Safety Information

For customers in the European Economic Area

Contains SVHC: octyl/nonylphenol ethoxylates. For use in research and under controlled conditions only – acc. to Art. 56.3 and 3.23 REACH Regulation.

2.2. Protocols

Preparation of RT-PCR master mixes

Prepare two RT-PCR master mixes. Master Mix 2 contains enzyme and reaction buffer; Master Mix 1 contains all other reaction components. This circumvents the need for hot start and avoids that the enzyme interacts with primers or template during the reaction setup.

Preparation of master mix 1

Always wear gloves when handling RNA to avoid any contamination with RNases.

1 Thaw components listed below and place them on ice.

2 Briefly vortex and centrifuge all reagents before setting up the reactions.

3 To a sterile, nuclease-free reaction tube on ice, add the components in the order listed for each 50 µl reaction:

| Reagent | Control Volume [µl] | Probe Volume [µl] | Final conc. |
|---|---------------------|-----------------------------------|--------------------------|
| Water, PCR Grade (Vial 6) | 10.5 | add up to a final volume of 25 | _ |
| dNTP Mix (10 mM of each dNTP) (Vial 2) | 4 | 4 | 200 μ M of each dNTP |
| DTT solution, 100 mM | 2.5 | 2.5 | 5 mM |
| RNase inhibitor (Vial 3) | 1 | 1 | 5 U |
| Control primer mix (Vial 5) | 2 | - | 400 nM each |
| Forward primer 1 | - | variable | 400 nM |
| Reverse primer 2 | - | variable | 400 nM |
| Human control RNA (Vial 4) | 5 | - | 10 pg |
| Template RNA | - | variable | 1 µg – 1 pg total RNA |
| Final Volume | 25 | 25 | |

4 Mix and centrifuge briefly to collect the sample at the bottom of the tube.

Preparation of master mix 2

Always wear gloves when handling RNA to avoid any contamination with RNases.

1 Thaw components listed below and place them on ice.

2 Briefly vortex and centrifuge all reagents before setting up the reactions.

3 To a sterile, nuclease-free reaction tube on ice, add the components in the order listed for each 50 µl reaction:

| Reagent | Control Volume [µl] | Probe Volume [µl] | Final conc. |
|----------------------------------|---------------------|-------------------|--------------------------------|
| Water, PCR Grade (Vial 6) | 14 | 14 | - |
| RT-PCR buffer, 5x conc. (Vial 7) | 10 | 10 | 1x (1.5 mM MgCl ₂) |
| Enzyme mix (Vial 1) | 1 | 1 | _ |
| Final Volume | 25 | 25 | |

4 Mix and centrifuge briefly to collect the sample at the bottom of the tube.

RT-PCR

The cycle number is dependent on the abundance of the respective mRNA. For rare mRNA messages, 40 cycles or a second (nested) PCR could be necessary.

Reverse transcription temperature

A template denaturation step prior to initiation of the reverse transcription is not required. If desired, a denaturation step may be incorporated by incubating primers and template after addition of the required amount of water at +68°C for 2 minutes or at +94°C for 1 minute.

A Do not incubate AMV or RNase inhibitor at this temperature since these components will become inactivated.

Reverse transcription may be performed between +42 and +60°C, see section, **Results**, Figures 1 and 2. When establishing a new assay, start with +50°C. If the reaction is performed at temperatures >+50°C, only short RNA fragments can be amplified due to partial inactivation of AMV reverse transcriptase. To amplify long fragments >3 kb by RT-PCR, use a temperature range of +45 to +48°C.

PCR temperature

The annealing temperature in PCR depends on the melting temperature of the respective primer pair. Use an appropriate computer program to calculate the optimal temperature for your primers. The elongation temperature should always be +68°C. The denaturation temperature depends on the G + C content of the template. A high G + C content (>60%) of your RNA might require either higher denaturation temperatures or a longer denaturation time.

RT-PCR protocol

🛕 Always wear gloves when handling RNA to avoid any contamination with RNases.

🕖 The following thermal profiles are an example. Different thermal cyclers may require different profiles.

- For each reaction, combine 25 μl Master Mix 1 and 25 μl Master Mix 2 in a 0.2 ml thin-walled PCR tube on ice.
 Gently vortex the mixture to produce a homogeneous reaction, then centrifuge briefly to collect the solution at the bottom of the tube.
 - Overlay the reaction carefully with 30 μ l mineral oil if required by the thermal cycler.

Place your samples in a thermal block cycler equilibrated to +50°C and incubate for 30 minutes.
 i Temperatures up to +60°C can be used for the reverse transcription step.

3 Start cycling using the following thermal profile.

| Step | Temperature [°C] | Time | Number of Cycles |
|------------------|------------------------|---|-------------------|
| Pre-Incubation | 94 | 2 min | 1 |
| Denaturation | 94 | 30 sec | 10 |
| Annealing | 45 – 65 ⁽¹⁾ | 30 sec | |
| Elongation | 68 | 45 sec – 4 min ⁽²⁾ | |
| Denaturation | 94 | 30 sec | 25 ⁽³⁾ |
| Annealing | 45 - 65 ⁽¹⁾ | 30 sec | |
| Elongation | 68 | 45 sec – 4 min ⁽²⁾ + 5 sec cycle | |
| - | | elongation for each successive cycle ⁽³⁾ | |
| Final Elongation | 68 | 7 min | 1 |

4 Analyze the samples on a 1 to 2% agarose gel.

⁽¹⁾ Optimal annealing temperature depends on the melting temperature of the primers and on the experimental system. Use an annealing temperature of +66°C for the human β -actin control primers supplied with the kit.

⁽²⁾ Elongation time depends on fragment length as shown in the table. Cycle number 11 is 5 seconds longer than cycle 10. Cycle number 12 is 10 seconds longer than cycle 10. Cycle number 13 is 15 seconds longer than cycle 10, etc.

| PCR fragment length [kb] | <1 | 1.5 | 3 | 4.5 | 6 |
|------------------------------|--------|-------|-------|-------|---------|
| Elongation time | 45 sec | 1 min | 2 min | 3 min | 4 min |
| Temperature for RT-step [°C] | ≤60 | 50 | 50 | 48 | 45 - 48 |

⁽³⁾ The cycle number is dependent on the abundance of the respective mRNA. For rare messages, 40 cycles or a second nested PCR may be necessary.

2.3. Parameters

Incorporation of Modified Nucleotides

AMV reverse transcriptase and the Expand High Fidelity enzyme mix both accept modified nucleotides, such as Digoxigenin-11-dUTP*, Biotin-16-dUTP*, or Fluorescein-12-dUTP* at a concentration usually used:

• For probe synthesis, for example, 1:3 to 1:6 DIG-dUTP:dTTP.

• For labeling PCR products to detect products in a ELISA format.

Labeling with DIG-dUTP for ELISA

Follow the protocol described in section, **Protocols, Preparation of RT-PCR master mixes** with one exception. Use instead of the 4 μ l supplied in the dNTP mix, use 5 μ l of the PCR DIG Labeling Mix* or the PCR ELISA (DIG Labeling)*. Both products use dNTPs with the following final concentration per reaction: 200 μ M dATP, dGTP, dCTP each; 190 μ M dTTP and 10 μ M DIG-11-dUTP.

Maximum Fragment Size

Up to 6 kb.

PCR Cloning

TA cloning or blunt-end cloning.

The obtained fragments are a mixture of fragments with A overhangs at the 3' end and blunt-ended fragments. Without any treatment of the ends, TA cloning will yield in general better results than blunt-end cloning.

Sensitivity

10 pg

1,000 copies of RNA targets can be visualized on an agarose gel and ethidium bromide staining, by analyzing 10 μ l of the 50 μ l reaction after 35 cycles. Sensitivity can easily be improved to the lowest level of RNA targets by using more cycles and/or using labeled nucleotides, such as DIG-dUTP during PCR, and analyzing the reaction in an ELISA format.

Specificity

Specificity of the reaction strongly depends on primer design and DNA contamination. See section, **Control Reactions** for additional information.

Temperature Optimum

+42°C to +60°C Reverse transcription

3. Results

Effect of varying cDNA synthesis reaction temperatures

Figure 1 shows the effect of varying cDNA synthesis reaction temperatures on sensitivity of RT-PCR performed with the Titan One Tube RT-PCR Kit and a 324 bp mouse β -actin template.

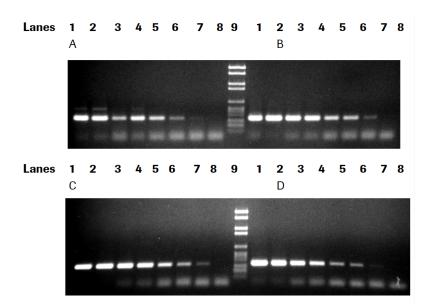


Fig. 1: Varying amounts of total mouse liver RNA as template was used. cDNA synthesis was performed at +45°C (panel A), +50°C (panel B), +55°C (panel C), and +60°C (panel D) for 30 minutes, followed by 35 cycles of PCR. A 15 µl aliquot of the PCR product was analyzed on a 1% agarose gel.

Lane 1: 10 ng total mouse liver RNA

Lane 2: 5 ng total mouse liver RNA

Lane 3: 1 ng total mouse liver RNA

Lane 4: 500 pg total mouse liver RNA

Lane 5: 100 pg total mouse liver RNA

Lane 6: 50 pg total mouse liver RNA

Lane 7: 10 pg total mouse liver RNA

Lane 8: negative control containing no template

Lane 9: DNA Molecular Weight Marker VI

Figure 2 shows the effect of different cDNA synthesis reaction temperatures on the sensitivity on RT-PCR performed with the Titan One Tube RT-PCR Kit and a 1.9 kb dystrophin template.

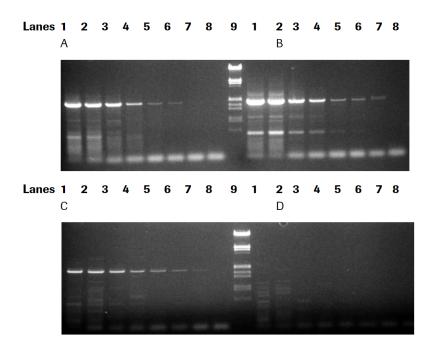


Fig. 2: Varying amounts of skeletal muscle liver RNA as template was used. cDNA synthesis was performed at +45°C (panel A), +50°C (panel B), +55°C (panel C), and +60°C (panel D) for 30 minutes, followed by 35 cycles of PCR. A 15 µl aliquot of the PCR product was analyzed on a 1% agarose gel.

Lanes 1 to 8: same as in Figure 1 but total skeletal muscle RNA was used as template. Lane 9: DNA Molecular Weight Marker III

Amplification of RT-PCR products

Figure 3 shows the amplification of various RT-PCR products with the Titan One Tube RT-PCR Kit.

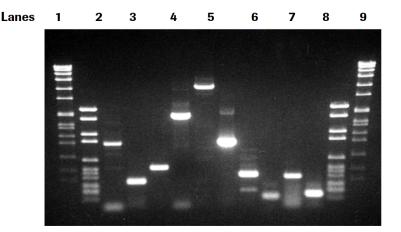


Fig. 3: RT-PCR was performed according to the Instructions for Use. cDNA synthesis was performed at +50°C, 35 cycles of PCR. 15 µl aliquots of the RT-PCR product were analyzed on a 1% agarose gel.

Lanes 1 and 9: DNA Molecular Weight Marker VII

Lanes 2 and 8: DNA Molecular Weight Marker VI

Lane 3: 324 bp ß-actin (1 ng human liver total RNA)

Lane 4: 980 bp GAPDH (10 ng human liver total RNA)

Lane 5: 2.45 kb ApoB (10 ng human liver total RNA)

Lane 6: 2 kb Dystrophin (10 ng human skeletal muscle total RNA)

Lane 7: 4 kb Dystrophin (10 ng human skeletal muscle total RNA)

4. Troubleshooting

| Observation | Possible cause | Recommendation |
|------------------------------------|--|---|
| No or very little PCR product. | Insufficient amount of template RNA. | Increase amount of RNA template in cDNA |
| | insufficient amount of template nin/t. | reaction. |
| | | Use poly(A)+ mRNA rather than total RNA as template. |
| | Template RNA degraded. | Prepare fresh RNA template, being careful to prevent RNase activity. |
| | | Check RNA preparation by gel electrophoresis. |
| | Too much template RNA. | Decrease amount of RNA template. <i>i</i> A too high amount of template RNA may affect/inhibit performance of RT-PCR. |
| | Reaction not optimized. | Increase the concentration of dNTPs in the PCR reaction up to 500 μ M maximum. At the same time, raise the final concentration of MgCl ₂ in the reaction to 3 mM to compensate for the additional dNTPs. |
| | | Increase primer concentration up to 1 μM maximum. |
| | | Synthesize the cDNA at a higher temperature. |
| | Template secondary structure prevented effective first strand cDNA | Optimize DMSO concentration up to 10% maximum. |
| | synthesis. | Briefly denature the RNA template at +94°C for 1 minute before adding reverse transcriptase. ▲ Do not incubate reverse transcriptase or RNase inhibitor at this elevated temperature as they will be inactivated. |
| | Template secondary structure inhibits effective formation of full-length products. | If GC content of RNA is high (>60%), increase denaturation temperature or denaturation time in PCR cycles. |
| Product is smeared. | Secondary amplification product(s). | Check reagent concentrations and cycling conditions: Check Mg²+ concentration. Optimize temperature of cDNA synthesis step. Optimize primer concentration. Decrease number of cycles. Check and perhaps decrease concentration of template. Titrate the amount of enzyme(s). |
| Nonspecific product bands present. | Annealing temperature too low. | Increase annealing temperature during PCR to increase specificity of amplification. |
| | Contaminating DNA in sample. | Perform a no-RT control, see section, Control Reactions for additional information. |

5. Additional Information on this Product

5.1. Test Principle

Background information

RT-PCR is a sensitive technique to determine the mere presence or relative quantity of specific RNA templates, for example, in gene expression studies. Furthermore, RT-PCR allows cloning of rare messages without the necessity to construct cDNA libraries.

- In two-step RT-PCR, reverse transcription of RNA into cDNA is performed prior to amplification of cDNA by PCR in a separate reaction. This reaction setup requires to open the reaction tube after cDNA synthesis and to add the reaction components required for the PCR step of the procedure. Not only is this handling inconvenient, but it also increases the risk of contamination.
- In one-step RT-PCR, cDNA synthesis as well as PCR are performed using an optimized buffer system in one reaction tube in a combined reaction without the requirement to add further reagents between cDNA synthesis and PCR.

Two basic techniques for one-step RT-PCR are described:

- Tth DNA polymerase can be used for both reverse transcription and PCR due to its intrinsic RT-activity. The thermostability of this enzyme can overcome secondary structures in RNAs by performing cDNA synthesis at +60 to +70°C. However, the length of the RT-PCR products is limited to less than 1 kb. Furthermore the polymerase possesses an increased error rate due to the use of manganese ions.
- The combination of AMV reverse transcriptase and Taq DNA polymerase allows amplification of fragments up to 2 kb with decreased error rates compared to the use of Tth DNA polymerase. However, this reaction has to be performed during the cDNA synthesis step at +42°C.

By combining AMV Reverse Transcriptase and the Expand High Fidelity PCR System together with an optimized buffer system in the Titan One Tube RT-PCR Kit, the limitations of this approach are now overcome allowing an RT reaction temperature of up to +60°C and fragment lengths of up to 6 kb.

Reaction mechanism

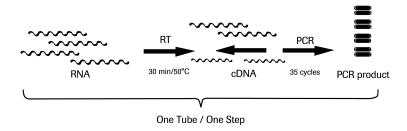


Fig. 4: Reaction mechanism of Titan One Tube RT-PCR Kit.

5.2. Quality Control

For lot-specific certificates of analysis, see section, Contact and Support.

6. Supplementary Information

6.1. Conventions

To make information consistent and easier to read, the following text conventions and symbols are used in this document to highlight important information:

| Text convention and symbols | | | | |
|---|--|--|--|--|
| <i>i</i> Information Note: Additional information about the current topic or procedure. | | | | |
| ▲ Important Note: Information critical to the success of the current procedure or use of the product. | | | | |
| 123 etc. | Stages in a process that usually occur in the order listed. | | | |
| 1 2 3 etc. | Steps in a procedure that must be performed in the order listed. | | | |
| * (Asterisk) | The Asterisk denotes a product available from Roche Diagnostics. | | | |

6.2. Changes to previous version

Layout changes. Editorial changes. New information added related to the REACH Annex X.

6.3. Ordering Information

| Product | Pack Size | Cat. No. |
|-------------------------------------|--|----------------|
| Reagents, kits | | |
| Digoxigenin-11-dUTP, alkali-labile | 25 nmol, 25 μl, 1 mM | 11 573 152 910 |
| | 125 nmol, 125 μl, 1 mM | 11 573 179 910 |
| Digoxigenin-11-dUTP, alkali-stable | 25 nmol, 25 μl, 1 mM | 11 093 088 910 |
| | 125 nmol, 125 μl, 1 mM | 11 558 706 910 |
| | 5 x 125 nmol, 5x 125 μl, 1 mM | 11 570 013 910 |
| Biotin-16-dUTP | custom fill | 11 093 711 103 |
| Fluorescein-12-dUTP | custom fill | 11 375 601 103 |
| Uracil-DNA Glycosylase, heat-labile | custom fill | 11 780 565 103 |
| Expand High Fidelity PCR System | 100 U, 1 x 100 U, 40 reactions in a final volume of 50 μl | 11 732 641 001 |
| | 500 U, 2 x 250 U, 200 reactions in a final volume of 50 μl | 11 732 650 001 |
| | 2,500 U, 10 x 250 U, 1,000 reactions in a final volume of 50 μl | 11 759 078 001 |
| Expand Long Template PCR System | 150 U, 1 x 150 U, 38 reactions in a final volume of 50 μl | 11 681 834 001 |
| | 720 U, 2 x 360 U, 190 reactions in a final volume of 50 μl | 11 681 842 001 |
| | 3,600 U, 10 x 360 U, 950 reactions in a final volume of 50 μl | 11 759 060 001 |
| Taq DNA Polymerase, 5 U/µl | 100 U, 5 U/µl, 80 reactions | 11 146 165 001 |
| | 500 U, 5 U/µl, 400 reactions | 11 146 173 001 |
| | 4 x 250 U, 5 U/µl, 800 reactions | 11 418 432 001 |
| | 10 x 250 U, 5 U/µl, 2,000 reactions | 11 596 594 001 |
| | 20 x 250 U, 5 U/µl, 4,000 reactions | 11 435 094 001 |

6.4. Trademarks

EXPAND is a trademark of Roche. All other product names and trademarks are the property of their respective owners.

6.5. License Disclaimer

For patent license limitations for individual products please refer to: **List of LifeScience products**.

6.6. Regulatory Disclaimer

For life science research only. Not for use in diagnostic procedures.

6.7. Safety Data Sheet

Please follow the instructions in the Safety Data Sheet (SDS).

6.8. Contact and Support

To ask questions, solve problems, suggest enhancements or report new applications, please visit our **Online Technical Support Site**.

To call, write, fax, or email us, visit **sigma-aldrich.com**, and select your home country to display country-specific contact information.



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