

NZY Reverse Transcriptase

Catalogue number:

MB12401, 20000 U
MB12402, 100000 U

Description

NZY Reverse Transcriptase is a modified recombinant form of the Moloney Murine Leukemia Virus (M-MuLV) Reverse Transcriptase purified from *Escherichia coli*. The enzyme has been modified in order to promote stability. NZY Reverse Transcriptase synthesizes the complementary DNA strand in the presence of a primer using either RNA (cDNA synthesis) or single-stranded DNA as a template in a wide range of temperatures (37-50°C). High reaction temperatures are beneficial to improve specificity and to allow synthesis of cDNA from complex templates, including those with high GC-content or with high degree of secondary structure. The enzyme lacks 3'→5' exonuclease activity and has no RNase H activity, enabling improved synthesis of full-length cDNA even for long mRNA, using random priming. Thus, NZY Reverse Transcriptase gives high yields of cDNA up to 7 kb. NZY Reverse Transcriptase can be used in first-strand cDNA synthesis experiments, RT-PCR, RT-qPCR, DNA labelling and analysis of RNA by primer extension.

Storage conditions

NZY Reverse Transcriptase should be stored at -20 °C in a freezer without defrost cycles. The protein will remain stable till the expiry date if stored as specified.

Reaction buffer (10×)

500 mM Tris-HCl, pH 8.3, 750 mM KCl, 30 mM MgCl₂, 100 mM DTT. Upon thawing, if any precipitate is observed, pulse vortex until the precipitate is completely resuspended.

Unit definition

One unit is defined as the amount of enzyme necessary to catalyse the incorporation of 1 nmol of dTTP into acid-insoluble material in 10 min at 37 °C, using poly(A)×oligo(dT)₁₂₋₁₈ as a template-primer.

Enzyme concentration: 200 U/μL

Inhibition and Inactivation

NZY Reverse Transcriptase is inhibited in the presence of metal chelators (e.g. EDTA), inorganic phosphate, pyrophosphate and polyamines. The enzyme is inactivated at 98 °C for 5 min.

Protocol for first-strand cDNA synthesis

Since RNA is very susceptible to degradation, a correct handling and storage of this nucleic acid is essential. Special precautions should be taken to avoid RNase contamination.

1. On ice, in a sterile, nuclease-free microcentrifuge tube, prepare a reaction mixture, combining the following components:

total RNA; or mRNA / poly(A) RNA	10 pg – 5 μg 10 pg – 0.5 μg
Oligo(dT) ₁₂₋₁₈ (50-60 μM); or random hexamer (50-250 ng/μL); or gene-specific primer (2 μM)	1 μL (2.5-3 μM final conc.); (or 2.5-12.5 ng/μL final conc.); (or 0.1 μM final conc.)
dNTP Mix (10 mM each)	1 μL (0.5 mM final conc.)
Nuclease-free water	up to 16 μL

2. For some GC-rich RNAs or nucleic acids with high degree of secondary structure, a denaturation step may be required. If so, centrifuge briefly and incubate the mix at 65 °C for 5 min. Chill on ice for at least 1 min, briefly centrifuge again and place on ice.

3. Add the following reaction components:

10× Reaction Buffer	2 μL
NZY Ribonuclease Inhibitor (not provided)*	1 μL (40 units)
NZY Reverse Transcriptase	1 μL (200 units)
FINAL Volume	20 μL

(* NZY Ribonuclease Inhibitor (Cat. No. MB084) will protect RNA, improve total cDNA yields and increase the percentage of full-length cDNA. Its addition is indispensable when amount of RNA template is ≤ 80 ng.

4. Mix gently and centrifuge briefly.

5. Incubate at 50 °C for 30-50 min.

Note: When using random-hexamer primers, incubate first at 25 °C for 10 min and then proceed with the 30-50 min reaction step (step 5).

6. Inactivate the reaction by heating at 85 °C for 5 min, and then chill on ice.

7. Store cDNA product at -20 °C or proceed to downstream applications.

Important notes

- cDNA can be stored at -20 °C or at 4 °C for up to one week. If long-term storage is required, -70 °C is recommended. Avoid freeze/thaw cycles of the cDNA.
- The resulting cDNA can be used for cloning or as a template in PCR or qPCR reactions. Typically, 10% (2 μL) of the first-strand reaction is enough for most PCR applications. Optionally, the cDNA can be diluted in TE buffer.
- When using cDNA in PCR amplification, some targets (> 1 kb) may require RNA-free DNA as template. To remove RNA bond to cDNA, add 1 μL (5 U) of NZY RNase H (Cat. No. MB085) and incubate at 37 °C for 20 min. This procedure will increase the sensitivity of the PCR step.

Quality control assays

Purity

NZY Reverse Transcriptase is >90% pure as judged by SDS polyacrylamide gel electrophoresis followed by BlueSafe staining.

Nucleases assays

To test for DNase contamination, 0.2-0.3 μg of pNZY28 DNA are incubated with 200 U of NZY Reverse Transcriptase for 14-16 h at 37 °C. To test for RNase contamination, 1 μg of RNA is incubated with 200 U of the enzyme for 1 h at 37°C. Following incubation, the nucleic acids are visualized on a GreenSafe-stained agarose gel. There must be no visible nicking or cutting of the nucleic acids. Similar tests are performed with reaction buffer.

Functional assay

NZY Reverse Transcriptase and respective buffer are tested for performance in a RT-qPCR experiment: a 10-fold serial dilution of total RNA from mouse liver (1 µg to 0.1 ng) is reverse transcribed using 200 units of enzyme; the resultant cDNA is then used as template in a quantitative real-time PCR assay using specific primers to amplify the mouse GAPDH gene.

Related products

Product name	Cat. No.
NZY Ribonuclease Inhibitor	MB084
NZY RNase H (<i>E. coli</i>)	MB085
Oligo (dT) ₁₈ primer mix	MB12801
Random hexamer mix	MB12901
dNTPs NZYMix	MB086
Agarose (routine grade)	MB144
NZY qPCR Green Master Mix (2x), no ROX/ ROX / ROX plus	MB221 / MB220 / MB219

Troubleshooting

Little or no RT-PCR/RT-qPCR amplification product

- RNA damage or degradation

Analyse RNA on a denaturing gel to verify integrity. Use aseptic conditions while working with RNA to prevent RNase contamination. Ensure the use of NZY Ribonuclease Inhibitor; the addition of this inhibitor is essential when using less than 50 ng of RNA in order to safeguard the template against degradation due to ribonuclease contamination. Replace RNA if necessary.

- Presence of RT inhibitors

Some inhibitors of RT enzymes include: SDS, EDTA, glycerol, sodium phosphate, spermidine, formamide and guanidine salts. They can be problematic in smaller reaction volumes. If necessary, remove inhibitors by ethanol precipitation of the RNA preparation before use; wash the pellet with 70% (v/v) ethanol.

- Not enough starting RNA

Increase the concentration of starting RNA.

Unexpected bands after electrophoretic analysis of amplified products

- Genomic DNA contamination

DNase I may be used to eliminate genomic DNA contamination from the starting RNA (pre-treatment RNA). The enzyme volume should not exceed 10% of the total reaction volume.

V2001

Certificate of Analysis

Test	Result
Enzyme purity	Pass
Nucleases assays	Pass
Functional assay	Pass

Approved by:



Patrícia Ponte
Senior Manager, Quality Systems

For research use only.

