



roboklon

# T7 RNA Polymerase

(Bacteriophage T7 of Escherichia coli)

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Modified T7 RNA Polymerase with higher tolerance towards modified nucleotides. Extremely useful for radioactive and non radioactive labeling as well as for RNA synthesis for preparative scale.

#### Description:

 Cat. No.
 Size

 E1290-01
 5 000 units

 E1290-02
 25 000 units

#### Unit Definition:

One unit is the amount of enzyme required to incorporate 1 nmol of labeled UTP into acid-soluble material in 1 hr at  $37^{\circ}$ C.

#### Storage Conditions:

Store at -20°C



#### Fig. T7 RNA transcription of 400 nt RNA.

5  $\mu$ l of T7 transcription were mixed with 5  $\mu$ l of 2 x RNA loading buffer and loaded on a 7 % polyacrylamide gel supplemented with 8 M urea. The gel was ethidium bromide stained. 1-5, respectively 50, 100, 200, 400, 800 U of T7 RNA Polymerase used for T7 transcription.

## ption:

- ➔ DNA-dependent RNA polymerase which has stringent specificity for T7 phage promoters sequence (1).
- → Ultrapure recombinant enzyme.
- ➔ Efficiently synthesizes *in vitro* transcripts from almost any DNA that is downstream from a T7 promoter (2).
- Suitable for preparing labeled single-stranded RNA probes of high specific activity(3).
- ➔ Transcripts can be used as hybridization probes, templates for in vitro translation, substrates in RNA processing systems, or exon and intron mapping of genomic DNA.

#### T7 in vitro transcription, Example reaction protocol:

Component	Final Concentr. / amount	Add Per Reaction
5 x Reaction Buffer	1 x	10 µl
NTP mix [25 mM each]	1.875 mM per NTP	3.75 μl*
DTT [100 mM]	2.5 mM	1.25 µl*
Thermostable Pyrophosphatase [20 U/µI] (Cat. No. E1267)	2.4 U	0.12 µl*
DNA template for T7 transcription	2 µg	Variable
T7 RNA Polymerase	50 - 800 U**	Variable
RNase-free H₂O		@ 50 µl

\* dependent on concentration of stock solution

\*\*50U is most efficient for labeling, more units are recommended for preparative scale

Incubate up to 2 hours at 37°C and check transcription on appropriate denaturing polyacrylamide gel. Load 5  $\mu$ l of reaction mixed with 5  $\mu$ l of 2 x RNA loading buffer (2.6 M urea, 2 x TBE, 0.02 % (w/v) BPB, 0.02 % (w/v) XCB, 66 % (v/v) formamid).

#### Storage Buffer:

20 mM potassium phosphate (pH 7.5), 0.1 M NaCl, 1 mM EDTA, 1 mM dithiothreitol, 100  $\mu g/ml$  bovine serum albumin and 50 % (v/v) glycerol.

#### 5 x Reaction Buffer:

0.4~M HEPES (pH 7.5), 16~mM MgCl2, 5~mM spermidine, 0.6~mg/ml bovine serum albumin.

#### Assay Conditions:

40 mM Tris-HCl (pH 7.9 at 22°C), 8.0 mM MgCl<sub>2</sub>, 5 mM dithiothreitol, 4 mM spermidine-(HCl)<sub>3</sub>,2.5 µg T7 DNA, 0.4 mM each of ATP, CTP, GTP and 0.4 mM [ $\alpha$ -<sup>32</sup>P]UTP. Incubation is at 37°C for 10 min in a reaction volume of 50 µl.

#### Quality Control:

All preparations are assayed for contaminating exonuclease, endonuclease and nonspecific RNase and single- and double-stranded DNase activities. Typical preparations are greater than 90 % pure, as judged by SDS polyacrylamide gel electrophoresis.

#### References:

- 1. Chamberlin, M. and Ring, J. (1973) J. Biol. Chem. 248, 2235-2244.
- 2. Tabor, s and Richardson, C.C. (1985) Proc. Natl. Acad. Sci. U.S.A. 82, 1074-1078.
- Sambrook, J., Fritsch, E. F. and Maniatis, T. (1989) Molecular Cloning: A Laboratory Manual, second edition, pp. 10.27-10.37, Cold Spring Harbour Laboratory, Cold Spring Harbour.

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