CIRCORRI
GIBCODIC

M-MLV Reverse Transcriptase

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Cat. No. 28025 -013 Lot No. _____40,000 units; 200 U/ μl Exp. Date: _____. Store at -20 °C (not frost-free).

Description :

 Moloney
 Murine Leukemia Virus Reverse
 Transcriptase (M-MLV RT) uses singlestranded RNA or DNA in the presence of a primer to synthesize a complementary

 DNA strand. This enzyme is isolated (1) from gene of M-MLV on a plasmid (2, 3).
 E. coli expressing a portion of the *pol pol*

 Components :
 28025 - 013 M-MLV RT
 Lot No.
 Lot No.

 Y00146
 5X First Strand Buffer
 Lot No.
 Y00147

Unit Defin ition :

<u>Storage Buffer</u>: 20 mM Tris-HCl (pH 7.5) 1 mM DTT 0.01% (v/v) Nonidet-P40 0.1 mM Na 2EDTA 0.1 M NaCl 50% (v/v) glycerol Quality Control Assays : See Back Page

Doc. Rev.: 011397

This product is distributed for laboratory research use only. CAUTION: Not for diagnostic use. The safety and efficacy of this product in diagnostic or other clinical uses has not been established.

For technical questions about this product, call the Life Technologies TECH-LINE^{sed} (800) 828-6686].

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 Quality Control Assays
 :

 This product has passed the following quality control assays:

 SDS-polyacrylamide gel analysis for purity; functional absence of endodeoxyribonuclease, 3 ' and 5 ' exodeoxyribonuclease, and ribonuclease activities; yield and length of cDNA product.

 $\label{eq:stress} Store the 5X First Strand Buffer and 0.1M DTT at -20 $$ ^C. Thaw the solutions at room temperature just prior to use and refreeze immediately. $$$

The enclosed buffers were assayed with the enzyme and met quality control specifications.

 Functional Assav Conditions
 :

 50 mM Tris-HCl (pH 8.3)
 75 mM KCl

 10 mM DTT
 3 mM MgCl 2

 0.5 mM each dGTP, dATP, dTTP, dCTP

 (1-10 µCi of [Ct⁻³²P] dCTP added as a tracer)

 10 µg/ml oligo (dT) 12-18

 20 µg/ml mRNA

 200 units M-MLV RT

 Reaction Volume: 20 µl

 Incubation: 60 minutes at 37 °C

Use 200 units of M-MLV RT per $\mbox{$\mu$}$ g of RNA (5) in a standard assay. However, reaction volume and amounts of enzyme and \$\$mRNA\$ should be tailored to the second-strand method.

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References:

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 Houts, G. E., Miyagi, M., Ellis, C., Beard, A., and Beard, J. W. (1979) J. Virol. 29, 517.
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