



Mbo II

Recognition Sequence:

S E471 200 units 5.000 u/ml GAAGA(N)₈↓
CTTCT(N)₇↑

Lot: Exp:

Store at -20C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

 %Activity
 50-75
 75-100
 25-50
 50-75
 100
 50

 37°C
 65°C
 Y
 λ
 RR
 Dam

For more details scen the code



CERTIFICATE OF ANALYSIS

<u>Source</u>: An E.coli strain that carries the cloned Mbo II gene from Moraxella bovis.

Supplied in:

10 mM Tris-HCl (pH 7.5), 50 mM NaCl, 0.1 mM EDTA, 1 mM DTT, 200 μg/ml BSA, 50% glycerol.

Reaction Conditions:

1X SE-Buffer Y. Incubate at 37° C.

1X SE-Buffer Y (pH 7.9 @ 25° C):

33 mM Tris-Ac 66 mM KAc 10 mM MgAc 1 mM DTT

Heat Inactivation:

Enzyme is inactivated by incubation at 65°C for 20 minutes.

<u>Unit Definition</u>:One unit is defined as the amount of enzyme required to digest 1 μ g of Lambda DNA (Dam-) in 1 hour at 37° C in a total reaction volume of 50 μ l.

Quality Control Assays

 $\frac{16 \hbox{-Hour Incubation:}}{\text{DNA and 5 Units of enzyme incubated for 16 hours}} \text{ and 5 Units of enzyme incubated for 16 hours} \\ \text{resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.}$

Oligonucleotide Assay:No detectable degradation of a single-stranded and double-stranded oligonucleotide was observed after incubation with 5 units of restriction endonuclease for 3 hours.

Enzyme Properties:

When using a buffer other than the optimal (Supplied) SE-Buffer, it may be necessary to add more enzymes to achieve complete digestion.

Reagents Supplied with Enzyme: 10X SF Buffer Y.

Blocked by overlapping Dam methylation ($G^{m}ATC$): $GAA\underline{GATC}$.