



Mab I

Recognition Sequence:

S E12

200 units 5,000 u/ml

ALCCWGGT TGGWCC1A

> Lot: Exp:

Store at -20°C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

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

37°C





For more details scen the code



CERTIFICATE OF ANALYSIS

Source: Microbacterium arborescens SE.

Supplied in:

10 mM Tris-HCl (pH 7.5), 100 mM NaCl, 0.1 mM EDTA, 7 mM 2-mercaptoethanol, $200~\mu g/ml$ BSA, 50% glycerol.

Reaction Conditions:

1X SE-Buffer W, BSA (100 $\mu g/ml).$ Incubate at 37° C.

1X SE-Buffer W (pH 8.5 @ 25° C):

10 mM Tris-HCl 100 mM NaCl 10 mM MgCl, 1 mM DTT

Heat Inactivation:

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

Quality Control Assays

<u>Ligation</u>:After 5-fold overdigestion with Mab I, > 90% of the DNA fragments can be ligated and recut.

16-Hour Incubation: A 50 µl reaction containing 1 µg of DNA and 10 Units of enzyme incubated for 16 hours resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.

No using BSA for long incubation.

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 SE Buffer W, BSA (10 mg/ml).

Blocked by overlapping Dcm methylation (G^mCWGG): ACCWGGT.