



BstMB I

Recognition Sequence:

S

500 units 10,000 u/ml

E119

↓GATC CTAG↑

Lot: Exp:

Store at -20°C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

 %Activity
 10-25
 25-50
 100
 75-100
 10-25
 100

For more details scen the code



CERTIFICATE OF ANALYSIS

Source: Bacillus stearothermophillus MB.

Supplied in:

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

Reaction Conditions:

1× SE-Buffer O. Incubate at 65° C.

 $\underline{\text{1X SE-Buffer O (pH 7.6 @ 25°C):}}$

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

Heat Inactivation:

Enzyme is inactivated by incubation at 80°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 65° C in a total reaction volume of 50 μ l.

Quality Control Assays

<u>Ligation</u>:After 5-fold overdigestion with BstMB I, more than 95% of the DNA fragments can be ligated and recut.

 $\underline{16\text{-Hour Incubation:}}A~50~\mu I$ reaction containing 1 μg of DNA and 20 Units of enzyme incubated for 16 hours 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 10 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 O.

Blocked by overlapping Dam methylation ($G^{m}ATC$): <u>GATC</u>.

Not blocked by CG methylation.

Not cut hemi methylated site: 5` - G(6mA) TC-3` /
3` -CTAG-5`