



# **Bmt I**

Recognition Sequence:

S E4

**E457**1,000 units
20,000 u/ml

GCTAG↓C C↑GATCG

Lot: Exp:

Store at -20C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

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

37°C





For more details scen the code



## **CERTIFICATE OF ANALYSIS**

<u>Source</u>: An E.coli strain that carries the cloned Bmt I gene from Bacillus megaterium S2.

#### Supplied in:

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

## **Reaction Conditions:**

1X SE-Buffer W. Incubate at 37° C.

## <u>1X SE-Buffer W(pH 8.5 @ 25° C)</u>:

10 mM Tris-HCl 100 mM NaCl 10 mM MgCl<sub>2</sub> 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  $\mu$ g of Lambda DNA/HindIII in 1 hour at 37° C in a total reaction volume of 50  $\mu$ l.

### **Quality Control Assays**

<u>Ligation</u>:After 20-fold overdigestion with Bmt I, ~95% of the DNA fragments can be ligated and recut.

 $\underline{16\text{-Hour Incubation:}}A~50~\mu\text{l}$  reaction containing 1  $\mu\text{g}$  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 20 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 W.

Bmt Lis an isoschizomer of Nhe L.

The minimum number of units that resulted in complete digestion of 1  $\mu g$  of substrate DNA in 16 hours is 0.13. Bmtl cleaves linear plasmid DNA at a rate 5 times higher than supercoiled plasmid DNA.