



# BstSC I

Recognition Sequence:

S E307
100 units
3.000 u/ml

↓CCNGG GGNCC↑

Lot: Exp:

50-75

Store at -20C

100

60

SE-Buffers B G O W Y ROSE

50-75

55°C 80°C Υ λ

50-75

50-75

For more details scen the code



## **CERTIFICATE OF ANALYSIS**

Source: Bacillus stearothermophilus SC.

### Supplied in:

 $\overline{10}$  mM Tris-HCl (pH 7.5), 50 mM KCl, 0.1 mM EDTA, 7 mM 2-mercaptoethanol, 200 µg/ml BSA, 50% glycerol.

#### **Reaction Conditions:**

1× SE-Buffer Y. Incubate at 55 °C.

## <u>1X SE-Buffer Y (pH 7.9 @ 25° C)</u>:

33 mM Tris-Ac 66 mM KAc 10 mM MgAc 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  $\mu$ g of Lambda DNA (Dcm-) in 1 hour at55° C in a total reaction volume of 50  $\mu$ l.

#### Quality Control Assays

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

16-Hour Incubation: A 50 µl reaction containing 1 µg of DNA and 3 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 3 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 Y.

Blocked by Dcm-methylation (C<sup>m</sup>CWGG): CCWGG.

At37° C activity is 10% from maximum.