



Sma I

Recognition Sequence:

S E17

2,000 units 20.000 u/ml

GGGT CCC

Lot: Exp:

Store at -20C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

 %Activity
 0-10
 0-10
 0-10
 100
 50

25°C 65°C Y

For more details scen the code

Ph/F+7(383)333-6853 info@sibenzyme.com www.sibenzyme.com

minimal

RR

CERTIFICATE OF ANALYSIS

<u>Source</u>: An *E.coli* strain that carries the cloned Sma I gene from *Serratia marcescens*.

Supplied in:

10~mM Tris-HCl (pH 7.6), 50~mM NaCl, 0.1~mM EDTA, $200~\mu\text{g/ml}$ BSA, 1~mM DTT, 50% glycerol.

Reaction Conditions:

1x SE-Buffer Y. Incubate at 25° 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 $^{\circ}$ C for 20 minutes.

<u>Unit Definition</u>: One unit of the enzyme is the amount required to hydrolyze 1 μ g of λ DNA (Hind III– digest) in 1 hour at 25° C in a total reaction volume of 50 μ l.

Quality Control Assays

 $\frac{Ligation}{approximately 90\% of the DNA fragments can be ligated with T4 DNA Ligase and recut. In the presence of 10 \% PEG ligation is better.$

16-Hour Incubation: A 50 µl reaction containing in 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 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 Y.