



# Sma I

Recognition Sequence:

E177m XS

20.000 u/ml

500 units

CCCTGGG GGGT CCC

Lot:

Exp:

Store at -20C

minimal

SE-Buffers	В	G	0	w	Υ	ROSE
%Activity	0-10	0-10	0-10	0-10	100	50

For more details scen the code



## CERTIFICATE OF ANALYSIS

Source: 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 µg/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 1 mM DTT 10 mM MgAc

### Heat Inactivation:

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

Unit Definition: One unit of the enzyme is the amount required to hydrolyze 1  $\mu$ g of  $\lambda$  DNA (Hind III- digest) in 1 hour at 25° C in a total reaction volume of 50 µl.

#### Quality Control Assays

Ligation: After 20-fold overdigestion with Smal, 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.