



Set I

Recognition Sequence:

S E537 200 units 5.000 u/ml ASST↓ †TSSA

Lot: Exp:

Store at -20C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

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

55°C 80°C Y (dsDNA) R

For more details scen the code



CERTIFICATE OF ANALYSIS

<u>Source</u>: An E.coli strain that carries the cloned Set I gene from Streptomyces werraensis 37.

Supplied in:

10 mM Tris-HCl (pH 7.6), 100 mM NaCl, 0.1 mM EDTA, 1 mM DTT, 50% glycerol.

Reaction Conditions:

1X SE-Buffer Y. Incubate at 55° 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 80°C for 20 minutes.

Unit Definition: One unit is defined as the amount of enzyme required to cleave 1 pmol of the double-stranded oligonucleotide of the below indicated structure in 1 hour at 55° Cin a total reaction volume of 20 μ l.

Quality Control Assays

 $\underline{\text{Ligation}}. \text{After 5-fold overdigestion with enzyme, \sim50\%$ of the pBR322 DNA fragments can be ligated with T4 DNA Ligase and recut.}$

16-Hour Incubation: A 50 μl reaction containing 1 μg of DNA and 5 Units of enzyme incubated for 16 hours resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.

Note!

SetI cleaves a canonical site and several other sites with a weaker activity.

In the case of long incubation with SetI DNA can be digested to small oligos.

Oligonucleotide Assay:No detectable degradation of a single-stranded and double-stranded oligonucleotide was observed after incubation with 5 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.