



# Abs I

Recognition Sequence:

**S** E535 50 units

1.000 u/ml

75-100

CC\$TCGAGG GGAGCTTCC

> Lot: Exp:

Store at -20°C

G O W Y ROSE
10-25 0 50-75 0-10 50

37°C

For more details

scen the code

SE-Buffers



osi (puci

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# **CERTIFICATE OF ANALYSIS**

Source: Arthrobacter species 7M06.

### Supplied in:

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 Absl. Incubate at 37° C.

1X SE-Buffer Absl (pH 9.0 @ 25° C):

 $\begin{array}{ll} 10~\text{mM Tris-HCl} & 50~\text{mM KCl} \\ 10~\text{mM MgCl}_2 & 1~\text{mM DTT} \end{array}$ 

#### **Heat Inactivation:**

Enzyme is inactivated by incubation at 65  $^{\circ}\text{C}$  for 20 minutes.

<u>Unit Definition</u>: One unit is defined as the amount of enzyme required to digest  $1 \mu g$  of pUC19SE/Dril DNA in 1 hour at  $37^{\circ}$  C in a total reaction volume of  $50 \mu l$ .

#### **Quality Control Assays**

 $\frac{Ligation}{Ligation}: After 2-fold overdigestion with Abs I, \sim 90\% of the DNA fragments can be ligated with T4 DNA Ligase and recut.$ 

 $\underline{16\text{-Hour Incubation}}$ : A 50  $\mu l$  reaction containing 1  $\mu g$  of DNA and 2 Units of enzyme incubated for 16 hours resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.

A long incubation time may result in star activity.

Oligonucleotide Assay: No detectable degradation of a single-stranded and double-stranded oligonucleotide was observed after incubation with 1 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 Absl.