



# Xba I

Recognition Sequence:

S

2,000 units 20.000 u/ml

E141

T‡CTAGA AGATC†T

Lot: Exp:

Exp:

Store at -20C

SE-Buffers	В	G	0	w	Υ	ROSE
%Activity	75-100	75-100	100	50-75	75-100	25
37°C 65°C O (\(\lambda\)/HindIII (\(\mathbb{RR}\)(\mathbb{RSA})(\(\mathbb{Dam}\)						

For more details scen the code



## **CERTIFICATE OF ANALYSIS**

<u>Source</u>: An E.coli strain that carries the cloned Xba I gene from Xanthomonas badrii.

#### Supplied in:

10 mM Tris-HCl (pH 7.5), 50 mM NaCl, 0.1 mM EDTA, 1 mM DTT, 50% glycerol.

### **Reaction Conditions:**

1X SE-0, BSA (100 μg/ml). Incubate at 37° C.

 $\underline{\text{1X SE-Buffer 0 (pH 7.6 @ 25°C)}}:$ 

 $\begin{array}{ccc} 50~\text{mM Tris-HCl} & 100~\text{mM NaCl} \\ 10~\text{mM MgCl}_2 & 1~\text{mM DTT} \end{array}$ 

#### **Heat Inactivation:**

Enzyme is inactivated by incubation at 65°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 (Dam-)/HindIII in 1 hour at 37° C in a total reaction volume of 50  $\mu$ l. To obtain 100% activity, BSA should be added to the 1 x reaction mix to a final concentration of 100  $\mu$ g/ml.

#### **Quality Control Assays**

endonuclease for 3 hours.

 $\label{eq:Ligation} \begin{tabular}{ll} $Ligation: After 20-fold overdigestion with Xba I, $\sim\!90\%$ of the DNA fragments can be ligated and recut. \end{tabular}$ 

16-Hour Incubation: A 50 µl reaction containing 1 µg of DNA and 40 Units of enzyme incubated for 16 hours resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.

Do not use BSA for long incubation.

Oligonucleotide Assay: No detectable degradation of a single-stranded and double-stranded oligonucleotide was observed after incubation with 20 units of restriction.

#### **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 O. BSA (10mg/ml).

Blocked by overlapping Dam-methylation (G  $^{\rm m}$  ATC): TCTA  $\underline{\rm GATC}$