



## Fau I

Recognition Sequence:

E209

100 units 2,000 u/ml

100

CCCGC(N)41
GGGCG(N)61

Lot: Exp:

Store at -20°C

G O W Y ROSE

RR

**55°C 65°C B** λ

For more details scen the code

SE-Buffers

%Activity



### **CERTIFICATE OF ANALYSIS**

<u>Source</u>: An E.coli strain that carries the cloned Fau I gene from Flavobacterium aquatili.

#### Supplied in:

10 mM Tris-HCl (pH 7.5), 50 mM NaCl, 0,1 mM EDTA, 1 mM DTT, 200 µg/ml BSA, and 50% glycerol.

#### **Reaction Conditions:**

1X SE-Buffer B. Incubate at 55° C.

<u>1X SE-Buffer B (pH 7.6 @ 25° C</u>):

10 mM Tris-HCl 10 mM MgCl<sub>2</sub> 1 mM DTT

#### **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 in 1 hour at 55°C in a total reaction volume of 50  $\mu$ l.

#### **Quality Control Assays**

<u>Ligation</u>:After 2-fold overdigestion with Fau I, more than 90% of the DNA fragments can be ligated and of these 95% can be recut.

<u>16-Hour Incubation:</u> A 50  $\mu$ l reaction containing 1  $\mu$ g of DNA and 4 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 2 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 B.

Blocked by CG methylation.

The minimum number of units that resulted in complete digestion of 1  $\mu$ g of substrate DNA in 16 hours is 0.5.