



HspA I

Recognition Sequence:

E070 5,000 units

CGCTG CJCGC

Lot: Exp:

Store at -20C

 SE-Buffers
 B
 G
 O
 W
 Y
 ROSE

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

20.000 u/ml

For more details scen the code



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CERTIFICATE OF ANALYSIS

<u>Source</u>: An E.coli strain that carries the cloned HspA I gene from Haemophilus species A1.

Supplied in:

10 mM Tris-HCl (pH 7.5), 50 mM NaCl, 0.1 mM EDTA, 7 mM 2-mercaptoethanol, 200 µg/ml BSA, 50% glycerol.

Reaction Conditions:

1X SE-Buffer Y. Incubate at 37° 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.

Quality Control Assays

<u>Ligation</u>:After 5-fold overdigestion with HspA I, \sim 90% of the DNA fragments can be ligated and recut.

 $\underline{16\text{-Hour Incubation}}\text{:A 50}\ \mu\text{I}\ reaction containing 1}\ \mu\text{g of DNA and 20 Units of enzyme incubated for 16 hours resulted in the same pattern of DNA bands as a reaction incubated for 1 hour.}$

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.

Blocked by CG methylation 5'-G(5mC) GC-3'/3'-CG (5mC) G-5'.

Not blocked by methylation 5'-GCG(5mC)-3'/3'-CGCG-5'.

Cut hemi methylated site: 5'- G(5mC) GC-3'/3'- CGCG-5`

HspA I is an neoschizomer of Hha I.