

BamH I

Restriction

Endonuclease

GLGATCC Recognition Sequence: CCTAGTG E949m XS

2000 units 20.000 u/ml Lot: Exp: Store at -20C

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SE-Buffers ROSE 25-50 25-50 10-25 10-25 100 40

BSA

Ph/F+7(383)333-6853 For more details info@sibenzyme.com scen the code

CERTIFICATE OF ANALYSIS

Source: An E.coli strain that carries the cloned BamH I gene from Bacillus amyloliquefaciens H.

Supplied in:

10 mM Tris-HCl (pH 7.4), 50 mM KCl, 0.1 mM EDTA, 1 mM DTT, 200 µg/ml BSA, 50% glycerol.

Reaction Conditions:

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

1X SE-Buffer Y (pH 7.9 @ 25° C): 33 mM Tris-Ac 66mM KCl

10 mM MqAc 1 mM DTT

Heat Inactivation:

NO (80 °C for 20 minutes).

Unit Definition: One unit is defined as the amount of enzyme required to digest 1 µg of Lambda DNA in 1 hour at 37°C in a total reaction volume of 50 µl.

To obtain 100% activity, BSA should be added to the 1 x reaction mix to a final concentration of 100 μg/ml.

Quality Control Assays Ligation: After 20-fold overdigestion with BamH I, ~90%

of the DNA fragments can be ligated with high -activity T4 DNA Ligase and recut.

incubated for 1 hour.

endonuclease for 3 hours.

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

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 Y, BSA (10 mg/ml).

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