Restriction Endonuclease

Gla I

For more details

scen the code

R

| ecognition | Pu(5mC)↓GPy | | |
|-------------|--------------|--|--|
| equence: | PyG1(5mC)Pu | | |
| E494 | Lot: | | |
| 5,000 units | Exp: | | |
| 50,000 u/ml | Store at -20 | | |

| SE-Buffers | В | G | 0 | w | Y | ROS |
|------------|--------|--------|-------|-------|-----|-----|
| %Activity | 75-100 | 75-100 | 25-50 | 25-50 | 100 | 100 |

pHspAI2/

Gsal

CERTIFICATE OF ANALYSIS

Source: An E.coli strain that carries the cloned Gla I gene from Glacial ice bacterium Gl29.

Supplied in:

SibEnzyme®

Store at -20°C

RR

Ph/F+7(383)333-6853

info@sibenzyme.com

www.sibenzvme.com

10 mM Tris-HCl (pH 7.6), 250 mM NaCl, 0.1 mM EDTA, 0.1% Triton X-100, 7 mM 2-mercaptoethanol, 50 µg/ml BSA, 50% glycerol.

Reaction Conditions: 1X SF-Buffer Y. Incubate at 30°C.

1X SE-Buffer Y (pH 7.9 @ 25° C): 33 mM Tris-Ac 66 mM KAc 1 mM DTT 10 mM MqAc

Heat Inactivation:

Enzyme is inactivated by incubation at 65°C for 20 minutes.

Unit Definition: One unit is defined as the amount of enzyme required to digest completely a unique 5`-G (5mC)G(5mC)-3³/3⁻(5mC)G(5mC)G-5^{*} site in 1 µg of pHspAI2 plasmid DNA, which is linearized with Gsal, in 1 hour at 30° C in a total reaction volume of 50 µl. DNA pHspAI2/Gsal is a linearized plasmid pHspAI2, which carries a gene of DNA-methyltransferase M. HspAI (recognition sequence 5'-GCGC-3') and includes a unique Glal recognition site 5`-G(5mC)G(5mC)-3`/3`-(5mC)G(5mC)G-5`.

Quality Control Assays

16-Hour Incubation: No detectable degradation of 1µg of Lambda DNA was observed after incubation with 100 units of enzyme for 16 hours at 30°C in a total reaction volume of 50 µl.

Oligonucleotide Assay:No detectable degradation of a single-stranded and double-stranded oligonucleotide was observed after incubation with 50 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 diaestion.

Substrate specificity:

The enzyme cleaves only C5-methylated DNA and does not cut unmodified DNA and DNA with N4-methylcytosines The enzyme activity depends on number and position of methylated nucleotides in the recognition sequence: Optimal substrate (100% activity) 5`-G(5mC)G(mC)-3`/3`-(5mC)G(5mC)G-5` Good substrates (> 25% activity) 5`-R(5mC)G(5mC)-3`/3`-YG(5mC)G-5` 5`-R(5mC)GY-3`/3`-YG(5mC)R-5` Medium substrates (> 6% activity) 5`-G(5mC)R(5mC)-3`/3`-(5mC)GYG-5` 5`-R(5mC)GY-3`/3`-YG(5mC)R-5` Bad substrates (6% activity) 5`-G(5mC)GC-3`/3`-CG(5mC)G-5`

Reagents Supplied with Enzyme: 10X SE Buffer Y, DNA pHspAl2/Gsal.