





DpnI

Dpn I

Restriction Endonuclease

Recognition Sequence:

5'-GATC-3' 3'-CTAG-5'

 Cat. No.
 Size

 E2135-01
 500 units

 E2135-02
 2 500 units

Reaction Temperature: 37°C

Inactivation Temperature (20 min): 80°C

Prototype / Isoschizomer: Dpnl

Source: Diplococcus pneumoniae Purified from *E.coli* strain that carries the DpnI gene

from Diplococcus pneumoniae.

Package Contents:

→ Dpnl

→ 10x Reaction Buffer ONE

Note 1: DpnI cleaves only methylated GATC sites.

Storage Conditions: Store at -20°C

Double Digestion - Buffer Compatibility:

ONE Buffer is compatible with most EURx restriction

The relative activity of DpnI in Pfu (Cat.No. E1114) and PfuPlus! (Cat.No. E1118) DNA Polymerase buffer is approx. 50 %.

Recommended Buffer: ONE (or compatible third party buffers)

100 μ g/ml BSA neither inhibits nor promotes Dpnl cleavage.

DNA Methylation Impact on Digestion:

No inhibition: All GATC sites with N⁶-Methyladenine.

Standard Reaction Protocol:

Mix the following reaction components: 1-2 µg pure DNA with methylated GATC sites 3 µl 10x Buffer ONE

1-2 U DpnI (use 1 U / µg DNA, < 10 % React. Volume!)
Tips: Add enzyme as last component. Mix components well before adding enzyme. After enzyme addition, mix gently by pipetting. Do not vortex. High (excess) amounts of enzyme can greatly speed up the reaction.

@ 30 μI $H_2O,$ DNA and DNase free

Incubate for 1 h at 37°C

(Agarose-embedded, RE buffer equilibrated DNA: 16 h)

Stop reaction by alternatively

(b) Heat Inactivation 20 min at 80°C or

(c) Spin Column DNA Purification

(e.g. EURx PCR/DNA CleanUp Kit, Cat.No. E3520) or (d) Gel Electrophoresis and Single Band Excision (e.g. EURx AgaroseOut DNA Kit, Cat.No. E3540) or (e) Phenol-Chloroform Extraction or Ethanol Precipitation.

Non-optimal buffer conditions:

To compensate for the lack of enzyme activity, increase the amount of enzyme and $\ / \$ or reaction time accordingly. The following values may serve as orientation:

- → Enzyme amount: Instead of 1 U enzyme, use ~4 U of enzyme in buffers providing 25 % rel. activity, ~2 U in 50 %, ~1.5 U in 75 % or ~1 U in 100 %, respectively.
- → Reaction time: Increase by ~1.3-fold (75 % rel. activity), ~2 fold (50 %) or ~4 fold (25 %), respectively.

Note 2: It may be necessary to add more enzyme to obtain complete digestion when using other buffer than optimal (ONE).

Note 3: It is recommended to add 10 U of the enzyme to obtain complete digestion of the methylated template after standard site-directed mutagenesis (total reaction volume is 50 μ l).

Unit Definition:

One unit is the amount of enzyme required to completely digest 1 μg of pBR322 \emph{dam} methylated DNA in 1 hr. Total reaction volume is 30 $\mu l.$ Enzyme activity was determined in the recommended reaction buffer.

Reaction Buffer:

1 x ONE Buffer

Restriction Enzyme Compatibility:

Both, enzyme and buffers are fully compatible to restrictases and buffer systems of other manufacturers and can be used along in double digestions. To obtain best results, consult the corresponding manuals from all involved products.

Storage Buffer:

10~mM Tris-HCl (pH 7.4 at $4\,^{\circ}$ C), 1~mM dithiothreitol, 400~mM NaCl, 0.1 mM EDTA, 0.1% [v/v] Tergitol m TMN, 200 $\mu g/ml$ bovine serum albumin and 50% [v/v] glycerol.

Quality Control:

All preparations are assayed for contaminating endonuclease, 3'-exonuclease, and nonspecific single- and double-stranded DNase activities.