

Nt.BbvCI

REF: EG25520S

5'...C C T C A G C...3'
3'...G G A G T C G...5'



Storage Condition

Store at -20°C for 2 years.

Components

Component	Amount
Nt.BbvCl (5 U/µl)	30 µl
10× CutOne [®] Buffer	1 ml
10× CutOne® Color Buffer	1 ml

Description

Nt.BbvCl is a nicking endonuclease that cleaves only one strand of DNA on a double-stranded DNA substrate. It creates nicks on dsDNA substrates without cutting through the dsDNA. Nt.BbvCl is commonly used in isothermal nucleic acid amplification (e.g., Strand Displacement Amplification [SDA], Rolling Circle Amplification [RCA]). DNA nicks generated by Nt.BbvCl trigger the strand displacement reaction of DNA polymerase, and the repeated processes of cleavage, displacement, and extension enable exponential nucleic acid amplification.

Recommended Reaction Conditions

1× CutOne® Buffer;

Incubate at 37°C;

Refer to "Protocol for DNA Digestion" for reaction setup.

Heat Inactivation

Incubation at 80°C for 20 minutes.

Definition of Activity Unit

One unit is defined as the amount of enzyme required to completely convert 1 μg of supercoiled p615 DNA into the open-circular form in a 50 μl reaction system at 37°C for 1 hour.

Quality Control

Function

A 20 μ l CutOne $^{\circ}$ reaction system containing 5 U of Nt.BbvCl can convert 1 μ g of supercoiled plasmid DNA p615 into an open circular form at 37 $^{\circ}$ C for 15 minutes.

Prolonged Incubation

A 10 μ l reaction containing 5 U of Nt.BbvCI and the supercoiled p615 DNA substrate were incubated at 37°C for 16 hours, and no change in the open-circular DNA was detected by agarose gel electrophoresis.

Icon Descriptions

- [37] The enzyme's optimum reaction temperature is 37°C.
- Cleavage with this restriction enzyme may be blocked or impaired when the substrate DNA is methylated by the CpG methylase.
- EB Cleavage with this restriction enzyme may be blocked or impaired when the substrate DNA is methylated by the EcoBI methylase.
- The enzyme can be heat inactivated at by incubation 80°C for 20 minutes.

Method of application

1. Protocol for DNA Digestion

① Combine the following reaction components on ice in the order indicated:

ddH_2O	up to 50 μl
10× CutOne® Buffer or 10× CutOne® Color Buffer	5 µl
DNA ^a	1 µg
Nt.BbvCI (5 U/µI)	1 µl
Total	50 µl

- a. DNA substrates should contain no phenol, chloroform, ethanol, EDTA, detergents, or high salt concentrations, otherwise enzyme activity will be affected;
- ② Mix gently and spin down.
- 3 Incubate at 37°C for 15 min~3 h.
- ④ Optional: Inactivate the enzyme by heating at 80°C for 20 minutes, or by adsorption column or phenol/chloroform purification to terminate the reaction

2. Notice

- ① The volume of enzyme added to the reaction mixture should not exceed 10% of the total volume to avoid star activity caused by excessive glycerol in the enzyme storage buffer.
- ② The additives (e.g., glycerol, salt) in the enzyme storage buffer are the same as the contaminants in the substrate solution (e.g., salt, EDTA, or ethanol, etc.). Therefore, the smaller the reaction volume, the stronger the digestion inhibition effect.



Number of Recognition Sites in DNA

λDNA	ФХ174	pBR322	pUC57	pUC18/19	SV40	M13mp18/19	Adeno2
7	3	0	0	0	0	2	9

Methylation Effects on Digestion

Dam	Dcm	CpG	EcoKI	EcoBI
No effect	No effect	No effect	No effect	Blocked

Activity in Different Buffers*

CutOne [®] Buffer		Thermo Scientific FastDigest Buffer	NEB rCutSmart™ Buffer	Takara QuickCut™ Buffer	
Activity	100%	100%	100%	100%	

^{*}The activity data come from the functional test described above.