

BbvCl

REF: EG25516S

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







Storage Condition

Components

Component	Amount	
BbvCl (2 U/μl)	25 μΙ	
10× CutOne [®] Buffer	1 ml	
10× CutOne [®] Color Buffer	1 ml	

Description

BbvCl is derived from the BbvCl gene of Bacillus brevis (L. Ge) and is obtained through recombinant expression in Escherichia coli. BbvCI is a Type IIA restriction enzyme existing as a heterodimer composed of two distinct catalytic subunits, each with an independent cleavage site. It recognizes and cleaves the non-palindromic sequence CCTCAGC(-5/-2), generating 5' cohesive ends with 3-base overhangs. BbvCl uses CutOne® reaction buffer and is compatible with double digestion with other LightNing® restriction enzymes. However, it is advisable to avoid extended digestion (exceeding 3 hours) to prevent star activity.

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 of activity refers to the amount of enzyme required to completely digest 1 μg of λDNA in a 50 μl reaction system at 37°C for 1 hour

Quality Control Assays

Function

2 U of BbvCl can completely digest 1 μg of λDNA within 15 minutes at 37°C.

Prolonged Incubation / Star Activity Assay

Under optimal reaction temperature, incubate 2 U BbvCl with 1 μg λDNA for 3 hours. No contamination from other nucleases or nonspecific substrate degradation caused by star activity was detected. Longer incubation may result in star activity.

Ligation and Recuting

Under optimal reaction temperature, digest the substrate using 2 U BsrDI and recover the digested products. <10% of the DNA fragments can be ligated with T4 DNA Ligase at 22°C. Of these ligated fragments, >95% can be recut with BsrDI as determined 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.
- The enzyme can be heat inactivated at by incubation 80°C for 20
- 3 hours incubation did not show star activity, but delayed enzyme digestion might show star activity.

Protocol

1. Protocol for DNA Digestion

① Combine the following components on ice in the following order:			
ddH_2O	up to 50 μl		
10× CutOne [®] Buffer or 10× CutOne [®] Color Buffer	5 μΙ		
DNA ^a	1 µg		
BbvCl (2 U/µl)	1 μΙ		
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 1~3 hours.
- 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	Impaired	No effect	No effect

Activity in Different Buffers*

	CutOne [®] Buffer		NEB	Takara	
Cutoffe Bullet	FastDigest Buffer	rCutSmart™ Buffer	QuickCut™ Buffer		
Activity	100%	100%	100%	100%	

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