

# **Product Components**

Components	Component Number	2,000 U	10,000 U
Smal (20,000 U/mL)	RM21618	100 μL	500 μL
10X Buffer CutS	RM20103	1.25 mL	2 × 1.25 mL

# **Product Description**

#### **Restriction Site**

5'...C C C G G G...3' 3'...G G G C C C...5'

## **Unit Definition**

One unit is defined as the amount of enzyme required to digest 1  $\mu g$  of  $\lambda DNA$  (HindIII digest) in 1 hour at 25°C in a total reaction volume of 50  $\mu L$ .

## Storage

-20°C

#### **Reaction Conditions**

1X Buffer CutS, incubate at 25°C.

## **Activity at Temperature**

@37°C: 50%

## **1X Buffer CutS**

50 mM KAc, 20 mM Tris-HAc, 10 mM MgAc<sub>2</sub>, 100 μg/mL rHSA, pH 7.9 @ 25°C

## **Quick Cut**

Yes. This enzyme will digest unit substrate in 5-15 minutes under recommended reaction conditions.

### **Heat Inactivation**

65°C for 20 min.

## **Methylation Sensitivity**

dam methylation	dcm methylation	CpG Methylation
not sensitive	not sensitive	Blocked

### **Instructions**

## **Recommended Protocol for Digestion**

Components	50 μL		
$ddH_2O$	Up to 50 μL		
10X Buffer CutS	5 μL		
DNA*	1 μg		
Smal	1 μL		

- \* Note: DNA substrates should be free of phenol, chloroform, ethanol, EDTA, detergents or high concentrations of salt, otherwise it will affect the enzyme activity.
- ◆ The substrates are completely digested in 5-15 min incubate at 25°C.
- ◆ Smal has a half-life of 15 minutes at 37°C.
- Smal is an isoschizomer of Xmal. Smal produces bluntended fragments whereas Xmal produces a 5' extension.

## **OC Process**

- ◆ Purity is above 95% detected by SDS-PAGE.
- ◆ No exonuclease, nuclease contamination.
- ◆ No residual host genomic DNA detected by PCR.

## **Optimizing Restriction Endonuclease Reactions**

There are several key factors to consider when setting up a restriction endonuclease digest. Using the proper amounts of DNA, enzyme and buffer components in the correct reaction volume will allow you to achieve optimal digestion. Most researchers follow the "typical"



reaction conditions listed, where a 5 - 10 fold overdigestion is recommended to overcome variability in DNA source, quantity and purity.

### A "Typical" Restriction Digest

Restriction Enzyme	10 units is sufficient,	
	generally 1 µL is used	
DNA	1 μg	
10X ABclonal Buffer	5 μL (1X)	
Total Reaction Volume	50 μL	
Incubation Time	1 hr*	
Incubation Temperature	Enzyme dependent	

<sup>\*</sup> Note: Can be decreased to 5-15 minutes by using a "Quick Cut" Restriction Enzyme.

#### 1. Enzyme

 In general, we recommend 5-10 units of enzyme per μg DNA, and 10-20 units for genomic DNA in a 1 hour digest.

#### 2. DNA

 Methylation of DNA can inhibit digestion with certain enzymes.

### 3. Buffer

- Use at a 1X concentration.
- Supplement with SAM (S-Adenosyl methionine) to the recommended concentration if required.

#### 4. Reaction Volume

- A 50 μL reaction volume is recommended for digestion of 1 μg of substrate.
- Enzyme volume should not exceed 10% of the total reaction volume to prevent star activity due to excess glycerol.
- Additives in the restriction enzyme storage buffer (e.g., glycerol, salt) as well as contaminants found in the substrate solution (e.g., salt, EDTA, or alcohol) can be problematic in smaller reaction volumes.
  The following guidelines can be used for techniques that require smaller reaction volumes.

Reaction	Enzyme	DNA	10X ABclonal
System	Amount*		Buffer
10 μL**	1 U	0.1 μg	1 μL
25 µL	5 U	0.5 μg	2.5 μL
50 μL	10 U	1 µg	5 μL

<sup>\*</sup> Note: Restriction Enzymes should be diluted when smaller amounts are needed.

#### 5. Incubation Time

- Incubation time is typically 1 hour.
- Can often be decreased by using an excess of enzyme, or by using one of our "Quick Cut" restriction enzymes.
- It is possible, with many enzymes, to use fewer units and digest for up to 16 hours.

### 6. Stopping a Reaction

If no further manipulation of DNA is required:

Terminate with a stop solution (10 μL per 50 μL rxn)
[1x: 2.5% Ficoll-400, 10 mM EDTA, 3.3 mM Tris-HCl,
0.08% SDS, 0.02% Tartrazine, 0.001% Xylene Cyanol
FF, pH 8.0 @ 25°C].

When further manipulation of DNA is required:

- Heat inactivation can be used.
- Remove enzyme by using a spin column or phenol/chloroform extraction.

## 7. Control Reactions

If you are having difficulty cleaving your DNA substrate, we recommend the following control reactions:

- Control DNA (DNA with multiple known sites for the enzyme, e.g. lambda or pUC19 DNA) with restriction enzyme to test enzyme viability.
- If the control DNA is cleaved and the experimental DNA resists cleavage, the two DNAs can be mixed to determine if an inhibitor is present in the experimental sample. If an inhibitor (often salt, EDTA or phenol) is present, the control DNA will not cut after mixing.

<sup>\*\*</sup> Note: 10  $\mu L$  rxns should not be incubated for longer than 1 hour to avoid evaporation.