

# Hinfl

Catalogue number	Presentation
MB23901	500 U
MB23902	2500 U

5'...G↓ANTC...3'  
 3'...CTNA↓G...5' (N= G, C, A or T)

## Description

Hinfl is a Restriction Endonuclease purified from an Escherichia coli strain that carries the Hinfl gene from *Haemophilus influenzae*.

## Shipping & Storage Conditions

This product is shipped at dry ice. After delivery, product should be stored at -85°C to -15°C in a constant temperature freezer. Minimize exposure of enzyme to temperatures higher than -15°C. To reduce freeze-thaw cycles, we recommend making small aliquots of the enzyme. Hinfl will remain stable till the expiry date if stored as specified.

## Components

COMPONENT	MB23901 (500 U)		MB23902 (2500 U)	
	TUBES	VOLUME	TUBES	VOLUME
Hinfl (500 U)	1	50 µL	5	50 µL
10x NZYBuffer C	1	1 mL	1	1 mL

## Specifications

**Unit Definition:** One unit is defined as the amount of this enzyme required to digest completely 1 µg of plasmid DNA in 50 µL of the reaction mixture at 37°C for one hour.

**Inhibition & Inactivation:** Hinfl is inhibited in the presence of metal chelators (e.g. EDTA), phenol, chloroform, alcohol, detergents or excessive salts. The enzyme's activity can also be inhibited by DNA methylation (see below). The enzyme is heat inactivated at 80°C for 20 min.

**Methylation Sensitivity:** Hinfl activity is blocked by CpG methylation.

**Optimal Activity temperature:** The optimum reaction temperature is at 37°C.

**Activity in NZYtech Buffers:**

NZYtech Buffers	A	B	C	U
Activity in NZYtech buffers (% of max)	(90)	(90)	100	100

( ) weak star activity is detected

## Standard Protocol

### Recommendations before starting

- **Nucleic acid manipulation:** Ensure that DNA is devoid of any potential inhibitors of the enzyme. Additional wash steps during purification are advised. The recommended A230/260 ratio is typically close to 2.0. Achieving this level of purity ensures minimal contamination, which could otherwise interfere with the enzymatic reaction. Store the purified DNA sample properly to prevent degradation. Typically, DNA can be stored at -30°C to -15°C or -80°C to -65°C for long-term stability. We recommend using nuclease-free plasticware and reagents throughout all steps of the procedures.
- **Handling instructions:** It is preferable to keep the enzyme frozen while working at the bench and only remove it at the moment of addition to the reaction. Do not leave the enzyme on ice for an extended period of time. During setup, the enzyme should be the last component added to the reaction.

- **Controls:** A negative control without the addition of enzyme is highly recommended for verifying the effectiveness of the digestion process and distinguishing between enzymatic and non-enzymatic sources of DNA fragmentation. This enhances the reliability and interpretability of the experimental results.

## Procedure

The recommended protocol includes a 10-fold overdigestion, which generally is sufficient to overcome variations than can occur in DNA type, quantity and purity, as well as on frequency of recognition sites. In general, we recommend using 10 units of enzyme to digest 1 µg substrate DNA (or 10-20 units for genomic DNA) in 1 hour at appropriate temperature.

1. On ice, add the following reaction components into a sterile, nuclease-free microcentrifuge tube

### Notes:

- *If setting up more than one reaction, prepare a reaction mixture volume 10% greater than the total required for the number of reactions to be performed. If required, include sufficient reactions for the controls.*
- *Enzyme should be the last component added to reaction.*

	1 REACTION VOLUME
Substrate DNA	≤ 1 µg
10× NZYBuffer C <sup>(1)</sup>	2 µL
Hinfl	1 µL (10 units)
Nuclease-free water	up to 20 µL

(1) For double digestions, use NZYBuffer U (not provided, Cat. No. MB110)

2. Mix reaction components gently by pipetting or by “flicking” the tube (do not vortex) and spin down.

3. Incubate at 37°C for one hour.

**Note:** *In some situations, digestion may be improved by increasing the incubation time.*

4. Stop the reaction. Depending on the downstream application, use one of the following inactivation procedures:

- Heat inactivation (20 min. at 80°C)
- Addition of 20-30 mM EDTA pH 8.0 (\*)
- Gel Electrophoresis and Band Excision
- Spin Column DNA Purification
- Phenol-Chloroform Extraction or Ethanol Precipitation

(\*) **Note:** *the chelating property of EDTA may inhibit some downstream applications.*

## Technical Notes

**Reaction Volume:** The recommended final volume is 20 µL but reaction volumes from 10 to 50 µL per µg of substrate DNA can be tested. Enzyme should not exceed 10% of total reaction volume. Variation on final volume has influence on the reaction. In some situations, small reaction volumes may be beneficial; however, caution should be taken when reducing reaction volume because it may lead to star activity by concentrating glycerol (should not exceed 5-8%), enzyme or salts, as well any contaminant present in the reaction.

**Incubation Time:** Care must be taken during reaction incubation. Keep the temperature constant and avoid sample evaporation. This is special critical for long incubation periods (more than 1 hour) and small reaction volumes (less than 15 µL).

## Quality control assays

### Purity

Hinfl purity is >90% as judged by SDS polyacrylamide gel electrophoresis followed by Coomassie Blue staining.

### Nuclease assays

To test for DNase contamination, 1 µg of pNZY28-derived plasmid DNA are incubated with 10 U of enzyme for 14-16 h at 37 °C. Following incubation, the nucleic acid is visualized on a GreenSafe-stained agarose gel. There must be no visible nicking or cutting of the nucleic acid.

### Functional assay

Hinfl was tested for performance in a digestion of 1 µg of a recombinant pNZY28 derivative using 10 U, 5 U and 2 U of enzyme. The resulting product was visualized in an agarose gel.

For life science research only. Not for use in diagnostic procedures.