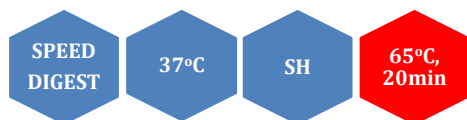


# Sal I



5' ...G▼TCGAC...3'  
3' ...CAGCT▲G...5'

Sal I is a restriction enzyme purified from *Streptomyces albus* G.

Catalogue No            130-1, 3000 U  
                                 130-2, 3x3000 U

Concentration            10-12u/μl and 40-60u/μl\*

\*Add an H to cat.# to order the high concentration

**Reagents supplied:** 10x SH buffer.

**Unit substrate:** Lambda DNA (HindIII digest).

**Unit calculation assay conditions:** 150 mM NaCl, 10 mM Tris-HCl (pH 7.9 @ 25°C), 10 mM MgCl<sub>2</sub>, 1 mM dithiothreitol, 100 μg/ml BSA. Incubate at 37°C.

**Absence of contaminants:** 400 units of Sal I incubated for 16 hours at 37°C with 1 μg of λ DNA (HindIII digest) resulted in a DNA pattern free of detectable nuclease degradation as determined by agarose gel electrophoresis. After 50-fold overdigestion with Sal I, greater than 95% of the DNA fragments can be ligated and recut with this enzyme.

**Storage buffer:** 50 mM KCl, 10 mM Tris-HCl (pH 7.4), 0.1 mM EDTA, 1 mM dithiothreitol, 300 μg/ml bovine serum albumin and 50% glycerol. Store at -20°C.

**Heat inactivation:** 65°C for 20 minutes.

**Methylation Sensitivity:**

dam methylation: Not sensitive

dcm methylation: Not sensitive

CpG methylation: Blocked

**Star activity:** Large excess of the enzyme results in the appearance of star activity.

**Percent Activity in MINOTECH Buffers**

L	M	H	SH	A	K
<10	25-50	50	100	<10	50

**General reaction mixture:**

10U Sal I                            1μl  
10x SH buffer \*                    2μl  
DNA substrate                    <1μg  
Sterile ultrapure water            Up to 20 μl  
*Incubate for 15 min at 37°C*

\*We recommend the addition of BSA to a final concentration of 100 μg/ml.

**Frequency of Cutting**

λ	Ad-2	Φx174	pUC18	M13mp18	pBR322
2	3	0	1	1	1



Lambda DNA 0.7 % agarose