Restriction Endonuclease *Nar* **I**

From Nocardia argentinensis

Cat. No. 11 103 024 001

1000 units (10 U/ μ l)



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Store at -15 to -25°C Stability/Storage The undiluted enzyme solution is stable when stored Number of cleavage sites on different DNAs (2): at -15 to -25°C until the control date printed on the Ad2 SV40 ΦX174 M13mp7 pBR322 pBR328 pUC18 λ label. Do not store below -25°C to avoid freezing. 1 20 5 Note: Product is shipped on dry ice. **PFGE** tested Nar I has been tested in Pulsed Field Gel Electro-Nar I recognizes the sequence GG/CGCC and gener-Sequence specificity ates fragments with 5'-cohesive ends (1). Nar I is simiphoresis (test system bacterial chromosomes). For cleavage of genomic DNA (E. coli C600) embedded in lar to Nae I in that it demonstrates marked site preferences on different substrate DNAs, e.g. λ, Ad2 agarose for PFGE analysis 10 units of enzyme/µg DNA pBR322. Several cleavage sites on these DNAs are and 4 h incubation time are recommended. cleaved at extremely slow rates and complete digestion is obtained only with large excess of enzyme (2). Ligation and Nar I fragments obtained by complete digestion of 1 μg recutting assay Ad2 DNA are ligated with 1 U T4 DNA ligase in a volume of 10 µl by incubation for 16 h at +4 °C in 66 mM **Compatible ends** Nar I generates compatible ends to Acy I, Cla I, Hpa II, Mae II, Msp I, Sfu I and Taq I. Tris-HCl, 5 mM MgCl₂, 5 mM Dithiothreitol, 1 mM ATP, pH 7.5 (at +20 °C), resulting in >90 % recovery of Ad2 DNA. Nar I is an isoschizomer to Bbe I, Ehe I, Kas I, Nun II. Isoschizomers Subsequent re-cutting with *Nar* I yields > 90% of the typical pattern of Ad 2 × *Nar* I fragments. Methylation Nar I is inhibited by 5-methylcytosine as indicated (*). sensitivity In addition Nar I is inhibited by 4-methylcytosine at the 3'-position and by 5'-hydroxymethylcytosin in all posi-Troubleshooting A critical component is the DNA substrate. Many comtions pounds used in the isolation of DNA such as phenol, chloroform, ethanol, SDS, high levels of NaCl, metal ions (e.g., Hg²⁺, Mn²⁺) inhibit or alter recognition 20 mM Tris-HCl, 50 mM NaCl, 0.1 mM EDTA, 1 mM Storage buffer dithioerythritol, 0.05% polydocanol, 50% glycerol (v/v), pH approx. 8.0 (at +4°C). specificity of many restriction enzymes. Such compounds should be removed by ethanol precipitation 330 mM Tris-acetate, 660 mM K-acetate, 100 mM followed by drying, before the DNA is added to the Incubation buffer Mg-acetate, 5 mM dithiothreitol, pH 7.9 (at +37°C), restriction digest reaction. Appropriate mixing of the (10x, included) enzyme is recommended. $(\triangle SuRE/Cut Buffer A)$ Activity in A В L Μ Н SuRE/Cut Buffer **Quality control** 100% 75-100% 75-100% 50-75% 0-10% System Lot-specific certificates of analysis are available at +37 °C www.roche-applied-science.com/certificates. Incubation temperature Absence of 1 µg Ad2-DNA is incubated for 16 h in 50 µl SuRE/Cut unspecific buffer A with excess of Nar I. The number of enzyme Unit definition One unit is the enzyme activity that completely cleaves units which do not change the enzyme-specific pattern 1 µg pBR322 DNA in 1 h at +37°C in the SuRE/Cut endonuclease activities is stated in the certificate of analysis. buffer A in a total volume of 25 µl. Approx. 5 µg [³H] labeled calf thymus DNA are incubated Absence of Heat inactivation The enzyme can be heat-inactivated by heating to exonuclease with 3 μ l Nar I for 4 h at +37°C in a total volume of 100 +65°C for 15 min. activity $\mu l~$ 50 mM Tris-HCl, 10 mM MgCl_2, 1 mM dithioerythritol, pH approx. 7.5. Under these conditions no released radio-Typical activity is detectable, as stated in the certificate of analy-**Final concentration** experiment Component sis. DNA 1 μg 10 × SuRE/Cut Buffer A 2.5 μl Up to a total volume of 25 μl Repurified water References Restriction enzyme 1 unit Incubate at +37°C for 1 h. Kessler, C. & Manta, V. (1990) Gene 92, 1-248.

- 2 Glück, B., Sagmeister, C., Rexer, B., unpublished observations.
- 3 Cooney, C.A. (1990) *Nucl. Acids Res.* **18,** 3667.
- 4 Lindsay, S & Bird, A.P. (1987) *Nature* **327**, 336-338.
- 5 Rina, M. et al. (1991) *Nucl. Acids Res.* **19**, 6341.
- 6 Rebase The Restriction Enzyme Database
- http://rebase.neb.com

7 Benchmate: http://www.roche-applied-science.com/benchmate



Ordering Information

Roche Applied Science offers a large selection of reagents and systems for life science research. For a complete overview of related products and manuals, please visit and bookmark our home page, <u>www.roche-applied-science.com</u>, and our Special Interest Sites, including "Mapping & Cloning": <u>http://www.restriction-enzymes.com</u>.

The convenient RE Finder Program located on our Bench Mate website, <u>http://www.roche-applied-science.com/benchmate</u> helps you identify the enzymes that will cut your DNA sequence, and displays the names and recognition sequences of enzymes and isoschizomers as well as links to detailed infor-

mation (*e.g.* instructions for use) of the selected restriction enzyme.

Product	Application	Packsize	Cat. No.
Restriction Enzymes	DNA restriction digestion	Please refer to website or catalogue	
Rapid DNA Liga- tion Kit	Ligation of sticky- or blunt-ended DNA fragments in just 5 min at +15 to +25 °C.	Kit (40 DNA ligations)	11 635 379 001
T4 DNA Ligase	Ligation of sticky- and blunt- ended DNA fragments.	100 U 500 units (1 U/μl)	10 481 220 001 10 716 359 001
rAPid Phosphatase	Dephosphorylation of 5´-phosphate residues from nucleic acids	1000 U 5000 U	04 898 133 001 04 898 141 001
rAPid Dephos and Ligation Kit	Dephosphorylation of nucleic acids.	40 reactions 160 reactions	04 898 117 001 04 898 125 001
Alkaline Phospha- tase (AP), special quality for molecu- lar biology	Dephosphorylation of 5'-phosphate residues from nucleic acids.	1000 U (20 U/µl)	11 097 075 001
Agarose MP	Multipurpose agarose for analytical and prepara- tive electrophoresis of nucleic acids	100 g 500 g	11 388 983 001 11 388 991 001
Agarose LE	Separation of nucleic acids in the range 0.2 - 1.5 kbp	100 g 500 g	11 685 660 001 11 685 678 001
Agarose Gel DNA Extraction Kit	For the elution of DNA fragments from agarose gels.	1 Kit (max. 100 reac- tions)	11 696 505 001
High Pure PCR Product Purifica- tion Kit	Purification of PCR or enzymatic modification reaction (<i>e.g.</i> restriction digest)	50 purifications 250 purifications	11 732 668 001 11 732 676 001
SuRE/Cut Buffer Set for Restriction Enzymes	Incubation buffers A, B, L, M and H for restriction enzymes	1 ml each (10× conc. solutions)	11 082 035 001
SuRE/Cut Buffer A	Restriction enzyme incubation	5×1 ml (10× conc. solution)	11 417 959 001
SuRE/Cut Buffer B	Restriction enzyme incubation	5×1 ml (10× conc. solution)	11 417 967 001
SuRE/Cut Buffer H	Restriction enzyme incubation	5×1 ml (10× conc. solution)	11 417 991 001
SuRE/Cut Buffer L	Restriction enzyme incubation	5×1 ml (10× conc. solution)	11 417 975 001
SuRE/Cut Buffer M	Restriction enzyme incubation	5×1 ml (10× conc. solution)	11 417 983 001
Water, PCR Grade	Specially purified, double-distilled, deionized, and autoclaved	100 ml (4 vials of 25 ml)	03 315 843 001
		25 ml (25 vials of 1 ml) 25 ml	03 315 932 001 03 315 959 001
		(1 vial of 25 ml)	
BSA, special qual- ity for molecular biology	Maintaining enzyme stability	20 mg (1 ml)	10 711 454 001

Printed Materials	You can view the following manuals on our website:	
	Lab FAQS "Find a Quick Solution"	
	Restriction Enzyme Ordering Guide	
	Molecular Weight Markers for Nucleic Acids	
Changes to previous version	Lot-specific information is no longer printed on the label of the product. Instead, the address for certificates of analysis is pro- vided (www.roche-applied-science.com/certificates). Update of unit definition and quality control.	
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Regulatory Disclaimer	For life science research only. Not for use in diagnostic procedures.	

Commonly used bacterial strains

Strain	Genotype
BL21	<i>E. coli B F</i> ⁻ <i>dcm ompT hsdS</i> ($r_B^-m_B^-$) <i>gal</i> (Studier, F.W. <i>et al</i> (1986) <i>J. Mol. Biol.</i> , 189 , 113.)
C600 ^e	<i>supE44 hsdR2 thi-1 thr-1 leuB6 lacY1 tonA21</i> ; (Hanahan, D. (1983) <i>J. Mol. Biol.</i> 166 , 557.)
DH5α	<i>supE</i> 44 Δ(<i>lac</i> U169 (φ80d <i>lac</i> ZΔM15) <i>hsd</i> R17 <i>rec</i> A1 <i>end</i> A1 <i>gyr</i> A96 <i>thi</i> -1 <i>rel</i> A1; (Hanahan, D. (1983) <i>J. Mol. Biol.</i> 166 , 557.)
HB101	supE44 hsdS20 recA13 ara-14 proA2 lacY1 galK2 rpsL20 xyl-5 mtl-1; (Hanahan, D., (1983) J. Mol. Biol. 166 , 557.)
JM108	recA1 supE44 endA1 hsdR17 gyrA96 relA1 thi ∆(lac-proAB); (Yanisch- Perron, C. et al., (1985) Gene 33 , 103.)
JM109	recA1 supE44 endA1 hsdR17 gyrA96 relA1 thi Δ (lac-proAB) F'[traD36proAB ⁺ , lacl ^q lacZ Δ M15]; (Yanisch- Perron, C. et al., (1985) Gene 33 , 103.)
JM110	<i>rpsL</i> (Str ⁷) thr leu thi-I lacY galK galT ara tonA tsx dam dcm supE44 Δ (lac-proAB) F[traD36proAB ⁺ , lacl ^q lacZ Δ M15]; (Yanisch- Perron, C. et al., (1985) Gene 33 , 103.)
K802	<i>supE hsdR gal metB;</i> (Raleigh, E. et al., (1986) Proc.Natl. Acad.Sci USA, 83, 9070.; Wood, W.B. (1966) J. Mol. Biol., 16 , 118.)
SURE ^r	recB recJ sbc C201 uvrC umuC::Tn5(kan ^r) lac , Δ(hsdRMS) endA1 gyrA96 thi relA1 supE44 F [*] [proAB ⁺ lacI ^q lacZΔM15 Tn10 (tet ^r); (Greener, A. (1990) Stratagies, 3 , 5.)
TG1	supE hsd $\Delta 5$ thi Δ (lac-proAB) F[traD36proAB ⁺ , lacl ^q lacZ Δ M15]; (Gibson, T.J. (1984) PhD Theses. Cambridge University, U.K.)
XL1-Blue ^r	supE44 hsdR17 recA1 endA1 gyrA46 thi relA1 lac $F'[proAB^+, lacl^q lacZ\DeltaM15 Tn10 (tet')]$; (Bullock et al., (1987) BioTechniques, 5, 376.)

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