## A general method for the purification of restriction enzymes

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#### ABSTRACT

An abbreviated procedure has been developed for the purification of restriction endonucleases. This procedure uses chromatography on phosphocellulose and hydroxylapatite and results in enzymes of sufficient purity to permit their use in the sequencing, molecular cloning, and physical mapping of DNA.

#### INTRODUCTION

The widespread use of restriction endonucleases has revolutionized DNA biochemistry (3, 4). We have devised a simple procedure for the preparation of several such enzymes sufficiently free from other nucleases and phosphatases to permit their use in sequencing, molecular cloning, and physical mapping of DNA. The following enzymes have been purified by this procedure: <u>AluI, BamI, BglI, BglII, EcoRI, EcoRII, HaeII, and HaeIII, HgaI, HincII, HsuI</u> (isoschizomer of <u>HindIII</u>), <u>PstI, SalI, TaqI, HpaI</u> and <u>HpaII</u>.

# BACTERIAL STRAINS AND GROWTH CONDITIONS

Strain sources, culture conditions and yields of cells are summarized in Table 1. Haemophilus strains were grown in a New Brunswick 14 liter microferm in the following medium: 37 g Brain Heart Infusion (BBL), 0.2 ml stock solution of nicotinamide adenine dinucleotide (NAD), and 10 ml stock solution of Hemin per liter. The NAD stock (10 mg/ml) was filter-sterilized and stored at -20°C. The Hemin stock (0.1% in 4% triethanolamine) was sterilized at  $65^{\circ}$ C for 15 min and stored at 4°C. A 500 ml culture, inoculated with a single colony and grown 16 hours, was used to inoculate 12 liters of medium in the microferm. This was grown 5-7 hours at  $37^{\circ}$ C with 11 1/min aeration and 3-500 rpm mixing. Increase in cell density was monitored on a Klett-Summerson photometer with a green filter, and the cells were harvested by centrifugation when the cell density stopped increasing (approximately 400-600 Klett units).

<u>Streptomyces albus</u> was grown in a New Brunswick microferm in Zubay's medium, which contains 28.9 g  $K_2$ HPO<sub>4</sub>, 5.6 g KH<sub>2</sub>PO<sub>4</sub>, 10 g yeast extract, 10 mg

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<b></b>			Table 1		
Enzyme		Strain	Source of Strain	<sup>a</sup> Fementor/Medium	Yield
AluI		Arthrobacter luteus	R. J. Roberts	Hi density/SLBH	280 g
BamI		Bacillus amyloliquifaciensH	G. Wilson	Hi density/SLBH	55 g
<u>Bgl</u> I BglII	}	Bacillus globigii	G. Wilson	Hi density/SLBH	60 g
EcoRI		Escherichia coli pMB4	H. W. Boyer	Hi density/SLBH	250 g
<u>Eco</u> RII		Escherichia coli RY22	H. W. Boyer	Hi density/SLBH	250 g
<u>Hae</u> II <u>HaeIII</u>	}	Haemophilus aegypticus	R. J. Roberts	Microferm/BHI	55 g
Hga		Haemophilus gallinarium	R. J. Roberts	Microferm/BHI	60 g
<u>Hin</u> cII		Haemophilus influenza C	R. J. Roberts	Microferm/BHI	47 g
<u>Hpa</u> I <u>Hpa</u> II	}	Haemophilus parainfluenza	H. W. Boyer	Microferm/BHI	60 g
Hsu		Haemophilus suis	R. J. Roberts	Microferm/BHI	25 g
SalI		Streptomyces albus		Microferm/Zubay's	-
Taq		Thermus aquaticus	R. J. Roberts	Microferm/Castenholz	22 g

<sup>a</sup>Details of growth conditions and compositions of the media are given in the text. Strains grown in the Labline hi density fermentor could also be grown in Zubay's medium in the New Brunswick microferm or other comparable fermentation units.

vitamin B<sub>1</sub> per liter plus 1% glucose (autoclaved separately). Il liters of medium was inoculated with 1 liter of culture (started from a single colony and grown 36 hr at  $37^{\circ}$ C). This was grown 36 hr at  $37^{\circ}$ C.

<u>Thermus aquaticus</u> was grown in a New Brunswick microferm in Castenholz TYE medium (Bergey's manual). This medium was prepared by mixing 100 ml sterile 1% TYE, 1 ml Nitsch's trace element solution, 50 ml Castenholz salts A, 50 ml Castenholz salts B and 800 ml distilled water. 1% TYE is 10 g tryptone, 10 g yeast extract plus 1 liter distilled water. Nitsche's trace elements contains 0.5 ml  $H_2SO_4$ , 2.2 g  $MnSO_4$ , 0.5 g  $ZnSO_4$ , 0.5 g  $H_3PO_3$ , 0.016  $CuSO_4$ , 0.025 g  $Na_2MOO_2$ , 0.046 g  $CaCl_2 \cdot 6H_2O$  per liter. Castenholz salts A contains 2.0 g nitriloacetic acid, 20.0 ml of 0.03% FeCl<sub>3</sub>, 1.2 g  $CaSO_4 \cdot 2H_2O$ , 2.1 g  $KNO_3$  per liter and Castenholz salts B contains 2.0 g  $MgSO_4 \cdot 7H_2O$ , 0.16 g NaCl, 14.0 g  $NaNO_3$ , 2.2 g  $Na_2HPO_4$  g per liter. The optimum temperature for growth of <u>Thermus aquaticus</u> is 70°C; however the maximum temperature achieved by our microferm was 63°C. 11 liters of medium was inoculated with 1 liter of

			T	able 2			
	Sonicat	ion			Assay		
Enzyme	Lysozyme	Total Time	<sup>a</sup> Buffer	<sup>b</sup> # Sites in Substrate	<sup>C</sup> Crude Extract	P.C. Column Eluate	Time
<u>Alu</u> I	100µg/ml 30'	<b>ц</b> •	666	10		2 µl	30 <b>'</b>
BamI	100µg/ml 30'	12'	TM	1	-		
<u>Bgl</u> I <u>Bgl</u> II	100μg/ml 15' 100μg/ml 15'	3' 3'	666 666	l λDNA 5	_	2 µl	60'
EcoRI	_	4'	TM + 0.1M NaCl	l	0.05µl	.05µl	10'
<u>Eco</u> RII	-	4 '	TM	5	2 µl	2 µl	30 <b>'</b>
HaeII HaeIII	-	<u>4</u> ,	666 666	2 pMB8 >12	-	l μl l μl	30' 30'
Hga	-	<u></u> ч	666	pKB252 5	8 µl	μl	20'
<u>Hin</u> cII	_	4'	666 + 0.05M NaCl	2	_	_	-
<u>Hpa</u> I HpaII	-	չեւ Դեւ	666 + 0.05M NaCl	р <b>МВ9 &gt;</b> 10		2 µl 2 µl	15' 15'
Hsu	_	4 <b>'</b>	666 + 0.05M NaCl	l	-	l µl	15'
Pst	100µg/ml 15'	2'	TM	1	-	2 µl	30'
<u>Sal</u> I	_	5'	666 + 0.15M NaCl	1	_	l µl	60'
Taq	-	5'	666	pBR345 l	_	-	-

<sup>a</sup>TM buffer is tris-HCl 0.1M, MgCl<sub>2</sub> 5mM, pH 7.5; 666 buffer is tris-HCl 6mM, MgCl<sub>2</sub> 6mM, βME 6mM, pH 7.5.

<sup>b</sup>Substrate DNA was pBR322 except where noted otherwise.

<sup>C</sup>More careful adjustment of amount of extract and time of incubation might render specific restriction patterns visible in some of these strains.

culture grown overnight at 70°C and this was grown 6 hr at 63°C.

Other strains were grown in a high density fermentor (Lab-line Instruments) to late log phase in SLBH medium (ll g tryptone, 22.5 g yeast extract, 51 ml 1M  $K_2$ HPO<sub>4</sub> and 15.7 ml 1M  $KH_2$ PO<sub>4</sub> per liter plus 0.4% glycerol). ENZYME ASSAYS

Enzymes were assayed by electrophoresis of digested DNA on a 1% agarose gel Tris-borate buffer (10.8 g tris base, 0.93 g Na<sub>2</sub>EDTA and 5.5 g boric acid per liter) for 1-1/4 hr at 16 V/cm (2).

Any defined DNA with a convenient number of restriction sites can be used as a substrate. We have generally used pBR322 (1) except as noted in Table 2.

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The DNA for <u>Eco</u>RII assays was prepared from <u>E</u>. <u>coli</u> strain GM31 which is <u>dcm</u> (lacking DNA cytosine methylase). The buffers used for assaying the various enzymes are listed in Table 2. Because of interference by other nucleases, we were unable to assay some of the enzymes until after phosphocellulose chromatography (Table 2). For all except <u>Eco</u>RI, 0.5-2.0  $\mu$ l of extract or phosphocellulose column eluate is incubated in 30  $\mu$ l containing 0.2  $\mu$ g DNA at 37°C (60°C for <u>Taq</u>) for 15-60 min. <u>Eco</u>RI is coded for by a multicopy plasmid and therefore is present at a much higher concentration than the other enzymes; crude extract and phosphocellulose eluate were diluted approximately 20 fold. These variations are summarized in Table 2. The amounts of enzyme to add and time of incubation are approximate and are included only to serve as a guide. PREPARATION OF CELL EXTRACT

The following purification is standardized to 50 g of cells. Frozen cell pellets were thawed and suspended in 200 ml of 10mM  $K_2HPO_4-KH_2PO_4$  pH 7.0, 7mM mercaptoethanol, 1mM EDTA (extract buffer) containing 25 µg/ml phenylmethyl sulfonylfluoride (PMSF), 1mM NaN<sub>3</sub> and 0.4M NaCl. Several of the enzymes have been successfully prepared without NaN<sub>3</sub> and the proteolytic inhibitor (PMSF); however these are generally added as a precaution.

The cell suspension was sonicated with a Branson model W-350 set at maximum output using a 1 inch horn. Sonication was carried out in 30"-90" pulses with the cell suspension stirring on ice; the temperature was monitored and kept below  $10^{\circ}$ C. Several of the strains were resistant to sonication. These were stirred on ice with the addition of  $100 \ \mu\text{g/ml}$  lysozyme for 15-30 minutes before sonication. (See Table 2) Completeness of sonication was monitored by following increase in absorbance at 280 of centrifuged aliquots after periods of sonication. In the case of EcoRI, release of enzyme activity closely paralleled release of  $A_{280}$  material.

The sonicated cell suspension was centrifuged for 1 hour at 100,000 X g (Beckman type 35 rotor). The supernatant was decanted and the conductivity was diluted to the equivalent of 0.2M NaCl or 0.1M NaCl, depending on the strain (see Table 3) and the pH was adjusted to 7.0. This extract was applied directly to a column of phosphocellulose.

## PHOSPHOCELLULOSE CHROMATOGRAPHY

Careful preparation of phosphocellulose is essential to the success of this purification method. 125 g of Whatman Pll was suspended in 4 liters of 0.2N HCl diluted 1:1 with 95% ETOH and stirred gently for 30 min at room temperature. The slurry was allowed to settle and the supernatant aspirated or decanted to remove fines and other particulate matter. The resin was

				Table 3			
Enzyme	Phosphoce Molarity NaCl for Loading Enzyme	ellulose Chromat Molarity NaCl in Gradient	ography Molarity NaCl Where Enzyme Elutes	<u>Hydroxylapatite</u> Molarity of Pota Gradient	Chromatography ssium Phosphate <sup>a</sup> Elution	<sup>b</sup> rield Units of Enzyme from 50g of cells	Used for DNA Sequencing
IULA	0.2	0.2 - 1.0	0.35	.0125	.10	4,000	+
BamI	0.2	0.2 - 0.6	0.35	.0125	.08	60,000	+
Bg1I Bg1II	0.2	0.2 - 0.8	0.3	.015 .014	.3 .27	1 1	TN
EcoRI	0.2	0.2 - 0.8	0.6	.01500	.20	1,000,000	+
EcoRII	0.2	0.2 - 0.6	0.4	.0125	<b>.</b> 0h	15,000	+
HaeII HaeIII	0.2	0.2 - 1.0	0.9	.0150	.25	60,000 180,000	+ +
Hga	0.2	0.2 - 1.0	0.4	I	I	6,000	TN
HincII	0.1	0.1 - 0.6	0.35	.0125	.06	1	TN .
Hpa.I Hpa.II	0.2	0.2 - 1.0	0.45 0.30	.0150 .0150	.30	500 25 <b>,</b> 000	LN
HsuI	0.2	0.2 - 0.6	0.3	.0125	I	60,000	+
Pst	0.2	0.2 - 0.6	0.35	0† - IO.	.20	30,000	TN
Sall	0.2	0.2 - 0.8	0.4	I	I	25 <b>,</b> 000	LN
Taq	0.2	0.2 - 0.8	0.4	.0125	.12	8,000	+
b <sup>a</sup> The el bA unit for Ta	ution positions is defined as These are	s are approximat the amount of e all minimum est	e. nzyme required imates.	to digest l ug of	DNA to completi	on in l hr a	t 37°C (or 60°

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collected by vacuum filtration and washed 2 to 3 times by suspension in 4 liters of distilled water. The pH was adjusted to near neutral with 1M NaOH and the resin was suspended in 4 liter 0.1N NaOH, stirred for 30 min at room temperature, collected by filtration, and suspended in 4 liters of 1mM EDTA. It was stirred for 30 min at room temperature and washed 3 times with distilled water. The pH was adjusted to near neutral with 1M HCl and the resin was suspended in extract buffer plus 0.2M NaCl. The pH was carefully adjusted to pH 7.0 before pouring a column, since many column volumes of dilute buffer are required for equilibration of the pH in a packed column. 125 g of dry phosphocellulose provides enough resin for a 500 ml packed column bed volume. If this preparation is scaled up, or if the starting resin is very dirty (phosphocellulose lots vary), the volumes or the number of the acid-ethanol and NaOH washes should be increased. Fines should be removed by allowing the resin to settle and aspirating or decanting the supernatant after several of the washes. The resin should be stirred gently and not too long so that fines are not generated, and it should not be allowed to stand at the extremes of pH for more than two hours.

A phosphocellulose column of 150-200 ml bed volume (2.5 X 35 cm) is sufficient for 50 g cell. It should be washed with several volumes of extract buffer plus NaCl (see Table 3) prior to loading. The extract was loaded directly onto the column and the column washed with several column volumes of extract buffer plus NaCl (see Table 3 for NaCl conc.) or until the  $A_{280}$  has decreased to near zero. The column was developed with a linear gradient of NaCl in extract buffer, 1-1.5 liter in volume (approximately 7 times the column volume). The NaCl concentrations for the individual enzymes are given in Table 3 together with the approximate NaCl concentration at which the enzyme elutes. Ten ml fractions were collected and assayed for endonuclease activity. Peak fractions were pooled and subjected to hydroxylapatite chromatography.

### HYDROXYLAPATITE CHROMATOGRAPHY

A 20 ml bed volume column of hydroxylapatite (Clarkson Chemical Co.) was packed and equilibrated with extract buffer plus 0.2M NaCl. The phosphocellulose pool was applied directly to this column and the column was washed with about 100 ml of extract buffer plus 0.2M NaCl. Enzymes were eluted with a 300 ml gradient of potassium phosphate pH 7.0 containing lmM EDTA, 7mM  $\beta$ ME, lmM NaN<sub>3</sub> and 0.2M NaCl. (Table 3 lists potassium phosphate concentrations for the individual enzymes.) 4 ml fractions were collected and assayed for enzyme activity.

Peak fractions were pooled and dialyzed against storage buffer (20mM  $K_2HPO_4 \cdot KH_2PO_4$  pH7, 0.2M NaCl, lmM EDTA, 7mM ßME, l mM NaN<sub>3</sub> diluted l:1 with glycerol). This dialysis concentrated the enzyme about three fold. Enzymes in this buffer were stable for many months at -20°C and working aliquots are also kept at -20°C. A more concentrated enzyme preparation can be obtained by dialyzing the pooled fractions from the hydroxylapatite column against solid PEG 6000 prior to dialysis against storage buffer.

Three enzymes, <u>SalI</u>, <u>Eco</u>RII, and <u>HgaI</u>, did not bind well to hydroxylapatite. Approximately 70% of the <u>Eco</u>RII, 90% of the <u>Hga</u>I, and all of the <u>SalI</u> activity were found in the flow through from this column; however, in the cases of <u>Eco</u>RII and <u>SalI</u>, most of the contaminating activities bound, and this was still an efficient purification step. Flow-through fractions were concentrated by dialysis versus solid polyethylene glycol (PEG6000) followed by dialysis against storage buffer.

## BIOREX 70 CHROMATOGRAPHY

After phosphocellulose and hydroxylapatite chromatography, <u>Hga</u>I still contained a large amount of exonuclease. The hydroxylapatite flow-through was concentrated, dialyzed against EB, and further purified by chromatography on Biorex 70 (Biorad).

A 20 ml bed volume column was equilibrated with EB and the dialyzed hydroxylapatite pool applied to the column. The column was washed with EB and developed with a gradient of 0 to 0.5 M NaCl in EB. Much of the exonuclease was present in the flow-through, while the <u>Hga</u> eluted at approximately 0.35 M NaCl. <u>Eco</u>RII and <u>Hpa</u>II were also subjected to Biorex 70 chromatography. As in the case of <u>Hga</u>I, exonuclease was present in the flow-through, and the endonuclease eluted at 0.35 - 0.4 M NaCl.

In our laboratory, all of the enzymes except <u>Hga</u>I, <u>Eco</u>RII, and <u>Hpa</u>II have been used routinely after phosphocellulose and hydroxylapatite chromatography. If further purification is required, however, Biorex 70 may generally be an excellent step, since for the enzymes tested the major contaminants were present in the flow-through, and the enzyme eluted at a relatively high NaCl concentration.

### DISCUSSION

This procedure is a simplification of a previously published method for <u>Eco</u>RI purification (2). All enzymes listed bind to phosphocellulose at NaCl concentrations high enough to dissociate the enzyme from the nucleic acids in the crude extract. This permits direct loading of the crude extract on the phosphocellulose column; nucleic acids do not bind to the column and are

eliminated in the flow-through and wash. Streptomycin and ammonium sulfate precipitations are eliminated and dialysis steps are reduced. This shortens the purification procedure and has the additional benefit that some enzymes (i.e. <u>Hsu</u> and <u>SalI</u>) are more stable and are obtained in higher yield than when prepared by more lengthy procedures. The approximate yield of the enzymes obtained from 50 g of cells is listed in Table 3.

Digestion of DNA with all of the enzymes produces fragments which yield clear gel patterns. Except in the case of  $\underline{\text{Hga}}$ , which has not been tested, the fragments serve as substrates for DNA ligase, and ligation results in regeneration of the restriction sites. By these criteria the enzymes are sufficiently free of non-specific nucleases and phosphatases to be useful as tools in physical mapping and cloning. All of those tested (Table 3) are also pure enough to be used in DNA sequencing.

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