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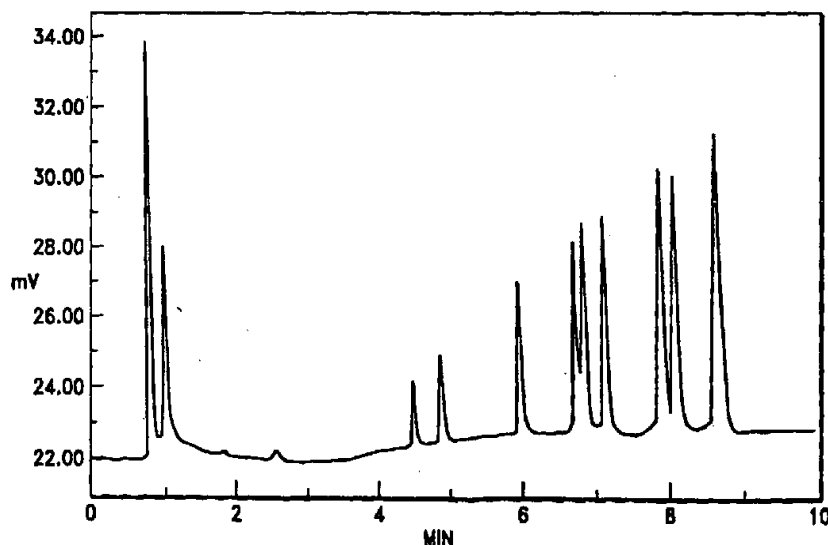


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(54) Title: POLYNUCLEOTIDE SEPARATIONS ON NONPOROUS POLYMER BEADS



(57) Abstract

The present invention describes a chromatographic method for separating polynucleotides with improved separation and efficiency. The method uses nonporous polymeric beads having a non-reactive, non-polar surface, and made from a variety of different polymerizable monomers.

TITLE OF THE INVENTION

POLYNUCLEOTIDE SEPARATIONS ON NONPOROUS POLYMER BEADS

FIELD OF THE INVENTION

The present invention is directed to the separation of polynucleotides
5 using nonporous polymeric beads.

BACKGROUND OF THE INVENTION

Separations of polynucleotides such as DNA have been traditionally
performed using slab gel electrophoresis or capillary electrophoresis.
However, liquid chromatographic separations of polynucleotides are
10 becoming more important because of the ability to automate the analysis and
to collect fractions after they have been separated. Therefore, columns for
polynucleotide separation by liquid chromatography (LC) are becoming more
important.

High quality materials for double stranded DNA separations previously
15 have been based on polymeric substrates disclosed in U.S. Patent No.
5,585,236, to Bonn et al., which showed that double-stranded DNA can be
separated on the basis of size with selectivity and performance similar to gel
electrophoresis using a process characterized as reverse phase ion pairing
chromatography (RPIPC). However, the chromatographic material described
20 was limited to nonporous beads substituted with alkyl groups having at least 3
carbons because Bonn et al. were unsuccessful in obtaining separations
using polymer beads lacking this substitution. Additionally, the polymer
beads were limited to a small group of vinyl aromatic monomers, and Bonn et
al. were unable to effect double stranded DNA separations with other
25 materials.

A need continues to exist for chromatographic methods for separating polynucleotides with improved separation efficiency and resolution.

Summary of the Invention

Accordingly, one object of the present invention is to provide a chromatographic method for separating polynucleotides with improved separation and efficiency.

Another object of the present invention is to provide a method for separating polynucleotides using nonporous polymer beads having a non-reactive, non-polar surface.

It is another object of this invention to provide the chromatographic separation of polynucleotides using nonporous polymeric beads made from a variety of different polymerisable monomers.

It is a further object of this invention to provide the chromatographic separation of polynucleotides using polymeric beads which can be unsubstituted, methyl-substituted, ethyl-substituted, hydrocarbon-substituted, or hydrocarbon polymer-substituted.

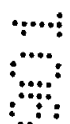
Yet another object of the present invention is to provide improved polymer beads by including steps to remove contamination occurring during the manufacturing process.

Still another object of the invention is to provide a method for separating polynucleotides using a variety of different solvent systems.

These and other objects which will become apparent from the following specification have been achieved by the present invention.

According to one aspect of the present invention there is provided a method for separating a mixture of polynucleotides, comprising flowing a mixture of polynucleotides having up to 1500 base pairs through a separation column containing polymer beads having an average diameter of 0.5 to 100 μ m, said beads having a surface composition essentially completely unsubstituted, or essentially completely substituted with a moiety selected from the group consisting of methyl, ethyl, hydrocarbon having from 23 to 1 000 000 carbons, and hydrocarbon polymer having from 23 to 1 000 000 carbons, and wherein said beads are characterised by being substantially free from multivalent cations which are free to bind with DNA and having a DNA Separation Factor of at least 0.5; and separating said mixture of polynucleotides.

In summary, this invention relates to a method for separating a mixture of polynucleotides comprising flowing a mixture of polynucleotides having up to



1500 base pairs through a separation column containing beads having an average diameter of 0.5 to 100 microns, and separating the mixture of polynucleotides using Matched Ion Polynucleotide Chromatography (as defined hereinbelow). The beads are preferably made from polymers, including mono- and di-vinyl substituted aromatic compounds such as styrene, substituted styrenes, alpha-substituted styrenes and divinylbenzene; acrylates and methacrylates; polyolefins such as polypropylene and polyethylene; polyesters; polyurethanes; polyamides; polycarbonates; and substituted polymers including fluorosubstituted ethylenes commonly known under the trademark TEFLON. The base polymer can also be mixtures of polymers, non-limiting examples of which include poly(styrene-divinylbenzene) and poly(ethylvinylbenzene-divinylbenzene). The polymer can be unsubstituted, or substituted with a moiety selected from the group consisting of methyl, ethyl, hydrocarbon, or hydrocarbon polymer. The hydrocarbon polymer optionally has from 23 to 1,000,000 carbons. The beads preferably have an average diameter of about 1 - 5 microns. In the preferred embodiment, precautions are taken during the production of the beads so that they are substantially free of multivalent cation contaminants and the beads are treated, for example by an acid wash treatment, to remove any residual surface metal contaminants. The beads of the invention are characterized by having a DNA Separation Factor (as defined hereinbelow) of at least 0.05. In a preferred embodiment, the beads are characterized by having a DNA Separation Factor of at least 0.5.

In addition to the beads themselves being substantially metal-free, we have also found that, to achieve optimum peak separation, the separation

column and all process solutions held within the column or flowing through the column should be substantially free of multivalent cation contaminants.

This can be achieved by supplying and feeding solutions entering the separation column with components which have process solution-contacting surfaces made of material which does not release multivalent cations into the process solutions held within or flowing through the column, in order to protect the column from multivalent cation contamination. The process solution-contacting surfaces of the system components are preferably material selected from the group consisting of titanium, coated stainless steel, and organic polymer.

For additional protection, multivalent cations in eluent solutions and sample solutions entering the column can be removed by contacting these solutions with multivalent cation capture resin before the solutions enter the column to protect the resin bed from multivalent cation contamination. The multivalent capture resin is preferably cation exchange resin and/or chelating resin.

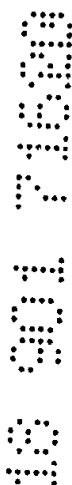
The method of the present invention can be used to separate double stranded polynucleotides having up to about 1500 to 2000 base pairs. In many cases, the method is used to separate polynucleotides having up to 600 bases or base pairs, or which have up to 5 to 80 bases or base pairs. The mixture of polynucleotides can be a polymerase chain reaction product.

The method is performed at a temperature within the range of 20°C to 90°C to yield a back-pressure not greater than 5000 psi. The method also preferably employs an organic solvent that is water soluble. The solvent is preferably selected from the group consisting of alcohols, nitriles,

dimethylformamide, esters, and ethers. The method also preferably employs a counter ion agent selected from trialkylamine acetate, trialkylamine carbonate, and trialkylamine phosphate. The most preferred counter ion agent is triethylammonium acetate or triethylammonium hexafluoroisopropyl alcohol.

8 As disclosed herein, the present invention also relates to a polymeric bead having an average bead diameter of 0.5-100 micron. Thus, according to another aspect of the invention there is provided a polymeric bead having an average bead diameter of 0.5-100µm and having a surface composition essentially completely unsubstituted, or essentially completely substituted with a moiety selected from the group consisting of
 10 methyl, ethyl, hydrocarbon having from 23 to 1 000 000 carbons, and hydrocarbon polymer having from 23 to 1 000 000 carbons, said bead being substantially free from multivalent cations which are free to bind with DNA and being characterised by having a DNA Separation Factor of at least 0.5.

Precautions are taken during the production of the beads so that they are
 15 substantially free of multivalent cation contaminants and the beads are treated, for example, by an acid wash treatment, to remove any residual surface metal contaminants. In one embodiment, the beads are characterised by having a DNA Separation Factor of at least 0.05. In a preferred embodiment, the beads are characterised by having a DNA Separation Factor of at least 0.5. The bead preferably has an average diameter of about
 20 1-10 microns, and most preferably has an average diameter of about 1-5 microns. The bead can be comprised of a copolymer of vinyl aromatic monomers. The vinyl aromatic monomers can be styrene, alkyl substituted styrene, alpha-methylstyrene or alkyl substituted alpha-methylstyrene. The bead can be a copolymer such as a copolymer of styrene, C₁₋₆ alkyl vinylbenzene and divinylbenzene. The bead can contain functional
 25 groups such as polyvinyl alcohol, hydroxy, nitro, halogen (c.g. bromo), cyano, aldehyde, or other groups that do not bind the sample. The bead can be unsubstituted, or substituted with a moiety such as methyl, ethyl, hydrocarbon, or hydrocarbon polymer. In preferred embodiments, the bead is octadecyl modified poly(ethylvinylbenzene-



divinylbenzene) or poly(styrene-divinylbenzene). The bead can also contain crosslinking divinylmonomer such as divinyl benzene or butadiene.

BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1 is a schematic representation of how the DNA Separation

5 Factor is measured.

FIG. 2 is a MIPC separation of pUC18 DNA-*Hae*III digestion fragments on a column containing alkylated poly(styrene-divinylbenzene) beads. Peaks are labeled with the number of base pairs of the eluted fragment.

FIG. 3 is a MIPC separation of pUC18 DNA-*Hae*III digestion fragments
10 on a column containing nonporous 2.1 micron beads of underivatized poly(styrene-divinylbenzene).

FIG. 4 is a Van't Hoff plot of $\ln k$ vs. $1/T [^{\circ}K^{-1}]$ with alkylated poly(styrene-divinylbenzene) beads showing positive enthalpy using acetonitrile as the solvent.

15 FIG. 5 is a Van't Hoff plot of $\ln k$ vs. $1/T [^{\circ}K^{-1}]$ with underivatized poly(styrene-divinylbenzene) beads showing positive enthalpy using acetonitrile as the solvent.

FIG. 6 is a Van't Hoff plot of $\ln k$ vs. $1/T [^{\circ}K^{-1}]$ with alkylated poly(styrene-divinylbenzene) beads showing negative enthalpy using
20 methanol as the solvent.

FIG. 7 is a separation using alkylated beads and acetonitrile as solvent.

FIG. 8 is a separation using alkylated beads and 50.0% methanol as the solvent.

FIG. 9 is a separation using alkylated beads and 25.0% ethanol as the solvent.

FIG. 10 is a separation using alkylated beads and 25.0% vodka (100 proof) as the solvent.

5 FIG. 11 is a separation using alkylated beads and 25.0% 1-propanol as the solvent.

FIG. 12 is a separation using alkylated beads and 25.0% 1-propanol as the solvent.

10 FIG. 13 is a separation using alkylated beads and 10.0% 2-propanol as the solvent.

FIG. 14 is a separation using alkylated beads and 10.0% 2-propanol as the solvent.

FIG. 15 is a separation using alkylated beads and 25.0% THF as the solvent.

15 FIG. 16 is a combination isocratic/gradient separation on a non-alkylated poly(styrene-divinylbenzene) beads.

DETAILED DESCRIPTION OF THE INVENTION

In its most general form, the subject matter of the present invention is the separation of polynucleotides utilizing columns filled with nonporous
20 polymeric beads having an average diameter of about 0.5 -100 microns; preferably, 1 - 10 microns; more preferably, 1 - 5 microns. Beads having an average diameter of 1.0 - 3.0 microns are most preferred.

In U.S. Patent No. 5,585,236, Bonn et al. had characterized the nucleic acid separation process as reverse phase ion pairing chromatography
25 (RPIPC). However, since RPIPC does not incorporate certain essential

characteristics described in the present invention, another term, Matched Ion Polynucleotide Chromatography (MIPC), has been selected. MIPC as used herein, is defined as a process for separating single and double stranded polynucleotides using non-polar beads, wherein the process uses a counter
5 ion agent, and an organic solvent to desorb the nucleic acid from the beads, and wherein the beads are characterized as having a DNA Separation Factor of at least 0.05. In a preferred embodiment, the beads have a DNA Separation Factor of at least 0.5. In an optimal embodiment, the beads have a DNA Separation Factor of at least 0.95.

10 The performance of the beads of the present invention is demonstrated by high efficiency separation by MIPC of double stranded and single stranded DNA. We have found that the best criterion for measuring performance of the beads is a DNA Separation Factor. This is measured as the resolution of 257- and 267-base pair double stranded DNA fragments of a pUC18 DNA-
15 *HaeIII* restriction digest wherein the distance from the valley between the peaks to the top of one of the peaks, over the distance from the baseline to the valley. Referring to the schematic representation of FIG. 1, the DNA Separation Factor is determined by measuring the distance "a" from the baseline to the valley "e" between the peaks "b" and "c" and the distance "d"
20 from the valley "e" to the top of one of the peaks "b" or "c". If the peak heights are unequal, the highest peak is used to obtain "d." The DNA Separation Factor is the ratio of $d/(a+d)$. The peaks of 257- and 267-base pairs in this schematic representation are similar in height. Operational beads of the present invention have a DNA Separation Factor of at least 0.05. Preferred
25 beads have a DNA Separation Factor of at least 0.5.

Without wishing to be bound by theory, we believe that the beads which conform to the DNA Separation Factor as specified herein have a pore size which essentially excludes the polynucleotides being separated from entering the bead. As used herein, the term "nonporous" is defined to denote
5 a bead which has surface pores having a diameter that is less than the size and shape of the smallest DNA fragment in the separation in the solvent medium used therein. Included in this definition are polymer beads having these specified maximum size restrictions in their natural state or which have been treated to reduce their pore size to meet the maximum effective pore
10 size required. Preferably, all beads which provide a DNA Separation Factor of at least 0.05 are intended to be included within the definition of "nonporous" beads.

The surface conformations of nonporous beads of the present invention can include depressions and shallow pit-like structures which do not
15 interfere with the separation process. A pretreatment of a porous bead to render it nonporous can be effected with any material which will fill the pores in the bead structure and which does not significantly interfere with the MIPC process.

Pores are open structures through which eluent and other materials
20 can enter the bead structure. Pores are often interconnected so that fluid entering one pore can exit from another pore. We believe that pores having dimensions that allow movement of the polynucleotide into the interconnected pore structure and into the bead impair the resolution of separations or result separations that have very long retention times. In MIPC, however, the beads
25 are "nonporous" and the polynucleotides do not enter the bead structure.

The term polynucleotide is defined as a linear polymer containing an indefinite number of nucleotides, linked from one ribose (or deoxyribose) to another via phosphoric residues. The present invention can be used in the separation of RNA or of double- or single-stranded DNA. For purposes of
5 simplifying the description of the invention, and not by way of limitation, the separation of double-stranded DNA will be described in the examples herein, it being understood that all polynucleotides are intended to be included within the scope of this invention.

Chromatographic efficiency of the column beads is predominantly
10 influenced by the properties of surface and near-surface areas. For this reason, the following descriptions are related specifically to the close-to-the-surface region of the polymeric beads. The main body and/or the center of such beads can exhibit entirely different chemistries and sets of physical properties from those observed at or near the surface of the polymeric beads
15 of the present invention.

The nonporous polymeric beads of the present invention are prepared by a two-step process in which small seed beads are initially produced by emulsion polymerization of suitable polymerizable monomers. The emulsion polymerization procedure of the invention is a modification of the procedure of
20 Goodwin et al. (J. W. Goodwin, J. Hearn, C. C. Ho and R. H. Ottweill, Colloid & Polymer Sci., (1974), 252:464-471). Monomers which can be used in the emulsion polymerization process to produce the seed beads include styrene, alkyl substituted styrenes, alpha-methyl styrene, and alkyl substituted alpha-methyl styrene. The seed beads are then enlarged and, optionally, modified

by substitution with various groups to produce the nonporous polymeric beads of the present invention.

The seed beads produced by emulsion polymerization can be enlarged by any known process for increasing the size of the polymer beads. For example, polymer beads can be enlarged by the activated swelling process disclosed in U.S. Patent No. 4,563,510. The enlarged or swollen polymer beads are further swollen with a crosslinking polymerizable monomer and a polymerization initiator. Polymerization increases the crosslinking density of the enlarged polymeric bead and reduces the surface porosity of the bead.

10 Suitable crosslinking monomers contain at least two carbon-carbon double bonds capable of polymerization in the presence of an initiator. Preferred crosslinking monomers are divinyl monomers, preferably alkyl and aryl (phenyl, naphthyl, etc.) divinyl monomers and include divinyl benzene, butadiene, etc. Activated swelling of the polymeric seed beads is useful to

15 produce polymer beads having an average diameter ranging from 1 up to about 100 microns.

Alternatively, the polymer seed beads can be enlarged simply by heating the seed latex resulting from emulsion polymerization. This alternative eliminates the need for activated swelling of the seed beads with

20 an activating solvent. Instead, the seed latex is mixed with the crosslinking monomer and polymerization initiator described above, together with or without a water-miscible solvent for the crosslinking monomer. Suitable solvents include acetone, tetrahydrofuran (THF), methanol, and dioxane. The resulting mixture is heated for about 1 - 12 hours, preferably about 4 - 8

25 hours, at a temperature below the initiation temperature of the polymerization

initiator, generally, about 10°C - 80°C, preferably 30°C - 60°C. Optionally, the temperature of the mixture can be increased by 10 - 20% and the mixture heated for an additional 1 to 4 hours. The ratio of monomer to polymerization initiator is at least 100:1, preferably about 100:1 to about 500:1, more
5 preferably about 200:1 in order to ensure a degree of polymerization of at least 200. Beads having this degree of polymerization are sufficiently pressure-stable to be used in high pressure liquid chromatography (HPLC) applications. This thermal swelling process allows one to increase the size of the bead by about 110 - 160% to obtain polymer beads having an average
10 diameter up to about 5 microns, preferably about 2 - 3 microns. The thermal swelling procedure can, therefore, be used to produce smaller particle sizes previously accessible only by the activated swelling procedure.

Following thermal enlargement, excess crosslinking monomer is removed and the particles are polymerized by exposure to ultraviolet light or
15 heat. Polymerization can be conducted, for example, by heating of the enlarged particles to the activation temperature of the polymerization initiator and continuing polymerization until the desired degree of polymerization has been achieved. Continued heating and polymerization allows one to obtain beads having a degree of polymerization greater than 500.

20 In the present invention, the packing material disclosed by Bonn et al. or U.S. Patent No. 4,563,510 can be modified through substitution of the polymeric beads with methyl or ethyl groups or with a hydrocarbon polymer group or can be used in its unmodified state. The polymer beads can be alkylated with 1 or 2 carbon atoms by contacting the beads with an alkylating
25 agent, such as methyl iodide or ethyl iodide. Alkylation is achieved by mixing

the polymer beads with the alkyl halide in the presence of a Friedel-Crafts catalyst to effect electrophilic aromatic substitution on the aromatic rings at the surface of the polymer blend. Suitable Friedel-Crafts catalysts are well-known in the art and include Lewis acids such as aluminum chloride, boron trifluoride, tin tetrachloride, etc. The beads can be hydrocarbon substituted by substituting the corresponding hydrocarbon halide for methyl iodide in the above procedure, for example.

The term "hydrocarbon" as used herein is defined to include alkyl and alkyl substituted aryl groups, having from 23 to 1,000,000 carbons, the alkyl groups including straight chained, branch chained, cyclic, saturated, unsaturated nonionic functional groups of various types including aldehyde, ketone, ester, ether, alkyl groups, and the like, and the aryl groups including as monocyclic, bicyclic, and tricyclic aromatic hydrocarbon groups including phenyl, naphthyl, and the like. Methods for hydrocarbon substitution are conventional and well-known in the art and are not an aspect of this invention. The substitution can also contain hydroxy, cyano, nitro groups, or the like which are considered to be non-polar, reverse phase functional groups. The term "hydrocarbon polymer" as used herein is defined to be a polymer having a hydrocarbon composition and from 23 to 1,000,000 carbons.

The chromatographic material reported in the Bonn patent was limited to nonporous beads substituted with alkyl groups having at least 3 carbons because Bonn et al. were unsuccessful in obtaining separations using polymer beads lacking this substitution. Additionally, the polymer beads were limited to a small group of vinyl aromatic monomers, and Bonn et al. were unable to effect double stranded DNA separations with other materials.

In the present invention, it has now been surprisingly discovered that successful separation of double stranded DNA can be achieved using underivatized nonporous beads as well as using beads derivatized with methyl, ethyl, hydrocarbon, or hydrocarbon polymer substitution.

5 The base polymer of the invention can also be other polymers, non-limiting examples of which include mono- and di-vinyl substituted aromatics such as styrene, substituted styrenes, alpha-substituted styrenes and divinylbenzene; acrylates and methacrylates; polyolefins such as polypropylene and polyethylene; polyesters; polyurethanes; polyamides; 10 polycarbonates; and substituted polymers including fluorosubstituted ethylenes commonly known under the trademark TEFLON. The base polymer can also be mixtures of polymers, non-limiting examples of which include poly(styrene-divinylbenzene) and poly(ethylvinylbenzene-divinylbenzene). Methods for making beads from these polymers are 15 conventional and well known in the art (for example, see U.S. Patent 4,906,378). The physical properties of the surface and near-surface areas of the beads are the predominant influence on chromatographic efficiency. The polymer, whether derivatized or not, must provide a nonporous, non-reactive, and non-polar surface for the Matched Ion Polynucleotide Chromatographic 20 separation.

The beads of the invention are also characterized by having low amounts of metal contaminants or other contaminants that can bind DNA.

The preferred beads of the present invention are characterized by having been subjected to precautions during production, including a decontamination 25 treatment, such as an acid wash treatment, designed to substantially

eliminate any multivalent cation contaminants (e.g. Fe(III), Cr(III), or colloidal metal contaminants). Only very pure, non-metal containing materials should be used in the production of the beads in order that the resulting beads will have minimum metal content.

5 In addition to the beads themselves being substantially metal-free, we have also found that, to achieve optimum peak separation during MIPC, the separation column and all process solutions held within the column or flowing through the column should be substantially free of multivalent cation contaminants. As described in commonly owned, copending U.S. application
10 Ser. No. 08/748,376 (filed November 13, 1996), this can be achieved by supplying and feeding solutions entering the separation column with components which have process solution-contacting surfaces made of material which does not release multivalent cations into the process solutions held within or flowing through the column, in order to protect the column from
15 multivalent cation contamination. The process solution-contacting surfaces of the system components are preferably material selected from the group consisting of titanium, coated stainless steel, passivated stainless steel, and organic polymer.

For additional protection, multivalent cations in eluent solutions and
20 sample solutions entering the column can be removed by contacting these solutions with multivalent cation capture resin before the solutions enter the column to protect the resin bed from multivalent cation contamination. The multivalent capture resin is preferably cation exchange resin and/or chelating resin.

To achieve high resolution chromatographic separations of polynucleotides, it is generally necessary to tightly pack the chromatographic column with the solid phase polymer beads. Any known method of packing the column with a column packing material can be used in the present invention to obtain adequate high resolution separations. Typically, a slurry of the polymer beads is prepared using a solvent having a density equal to or less than the density of the polymer beads. The column is then filled with the polymer bead slurry and vibrated or agitated to improve the packing density of the polymer beads in the column. Mechanical vibration or sonification are typically used to improve packing density.

For example, to pack a 4.6 x 50 mm i.d. column, 2.0 grams of beads can be suspended in 10 mL of methanol with the aid of sonification. The suspension is then packed into the column using 50 mL of methanol at 8,000 psi of pressure. This improves the density of the packed bed.

The separation method of the invention is generally applicable to the chromatographic separation of single stranded and double stranded polynucleotides of DNA and RNA. Samples containing mixtures of polynucleotides can result from total synthesis of polynucleotides, cleavage of DNA or RNA with restriction endonucleases or with other enzymes or chemicals, as well as nucleic acid samples which have been multiplied and amplified using polymerase chain reaction techniques.

The method of the present invention can be used to separate double stranded polynucleotides having up to about 1500 to 2000 base pairs. In many cases, the method is used to separate polynucleotides having up to 600 bases or base pairs, or which have up to 5 to 80 bases or base pairs.

In a preferred embodiment, the separation is by Matched Ion Polynucleotide Chromatography (MIPC). The nonporous beads of the invention are used as a reverse phase material that will function with counter ion agents and a solvent gradient to produce the DNA separations. In MIPC, 5 the polynucleotides are paired with a counter ion and then subjected to reverse phase chromatography using the nonporous beads of the present invention. Counter ion agents that are volatile, such as trialkylamine acetate, trialkylamine carbonate, trialkylamine phosphate (TEAAHPO_4), etc., are preferred for use in the method of the invention, with triethylammonium 10 acetate (TEAA) and triethylammonium hexafluoroisopropyl alcohol being most preferred.

To achieve optimum peak resolution during the separation of DNA by MIPC using the beads of the invention, the method is preferably performed at a temperature within the range of 20°C to 90°C; more preferably, 30°C to 15 80°C; most preferably, 50°C to 75°C. 14. The temperature is selected to yield a back pressure not exceeding 5000 psi. In general, separation of single-stranded fragments should be performed at higher temperatures.

We have found that the temperature at which the separation is performed affects the choice of solvents used in the separation. One reason 20 is that the solvents affect the temperature at which a double stranded DNA will melt to form two single strands or a partially melted complex of single and double stranded DNA. Some solvents can stabilize the melted structure better than other solvents. The other reason a solvent is important is because it affects the transport of the DNA between the mobile phase and the 25 stationary phase. Acetonitrile and 1-propanol are preferred solvents in these

cases. Finally, the toxicity (and cost) of the solvent can be important. In this case, methanol is preferred over acetonitrile and 1-propanol is preferred over methanol.

When the separation is performed at a temperature within the above
 5 range, an organic solvent that is water soluble is preferably used, for example, alcohols, nitriles, dimethylformamide (DMF), tetrahydrofuran (THF), esters, and ethers. Water soluble solvents are defined as those which exist as a single phase with aqueous systems under all conditions of operation of the present invention. Solvents which are particularly preferred for use in the
 10 method of this invention include methanol, ethanol, 2-propanol, 1-propanol, tetrahydrofuran (THF), and acetonitrile, with acetonitrile being most preferred overall.

We have determined that the chromatographic separations of double stranded DNA fragments exhibit unique Sorption Enthalpies (ΔH_{sorp}). Two
 15 compounds (in this case, DNA fragments of different size) can only be separated if they have different partition coefficients (K). The Nernst partition coefficient is defined as the concentration of an analyte (A) in the stationary phase divided by its concentration in the mobile phase:

$$20 \quad K = \frac{[A]_s}{[A]_m}$$

The partition coefficient (K) and the retention factor (k) are related through the following equations:

$$K = \frac{n(A)_s V_m}{n(A)_m V_s} \quad \text{and} \quad k = \frac{n(A)_s}{n(A)_m}$$

25 the quotient V_m/V_s is also called phase volume ratio (Φ). Therefore:

$$k = K \Phi$$

To calculate the sorption enthalpies, the following fundamental thermodynamic equations are necessary:

$$\ln K = -\frac{\Delta G_{\text{sorp}}}{RT}, \quad \ln k = -\frac{\Delta G_{\text{sorp}}}{RT} + \ln \Phi \quad \text{and} \quad \Delta G_{\text{sorp}} = \Delta H_{\text{sorp}} - T\Delta S_{\text{sorp}}$$

- 5 By transforming the last two equations, we obtain the Van't Hoff equation:

$$\ln k = -\frac{\Delta H_{\text{sorp}}}{RT} + \frac{\Delta S_{\text{sorp}}}{R} + \ln \Phi$$

- From a plot $\ln k$ versus $1/T$, the sorption enthalpy ΔH_{sorp} can be obtained from
 10 the slope of the graph (if a straight line is obtained). ΔS_{sorp} can be calculated if the phase volume ratio (Φ) is known.

- The Sorption Enthalpy ΔH_{sorp} is positive ($\Delta H_{\text{sorp}} > 0$) showing the separation is endothermic using acetonitrile as the solvent (Figs. 3 and 4), and using methanol as the solvent, the Sorption Enthalpy ΔH_{sorp} is negative
 15 ($\Delta H_{\text{sorp}} < 0$), showing the separation is exothermic (FIG. 5).

- The thermodynamic data (as shown in the Examples hereinbelow) reflect the relative affinity of the DNA-counter ion agent complex for the beads of the invention and the elution solvent. An endothermic plot indicates a preference of the DNA complex for the bead. An exothermic plot indicates a
 20 preference of the DNA complex for the solvent over the bead. The plots shown herein are for alkylated and non-alkylated surfaces as described in the Examples. Most liquid chromatographic separations show exothermic plots.

- Other features of the invention will become apparent in the course of the following descriptions of exemplary embodiments which are given for
 25 illustration of the invention and are not intended to be limiting thereof.

All references cited herein are hereby incorporated by reference in their entirety.

5

Preparation of nonporous poly(styrene-divinylbenzene) particles

20

Beads resulting from the first step are still generally too small and too soft (low pressure stability) for use as chromatographic packings. The softness of these beads is caused by an insufficient degree of crosslinking. In a second step, the beads are enlarged and the degree of crosslinking is increased.

The protocol for the second step is based on the activated swelling method described by Ugelstad et al. (*Adv. Colloid Interface Sci.*, 13:101-140 (1980)). In order to initiate activated swelling, or the second synthetic step, the aqueous suspension of polystyrene seeds (200 ml) from the first step was mixed first with 60 mL of acetone and then with 60 mL of a 1-chlorododecane emulsion. To prepare the emulsion, 0.206 g of sodium dodecylsulfate, 49.5 mL of deionized water, and 10.5 mL of 1-chlorododecane were brought together and the resulting mixture was kept at 0°C for 4 hours and mixed by sonication during the entire time period until a fine emulsion of < 0.3 microns was obtained. The mixture of polystyrene seeds, acetone, and 1-chlorododecane emulsion was stirred for about 12 hours at room temperature, during which time the swelling of the beads occurred. Subsequently, the acetone was removed by a 30 minute distillation at 80°C.

Following the removal of acetone, the swollen beads were further grown by the addition of 310 g of a ethyldivinylbenzene and divinylbenzene (DVB) (1:1.71) mixture also containing 2.5 g of dibenzoylperoxide as an initiator. The growing occurred with stirring and with occasional particle size measurements by means of light microscopy.

After completion of the swelling and growing stages, the reaction mixture was transferred into a separation funnel. In an unstirred solution, the

excess amount of the monomer separated from the layer containing the suspension of the polymeric beads and could thus be easily removed. The remaining suspension of beads was returned to the reactor and subjected to a stepwise increase in temperature (63°C for about 7 hours, 73°C for about 2
5 hours, and 83°C for about 12 hours), leading to further increases in the degree of polymerization (> 500). The pore size of beads prepared in this manner was below the detection limit of mercury porosimetry (< 30Å).

After drying, the dried beads (10 g) from step two were washed four times with 100 mL of n-heptane, and then two times with each of the
10 following: 100 mL of diethylether, 100 mL of dioxane, and 100 mL of methanol. Finally, the beads were dried.

EXAMPLE 2

Acid Wash Treatment

The beads prepared in Example 1 were washed three times with
15 tetrahydrofuran and two times with methanol. Finally the beads were stirred in a mixture containing 100 mL tetrahydrofuran and 100 mL concentrated hydrochloric acid for 12 hours. After this acid treatment, the polymer beads were washed with a tetrahydrofuran/water mixture until neutral (pH = 7). The beads were then dried at 40°C for 12 hours.

20 EXAMPLE 3

Standard Procedure for Testing the Performance of Separation Media

Separation particles are packed in an HPLC column and tested for their ability to separate a standard DNA mixture. The standard mixture is a pUC18 DNA-*Hae*III digest (Sigma-Aldrich, D6293) which contains 11
25 fragments having 11, 18, 80, 102, 174, 257, 267, 298, 434, 458, and 587

base pairs, respectively. The standard is diluted with water and five μL , containing a total mass of DNA of 0.25 μg , is injected.

Depending on the packing volume and packing polarity, the procedure requires selection of the driving solvent concentration, pH, and temperature.

- 5 The separation conditions are adjusted so that the retention time of the 257, 267 peaks is about 6 to 10 minutes. Any one of the following solvents can be used: methanol, ethanol, 2-propanol, 1-propanol, tetrahydrofuran (THF), or acetonitrile. A counter ion agent is selected from trialkylamine acetate, trialkylamine carbonate, trialkylamine phosphate, or any other type of cation
- 10 that can form a matched ion with the polynucleotide anion.

As an example of this procedure, FIG. 2 shows the high resolution of the standard DNA mixture using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads. The separation was conducted under the following conditions: Eluent A: 0.1 M TEAA, pH 7.0;

- 15 Eluent B: 0.1 M TEAA, 25% acetonitrile; Gradient:

Time (min)	%A	%B
0.0	65	35
3.0	45	55
10.0	35	65
13.0	35	65
14.0	0	100
15.5	0	100
16.5	65	35

The flow rate was 0.75 mL/min, detection UV at 260 nm, column temp. 50°C. The pH was 7.0.

As another example of this procedure using the same separation conditions as in FIG. 2, FIG. 3 is a high resolution separation of the standard DNA mixture on a column containing nonporous 2.1 micron beads of underivatized poly(styrene-divinylbenzene).

5

EXAMPLE 4

Sorption Enthalpy Measurements

Four fragments (174 bp, 257 bp, 267 bp, and 298 bp, found in 5 μ L pUC18 DNA-*Hae*III digest, 0.04 μ g DNA/ μ L) were separated under isocratic conditions at different temperatures using octadecyl modified, nonporous poly(styrene-divinylbenzene) polymer beads. The separation was carried out using a Transgenomic WAVE™ DNA Fragment Analysis System equipped with a DNASep™ column (Transgenomic, Inc., San Jose, CA) under the following conditions: Eluent: 0.1 M triethylammonium acetate, 14.25% (v/v) acetonitrile at 0.75 mL/min, detection at 250 nm UV, temperatures at 35, 40, 15 45, 50, 55, and 60°C, respectively. A plot of $\ln k$ versus $1/T$ shows that the retention factor k is increasing with increasing temperature (FIG. 4). This indicates that the retention mechanism is based on an endothermic process ($\Delta H_{\text{sorp}} > 0$).

The same experiments on non-alkylated poly(styrene-divinylbenzene) beads gave a negative slope for a plot of $\ln k$ versus $1/T$, although the plot is slightly curved (FIG. 5).

The same experiments performed on octadecyl modified, nonporous poly(styrene-divinylbenzene) beads but with methanol replacing the acetonitrile as solvent gave a plot $\ln k$ versus $1/T$ showing the retention factor

k is decreasing with increasing temperature (FIG. 6). This indicates the retention mechanism is based on an exothermic process ($\Delta H_{\text{sorp}} < 0$).

EXAMPLE 5

Separations with alkylated poly(styrene-divinylbenzene) beads

5 Eluents are chosen to match the desorption ability of the elution solvent to the attraction properties of the bead to the DNA-counter ion complex. As the polarity of the bead decreases, a stronger (more organic) or higher concentration of solvent will be required. Weaker organic solvents such as methanol are generally required at higher concentrations than
10 stronger organic solvents such as acetonitrile.

FIG. 7 shows the high resolution separation of DNA restriction fragments using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads. The experiment was conducted under the following conditions: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M TEAA, pH 7.2;
15 gradient: 33-55% acetonitrile in 3 min, 55-66% acetonitrile in 7 min, 65% acetonitrile for 2.5 min; 65-100% acetonitrile in 1 min; and 100-35% acetonitrile in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, column temp. 51°C. The sample was 5 μ L (= 0.2 μ g pUC18 DNA-*Hae*III digest).

20 Repeating the procedure of FIG. 7 replacing the acetonitrile with 50.0% methanol in 0.1 M (TEAA) gave the separation shown in FIG. 8.

Repeating the procedure of FIG. 7 replacing the acetonitrile with 25.0% ethanol in 0.1 M (TEAA) gave the separation shown in FIG. 9.

Repeating the procedure of FIG. 7 replacing the acetonitrile with 25% vodka (Stolichnaya, 100 proof) in 0.1 M (TEAA) gave the separation shown in FIG. 10.

The separation shown in FIG. 11 was obtained using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads as follows: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M tetraethylacetic acid (TEAA), pH 7.3; gradient: 12-18% 0.1 M TEAA and 25.0% 1-propanol (Eluent B) in 3 min, 18-22% B in 7 min, 22% B for 2.5 min; 22-100% B in 1 min; and 100-12% B in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, and column temp. 51°C. The sample was 5 μ L (= 0.2 μ g pUC18 DNA-*Hae*III digest).

The separation shown in FIG. 12 was obtained using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads as follows: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M TEAA, pH 7.3; gradient: 15-18% 0.1 M TEAA and 25.0% 1-propanol (Eluent B) in 2 min, 18-21% B in 8 min, 21% B for 2.5 min; 21-100% B in 1 min; and 100-15% B in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, and column temp. 51°C. The sample was 5 μ L (= 0.2 μ g pUC18 DNA-*Hae*III digest).

The separation shown in FIG. 13 was obtained using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads as follows: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M TEAA, pH 7.3; gradient: 35-55% 0.1 M TEAA and 10.0% 2-propanol (Eluent B) in 3 min, 55-65 % B in 10 min, 65% B for 2.5 min; 65-100% B in 1 min; and 100-35% B in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, and column temp. 51°C. The sample was 5 μ L (= 0.2 μ g pUC18 DNA-*Hae*III digest).

The separation shown in FIG. 14 was obtained using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads as follows: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M TEA₂HPO₄, pH 7.3; gradient: 35-55% 0.1 M TEA₂HPO₄ and 10.0% 2-propanol (Eluent B) in 3 min,
5 55-65% B in 7 min, 65% B for 2.5 min; 65-100% B in 1 min; and 100-65% B in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, and column temp. 51°C. The sample was 5 µL (= 0.2 µg pUC18 DNA-*Hae*III digest).

The separation shown in FIG. 15 was obtained using octadecyl modified, nonporous poly(ethylvinylbenzene-divinylbenzene) beads as
10 follows: Column: 50 x 4.6 mm i.d.; mobile phase 0.1 M TEAA, pH 7.3; gradient: 6-9% 0.1 M TEAA and 25.0% THF (Eluent B) in 3 min, 9-11 % B in 7 min, 11% B for 2.5 min; 11-100% B in 1 min; and 100-6% B in 1.5 min. The flow rate was 0.75 mL/min, detection UV at 260 nm, and column temp. 51°C. The sample was 5 µL (= 0.2 µg pUC18 DNA-*Hae*III digest).

15

EXAMPLE 6

Isocratic/gradient separation of ds DNA

The following is an isocratic/gradient separation of ds DNA using nonporous poly(styrene-divinylbenzene) beads. Isocratic separations have not been performed in DNA separations because of the large differences in
20 the selectivity of DNA/alkylammonium ion pair for beads. However, by using a combination of gradient and isocratic elution conditions, the resolving power of a system can be enhanced for a particular size range of DNA. For example, the range of 250-300 base pairs can be targeted by using an eluent of 0.1 M TEAA, and 14.25% acetonitrile at 0.75 mL/min at 40°C on 50 x 4.6

mm cross-linked poly(styrene-divinylbenzene) column, 2.1 micron. 5 μ L of pUC18 DNA-*Hae*III digest (0.2 μ g) was injected under isocratic conditions and 257, 267 and 298 base pairs DNA eluted completely resolved as shown in FIG. 16. Then the column was cleaned from larger fragments with 0.1M

5 TEAA/25% acetonitrile at 9 minutes. In other examples, there might be an initial isocratic step (to condition the column), then a gradient step (to remove or target the first group of DNA at a particular size), then an isocratic step (to separate the target material of a different size range) and finally a gradient step to clean the column.

10

EXAMPLE 7

Bromination of Remaining Double Bonds on the Surface of Poly(Styrene-Divinylbenzene) Polymer Beads

50.0 g of a poly(styrene-divinylbenzene) polymer beads were suspended in 500 g of tetrachloromethane. The suspension was transferred
15 into a 1000 mL glass reactor (with attached reflux condenser, separation funnel and overhead stirrer). The mixture was kept at 20°C. Bromine (100 mL) was added over a period of 20 minutes. After addition was completed, stirring continued for 60 minutes. The temperature was raised to 50°C to complete the reaction (2 hours).

20 The polymer beads were separated from the tetrachloromethane and excess bromine by means of centrifugation and cleaned with tetrahydrofuran (once with 100 mL) and methanol (twice with 100 mL). The polymer beads were dried at 40°C.

The polymer beads are packed into a 50 x 4.6 mm i.d column and the DNA Separation Factor is greater than 0.05 as tested by the procedure of Example 3.

EXAMPLE 8

5 *Nitration of a Poly(Styrene-Divinylbenzene) Polymer Beads*

In a 1000 mL glass reactor 150 mL of concentrated nitric acid (65%) were combined with 100 mL concentrated sulfuric acid. The acid mixture was cooled to 0-4°C. When the temperature had dropped to <4°C, 50 g of poly(styrene-divinylbenzene) polymer beads were added slowly under
10 continuous stirring. After addition was completed, 50 mL of nitric acid (65%) was added. The suspension was stirred for three hours, maintaining a temperature of 5-10°C.

On the next day the reaction was quenched by adding ice to the suspension. The polymer beads were separated from the acid by means of
15 centrifugation. The polymer beads were washed to neutrality with water, followed by washing steps with tetrahydrofurane (four times with 100 mL) and methanol (four times with 100 mL). The polymer beads were dried at 40°C.

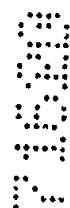
The polymer beads are packed into a 50 x 4.6 mm i.d column and the DNA Separation Factor is greater than 0.05 as tested by the procedure of
20 Example 3.

While the foregoing has presented specific embodiments of the present invention, it is to be understood that these embodiments have been presented by way of example only. It is expected that others will perceive and practice variations which, though differing from the foregoing, do not

depart from the spirit and scope of the invention as described and claimed herein.

The claims defining the invention are as follows:

1. A method for separating a mixture of polynucleotides, comprising flowing a mixture of polynucleotides having up to 1500 base pairs through a separation column containing polymer beads having an average diameter of 0.5 to 100µm, said beads having
 5 a surface composition essentially completely unsubstituted, or essentially completely substituted with a moiety selected from the group consisting of methyl, ethyl, hydrocarbon having from 23 to 1 000 000 carbons, and hydrocarbon polymer having from 23 to 1 000 000 carbons, and wherein said beads are characterised by being substantially free from multivalent cations which are free to bind with DNA and having a DNA
 10 Separation Factor of at least 0.5; and separating said mixture of polynucleotides.
2. The method of claim 1, wherein said beads are substituted with a moiety selected from the group consisting of methyl, ethyl, hydrocarbon having from 23 to 1 000 000 carbons, and hydrocarbon polymer having from 23 to 1 000 000 carbons.
3. The method of claim 2 wherein said moiety is substituted with a functional
 15 group selected from the group consisting of polyvinyl alcohol, hydroxy, nitro, bromo, cyano, and aldehyde.
4. The method of any one of claims 1 to 3, wherein said separation is by Matched Ion Polynucleotide Chromatography.
5. The method of any one of claims 1 to 4, wherein said beads are subjected to
 20 an acid wash treatment.
6. The method of any one of claims 1 to 5, wherein said beads are comprised of a polymer selected from mono- and di-vinyl substituted aromatic compounds, acrylates, methacrylates, polyolefins, polyesters, polyurethanes, polyamides, polycarbonates, and fluorosubstituted ethylenes.
7. A method for separating a mixture of polynucleotides, said method being
 25 substantially as hereinbefore described with reference to any one of the examples.
8. A polymeric bead having an average bead diameter of 0.5-100µm and having a surface composition essentially completely unsubstituted, or essentially completely substituted with a moiety selected from the group consisting of methyl, ethyl,
 30 hydrocarbon having from 23 to 1 000 000 carbons, and hydrocarbon polymer having from 23 to 1 000 000 carbons, said bead being substantially free from multivalent cations which are free to bind with DNA and being characterised by having a DNA Separation Factor of at least 0.5.



9. The bead of claim 8, wherein said bead is subjected to an acid wash treatment.

10. The bead of claim 8 or claim 9, wherein said bead is comprised of a copolymer of vinyl aromatic monomers and said vinyl aromatic monomers are selected from the group consisting of styrene, alkyl substituted styrene, α -methylstyrene and alkyl substituted α -methylstyrene.

11. The polymer bead of any one of claims 8 to 10, wherein said bead is substituted with a functional group selected from the group consisting of polyvinyl alcohol, hydroxy, nitro, bromo, cyano, and aldehyde.

12. The polymer bead of any one of claims 8 to 11, wherein said bead comprises a polymer selected from mono- and di-vinyl substituted aromatic compounds, acrylates, methacrylates, polyolefins, polyesters, polyurethanes, polyamides, polycarbonates, and fluorosubstituted ethylenes.

13. A polymeric bead, substantially as hereinbefore described with reference to any one of the examples.

Dated 17 September, 2001

Transgenomic, Inc.

Patent Attorneys for the Applicant/Nominated Person

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1/8

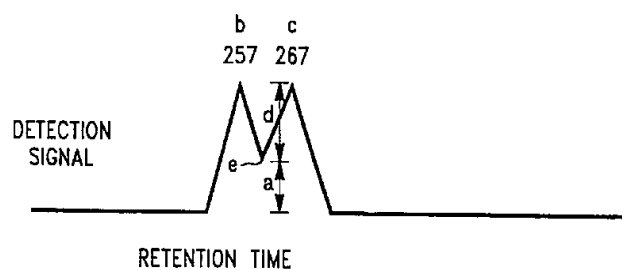
DNA SEPARATION FACTOR = $d/(a+d)$ 

FIG.-1

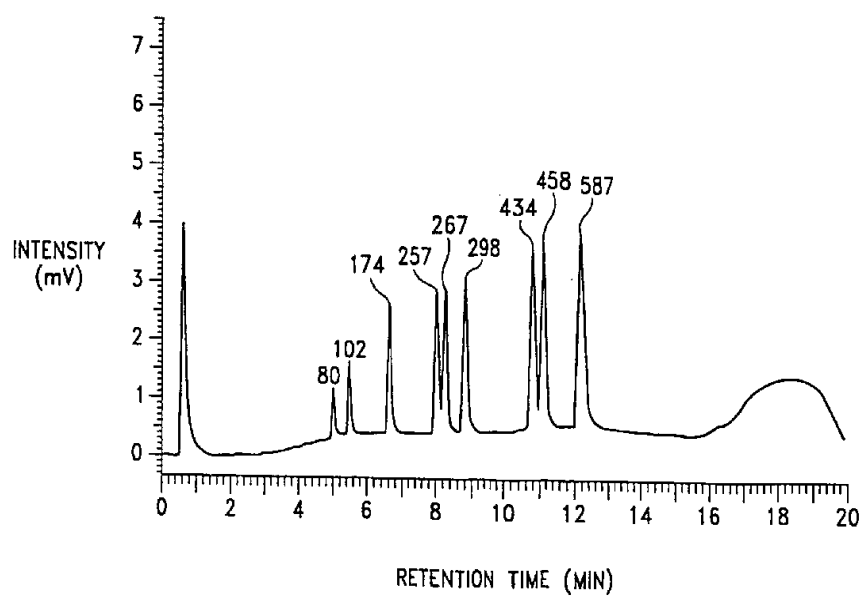


FIG.-2

2/8

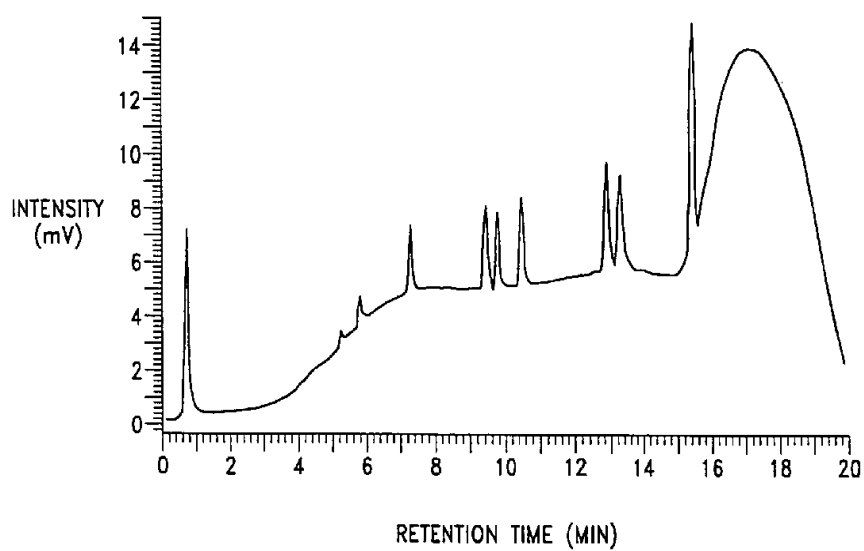


FIG.-3

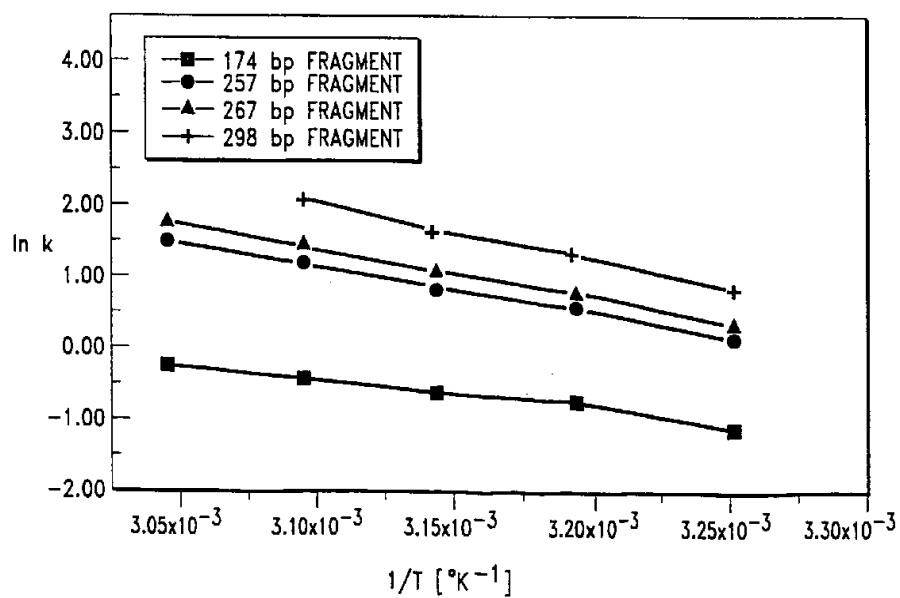


FIG.-4

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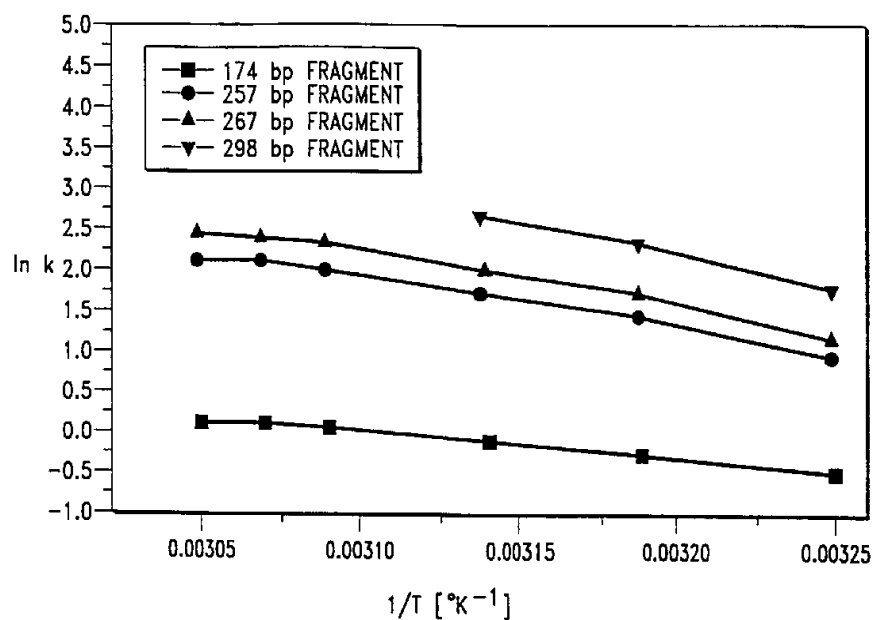


FIG.-5

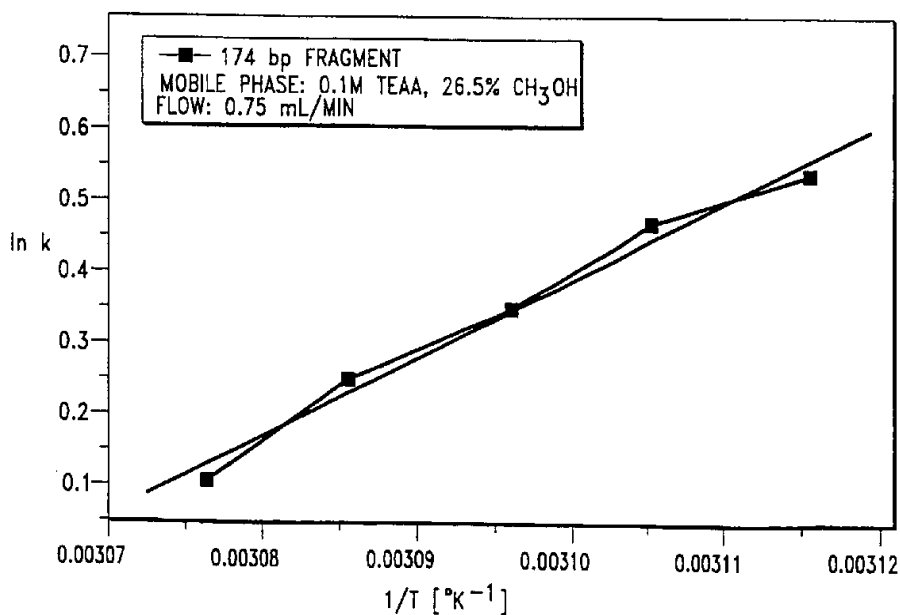
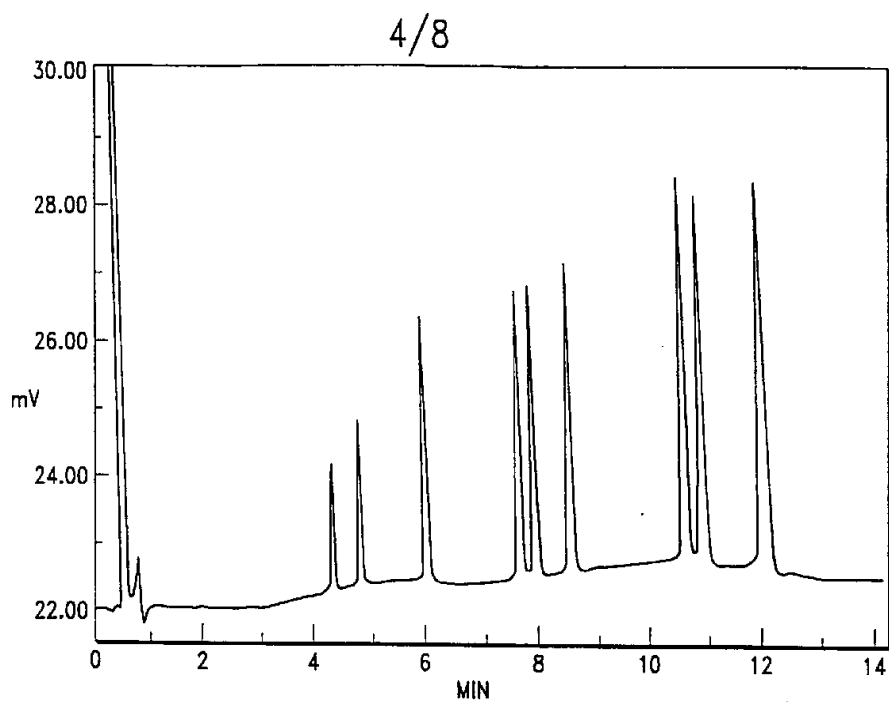
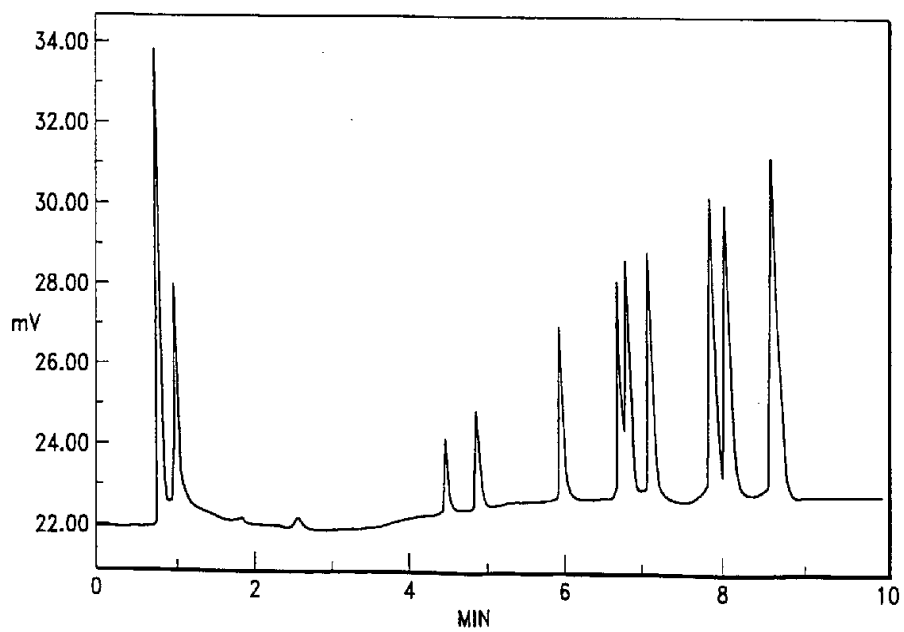
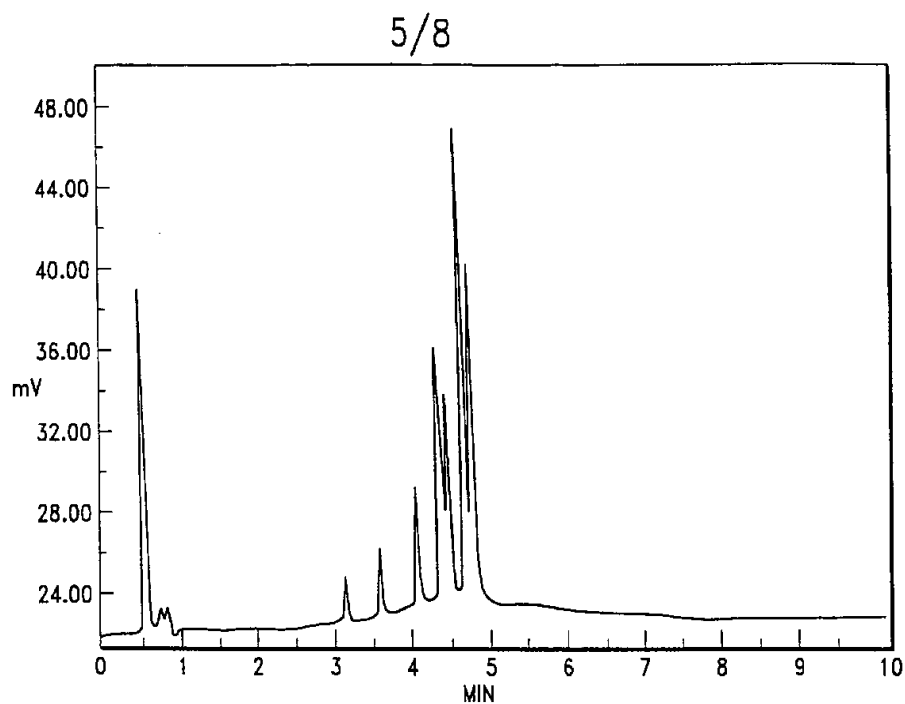
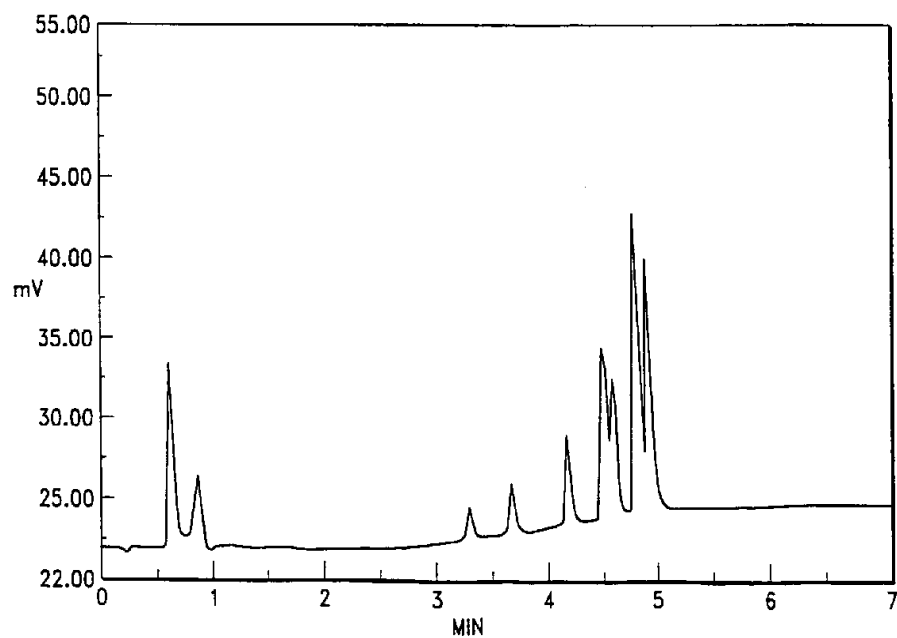
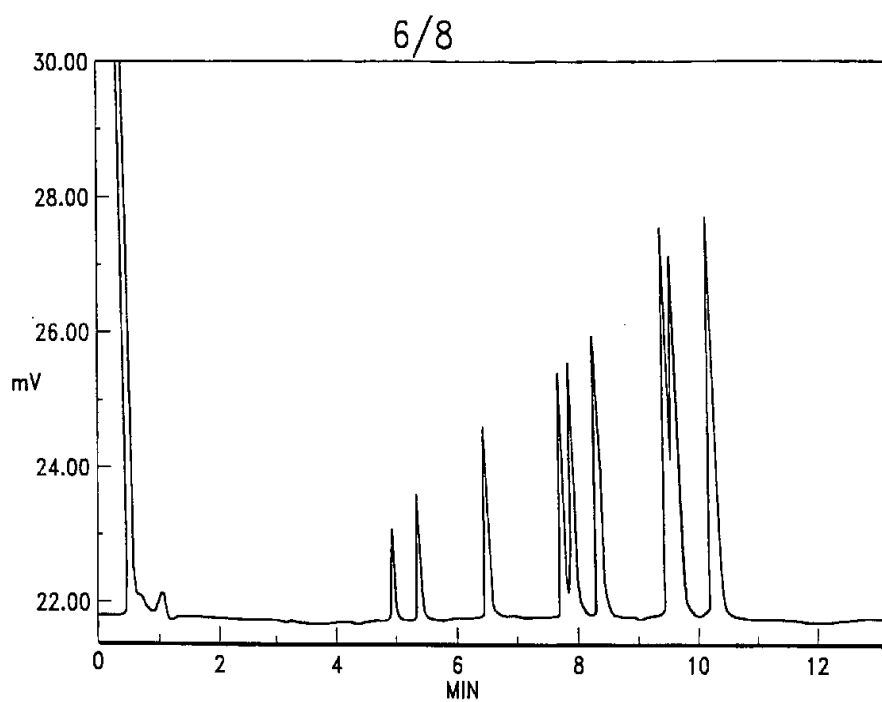
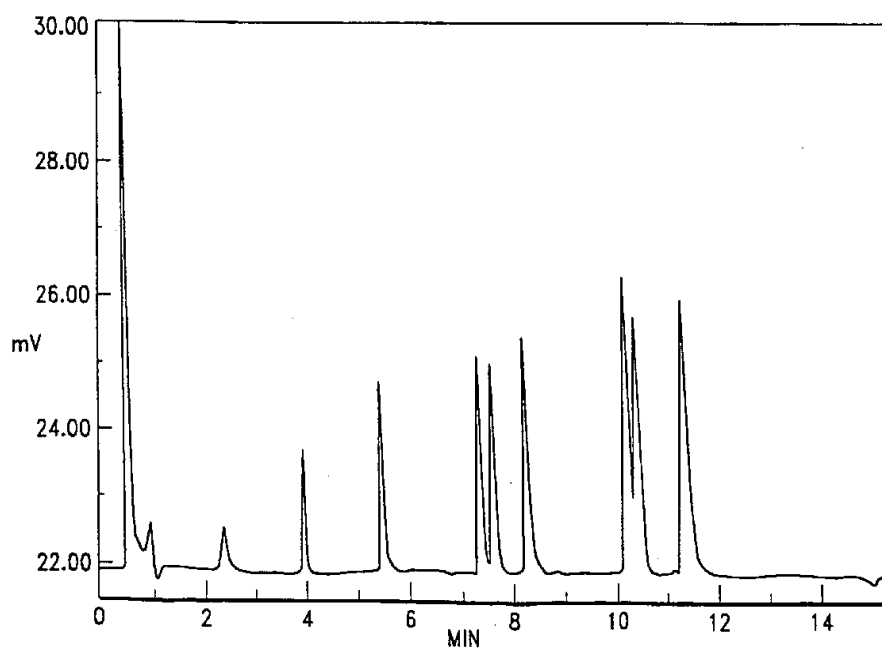


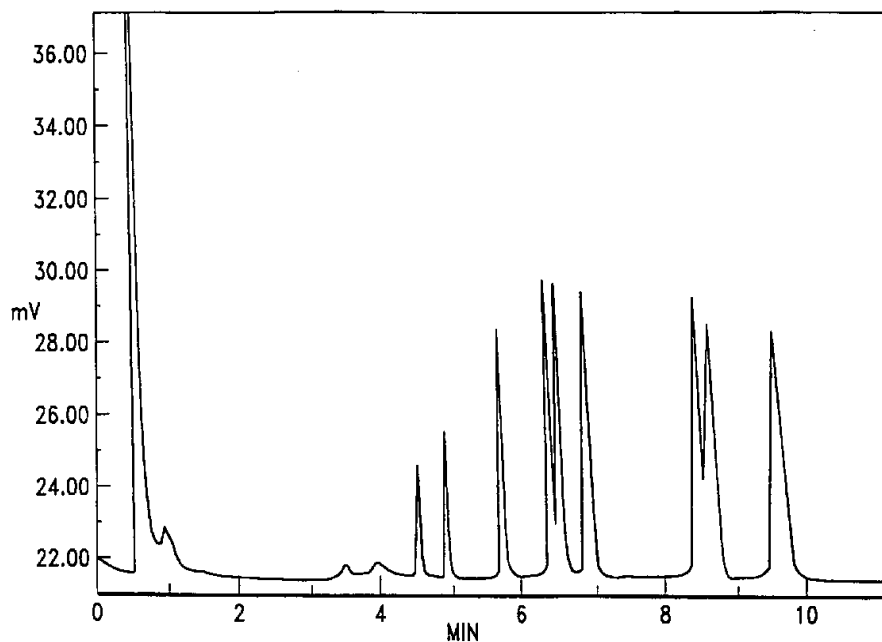
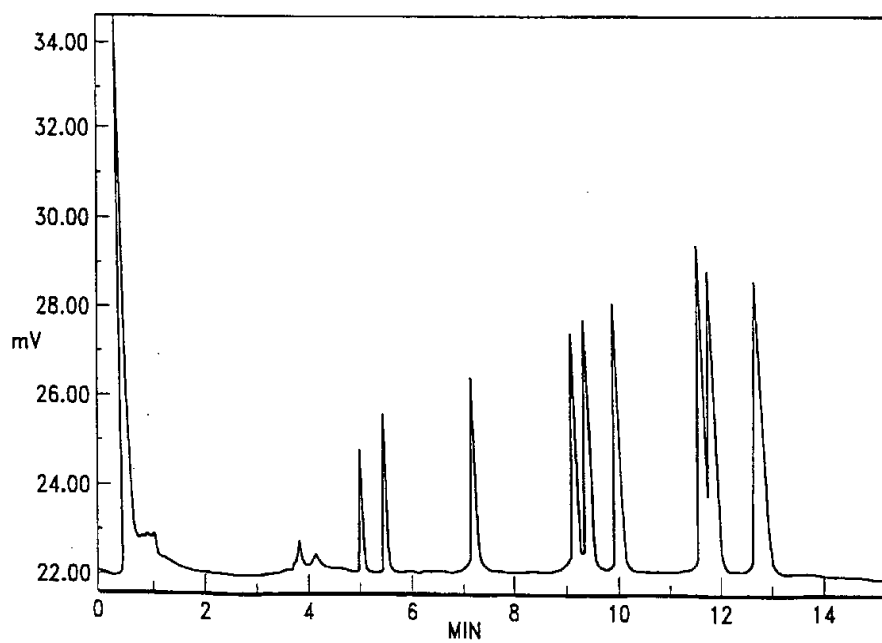
FIG.-6

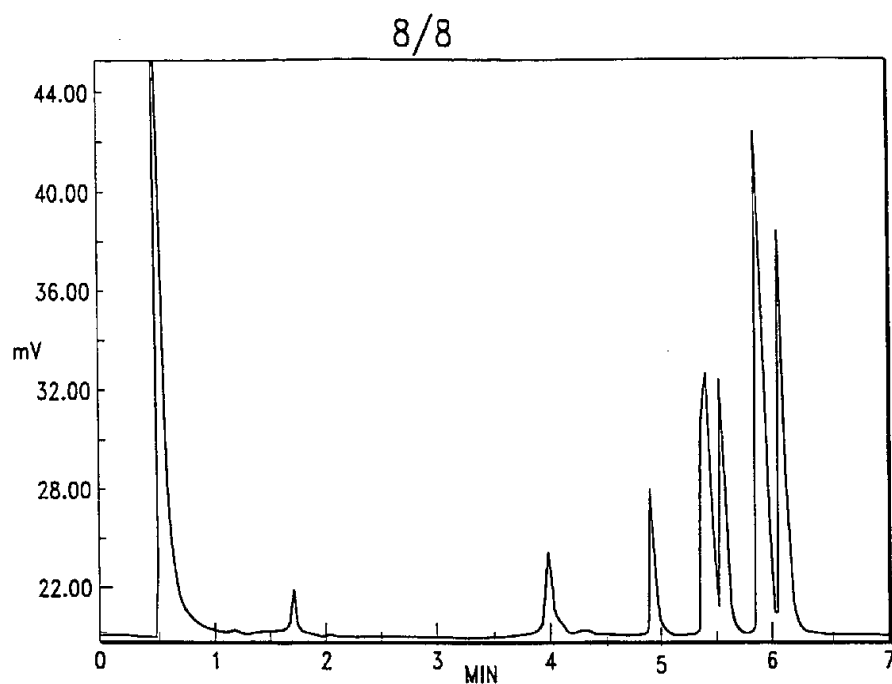
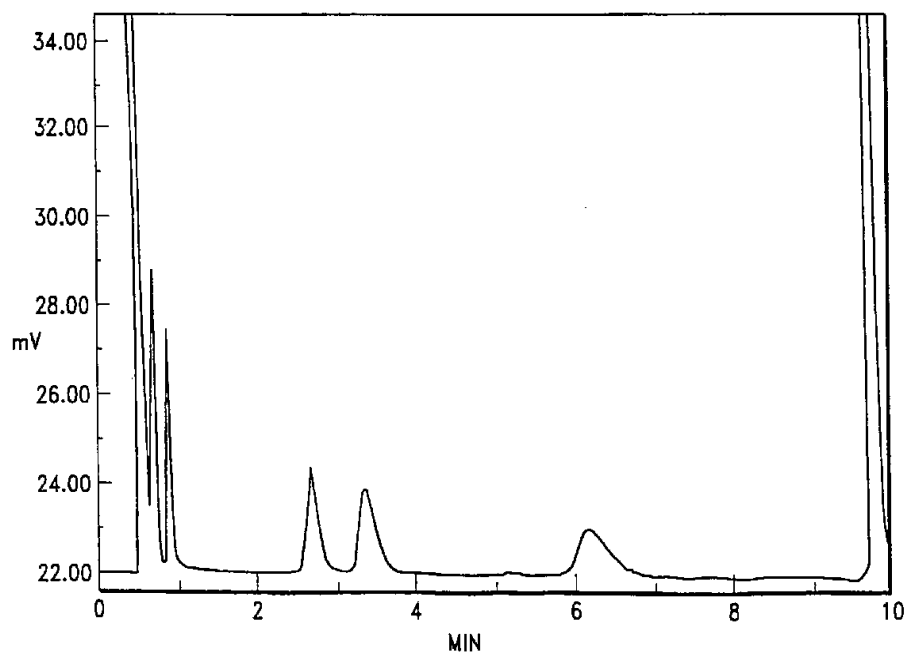
*FIG.-7**FIG.-8*

*FIG.-9**FIG.-10*

*FIG.-11**FIG.-12*

7/8

*FIG.-13**FIG.-14*

*FIG.-15**FIG.-16*