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(54) Title: HUMAN DNASE II

[illegible]

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#61  AAGAGGACCACTCCAAATGTCGTCGCTCCCAAGAGGAGCCCTGAGCTCGGTCGGTGCAC
277  E D E K S R N C V S F K G F W T C
      .
1021 GAATCGGAACCAAGGAGAGAGGACACCGGCTGCGGACACACTGTGTGCTCCACCTCCACG
297  K R T K A A C C A G G G A G A G G A C C G G C T G C G G A C A C T G T G T G C T C C A C C G
      .
1081 CCTCGGAAAGACCTTCCACCTGCTGTGGGAGAACTACGACGCTGATATATGATATGACCA
317  L M K A F Q D P L V K N Y Q P C N G N A R
      .
1161 GAGGCCACAGACAGACCTTATAGATGCTGGCCCTTATGCGCAGAGTGCAGTGTGCTCAGTTAT
337  K F S H A Y R E
      .
1261 TAATCTGACACACTTGGGAGACAGAGAGGAGAGTCACTTGAATCTCAGGATTTCCAGAG
1361 CAGCCTTGCGCTACATATGTAAGAGCAGACTCTACTCTAGAGACTTAAAGAAAGTTATGCCAGC
1381 ACGGTGTATATGCTGTATATGTCGACGACCACTGAGCGCAGAGGATGATTTAAGACAGAGAG
1381 ATCATGCGCAGATGAGGATCTAGATTTAGCCCAACCTGCGGTCACTAGCAGACACTCTGTT
1441 AAAAAAAGAGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG
1461 CTCTGGGAGAGCTGATATGAGCAGATCTAGTTAGAGTCAAGAGATTCGAGACAGAGCTGAGC
1501 ACATGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG
1521 ACATGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG

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(57) Abrégé/Abstract:

This invention relates to a novel human deoxyribonuclease, referred to as human DNase II. The invention provides nucleic acid sequences encoding human DNase II, thereby enabling the production of human DNase II by recombinant DNA methods in

(57) **Abrégé(suite)/Abstract(continued):**

quantities sufficient for clinical use. The invention also relates to pharmaceutical compositions and diagnostic and therapeutic uses of human DNase II.

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(57) Abstract

This invention relates to a novel human deoxyribonuclease, referred to as human DNase II. The invention provides nucleic acid sequences encoding human DNase II, thereby enabling the production of human DNase II by recombinant DNA methods in quantities sufficient for clinical use. The invention also relates to pharmaceutical compositions and diagnostic and therapeutic uses of human DNase II.

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1  GAATTTCGGGCAAGCTGTGGCTCTGATGTAAGCCGAGGCCCCGCAAGTCTCCGACACAGATT
61  CTGGGAGTCTCAGCCCTCAATGACAGCACTATGATCTCCGCTGTCTGTGGAGACAGCTCTGCTGCG
-16  H 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16
121  CCCCCCGGGGCGCCCTGACATCTGCTACGGGAGACTCCGGGCGAGCCTCTAGACATGTTTCTGTGT
-4  F A G A L T C T G D S Q Q P V D W V V V
181  CTACAGAGTGCACAGCTTTAGAGAGGCTCCGGAGGCGCGCCAGAGAGGCGCTCAGTTACAA
17  T K L P A L R G G A A G G A A G R G L G L A A
241  GTATTCTGACAGAGAGCTCCGAGAGAGCTCCGGGAGACAGCTCATCAGACAGCCGGA
17  Y L D E S G Q W R D D R A L I N S P E
321  GGGGGCGCTGGGGCGAGAGCTTCGAGGCCCTGTACCGGAGCAGACAGAGCAGGCTCTGCTCT
97  Q A Y V G R L A C C G C L V R S H M T S G L A F
381  CCTGCTCTACAAATGACCAAGCTCTCAACCAAGAGAGCTCTGCTGCTCTGCTGCTGGG
77  L Y L M A G G T P P P Q P S K A Q D S S M R G
441  GCACAGAGAGGTTCTGCTGCTCTTCTGACACAGATGGGGGCTCTGCTGCTCTGCTCAGACAT
97  H T R G V L L L L D N D D Q F W L V M S V
481  ACCTATGTTTGGCTCTACCGGCGCTCTCTGCTGCTATGACAGAGCGGCTCTATAGCGGCTGTAC
117  P H P T P P P A S A S A Y S H M S A C T
541  CTACGGGCGAGCCTCTGCTCTGGTGTGCTCTTCCGCTGCTGCTGCTCTCTCAGAGAGGGCGAA
137  V G Q T F L L C V S Y P P F A Q C F S K M G K
581  CGACGCTGACTACACTACCTACCTCTGGGCTCTATTAACCTACCGCTCGAAGGAGATCTTTGCCCA
167  G L T P T P T P W V Y A A G G G G G G G I F A G
641  GGAAATCCCGGAGCTTGGAGAGGTTGTGTCAGAGGCGACACAGTATGACCAAGAGAGCTCGAGAA
177  X P P D L E N V Y R G N B V S Q R P W M
701  CAGCAGATCTCTACTCATCATCTCGAGCGCGGGGCTGTTTTCAGACAGCTTTCAGCAAGTTTCAG
197  E 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16
761  CAATTTGAGATGAGCTGCTACTCCGCTGTGTTGGCAGACAGCTTTGTGACACAGCTGCA
217  H P Q D D L Y S G W L A A L G T D T N L Q
821  GGTTCAGTGTCTGACCAAAATCTGTAGGACATCTGCTGCTCTTACTGCTCTGGATCTATCGCA
237  V Q P W N X T V G I L P S H C S I D I W Q
881  GGTTCCTGAATGTGAACACAGATGATCTTCCCTCGAGACCGGCGCAGAGCTTTCAGACAGCA
257  V L N W Y Q I A P P O P A G C G S F F N M T
941  GGTTCCTGAATGTGAACACAGATGATCTTCCCTCGAGACCGGCGCAGAGCTTTCAGACAGCA
277  V L N W Y Q I A P P O P A G C G S F F N M T

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961 AAGAGGACGACCTCAAAATGTCGTCTGTCCCAAGAAAGGCGCTGGACCTGGGTGGGTGAAT
 277 K D N G S R X M C V S F R Q V W T C C V G D M
 1021 GAATCGGAACCAAGGAGGAGCAACCGGCGGCGGACACTGTGTGTCGACGCTGCCAAG
 327 N R N G A C C G G E X Q G O G O T L C C C A G L C T F A
 1081 CCTCTGGAAAGCCCTTCCACGCCCTCGGTGAAGAACTACCAAGCCCTGGTAATGCGATGCCAC
 319 L W K E A F Q P L V R M T Q P C M G N A R
 1161 GAAAGCCACAGCAAGCTTATAAGATCTTCCCTTATATGCGCAGGTGCATGCGCTCAGCTATG
 339 K P S R A Y R E
 1201 TAATGCCACACACTTTGGGAAAGCCAAAGAGGGAGGATCACTTGAACCTCAGGAATCTCGAAG
 1281 CAGCCTCTGAAATCAATATGAAAGGACCACTCTGACTAGAAACTTAAAGAAATTTAGCCGAGC
 1381 ACGGTGTATGATACCTCTGTATGCTGACCTGACCTGAGGAGGAGCATTTGAAGACCAAGAG
 1361 ATCATGGGTCAAGATGAATATGATTTACGCCAACTCGGTCACATAGCAAGACTCTGTCTT
 1441 AAAAATAAATAGGAGGAGCGAAGGACACGGTGGGTGCATGTCCTCAGCTTTGTAATACCCAGCA
 1501 CTTTGGGAGGCTGAGATGCGGACATCACTTGAAGTCAAGAGTTCGAGAACCAAGCTTGCCAC
 1581 ACATGTGGTGAAGCCCC

HUMAN DNASE II

Field of the Invention

The present invention relates to newly identified human deoxyribonuclease (DNase) protein, nucleic acid encoding such protein, the
5 use of such protein and nucleic acid, as well as the production of such protein and nucleic acid, for example, by recombinant DNA methods.

Background of the Invention

Deoxyribonuclease (DNase) is a phosphodiesterase capable of hydrolyzing polydeoxyribonucleic acid, and is known to occur in several
10 molecular forms. Based on their biochemical properties and enzymatic activities, DNase proteins have been classified as two types, DNase I and DNase II. DNase I proteins have a pH optimum near neutrality, an obligatory requirement for divalent cations, and produce 5'-phosphate nucleotides on hydrolysis of DNA. DNase II proteins exhibit an acid pH
15 optimum, do not require divalent cations for activity, and produce 3'-phosphate nucleotides on hydrolysis of DNA.

DNase from various species have been purified to a varying degree. For example, various forms of bovine DNase I have been purified and completely sequenced (Liao, et al., J. Biol. Chem. 248:1489-1495 (1973);
20 Oefner, et al., J. Mol. Biol. 192:605-632 (1986); Lahm, et al., J. Mol. Biol. 221:645-667 (1991)), and DNA encoding bovine DNase I has been cloned and expressed (Worrall, et al., J. Biol. Chem. 265:21889-21895 (1990)). Porcine and oricine DNase I proteins also have been purified and completely sequenced (Paudel, et al., J. Biol. Chem. 261:16006-16011 (1986); Paudel,
25 et al., J. Biol. Chem. 261:16012-16017 (1986)).

DNA encoding a human DNase I has been isolated and sequenced and the DNA has been expressed in recombinant host cells, thereby enabling the production of human DNase I in commercially useful quantities. Shak, et al., Proc. Natl. Acad. Sci. 87:9188-9192 (1990). The term "human DNase I"
30 will be used hereafter to refer to the mature polypeptide disclosed in Shak, et al.

DNA encoding other polypeptides having homology to human DNase I also have been identified. Rosen, et al., PCT Patent Publication No. WO 95/30428, published November 16, 1995; and Parrish, et al., Hum. Mol. Genet. 4:1557-1564 (1995).

5 DNase I has a number of known utilities and has been used for therapeutic purposes. Its principal therapeutic use has been to reduce the viscoelasticity of pulmonary secretions (including mucus) in such diseases as pneumonia and cystic fibrosis (CF), thereby aiding in the clearing of respiratory airways. See e.g., Lourenco, et al., Arch. Intern. Med.
10 142:2299-2308 (1982); Shak, et al., Proc. Natl. Acad. Sci. 87:9188-9192 (1990); Hubbard, et al., New Engl. J. Med. 326:812-815 (1992); Fuchs, et al., New Engl. J. Med. 331:637-642 (1994); Bryson, et al., Drugs 48:894-906 (1994). Mucus also contributes to the morbidity of chronic bronchitis, asthmatic bronchitis, bronchiectasis, emphysema, acute and chronic
15 sinusitis, and even the common cold. DNase I is effective in reducing the viscoelasticity of pulmonary secretions by hydrolyzing, or degrading, high-molecular-weight DNA that is present in such secretions. Shak, et al., Proc. Natl. Acad. Sci. 87:9188-9192 (1990); Aitken, et al., J. Am. Med. Assoc. 267:1947-1951 (1992).

20 Various forms of DNase II also have reportedly been purified, including bovine DNase II (Lesca, J. Biol. Chem. 251:116-123 (1976)), human DNase II (Yamanaku, et al., J. Biol. Chem. 249:3884-3889 (1974); Murai, et al., J. Biochem. 87:1097-1103 (1980); Harosh, et al., Eur. J. Biochem. 202:479-484 (1991); Yasuda, et al., Biochem. Biophys. Acta 1119:185-193
25 (1992)), porcine DNase II (Bernardi, et al., Biochemistry 4:1725-1729 (1965); Liao, et al., J. Biol. Chem. 260:10708-10713 (1990)), and rat DNase II (Dulaney, et al., J. Biol. Chem. 247:1424-1432 (1972)). The physical properties of the human DNase II proteins described in these reports vary considerably (e.g., reported molecular weights range from
30 32,000 to 45,000 Daltons), which leads to uncertainty whether there is one or multiple naturally occurring forms of the human protein.

Recent interest in human DNase II has arisen because of its possible role in the programmed cell death process of apoptosis (Barry, et al., Arch. Biochem. Biophys. 300:440-450 (1993); Barry, et al., Cancer Res.
35 53:2349-2357 (1993)). One of the events that is characteristic of that process is the degradation of nuclear DNA into nucleosomal fragments. The ability to prevent or inhibit the expression of human DNase II or its enzymatic activity within human cells may be important in preventing or

limiting such intracellular destruction of DNA, and thus may be an effective means of interrupting the process of apoptosis. In other instances, it may be useful to increase the expression of human DNase II within a certain population of cells within a human patient, such as cancer
5 cells, in order to induce apoptosis of those cells.

Summary of the Invention

The present invention provides human DNase II protein, as well as analogs and variants thereof, that have DNA-hydrolytic activity. As is
10 characteristic of DNase II proteins in general, the human DNase II of the present invention exhibits an acid pH optimum, and does not require divalent cation for activity.

The invention also provides nucleic acids encoding human DNase II, recombinant vectors comprising such nucleic acids, recombinant host cells
15 transformed with those nucleic acids or vectors, and processes for producing human DNase II by means of recombinant DNA technology. The invention includes the use of such nucleic acids and vectors for in vivo or ex vivo gene therapy.

The invention also provides complementary nucleic acids, including
20 so-called anti-sense oligonucleotides, that are capable of binding to and preventing the expression of nucleic acid within a cell that encodes human DNase II.

The invention also provides pharmaceutical compositions comprising human DNase II, optionally together with a pharmaceutically acceptable
25 excipient, as well as substantially purified antibodies that are capable of binding to human DNase II.

The invention also provides methods for reducing the viscoelasticity or viscous consistency of DNA-containing material in a patient, comprising administering a therapeutically effective dose of human DNase II to the
30 patient. The invention is particularly directed to a method of treating a patient having a disease such as cystic fibrosis, chronic bronchitis, pneumonia, bronchiectasis, emphysema, asthma, or systemic lupus erythematosus, that comprises administering a therapeutically effective amount of human DNase II to the patient. The invention also is directed
35 to the use of human DNase II in vitro, such as for degrading DNA that is present in a biological specimen or other material, and in diagnostic and other assays.

These and other aspects of the invention will be apparent to the ordinary skilled artisan upon consideration of the following detailed description.

Brief Description of the Figures

5 Figure 1 shows the nucleotide sequence (SEQ. ID. NO: 1) and deduced amino acid sequence (SEQ. ID. NO: 2) of human DNase II. The predicted leader (signal) amino acid sequence is underlined and the start of the mature protein is indicated by the arrowhead. The eight cysteine residues are indicated by asterisks and potential N-linked glycosylation sites are
10 boxed.

Detailed Description

The various aspects of the present invention are accomplished by first providing isolated DNA comprising the nucleotide coding sequence for human DNase II. By providing the full nucleotide coding sequence for human
15 DNase II, the invention enables the production of human DNase II by means of recombinant DNA technology, thereby making available for the first time sufficient quantities of substantially pure human DNase II protein for diagnostic and therapeutic uses.

As used herein, the term "human DNase II" refers to the polypeptide
20 having the amino acid sequence of the mature protein set forth in Figure 1 as well as modified and variant forms thereof as described herein. Modified and variant forms of human DNase II are produced in vitro by means of chemical or enzymatic treatment or in vivo by means of recombinant DNA technology. Such polypeptides differ from human DNase II, for example, by
25 virtue of one or more amino acid substitutions, insertions, and/or deletions, or in the extent or pattern of glycosylation, but in all cases will possess DNA-hydrolytic activity. A "variant" or "amino acid sequence variant" of human DNase II is a polypeptide that comprises an amino acid sequence different from that of human DNase II. Generally, a variant will
30 have at least 80% sequence identity, preferably at least 90% sequence identity, more preferably at least 95% sequence identity, and most preferably at least 98% sequence identity with human DNase II. Percentage sequence identity is determined, for example, by the Fitch, et al., Proc. Natl. Acad. Sci. USA 80:1382-1386 (1983), version of the algorithm
35 described by Needleman, et al., J. Mol. Biol. 48:443-453 (1970), after aligning the sequences to provide for maximum homology. Such variants

include naturally occurring allelic forms of human DNase II that are of human origin as well as naturally occurring homologs of human DNase II that are found in other animal species.

"DNA-hydrolytic activity" refers to the enzymatic activity of human DNase II in hydrolyzing (cleaving) substrate DNA to yield 3'-phosphorylated oligonucleotide end products. DNA-hydrolytic activity is readily determined by any of several different methods known in the art, including analytical polyacrylamide and agarose gel electrophoresis, hyperchromicity assay (Kunitz, J. Gen. Physiol. 33:349-362 (1950); Kunitz, J. Gen. Physiol. 33:363-377 (1950)), or methyl green assay (Kurnick, Arch. Biochem. 29:41-53 (1950); Sinicropi, et al., Anal. Biochem. 222:351-358 (1994)). As a routine matter, the pH and buffer used in these methods are varied so as to provide the conditions wherein the particular human DNase II will exhibit such activity, if any.

For convenience, substitutions, insertions, and/or deletions in the amino acid sequence of human DNase II are usually made by introducing mutations into the corresponding nucleotide sequence of the DNA encoding human DNase II, for example by site-directed mutagenesis. Expression of the mutated DNA then results in production of the variant human DNase II, having the desired amino acid sequence.

Whereas any technique known in the art can be used to perform site-directed mutagenesis, e.g. as disclosed in Sambrook, et al., Molecular Cloning: A Laboratory Manual, Second Edition (Cold Spring Harbor Laboratory Press, New York (1989)), oligonucleotide-directed mutagenesis is the preferred method for preparing the human DNase II variants of this invention. This method, which is well known in the art (Zoller, et al., Meth. Enzymol. 100:4668-500 (1983); Zoller, et al., Meth. Enzymol. 154:329-350 (1987); Carter, Meth. Enzymol. 154:382-403 (1987); Kunkel, et al., Meth. Enzymol. 154:367-382 (1987); Horwitz, et al., Meth. Enzymol. 185:599-611 (1990)), is particularly suitable for making substitution variants, although it may also be used to conveniently prepare deletion and insertion variants, as well as variants having multiple substitution, insertion, and/or deletion mutations.

Briefly, in carrying out site-directed mutagenesis of DNA encoding human DNase II (or a variant thereof), the DNA is altered by first hybridizing an oligonucleotide encoding the desired mutation to a single strand of the DNA. After hybridization, a DNA polymerase is used to synthesize an entire second strand, using the hybridized oligonucleotide

as a primer, and using the single strand of the DNA as a template. Thus, the oligonucleotide encoding the desired mutation is incorporated in the resulting double-stranded DNA.

Oligonucleotides may be prepared by any suitable method, such as by
5 purification of a naturally occurring DNA or by in vitro synthesis. For example, oligonucleotides are readily synthesized using various techniques in organic chemistry, such as described by Narang, et al., Meth. Enzymol. 68:90-98 (1979); Brown, et al., Meth. Enzymol. 68:109-151 (1979); Caruthers, et al., Meth. Enzymol. 154:287-313 (1985). The general approach
10 to selecting a suitable oligonucleotide for use in site-directed mutagenesis is well known. Typically, the oligonucleotide will contain 10-25 or more nucleotides, and will include at least 5 nucleotides on either side of the sequence encoding the desired mutation so as to ensure that the oligonucleotide will hybridize preferentially at the desired location to
15 the single-stranded DNA template molecule.

"Polymerase chain reaction," or "PCR," generally refers to a method for amplification of a desired nucleotide sequence in vitro, as described, for example, in U.S. Pat. No. 4,683,195. In general, the PCR method involves repeated cycles of primer extension synthesis, using
20 oligonucleotide primers capable of hybridizing preferentially to a template nucleic acid.

PCR mutagenesis (Higuchi, in PCR Protocols, pp.177-183 (Academic Press, 1990); Vallette, et al., Nuc. Acids Res. 17:723-733 (1989)) is also suitable for making the variants of human DNase II. Briefly, when small
25 amounts of template DNA are used as starting material in a PCR, primers that differ slightly in sequence from the corresponding region in the template DNA can be used to generate relatively large quantities of a specific DNA fragment that differs from the template sequence only at the positions where the primers differ from the template. For introduction of
30 a mutation into a plasmid DNA, for example, the sequence of one of the primers includes the desired mutation and is designed to hybridize to one strand of the plasmid DNA at the position of the mutation; the sequence of the other primer must be identical to a nucleotide sequence within the opposite strand of the plasmid DNA, but this sequence can be located
35 anywhere along the plasmid DNA. It is preferred, however, that the sequence of the second primer is located within 200 nucleotides from that of the first, such that in the end the entire amplified region of DNA bounded by the primers can be easily sequenced. PCR amplification using

a primer pair like the one just described results in a population of DNA fragments that differ at the position of the mutation specified by the primer, and possibly at other positions, as template copying is somewhat error-prone. Wagner, et al., in PCR Topics, pp.69-71 (Springer-Verlag, 5 1991).

If the ratio of template to product amplified DNA is extremely low, the majority of product DNA fragments incorporate the desired mutation(s). This product DNA is used to replace the corresponding region in the plasmid that served as PCR template using standard recombinant DNA methods. 10 Mutations at separate positions can be introduced simultaneously by either using a mutant second primer, or performing a second PCR with different mutant primers and ligating the two resulting PCR fragments simultaneously to the plasmid fragment in a three (or more)-part ligation.

Another method for preparing variants, cassette mutagenesis, is based 15 on the technique described by Wells et al., Gene, 34:315-323 (1985). The starting material is the plasmid (or other vector) comprising the DNA sequence to be mutated. The codon(s) in the starting DNA to be mutated are identified. There must be a unique restriction endonuclease site on each side of the identified mutation site(s). If no such restriction sites 20 exist, they may be generated using the above-described oligonucleotide-mediated mutagenesis method to introduce them at appropriate locations in the DNA. The plasmid DNA is cut at these sites to linearize it. A double-stranded oligonucleotide encoding the sequence of the DNA between the restriction sites but containing the desired 25 mutation(s) is synthesized using standard procedures, wherein the two strands of the oligonucleotide are synthesized separately and then hybridized together using standard techniques. This double-stranded oligonucleotide is referred to as the cassette. This cassette is designed to have 5' and 3' ends that are compatible with the ends of the linearized 30 plasmid, such that it can be directly ligated to the plasmid. The resulting plasmid contains the mutated DNA sequence.

The presence of mutation(s) in a DNA is determined by methods well known in the art, including restriction mapping and/or DNA sequencing. A preferred method for DNA sequencing is the dideoxy chain termination method 35 of Sanger, et al., Proc. Natl. Acad. Sci. USA 72:3918-3921 (1979).

DNA encoding human DNase II is inserted into a replicable vector for further cloning or expression. "Vectors" are plasmids and other DNAs that are capable of replicating within a host cell, and as such, are useful for

performing two functions in conjunction with compatible host cells (a vector-host system). One function is to facilitate the cloning of nucleic acid that encodes human DNase II, i.e., to produce usable quantities of the nucleic acid. The other function is to direct the expression of human DNase
5 II. One or both of these functions are performed by the vector in the particular host cell used for cloning or expression. The vectors will contain different components depending upon the function they are to perform.

The human DNase II of the present invention may be expressed in the
10 form of a preprotein wherein the DNase II includes a leader or signal sequence, or may be in the form of a mature protein which lacks a leader or signal sequence. The human DNase II also may be in the form of a fusion protein wherein additional amino acid residues are covalently joined to the amino- or carboxy-terminus of the preprotein or mature form of the DNase.

15 To produce human DNase II, an expression vector will comprise DNA encoding human DNase II, as described above, operably linked to a promoter and a ribosome binding site. The human DNase II then is expressed directly in recombinant cell culture, or as a fusion with a heterologous polypeptide, preferably a signal sequence or other polypeptide having a
20 specific cleavage site at the junction between the heterologous polypeptide and the human DNase II amino acid sequence.

"Operably linked" refers to the covalent joining of two or more DNA sequences, by means of enzymatic ligation or otherwise, in a configuration relative to one another such that the normal function of the sequences can
25 be performed. For example, DNA for a presequence or secretory leader is operably linked to DNA for a polypeptide if it is expressed as a preprotein that participates in the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding sequence if it affects the transcription of the sequence; or a ribosome binding site is operably
30 linked to a coding sequence if it is positioned so as to facilitate translation. Generally, "operably linked" means that the DNA sequences being linked are contiguous and, in the case of a secretory leader, contiguous and in reading phase. Linking is accomplished by ligation at convenient restriction sites. If such sites do not exist, then synthetic
35 oligonucleotide adaptors or linkers are used, in conjunction with standard recombinant DNA methods.

Prokaryotes (e.g., *E. coli*, strains of *Bacillus*, *Pseudomonas*, and other bacteria) are the preferred host cells for the initial cloning steps

of this invention. They are particularly useful for rapid production of large amounts of DNA, for production of single-stranded DNA templates used for site-directed mutagenesis, and for DNA sequencing of the variants generated. Prokaryotic host cells also may be used for expression of DNA
5 encoding human DNase II. Polypeptides that are produced in prokaryotic cells typically will be non-glycosylated.

In addition, human DNase II may be expressed in eukaryotic host cells, including eukaryotic microbes (e.g., yeast) or cells derived from an animal or other multicellular organism (e.g., Chinese hamster ovary
10 cells, and other mammalian cells), or in live animals (e.g., cows, goats, sheep). Insect cells and fungi also may be used.

Cloning and expression methodologies are well known in the art. Examples of prokaryotic and eukaryotic host cells, and starting expression vectors, suitable for use in producing human DNase II are, for example,
15 those disclosed in Shak, PCT Patent Publication No. WO 90/07572, published July 12, 1990. To obtain expression of human DNase II, an expression vector of the invention is introduced into host cells by transformation or transfection, and the resulting recombinant host cells are cultured in conventional nutrient media, modified as appropriate for inducing
20 promoters, selecting recombinant cells, or amplifying human DNase II DNA. The culture conditions, such as temperature, pH, and the like, are those previously used with the host cell, and as such will be apparent to the ordinarily skilled artisan.

"Transformation" and "transfection" are used interchangeably to refer
25 to the process of introducing DNA into a cell. Following transformation or transfection, the DNA may integrate into the host cell genome, or may exist as an extrachromosomal element. If prokaryotic cells or cells that contain substantial cell wall constructions are used as hosts, the preferred methods of transfection of the cells with DNA is the calcium
30 treatment method described by Cohen et al., Proc. Natl. Acad. Sci. 69:2110-2114 (1972) or the polyethylene glycol method of Chung et al., Nuc. Acids. Res. 16:3580 (1988). If yeast are used as the host, transfection is generally accomplished using polyethylene glycol, as taught by Hinnen, Proc. Natl. Acad. Sci. U.S.A., 75: 1929-1933 (1978). If mammalian cells
35 are used as host cells, transfection generally is carried out by the calcium phosphate precipitation method, Graham, et al., Virology 52:546 (1978), Gorman, et al., DNA and Protein Eng. Tech. 2:3-10 (1990). However, other known methods for introducing DNA into prokaryotic and eukaryotic

cells, such as nuclear injection, electroporation, or protoplast fusion also are suitable for use in this invention.

Particularly useful in this invention are expression vectors that provide for the transient expression in mammalian cells of DNA encoding human DNase II. In general, transient expression involves the use of an expression vector that is able to efficiently replicate in a host cell, such that the host cell accumulates many copies of the expression vector and, in turn, synthesizes high levels of a desired polypeptide encoded by the expression vector. Transient expression systems, comprising a suitable expression vector and a host cell, allow for the convenient positive identification of polypeptides encoded by cloned DNAs, as well as for the rapid screening of such polypeptides for desired biological or physiological properties. Wong, et al., Science 228:810-815 (1985); Lee, et al., Proc. Nat Acad. Sci. USA 82:4360-4364 (1985); Yang, et al., Cell 47:3-10 (1986). Thus, transient expression systems are conveniently used for expressing the DNA encoding amino acid sequence variants of human DNase II, in conjunction with assays to identify those variants that have such useful properties as increased half-life or decreased immunogenicity in vivo, or increased DNA hydrolytic activity at physiological pH.

Human DNase II preferably is secreted from the host cell in which it is expressed, in which case the variant is recovered from the culture medium in which the host cells are grown. In that case, it may be desirable to grow the cells in a serum free culture medium, since the absence of serum proteins and other serum components in the medium may facilitate purification of the variant. If it is not secreted, then the human DNase II is recovered from lysates of the host cells. When the human DNase II is expressed in a host cell other than one of human origin, the variant will be completely free of proteins of human origin. In any event, it will be necessary to purify the human DNase II from recombinant cell proteins in order to obtain substantially homogeneous preparations of the human DNase II. For therapeutic uses, the purified human DNase II preferably will be greater than 99% pure (i.e., any other proteins will comprise less than 1% of the total protein in the purified composition).

It is further contemplated that human DNase II may be produced by a method involving homologous recombination and amplification, for example, as described in PCT Patent Publication No. WO 91/06667, published May 16, 1991. Briefly, this method involves transforming cells containing an endogenous gene encoding human DNase II with a homologous DNA, which

homologous DNA comprises (1) an amplifiable gene (e.g., a gene encoding dihydrofolate reductase (DHFR)), and (2) at least one flanking sequence, having a length of at least about 150 base pairs, which is homologous with a nucleotide sequence in the cell genome that is within or in proximity to the gene encoding human DNase II. The transformation is carried out under conditions such that the homologous DNA integrates into the cell genome by recombination. Cells having integrated the homologous DNA then are subjected to conditions which select for amplification of the amplifiable gene, whereby the human DNase II gene amplified concomitantly. The resulting cells then are screened for production of desired amounts of human DNase II. Flanking sequences that are in proximity to a gene encoding human DNase II are readily identified, for example, by the method of genomic walking, using as a starting point the nucleotide sequence of human DNase II shown in Figure 1. Spoerel, et al., Meth. Enzymol. 152:598-603 (1987).

Generally, purification of human DNase II is accomplished by taking advantage of the differential physicochemical properties of the human DNase II as compared to the contaminants with which it may be associated. For example, as a first step, the culture medium or host cell lysate is centrifuged to remove particulate cell debris. The human DNase II thereafter is purified from contaminant soluble proteins and polypeptides, for example, by ammonium sulfate or ethanol precipitation, gel filtration (molecular exclusion chromatography), ion-exchange chromatography, hydrophobic chromatography, immunoaffinity chromatography (e.g., using a column comprising anti-human DNase II antibodies coupled to Sepharose), tentacle cation exchange chromatography (Frenz, et al., U.S. Patent No. 5,279,823, issued January 18, 1994), reverse phase HPLC, and/or gel electrophoresis.

In some host cells (especially bacterial host cells) the human DNase II may be expressed initially in an insoluble, aggregated form (referred to in the art as "refractile bodies" or "inclusion bodies") in which case it will be necessary to solubilize and renature the human DNase II in the course of its purification. Methods for solubilizing and renaturing recombinant protein refractile bodies are known in the art (see e.g., Builder, et al., U.S. Patent No. 4,511,502, issued April 16, 1985).

In another embodiment of this invention, covalent modifications are made directly to human DNase II to give it a desired property (for example, increased half-life or decreased immunogenicity in vivo, or increased DNA

hydrolytic activity at physiological pH), and may be made instead of or in addition to the amino acid sequence substitution, insertion, and deletion mutations described above.

Covalent modifications are introduced by reacting targeted amino acid
5 residues of human DNase II with an organic derivatizing agent that is capable of reacting with selected amino acid side-chains or N- or C-terminal residues. Suitable derivatizing agents and methods are well known in the art. Covalent coupling of glycosides to amino acid residues of the protein may be used to modify or increase the number or profile of
10 carbohydrate substituents.

The covalent attachment of agents such as polyethylene glycol (PEG) or human serum albumin to human DNase II may reduce immunogenicity and/or toxicity of the human DNase II and/or prolong its half-life, as has been observed with other proteins. Abuchowski, et al., J. Biol. Chem. 252:3582-
15 3586 (1977); Poznansky, et al., FEBS Letters 239:18-22 (1988); Goodson, et al., Biotechnology 8:343-346 (1990); Katre, J. Immunol. 144:209-213 (1990); Harris, Polyethylene Glycol Chemistry (Plenum Press, 1992). As another example, the variant or modified form of human DNase II may comprise an amino acid sequence mutation or other covalent modification
20 that reduces the susceptibility of the variant to degradation by proteases (e.g., neutrophil elastase) that may be present in sputum and other biological materials, as compared to human DNase II.

Antibodies to human DNase II are produced by immunizing an animal with human DNase II or a fragment thereof, optionally in conjunction with
25 an immunogenic polypeptide, and thereafter recovering antibodies from the serum of the immunized animals. Alternatively, monoclonal antibodies are prepared from cells of the immunized animal in conventional fashion. The antibodies also can be made in the form of chimeric (e.g., humanized) or single chain antibodies or Fab fragments, using methods well known in the
30 art. Preferably, the antibodies will bind to human DNase II but will not substantially bind to (i.e., cross react with) other DNase proteins (such as human and bovine DNase I). The antibodies can be used in methods relating to the localization and activity of human DNase II, for example, for detecting human DNase II and measuring its levels in tissues or
35 clinical samples. Immobilized anti-human DNase II antibodies are particularly useful in the detection of human DNase II in clinical samples for diagnostic purposes, and in the purification of human DNase II.

Purified human DNase II is used to reduce the viscoelasticity of DNA-containing material, such as sputum, mucus, or other pulmonary secretions. human DNase II is particularly useful for the treatment of patients with pulmonary disease who have abnormal viscous or inspissated secretions and
5 conditions such as acute or chronic bronchial pulmonary disease, including infectious pneumonia, bronchitis or tracheobronchitis, bronchiectasis, cystic fibrosis, asthma, tuberculosis, and fungal infections. For such therapies, a solution or finely divided dry preparation of the human DNase II is instilled in conventional fashion into the airways (e.g., bronchi)
10 or lungs of a patient, for example by aerosolization.

Human DNase II also is useful for adjunctive treatment of abscesses or severe closed-space infections in conditions such as empyema, meningitis, abscess, peritonitis, sinusitis, otitis, periodontitis, pericarditis, pancreatitis, cholelithiasis, endocarditis and septic
15 arthritis, as well as in topical treatments in a variety of inflammatory and infected lesions such as infected lesions of the skin and/or mucosal membranes, surgical wounds, ulcerative lesions and burns. Human DNase II may improve the efficacy of antibiotics used in the treatment of such infections (e.g., gentamicin activity is markedly reduced by reversible
20 binding to intact DNA).

Human DNase II also is useful for preventing the new development and/or exacerbation of respiratory infections, such as may occur in patients having cystic fibrosis, chronic bronchitis, asthma, pneumonia, or other pulmonary disease, or patients whose breathing is assisted by
25 ventilator or other mechanical device, or other patients at risk of developing respiratory infections, for example post-surgical patients.

Human DNase II also is useful for the treatment for systemic lupus erythematosus (SLE), a life-threatening autoimmune disease characterized by the production of diverse autoantibodies. DNA is a major antigenic
30 component of the immune complexes. In this instance, the human DNase II may be given systemically, for example by intravenous, subcutaneous, intrathecal, or intramuscular administration to the affected patient.

Finally, human DNase II is useful for the treatment of other non-infected conditions in which there is an accumulation of cellular debris
35 that includes cellular DNA, such as pyelonephritis and tubulo-interstitial kidney disease.

Human DNase II can be formulated according to known methods to prepare therapeutically useful compositions. Typically, the human DNase

II is formulated with a physiologically acceptable excipient (or carrier) for therapeutic use. Such excipients are used, for example, to provide liquid formulations and sustained-release formulations of human DNase II. The human DNase II formulation may be used with commercially-available
5 nebulizers including jet nebulizers and ultrasonic nebulizers for administration of the DNase II directly into the airways or lungs of an affected patient. Another preferred therapeutic composition is a dry powder of human DNase II, preferably prepared by spray-drying of a solution of the human DNase II. In all cases, it is desirable that the therapeutic
10 compositions of DNase II be sterile. Preferably, the therapeutic compositions are disposed in a container fabricated of plastic or other non-glass material.

In a further embodiment, the therapeutic composition comprises cells actively producing human DNase II. Such cells may be directly introduced
15 into the tissue of a patient, or may be encapsulated within porous membranes which are then implanted in a patient (see e.g., Aebischer, et al., U.S. Patent No. 4,892,538, issued January 9, 1990; Aebischer, et al., U.S. Patent No. 5,283,187, issued February 1, 1994), in either case providing for the delivery of the human DNase II into areas within the body
20 of the patient in need of increased concentrations of DNA-hydrolytic activity. In one embodiment of the invention, the patient's cells are transformed, either in vivo or ex vivo, with DNA encoding human DNase II, and then used to produce the human DNase II directly within the patient. This latter method is commonly referred to as gene therapy. In another
25 embodiment, the patient's cells are transformed with other DNA (such as a promoter, enhancer, or amplifiable gene) that is capable of activating or increasing expression of an endogenous human DNase II gene.

In certain circumstances, it may be desirable to decrease the amount of human DNase II expressed in a cell. For that purpose, human DNase II
30 anti-sense oligonucleotides can be made and a method utilized for diminishing the level of human DNase II within the cell comprising introducing into the cell one or more human DNase II anti-sense oligonucleotides. The term "human DNase II anti-sense oligonucleotide" refers to an oligonucleotide that has a nucleotide sequence that is capable
35 of interacting through base pairing with a complementary nucleotide sequence that is involved in the expression of human DNase II within a cell, and thereby interfering with such expression.

The therapeutically effective amount of human DNase II will depend, for example, upon the amount of DNA in the material to be treated, the therapeutic objectives, the route of administration, and the condition of the patient. Accordingly, it will be necessary for the therapist to titer the dosage and modify the route of administration as required to obtain the optimal therapeutic effect. Generally, the therapeutically effective amount of human DNase II will be a dosage of from about 0.1 μ g to about 5 mg of the variant per kilogram of body weight of the patient, administered within pharmaceutical compositions, as described herein.

Human DNase II optionally is combined with or administered in concert with one or more other pharmacologic agents used to treat the conditions listed above, such as antibiotics, bronchodilators, anti-inflammatory agents, mucolytics (e.g. n-acetyl-cysteine), actin binding or actin severing proteins (e.g., gelsolin; Matsudaira et al., Cell 54:139-140 (1988); Stossel, et al., PCT Patent Publication No. WO 94/22465, published October 13, 1994; protease inhibitors; or gene therapy product (e.g., comprising the cystic fibrosis transmembrane conductance regulator (CFTR) gene); Riordan, et al., Science 245:1066-1073 (1989)).

This invention also provides methods for determining the presence of a nucleic acid molecule encoding human DNase II in test samples prepared from cells, tissues, or biological fluids, comprising contacting the test sample with isolated DNA comprising all or a portion of the nucleotide coding sequence for human DNase II and determining whether the isolated DNA hybridizes to a nucleic acid molecule in the test sample. DNA comprising all or a portion of the nucleotide coding sequence for human DNase II is also used in hybridization assays to identify and to isolate nucleic acids sharing substantial sequence identity to the coding sequence for human DNase II, such as nucleic acids that encode naturally-occurring allelic variants of human DNase II.

Also provided is a method for amplifying a nucleic acid molecule encoding human DNase II that is present in a test sample, comprising the use of an oligonucleotide having a portion of the nucleotide coding sequence for human DNase II as a primer in a polymerase chain reaction.

The following examples are offered by way of illustration only and are not intended to limit the invention in any manner.

EXAMPLE 1

Cloning Human DNase II cDNA

Full-length cDNA encoding human DNase II was identified by screening a human placental cDNA library (in p-gt11, Clontech, Palo Alto, California USA) with a mixture of the following oligonucleotide probes, each of which
 5 had been end-labeled with T4 polynucleotide kinase and p-³²P-adenosine triphosphate to a high specific radioactivity:

5'-GCCCAGAGAGGGCTGCAGTACAAGTATCTGGACGAGAGCTCCGGAGGC-3' (SEQ. ID. NO: 3)

5'-CCCAGCGCCCCGCGAGTCCCAGACACAGATTCTGGATCTCAGCCC-3' (SEQ. ID. NO: 4)

10 5'-GAYCARGARGGNGGNTTYTGGCTNAT-3' (SEQ. ID. NO: 5)

5'-GAYCARGARGGNGGNTTYTGGTTRAT-3' (SEQ. ID. NO: 6)

5'-AAYCGNGGNCAYACNAARGGNGT-3' (SEQ. ID. NO: 7)

5'-AAYAGRGGNCAYACNAARGGNGT-3' (SEQ. ID. NO: 8)

The first two of the oligonucleotide probes listed above (SEQ. ID. NOS: 3 and 4) comprise portions of the EST sequence having accession code T53394, in the Genbank database.

Hybridization of the probes to the cDNA library was carried out under low stringency conditions (in 20% vol/vol formamide, 5X SSPE, 5X Denhardt's solution, 0.1% sodium dodecyl sulfate (SDS), and 100µg/ml sonicated salmon
 20 sperm DNA), at 42°C, for 20 hours. Post hybridization washes were carried out in 2X SSC, 0.1% SDS, at 42°C. 1X SSPE is 150mM NaCl, 10mM sodium phosphate, 1mM EDTA, pH 7.4. 1X Denhardt's solution is 0.02% Ficoll*, 0.02% bovine serum albumin, and 0.02% polyvinyl-pyrrolidone. 1X SSC is 0.15 M NaCl, 0.015 M sodium citrate, pH 7.0.

25 Hybridization-positive phage clones were isolated and their DNAs sequenced following standard procedures. A 1575 base-pair insert was identified amongst the hybridization-positive phage clones, including an open reading frame of 1080 base-pairs that encodes a predicted protein that is 360 amino acid residues in length. The nucleotide sequence of
 30 the 1575 base-pair insert (SEQ. ID. NO: 1) and the amino acid sequence of predicted protein (SEQ. ID. NO: 2) are shown in Figure 1.

The predicted protein includes a signal sequence that is 16 amino acid residues in length. Cleavage of the signal sequence releases the mature protein (human DNase II) that is 344 amino acid residues in
 35 length, and that has a predicted molecular weight of 38,000 Daltons and a predicted pI of 9.0.

EXAMPLE 2

*-trademark

Expression of Human DNase II cDNA

The cDNA encoding human DNase II was subcloned into a mammalian expression vector pRK5 (Gorman, et al., DNA and Protein Engineering Techniques 2:1 (1990); European Patent Publication EP 307,247, published 5 March 15, 1989). The resulting recombinant vector is referred to as pRK5/human DNase II. Human embryonic kidney 293 cells (American Type Culture Collection, CRL 1573) were grown in serum-containing Dulbecco Modified Eagle's medium (DMEM) to 70% confluency and then transiently transfected with pRK5/human DNase II, or as a control, pRK5 alone. 24 10 hours post-transfection, the cells were washed with phosphate buffered saline and transferred to serum-free medium containing insulin. 72-96 hours later, conditioned medium was collected from the cell cultures and concentrated approximately 10-fold. Proteins expressed in the cell cultures were analyzed by SDS-polyacrylamide gel electrophoresis (SDS- 15 PAGE).

Cells transfected with pRK-5/human DNase II were found to produce a unique protein of about 42,000 - 44,000 Daltons, that was not produced in cells transfected with pRK5 alone.

The amino-terminal sequence of that secreted protein was determined 20 by preparing a poly-His tagged version of human DNase II. DNA encoding the poly-His tagged version of human DNase II was prepared by joining a nucleotide sequence encoding the amino acid sequence

Met-Arg-Gly-Ser-His-His-His-His-His (SEQ. ID. NO: 9) to the 3' end of the nucleotide sequence encoding human DNase II that is 25 shown in Figure 1. Human embryonic kidney 293 cells were transiently transfected with the DNA, and Ni-NTA-silica (Qiagen, Inc., Chatsworth, California USA) was used to purify the secreted poly-His tagged human DNase II. The amino-terminal amino acid sequence of the secreted protein was determined to be Leu-Thr-Cys-Tyr-Gly-Asp-Ser-Gly-Gln, in agreement with the 30 predicted amino acid sequence of the mature human DNase II protein shown in Figure 1.

EXAMPLE 3Biological Activity of Human DNase II

Concentrated cell culture supernatants, prepared as described above, 35 were tested for DNA-hydrolytic activity in a hyperchromicity assay (Kunitz, J. Gen. Physiol. 33:349-362 (1950); Kunitz, J. Gen. Physiol. 33:363-377 (1950)), in which the buffer used was 0.1M sodium acetate, pH 4.6, 1 mM

magnesium chloride. Such activity was detected in the supernatants from cells transfected with pRK5/human DNase II, but not in the supernatants from cells transfected with pRK5 alone. By also assaying cell lysates, it was determined that approximately 20% - 30% of the total human DNase II activity in the cells transfected with pRK5/human DNase was secreted.

EXAMPLE 4

Pattern of Expression of Human DNase II in Human Tissue

Northern blots of various human tissues were performed using a radiolabeled probe comprising a portion of the coding sequence of the cloned human DNase II cDNA. Expression of human DNase II messenger RNA (mRNA) was found in all tissues examined (brain, colon, heart, small intestine, kidney, liver, lung, peripheral blood lymphocytes, skeletal muscle, ovary, pancreas, placenta, prostate, spleen, testis, and thymus).

Sequence Listing

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Genentech, Inc.
Baker, Kevin P.
Baron, Will F.

(ii) TITLE OF INVENTION: Human DNase II

(iii) NUMBER OF SEQUENCES: 9

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Dennison Associates
(B) STREET: 133 Richmond Street West, Suite 301
(C) CITY: Toronto
(D) PROVINCE: Ontario
(E) COUNTRY: Canada
(F) POSTAL CODE: M5H 2L7

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: WinPatin (Genentech)

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 2,250,433
(B) FILING DATE: 23-APR-1997
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 08/639294
(B) FILING DATE: 25-APR-1996

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Dennison Associates
(B) REFERENCE/DOCKET NUMBER: JJ-10265CA

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 416-368-8313
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(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 1575 base pairs
(B) TYPE: Nucleic Acid
(C) STRANDEDNESS: Single
(D) TOPOLOGY: Linear

(ii) MOLECULE TYPE: DNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

WO 97/40134

PCT/US97/06664

GAATTCGGGC CAGTCCTGGC CTCTGATGTA ACCCAGCGCC CCGCAGTCCC 50
 GACACAGATT CCTGGATCTC AGCCCCATAG CAGCTATGAT CCCGCTGCTG 100
 CTGGCAGCGC TGCTGTGCGT CCCC GCCGGG GCCCTGACCT GCTACGGGGA 150
 CTCCGGGCAG CCTGTAGACT GGTTCGTGGT CTACAAGCTG CCAGCTCTTA 200
 5 GAGGGTCCGG GGAGGCGGCG CAGAGAGGGC TGCAGTACAA GTATCTGGAC 250
 GAGAGCTCCG GAGGCTGGCG GGACGGCAGG GCACTCATCA ACAGCCCGGA 300
 GGGGGCCGTG GGCCGAAGCC TGCAGCCGCT GTACCGGAGC AACACCAGCC 350
 AGCTCGCCTT CCTGCTCTAC AATGACCAAC CGCCTCAACC CAGCAAGGCT 400
 CAGGACTCTT CCATGCGTGG GCACACGAAG GGTGTCCTGC TCCTTGACCA 450
 10 CGATGGGGGC TTCTGGCTGG TCCACAGTGT ACCTAACTTC CCTCCACCGG 500
 CCTCCTCTGC TGCATACAGC TGGCCTCATA GCGCCTGTAC CTACGGGCAG 550
 ACCCTGCTCT GTGTGTCTTT TCCCTTCGCT CAGTTCTCGA AGATGGGCAA 600
 GCAGCTGACC TACACCTACC CCTGGGTCTA TAACTACCAG CTGGAAGGGA 650
 TCTTTGCCCA GGAATTCCCC GACTTGGAGA ATGTGGTCAA GGGCCACCAC 700
 15 GTTAGCCAAG AACCTGGAA CAGCAGCATC AACTCACAT CCCAGGCCGG 750
 GGCTGTTTTT CAGAGCTTTG CCAAGTTCAG CAAATTTGGA GATGACCTGT 800
 ACTCCGGCTG GTTGGCAGCA GCCCTTGGTA CCAACCTGCA GGTCCAGTTC 850
 TGGCACAAAA CTGTAGGCAT CCTGCCCTCT AACTGCTCGG ATATCTGGCA 900
 GGTTCCTGAAT GTGAACCAGA TAGCTTTCCC TGGACCAGCC GGCCCAAGCT 950
 20 TCAACAGCAC AGAGGACCAC TCCAAATGGT GCGTGTCCCC AAAAGGGCCC 1000
 TGGACCTGCG TGGGTGACAT GAATCGGAAC CAGGGAGAGG AGCAACGGGG 1050
 TGGGGGCACA CTGTGTGCCC AGCTGCCAGC CCTCTGGAAA GCCTTCCAGC 1100
 CGCTGGTGAA GAACTACCAG CCCTGTAATG GCATGGCCAG GAAGCCCAGC 1150
 AGAGCTTATA AGATCTAACC CTTATGGCCA GGTGCAGTGG CTCACGTATG 1200
 25 TAATCCCAGC ACTTTGGGAA GCCAAGGAGG GAGGATCACT TGAATCAGG 1250
 AATTCGAGAC CAGCCTGGGC TACATAGTGA GACCACATCT CTACTAGAAC 1300
 TTAAAAAAG TTAGCCAGGC ACGGTGATAA ATGCCTGTAG TCCCAGCCAC 1350
 TGAAGCCAGA GGATCGATTG AACCAGGGAG ATCATGGTCA CAGTGA ACTA 1400

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TGATTACGCC AACCTGGGTC ACATAGCAAG ACTCTGTTTC AAAAAAAAAAG 1450
 GGGGGGCGGG GGACGGGTGG GTGCAGTGGC TCACATCTGT AACCCCAGCA 1500
 CTTTGGGAGG CTGAGATGGG CAGATCACTT GAGGTCAGGA GTTCGAGACC 1550
 AGCCTGGCCA ACATGGTGAA ACCCC 1575

5 (2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 360 amino acids
 (B) TYPE: Amino Acid
 (D) TOPOLOGY: Linear

10 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met	Ile	Pro	Leu	Leu	Leu	Ala	Ala	Leu	Leu	Cys	Val	Pro	Ala	Gly	1	5	10	15
Ala	Leu	Thr	Cys	Tyr	Gly	Asp	Ser	Gly	Gln	Pro	Val	Asp	Trp	Phe	20	25	30	
Val	Val	Tyr	Lys	Leu	Pro	Ala	Leu	Arg	Gly	Ser	Gly	Glu	Ala	Ala	35	40	45	
Gln	Arg	Gly	Leu	Gln	Tyr	Lys	Tyr	Leu	Asp	Glu	Ser	Ser	Gly	Gly	50	55	60	
Trp	Arg	Asp	Gly	Arg	Ala	Leu	Ile	Asn	Ser	Pro	Glu	Gly	Ala	Val	65	70	75	
Gly	Arg	Ser	Leu	Gln	Pro	Leu	Tyr	Arg	Ser	Asn	Thr	Ser	Gln	Leu	80	85	90	
Ala	Phe	Leu	Leu	Tyr	Asn	Asp	Gln	Pro	Pro	Gln	Pro	Ser	Lys	Ala	95	100	105	
Gln	Asp	Ser	Ser	Met	Arg	Gly	His	Thr	Lys	Gly	Val	Leu	Leu	Leu	110	115	120	
Asp	His	Asp	Gly	Gly	Phe	Trp	Leu	Val	His	Ser	Val	Pro	Asn	Phe	125	130	135	
Pro	Pro	Pro	Ala	Ser	Ser	Ala	Ala	Tyr	Ser	Trp	Pro	His	Ser	Ala	140	145	150	
Cys	Thr	Tyr	Gly	Gln	Thr	Leu	Leu	Cys	Val	Ser	Phe	Pro	Phe	Ala	155	160	165	
Gln	Phe	Ser	Lys	Met	Gly	Lys	Gln	Leu	Thr	Tyr	Thr	Tyr	Pro	Trp	170	175	180	
Val	Tyr	Asn	Tyr	Gln	Leu	Glu	Gly	Ile	Phe	Ala	Gln	Glu	Phe	Pro	185	190	195	

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	Asp	Leu	Glu	Asn	Val	Val	Lys	Gly	His	His	Val	Ser	Gln	Glu	Pro	
					200					205					210	
	Trp	Asn	Ser	Ser	Ile	Thr	Leu	Thr	Ser	Gln	Ala	Gly	Ala	Val	Phe	
					215					220					225	
5	Gln	Ser	Phe	Ala	Lys	Phe	Ser	Lys	Phe	Gly	Asp	Asp	Leu	Tyr	Ser	
					230					235					240	
	Gly	Trp	Leu	Ala	Ala	Ala	Leu	Gly	Thr	Asn	Leu	Gln	Val	Gln	Phe	
					245					250					255	
10	Trp	His	Lys	Thr	Val	Gly	Ile	Leu	Pro	Ser	Asn	Cys	Ser	Asp	Ile	
					260					265					270	
	Trp	Gln	Val	Leu	Asn	Val	Asn	Gln	Ile	Ala	Phe	Pro	Gly	Pro	Ala	
					275					280					285	
	Gly	Pro	Ser	Phe	Asn	Ser	Thr	Glu	Asp	His	Ser	Lys	Trp	Cys	Val	
					290					295					300	
15	Ser	Pro	Lys	Gly	Pro	Trp	Thr	Cys	Val	Gly	Asp	Met	Asn	Arg	Asn	
					305					310					315	
	Gln	Gly	Glu	Glu	Gln	Arg	Gly	Gly	Gly	Thr	Leu	Cys	Ala	Gln	Leu	
					320					325					330	
20	Pro	Ala	Leu	Trp	Lys	Ala	Phe	Gln	Pro	Leu	Val	Lys	Asn	Tyr	Gln	
					335					340					345	
	Pro	Cys	Asn	Gly	Met	Ala	Arg	Lys	Pro	Ser	Arg	Ala	Tyr	Lys	Ile	
					350					355					360	

(2) INFORMATION FOR SEQ ID NO:3:

- (i) SEQUENCE CHARACTERISTICS:
- 25 (A) LENGTH: 48 base pairs
- (B) TYPE: Nucleic Acid
- (C) STRANDEDNESS: Single
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

30 GCCCAGAGAG GGCTGCAGTA CAAGTATCTG GACGAGAGCT CCGGAGGC 48

(2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
- 35 (A) LENGTH: 45 base pairs
- (B) TYPE: Nucleic Acid
- (C) STRANDEDNESS: Single
- (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

WO 97/40134

PCT/US97/06664

CCCAGCGCCC CGCAGTCCCA GACACAGATT CCTGGATCTC AGCCC 45

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- 5 (A) LENGTH: 26 base pairs
 (B) TYPE: Nucleic Acid
 (C) STRANDEDNESS: Single
 (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GAYCARGARG GNGGNTTYTG GCTNAT 26

10 (2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- 15 (A) LENGTH: 26 base pairs
 (B) TYPE: Nucleic Acid
 (C) STRANDEDNESS: Single
 (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

GAYCARGARG GNGGNTTYTG GTTRAT 26

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS:

- 20 (A) LENGTH: 23 base pairs
 (B) TYPE: Nucleic Acid
 (C) STRANDEDNESS: Single
 (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

25 AAYCGNGGNC AYACNAARGG NGT 23

(2) INFORMATION FOR SEQ ID NO:8:

(i) SEQUENCE CHARACTERISTICS:

- 30 (A) LENGTH: 23 base pairs
 (B) TYPE: Nucleic Acid
 (C) STRANDEDNESS: Single
 (D) TOPOLOGY: Linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

AAYAGRGGNC AYACNAARGG NGT 23

(2) INFORMATION FOR SEQ ID NO:9:

WO 97/40134

PCT/US97/06664

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 10 amino acids
(B) TYPE: Amino Acid
(D) TOPOLOGY: Linear

5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Met	Arg	Gly	Ser	His	His	His	His	His	His
1				5					10

What is claimed is:

1. An isolated nucleic acid molecule comprising a nucleotide sequence encoding a polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2.
2. An expression vector comprising a nucleotide sequence encoding a polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II operably linked to a promoter recognized by a host cell transformed with the vector.
3. An isolated nucleic acid molecule comprising a nucleotide sequence that encodes an amino acid sequence having at least 95% identity with the polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II, wherein the amino acid sequence encoded by said isolated nucleic acid molecule has DNA-hydrolytic activity.
4. An isolated nucleic acid molecule comprising a nucleotide sequence that encodes an amino acid sequence that differs from the polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II by the substitution of one amino acid for another at only a single position within the polypeptide sequence, wherein the amino acid sequence encoded by said isolated nucleic acid molecule has DNA-hydrolytic activity.
5. An isolated nucleic acid molecule comprising a nucleotide sequence that encodes an amino acid sequence that differs from the polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II by the substitution of one amino acid for another at only two positions within the polypeptide sequence, wherein the amino acid sequence encoded by said isolated nucleic acid molecule has DNA-hydrolytic activity.
6. A host cell transformed with an expression vector comprising a nucleotide sequence encoding a polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II.
7. A method of culturing a host cell transformed with an expression vector comprising a nucleotide sequence encoding a polypeptide having an amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II, which comprises culturing the host cell under conditions such that expression vector is replicated.

8. A process for producing a mature DNase II polypeptide which comprises transforming a host cell with a nucleic acid molecule that encodes a polypeptide comprising the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II and culturing the host cell under conditions such that the polypeptide is produced in the host cell.
9. An isolated polypeptide comprising the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II.
10. An isolated polypeptide comprising an amino acid sequence having at least 95% identity with the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II, which polypeptide has DNA-hydrolytic activity.
11. An isolated polypeptide comprising an amino acid sequence that differs from the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II by the substitution of one amino acid for another at only a single position wherein the isolated polypeptide has DNA-hydrolytic activity.
12. A polypeptide of claim 11 wherein the amino acid substitution creates a glycosylation site within the polypeptide that is not present in naturally-occurring human DNase II.
13. A pharmaceutical composition comprising a polypeptide comprising the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II and a physiologically acceptable excipient.
14. A composition of claim 13 that is sterile.
15. An antibody that is capable of specifically binding to the amino acid sequence of residues 17-360 of SEQ ID NO.:2 for mature human DNase II.
16. The use of the polypeptide according to claim 10 for the manufacture of a medicament for the treatment of a patient having a pulmonary disease or disorder.
17. The use of claim 16 wherein the disease or disorder is cystic fibrosis.
18. The use of the polypeptide according to claim 10 for the manufacture of a medicament for the treatment of a patient having systemic lupus erythematosus.

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1 GAATTCGGGCCAGTCCTGGCCTCTGATG TAACCCGAGCGCCCCGCAGTCCCGACACAGATT
 61 CCTGGATCTCAGCCCCATAGCAGCTATGATCCCGCTGCTGCTGGCAGCGCTGCTGTGCGT
 -16 M I P L L L A A L L C V
 121 CCCC GCCGGGGCCCTGACCTGCTACGGGGACTCCGGGCAGCCTGTAGACTGGTTCGTGGT
 -4 P A G A L T C Y G D S G Q P V D W F V V
 *
 181 CTACAAGCTGCCAGCTCTTAGAGGGTCCGGGGAGGCGGCGCAGAGAGGGCTGCAGTACAA
 17 Y K L P A L R G S G E A A Q R G L Q Y K
 241 GTATCTGGACGAGAGCTCCGGAGGCTGGCGGGACGGCAGGGCACTCATCAACAGCCCGGA
 37 Y L D E S S G G W R D G R A L I N S P E
 301 GGGGGCCGTGGGCCGAAGCCTGCAGCCGCTGTACCGGAGCAACACCAGCCAGCTCGCCTT
 57 G A V G R S L Q P L Y R S N T S Q L A F
 361 CCTGCTCTACAATGACCAACCGCCTCAACCCAGCAAGGCTCAGGACTCTTCCATGCGTGG
 77 L L Y N D Q P P Q P S K A Q D S S M R G
 421 GCACACGAAGGGTGTCTGCTCCTTGACCACGATGGGGGCTTCTGGCTGGTCCACAGTGT
 97 H T K G V L L L D H D G G F W L V H S V
 481 ACCTAACTTCCTCCACCGGCCTCCTCTGCTGCATACAGCTGGCCTCATAGCGCCTGTAC
 117 P N F P P P A S S A A Y S W P H S A C T
 *
 541 CTACGGGCAGACCCTGCTCTGTGTGTCTTTTCCCTTCGCTCAGTTCTCGAAGATGGGCAA
 137 Y G Q T L L C V S F P F A Q F S K M G K
 *
 601 GCAGCTGACCTACACCTACCCCTGGGTCTATAACTACCAGCTGGAAGGGATCTTTGCCCA
 157 Q L T Y T Y P W V Y N Y Q L E G I F A Q
 661 GGAATTCCCCGACTTGGAGAATGTGGTCAAGGGCCACCACGTTAGCCAAGAACCCTGGAA
 177 E F P D L E N V V K G H H V S Q E P W N
 721 CAGCAGCATCACACTCACATCCCAGGCCGGGGCTGTTTTCCAGAGCTTTGCCAAGTTCAG
 197 S S I T L T S Q A G A V F Q S F A K F S
 781 CAAATTTGGAGATGACCTGTACTCCGGCTGGTTGGCAGCAGCCCTTGGTACCAACCTGCA
 217 K F G D D L Y S G W L A A A L G T N L Q
 841 GGTCCAGTTCTGGCACAAACTGTAGGCATCCTGCCCTCTAACTGCTCGGATATCTGGCA
 237 V Q F W H K T V G I L P S N C S D I W Q
 *
 901 GGTCTGAATGTGAACCAGATAGCTTTCCCTGGACCAGCCGGCCCAAGCTTCAACAGCAC
 257 V L N V N Q I A F P G P A G P S F N S T

FIG. 1A
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961 AGAGGACCACTCCAAATGGTGC GTGTCCCCAAAAGGGCCCTGGACCTGCGTGGGTGACAT
277 E D H S K W C V S P K G P W T C V G D M
*

1021 GAATCGGAACCAGGGAGAGGAGCAACGGGGTGGGGGCACACTGTGTGCCCAGCTGCCAGC
297 N R N Q G E E Q R G G G T L C A Q L P A
*

1081 CCTCTGGAAAGCCTTCCAGCCGCTGGTGAAGAACTACCAGCCCTGTAATGGCATGGCCAG
317 L W K A F Q P L V K N Y Q P C N G M A R
*

1141 GAAGCCCAGCAGAGCTTATAAGATCTAACCTTATGGCCAGGTGCAGTGGCTCACGTATG
337 K P S R A Y K I

1201 TAATCCCAGCACTTTGGGAAGCCAAGGAGGGAGGATCACTTGAACCTCAGGAATTCGAGAC

1261 CAGCCTGGGCTACATAGTGAGACCACATCTCTACTAGAACTTAAAAAAGTTAGCCAGGC

1321 ACGGTGATAAATGCCTGTAGTCCCAGCCACTGAAGCCAGAGGATCGATTGAACCAGGGAG

1381 ATCATGGTCACAGTGAACCTATGATTACGCCAACCTGGGTACATAGCAAGACTCTGTTTC

1441 AAAAAAAGGGGGGGCGGGGACGGGTGGGTGCAGTGGCTCACATCTGTAACCCCAGCA

1501 CTTTGGGAGGCTGAGATGGGCAGATCACTTGAGGTCAGGAGTTCGAGACCAGCCTGGCCA

1561 ACATGGTGAAACCCC

FIG._1B

