

APPENDIX B
(U.S. Patent No. 5,547,933)
Part 1 of 4

United States Patent

Lin

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[54] PRODUCTION OF ERYTHROPOIETIN

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0136490	4/1985	European Pat. Off.
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380, 397, 834, 835, 23.5, 23.51, 388.7;
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[56]

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ABSTRACT

Disclosed are novel polypeptides possessing part or all of the primary structural conformation and one or more of the biological properties of mammalian erythropoietin ("EPO") which are characterized in preferred forms by being the product of prokaryotic or eucaryotic host expression of an exogenous DNA sequence. Illustratively, genomic DNA, cDNA and manufactured DNA sequences coding for part or all of the sequence of amino acid residues of EPO or for analogs thereof are incorporated into autonomously replicating plasmid or viral vectors employed to transform or transfect suitable prokaryotic or eucaryotic host cells such as bacteria, yeast or vertebrate cells in culture. Upon isolation from culture media or cellular lysates or fragments, products of expression of the DNA sequences display, e.g., the immunological properties and in vitro and in vivo biological activities of EPO of human or monkey species origins. Disclosed also are chemically synthesized polypeptides sharing the biochemical and immunological properties of EPO. Also disclosed are improved methods for the detection of specific single stranded polynucleotides in a heterologous cellular or viral sample prepared from, e.g., DNA present in a plasmid or viral-borne cDNA or genomic DNA "library".

14 Claims, 27 Drawing Sheets

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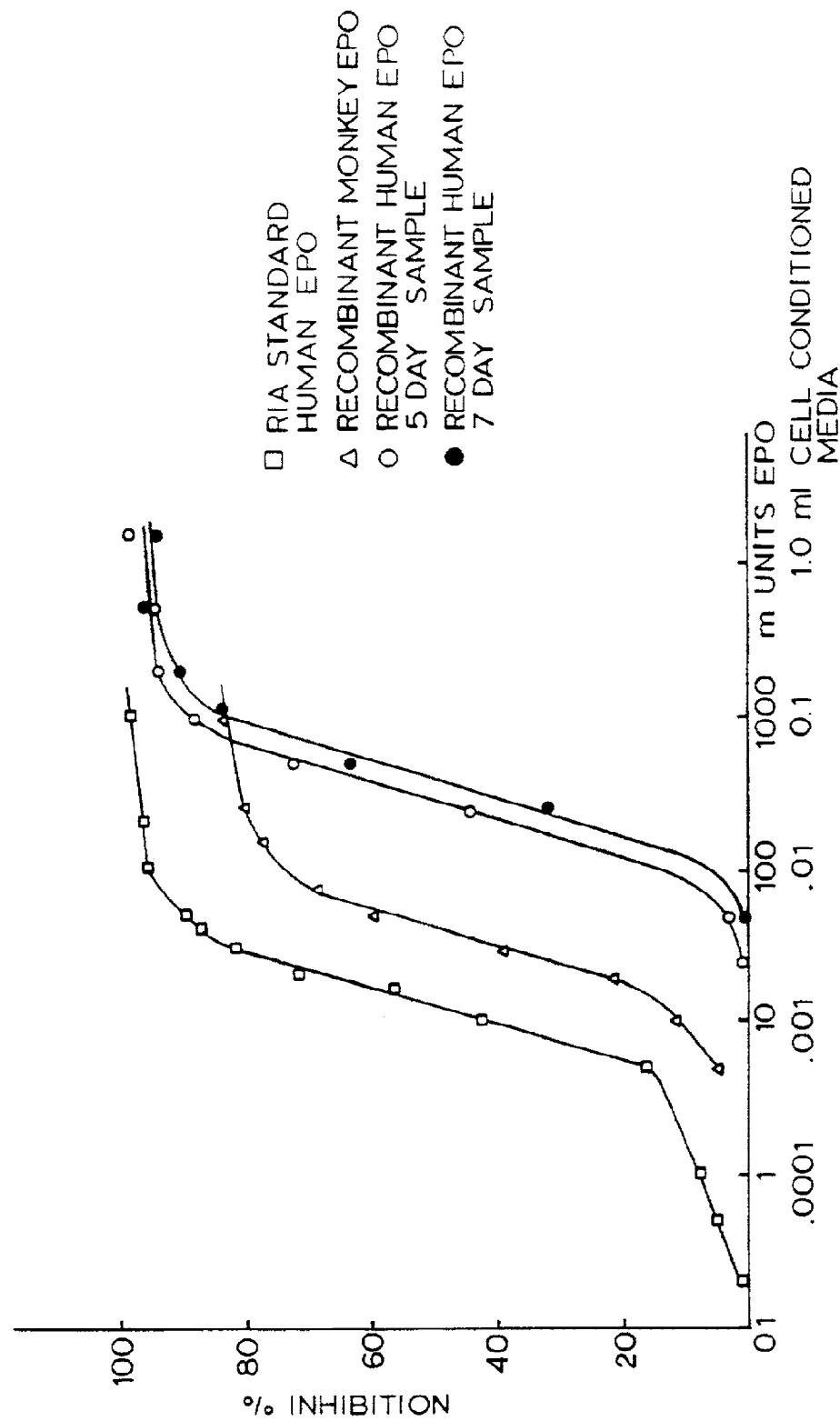
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FIG. 1

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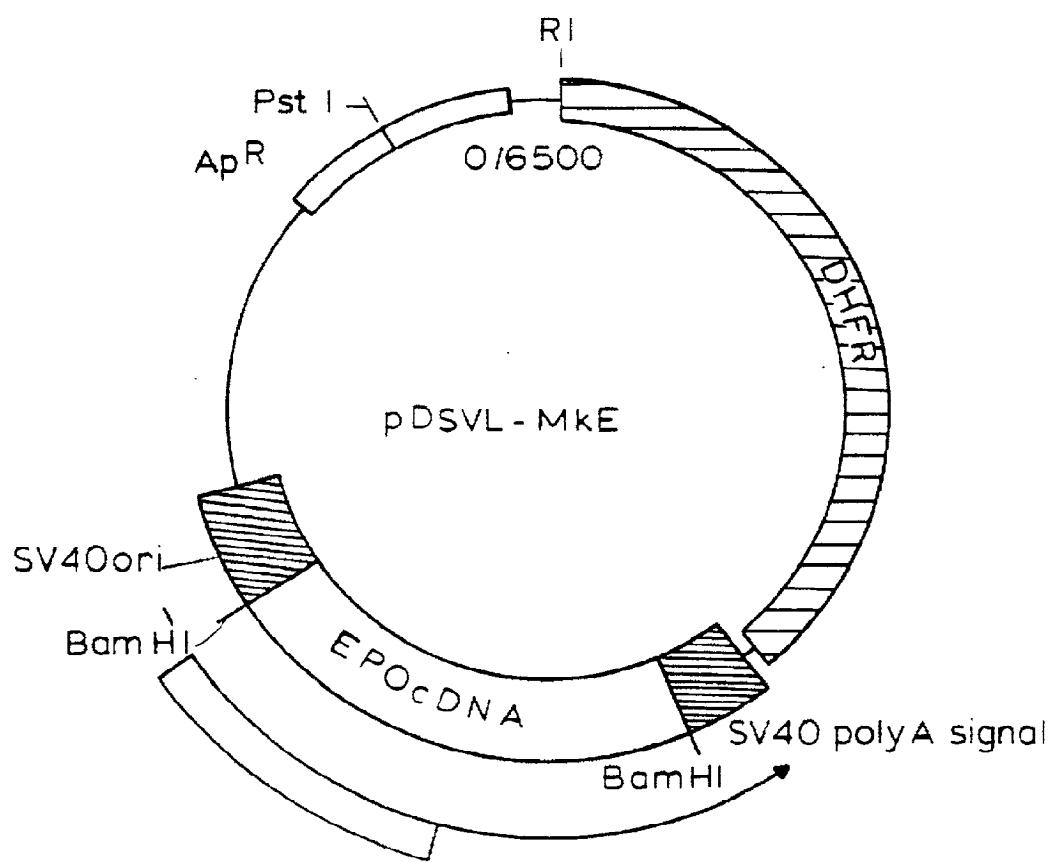
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FIG. 2



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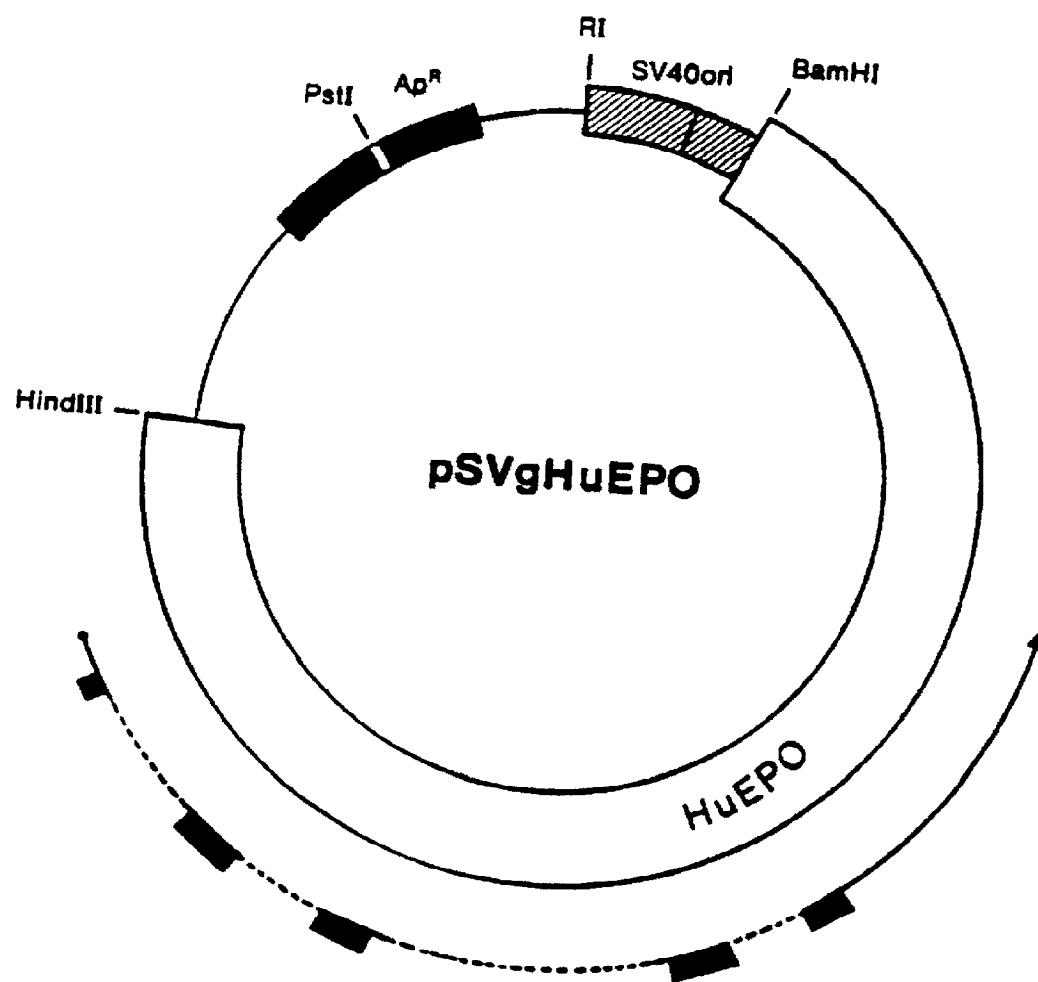
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FIG. 3



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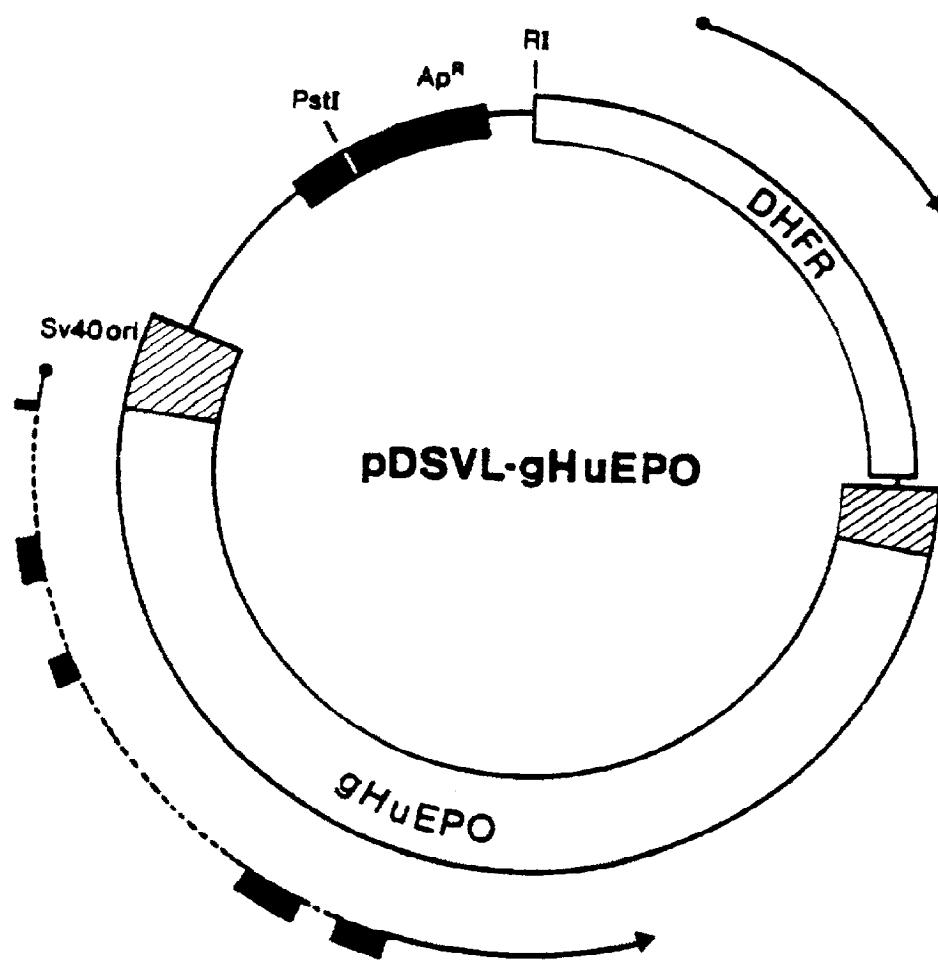
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FIG. 4



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FIG. 5A

Sequence 3A
GAATCCCGCCCCCTGAGCAACCCCTCTCCGACCCCTGGCTGCC
CGCTTAACTTCCGGATTCAGCTCCGGTGTGGCTAACCGCCCTAGGTGGCTGAG

-27	-20													
Met	Gly	Val	His	Glu	Cys	Pro	Ala	Trp						
GGACCCCGCCGCCCCGAGATG	GGG	ATG	CAC	GAA	TGT	CCT	GCC	TGG						
-10														
Leu	Trp	Leu	Leu	Ser	Leu	Val	Ser	Leu	Pro	Leu	Gly	Leu	Pro	
CTG	TGG	CTT	CTC	CTG	TCT	CTC	CTG	CTG	CTG	CTG	CTG	CTG	CTC	CCA
-1									-1	+1				
Val	Pro	Gly	Ala	Pro	Pro	Arg	Leu	Ile	Cys	Asp	Ser	Arg	Val	Leu
GTC	CCC	GGC	CCA	CCC	CCA	CCA	CTC	ATC	TCT	GAC	AGC	CGA	CTG	CTG
20									*					
Glu	Arg	Tyr	Leu	Glu	Ala	Lys	Glu	Ala	Glu	Asn	Val	Thr	Met	
GAG	AGG	TAC	CTC	TTC	GAG	GCC	AAG	GAG	GCC	GAG	AAT	GTC	ACG	ATG
30									*					
Gly	Cys	Ser	Glu	Ser	Cys	Ser	Leu	Asn	Glu	Asn	Ile	Thr	Val	Pro
GGC	TGT	TCC	GAA	AGC	TGC	AGC	TTC	ATG	GAG	ATG	ATC	ACC	GTC	CCA

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FIG. 5B

Asp	Thr	Lys	Val	Asn	Phe	Tyr	Ala	Trp	Lys	Arg	Met	Leu	Val	Gly
GAC	ACC	AAA	GTT	AAC	TTC	TAT	GCC	TGC	AGG	ATC	GGG	ATG	GTG	GGG
50														
Gln	Gln	Ala	Glu	Val	Trp	Gln	Gly	Leu	Ala	Leu	Ser	Glu		
CAG	CAG	CTG	GCT	CTA	AAA	CTC	TGG	CAC	GGC	CTG	GCC	CTG	TCA	GAA
60														
Ala	Val	Leu	Arg	Gly	Gln	Ala	Val	Leu	Ala	Asn	Ser	Ser	Gln	Pro
GCT	GTC	CTG	CGG	GGC	CAG	CCC	GTC	TTG	GCC	AMC	TCT	TCC	CAG	CCT
70														
Phe	Glu	Pro	Leu	Gln	Leu	His	Met	Asp	Lys	Ala	Ile	Ser	Gly	Leu
TTC	GAG	CCC	CTG	CAG	CTG	CAC	ATG	GAT	AAA	GCC	ATC	ATG	GGC	CTT
80														
Arg	Ser	Ile	Thr	Thr	Leu	Leu	Arg	Ala	Leu	Gly	Ala	Gln	Glu	Ala
CGC	AGC	ATC	ACC	ACT	CTG	CTT	CGG	GGC	CTG	GGA	GCC	CAG	GAA	GCC
90														
Ile	Ser	Leu	Pro	Asp	Ala	Ala	Ser	Ala	Ala	Pro	Leu	Arg	Thr	Ile
ATC	TCC	CTC	CCA	GAT	GCG	GGC	CCC	TCG	GCT	GCT	CCA	CTC	CGA	ACC
100														
Thr	Ala	Asp	Thr	Phe	Cys	Lys	Leu	Phe	Arg	Val	Tyr	Ser	Asn	Phe
ACT	GCT	GAC	ACT	TTC	TGC	AAA	CTC	TRC	CAA	CTC	TAC	TCC	AAU	TTC
110														
Ile	Ser	Ile	Thr	Leu	Leu	Arg	Ala	Leu	Gly	Ala	Gln	Glu	Ala	
ATC	TCC	CTC	CCA	GAT	GCG	GGC	CCC	TCG	GCT	GCT	CCA	CTC	CGA	ACC
120														
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Thr	Ala	Asp	Thr	Phe	Cys	Lys	Leu	Phe	Arg	Val	Tyr	Ser	Asn	Phe
ACT	GCT	GAC	ACT	TTC	TGC	AAA	CTC	TRC	CAA	CTC	TAC	TCC	AAU	TTC
140														

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FIG. 5C

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FIG. 6A

AGCTTCTGGCTTCGACCCAGCTTACCTTACGGAAACTCAGGCAACCCAGCTTCTGGCTTCGACCC
AGCCCGGGATGCCGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCG
ACGCTTGCG
GTGGCCCTTAACCCCTTGGCGAACCCCTTCAACCGACCGACCGACCGACCGACCGACCGACCGAC
CAATTAACCCCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCG
CGCCCGGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCG
CTTGGAGACCCGGCT
GAGCTTACGGCTGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCGGCG
-27 -24
Met Gly Val His
ATG AGG GTC CAC G

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