

# **EXHIBIT H-1**

## **Part 2 of 2**

and 567 mU/ml for the seven day growth sample. Estimated monkey EPO production levels in the Example 7B expression system were on the same order or better.

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EXAMPLE 9

P Culture fluids prepared according to Examples 6 and 7 were subjected to an in vitro assay for EPO activity according to the procedure of Goldwasser, et al., Endocrinology, 97, 2, pp. 315-323 (1975). Estimated monkey EPO values for culture fluids tested ranged from 3.2 to 4.3 U/ml. Human EPO culture fluids were also active in this in vitro assay and, further, this activity could be neutralized by anti-EPO antibody. The recombinant monkey EPO culture fluids according to Example 6 were also subjected to an assay for in vivo biological activity according to the general procedures of Cotes, et al., Nature, 191, pp. 1065-1067 (1961) and Hammond, et al., Ann.N.Y.Acad.Sci., 149, pp. 516-527 (1968) and activity levels ranged from 0.94 to 1.24 U/ml.

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EXAMPLE 10

P In the previous examples, recombinant monkey or human EPO material was produced from vectors used to transfect COS-1 cells. These vectors replicate in COS-1 cells due to the presence of SV40 T antigen within the cell and an SV40 origin of replication on the vectors. Though these vectors produce useful quantities of EPO in COS-1 cells, expression is only transient (7 to 14 days) due to the eventual loss of the vector. Additionally, only a small percentage of COS-1 became productively transfected with the vectors. The present example describes expression systems employing Chinese hamster ovary (CHO) DHFR<sup>-</sup> cells and the selectable marker, DHFR. [For discussion of related expression systems, see

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U.S. Letters Patent No. 4,399,216 and European Patent Applications 117058, 117059 and 117060, all published August 29, 1984.]

31 CHO DHFR<sup>-</sup> cells (DuX-B11) CHO K1 cells, Urlaub,  
5 et al., Proc. Nat. Acad. Sci. (U.S.A.), Vol. 77, 4461  
(1980) lack the enzyme dihydrofolate reductase (DHFR) due  
to mutations in the structural genes and therefore  
require the presence of glycine, hypoxanthine, and thymi-  
dine in the culture media. Plasmids pDSVL-MkE (Example  
10 6) or pDSVL-gHuEPO (Example 7B) were transfected along  
31 with carrier DNA into CHO DHFR<sup>-</sup> cells growing in media  
containing hypoxanthine, thymidine, and glycine in 60 mm  
culture plates. Plasmid pSVgHuEPO (Example 7A) was mixed  
15 with the plasmid pMG2 containing a mouse dihydrofolate  
reductase gene cloned into the bacterial plasmid vector  
pBR322 (per Gasser, et al., supra.) The plasmid mixture  
31 and carrier DNA was transfected into CHO DHFR<sup>-</sup> cells.  
(Cells which acquire one plasmid will generally also  
acquire a second plasmid). After three days, the cells  
20 were dispersed by trypsinization into several 100 mm  
culture plates in media lacking hypoxanthine and thymi-  
dine. Only those cells which have been stably trans-  
formed with the DHFR gene, and thereby the EPO gene,  
14 survive in this media. After 7-21 days, colonies of sur-  
25 viving cells became apparent. These transformant colo-  
nies, after dispersion by trypsinization can be  
continuously propagated in media lacking hypoxanthine and  
thymidine, creating new cell strains (e.g., CHO  
pDSVL-MkEPO, CHO pSVgHuEPO, CHO-pDSVL-gHuEPO).  
30 Culture fluids from the above cell strains were  
tested in the RIA for the presence of recombinant monkey  
or human EPO. Media for strain CHO pDSVL-MkEPO contained  
EPO with immunological properties like that obtained from  
COS-1 cells transfected with plasmid pDSVL-MkEPO. A  
35 representative 65 hour culture fluid contained monkey EPO  
at 0.60 U/ml.

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Culture fluids from CHO pSVgHuEPO and CHO pDSVL-gHuEPO contained recombinant human EPO with immunological properties like that obtained with COS-1 cells transfected with plasmid pSVgHuEPO or pDSVL-gHuEPO. A representative 3 day culture fluid from CHO pSVgHuEPO contained 2.99 U/ml of human EPO and a 5.5 day sample from CHO pDSVL-gHuEPO had 18.2 U/ml of human EPO as measured by the RIA.

The quantity of EPO produced by the cell strains described above can be increased by gene amplification giving new cell strains of greater productivity. The enzyme dihydrofolate reductase (DHFR) which is the product coded for by the DHFR gene can be inhibited by the drug methotrexate (MTX). More specifically, cells propagated in media lacking hypoxanthine and thymidine are inhibited or killed by MTX. Under the appropriate conditions, (e.g., minimal concentrations of MTX) cells resistant to and able to grow in MTX can be obtained. These cells are found to be resistant to MTX due to an amplification of the number of their DHFR genes, resulting in increased production of DHFR enzyme. The surviving cells can, in turn, be treated with increasing concentrations of MTX, resulting in cell strains containing greater numbers of DHFR genes. "Passenger genes" (e.g., EPO) carried on the expression vector along with the DHFR gene or transformed with the DHFR gene are frequently found also to be increased in their gene copy number.

As examples of practice of this amplification system, cell strain CHO pDSVL-MkE was subjected to increasing MTX concentrations (0 nM, 30 nM and 100 nM). Representative 65-hour culture media samples from each amplification step were assayed by RIA and determined to contain 0.60, 2.45 and 6.10 U/ml, respectively. Cell strain CHO pDSVL-gHuEPO was subjected to a series of increasing MTX concentrations of 30 nM, 50 nM, 100 nM,

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32 200 nM, 1  $\mu$ M, and 5  $\mu$ M MTX. A representative 3-day  
 culture media sample from the 100 nM MTX step contained  
 35 human EPO at  $3089 \pm 129$  u/ml as judged by RIA.  
 Representative 48 hour cultural medium samples from the  
 825 100 nM and 1  $\mu$ M MTX steps contained, respectively, human  
 EPO at 466 and 1352 U/ml as judged by RIA (average of  
 33 triplicate assays). In these procedures,  $1 \times 10^6$  cells  
 were plated in 5 ml of media in 60 mm culture dishes.  
 Twenty-four hours later the media were removed and  
 10 replaced with 5 ml of serum-free media (high glucose DMEM  
 supplemented with 0.1 mM non-essential amino acids and  
 L-glutamine). EPO was allowed to accumulate for 48 hours  
 in the serum-free media. The media was collected for RIA  
 assay and the cells were trypsinized and counted. The  
 15 average RIA values of 467 U/ml and 1352 U/ml for cells  
 92 grown at 100 nM and 1  $\mu$ M MTX, respectively, provided  
 actual yields of 2335 U/plate and 6750 U/plate. The  
 33 average cell numbers per plate were  $1.94 \times 10^6$  and  
 33  $3.12 \times 10^6$  cells, respectively. The effective production  
 20 rates for these culture conditions were thus 1264 and  
 2167 U/ $10^6$  cells/48 hours.

The cells in the cultures described immediately  
 above are a genetically heterogeneous population.  
 Standard screening procedures are being employed in an  
 25 attempt to isolate genetically homogeneous clones with  
 the highest production capacity. See, Section A, Part 2,  
 of "Points to Consider in the Characterization of Cell  
 Lines Used to Produce Biologics", June 1, 1984, Office of  
 Biologics Research Review, Center for Drugs and  
 30 Biologics, U.S. Food and Drug Administration.

The productivity of the EPO producing CHO cell  
 lines described above can be improved by appropriate cell  
 culture techniques. The propagation of mammalian cells  
 in culture generally requires the presence of serum in  
 35 the growth media. A method for production of erythro-  
 poietin from CHO cells in media that does not contain

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serum greatly facilitates the purification of erythropoietin from the culture medium. The method described below is capable of economically producing erythropoietin in serum-free media in large quantities sufficient for  
5 production.

Strain CHO pDSVL-gHuEPO cells, grown in standard cell culture conditions, are used to seed spinner cell culture flasks. The cells are propagated as a suspension cell line in the spinner cell culture flask in media consisting of a 50-50 mixture of high glucose DMEM and Ham's F12 supplemented with 5% fetal calf serum, L-glutamine, Penicillin and Streptomycin, 0.05 mM non-essential amino acids and the appropriate concentration of methotrexate. Suspension cell culture allows the EPO-producing  
15 CHO cells to be expanded easily to large volumes. CHO cells, grown in suspension, are used to seed roller bottles at an initial seeding density of  $1.5 \times 10^7$  viable  
33 cells per 850 cm<sup>2</sup> roller bottle in 200 ml of media. The cells are allowed to grow to confluency as an adherent  
20 cell line over a three-day period. The media used for this phase of the growth is the same as used for growth in suspension. At the end of the three-day growth period, the serum containing media is removed and  
13 replaced with 100 ml of serum-free media; 50-50 mixture  
3 25 of high glucose DMEM and Ham's F12 supplemented with 0.05 mM non-essential amino acids and L-glutamine. The roller bottles are returned to the roller bottle incubator for a period of 1-3 hours and the media again is removed and replaced with 100 ml of fresh serum-free  
14 30 media. The 1-3 hour incubation of the serum-free media reduces the concentration of contaminating serum proteins. The roller bottles are returned to the incubator for seven days during which erythropoietin accumulates in the serum-free culture media. At the end of the seven  
35 day production phase, the conditioned media is removed and replaced with fresh serum-free medium for a second

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production cycle. As an example of the practice of this production system, a representative seven-day, serum-free  
 35 media sample contained human erythropoietin at  $3892 \pm 409$   
 U/ml as judged by the RIA. Based on an estimated cell  
 33 5 density of  $0.9$  to  $1.8 \times 10^5$  cells/cm<sup>2</sup>, each 850  
 33 cm<sup>2</sup> roller bottle contained from  $0.75$  to  $1.5 \times 10^8$  cells  
 and thus the rate of production of EPO in the 7-day, 100  
 ml culture was 750 to 1470 U/10<sup>6</sup> cells/48 hours.

Culture fluids from cell strain CHO pDSVL-MKEPO  
 10 carried in 10 nM MTX were subjected to RIA in vitro and  
in vivo EPO activity assays. The conditioned media  
 35 sample contained  $41.2 \pm 1.4$  U/ml of MKEPO as measured by  
 35 the RIA,  $41.2 \pm 0.064$  U/ml as measured by the in vitro  
 35 biological activity assay and  $42.5 \pm 5$  U/ml as measured  
 15 by the in vivo biological activity assay. Amino acid  
 sequencing of polypeptide products revealed the presence  
 of EPO products, a principle species having 3 residues of  
 the "leader" sequence adjacent the putative amino ter-  
 20 minant alanine. Whether this is the result of incorrect  
 membranc processing of the polypeptide in CHO cells or  
 reflects a difference in structure of the amino terminus  
 of monkey EPO vis-a-vis human EPO, is presently unknown.

Culture fluids from cell strain CHO pDSVL-gHuEPO  
 were subjected to the three assays. A 5.5 day sample  
 25 contained recombinant human EPO in the media at a level  
 35 of  $18.2$  U/ml by RIA assay,  $15.8 \pm 4.6$  U/ml by in vitro  
 35 assay and  $16.8 \pm 3.0$  U/ml by in vivo assay.

Culture fluid from CHO pDSVL-gHuEPO cells pre-  
 pared amplified by stepwise 100 nM MTX were subjected to  
 30 the three assays. A 3.0 day sample contained recombinant  
 35 human EPO at a level of  $3089 \pm 129$  U/ml by RIA,  $2589 \pm$   
 35  $71.5$  U/ml by in vitro assay, and  $2040 \pm 160$  U/ml by in  
vivo assay. Amino acid sequencing of this product  
 reveals an amino terminal corresponding to that  
 35 designated in <sup>Figure 6</sup> ~~Table VI~~.

Cell conditioned media from CHO cells trans-  
 fected with plasmid pDSVL-MKE in 10 nM MTX were pooled,

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and the MTX dialyzed out over several days, resulting in  
35 media with an EPO activity of  $221 \pm 5.1$  U/ml (EPO-CCM).  
To determine the in vivo effect of the EPO-CCM upon hema-  
tocrit levels in normal Balb/C mice, the following  
5 experiment was conducted. Cell conditioned media from  
untransfected CHO cells (CCM) and EPO-CCM were adjusted  
with PBS. CCM was used for the control group (3 mice)  
13 and two dose levels of EPO-CCM -- 4 units per injection  
13 and 44 units per injection -- were employed for the  
10 experimental groups (2 mice/group). Over the course of 5  
weeks, the seven mice were injected intraperitoneally, 3  
times per week. After the eighth injection, average  
hematocrit values for the control group were determined  
to be 50.4%; for the 4U group, 55.1%; and, for the 44U  
15 group, 67.9%.

Mammalian cell expression products may be  
readily recovered in substantially purified form from  
culture media using HPLC (C<sub>4</sub>) employing an ethanol gra-  
dient, preferably at pH7.

20 A preliminary attempt was made to characterize  
recombinant glycoprotein products from conditioned medium of  
COS-1 and CHO cell expression of the human EPO gene in  
comparison to human urinary EPO isolates using both  
Western blot analysis and SDS-PAGE. These studies indi-  
25 cated that the CHO-produced EPO material had a somewhat  
higher molecular weight than the COS-1 expression product  
which, in turn, was slightly larger than the pooled  
source human urinary extract. All products were somewhat  
heterogeneous. Neuraminidase enzyme treatment to remove  
30 sialic acid resulted in COS-1 and CHO recombinant pro-  
ducts of approximately equal molecular weight which were  
both nonetheless larger than the resulting asialo human  
urinary extract. Endoglycosidase F enzyme (EC 3.2.1)  
treatment of the recombinant CHO product and the urinary  
35 extract product (to totally remove carbohydrate from

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both) resulted in substantially homogeneous products having essentially identical molecular weight characteristics.

Purified human urinary EPO and a recombinant,  
 5 CHO cell-produced, EPO according to the invention were subjected to carbohydrate analysis according to the procedure of Ledeen, et al. Methods in Enzymology,  
 14 83(Part D), 139-191 (1982) as modified through use of the hydrolysis procedures of Nesser, et al., Anal. Biochem.,  
 1410 142, 58-67 (1984). Experimentally determined carbohydrate constitution values (expressed as molar ratios of carbohydrate in the product) for the urinary isolate were as follows: Hexoses, 1.73; N-acetylglucosamine, 1; N-acetylneuraminic acid, 0.93; Fucose, 0; and N-acetyl-  
 15 galactosamine, 0. Corresponding values for the recombinant product (derived from CHO pDSVL-gHuEPO 3-day culture media at 100 nM MTX) were as follows: Hexoses, 15.09; N-acetylglucosamine, 1; N-acetylneuraminic acid, 0.998; Fucose, 0; and N-acetylgalactosamine, 0. These  
 20 findings are consistent with the Western blot and SDS-PAGE analysis described above.

Glycoprotein products provided by the present invention are thus comprehensive of products having a primary structural conformation sufficiently duplicative  
 25 of that of a naturally-occurring erythropoietin to allow possession of one or more of the biological properties thereof and having an average carbohydrate composition which differs from that of naturally-occurring erythropoietin.

30 CL EXAMPLE 11

P The present example relates to the total manufacture by assembly of nucleotide bases of two structural genes encoding the human species EPO sequence of <sup>Figure 6</sup> ~~Table VI~~  
 35 and incorporating, respectively "preferred" codons for expression in E.coli and yeast (S.cerevisiae) cells.

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Also described is the construction of genes encoding analogs of human EPO. Briefly stated, the protocol employed was generally as set out in the previously noted disclosure of Alton, et al. (WO 83/04053). The genes were  
5 designed for initial assembly of component oligonucleotides into multiple duplexes which, in turn, were assembled into three discrete sections. These sections were designed for ready amplification and, upon removal from the amplification system, could be assembled sequentially  
10 or through a multiple fragment ligation in a suitable expression vector.

*10*  
*Figures 10 through 15 and 7*  
~~Tables VIII through XIV below~~ illustrate the design and assembly of a manufactured gene encoding a human EPO translation product lacking any leader or pre-  
15 sequence but including an initial methionine residue at position -1. Moreover, the gene incorporated in  
21 substantial part E.coli preference codons and the construction was therefore referred to as the "ECEPO"  
gene.

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TABLE VIII

ECEPO SECTION 1 OLIGONUCLEOTIDES

1. AATTCTAGAAACCATGAGGGTAATAAAAATA
2. CCATTATTTTATTACCCTCATGGTTCTAG
- 5 3. ATGGCTCCGCCGCGTCTGATCTCCGAC
4. CTCGAGTCGCAGATCAGACGCCGCCGAG
5. TCGAGAGTTCTGGAACGTTACCTGCTG
6. CTTCCAGCAGGTAACGTTCCAGAACT
7. GAAGCTAAAGAAGCTGAAAACATC
- 10 8. GTGGTGATGTTTTAGCTTCTTTAG
9. ACCACTGGTTGTGCTGAACACTGTTC
10. CAAAGAACAGTGTTCCAGCACAAACCA
11. TTTGAACGAAAACATTACGGTACCG
12. GATCCGGTACCGTAATGTTTTTCGTT

TABLE IX

ECEPO SECTION 1

<sup>XbaI</sup>  
<sup>EcoRI</sup>  
 AATTCTAG AAACCATGAG<sup>1</sup> GGTAATAAAA TAATGGCTCC<sup>3</sup> GCCGCGTCTG  
 GATC TTTGGTACTC<sup>2</sup> CCATTATTTT ATTACCGAGG<sup>4</sup> CGGCGCAGAC  
 20  
 ATCTGCGACT<sup>5</sup> CGAGAGTTCT GGAACGTTAC CTGCTGGAAG<sup>6</sup> CTAAAGAAGC  
 TAGACGCTGA GCTCTCAAGA CCTTGCAATG GACGACCTTG<sup>7</sup> GATTTCTTCG  
 25  
 TGAAAACATC<sup>8</sup> ACCACTGGTT<sup>9</sup> GTGCTGAACA CTGTTCTTTG<sup>10</sup> AACGAAAACA<sup>11</sup>  
 ACTTTTGTAG TGGTGACCAA CACGACTTGT GACAAGAAAC<sup>12</sup> TTGCTTTTGT  
<sup>KpnI</sup> <sup>BamHI</sup>  
 TTACGGTACC<sup>12</sup> CCTAG

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TABLE X  
ECEPO SECTION 2 OLIGONUCLEOTIDES

- 1. AATTGGGTACCAGACACCAAGT
- 2. GTTAACCTTGGTGTCTGGTACCG
- 5 3. TAACTTCTACGCTTGGAAACGTAT
- 4. TTCCATACGTTTCCAAGCGTAGAA
- 5. GGAAGTTGGTCAACAAGCAGTTGAAGT
- 6. CCAAACCTCAACTGCTTGTGACCAAC
- 7. TTGGCAGGGTCTGGCACTGCTGAGCG
- 10 8. GCCTCGCTCAACAGTGCCAGACCCTG
- 9. AGGCTGTACTGCCTGGCCAGGCA
- 10. GCAGTGCCCTGGCCACGCAGTACA
- 11. CTGCTGGTAAACTCCTCTCAGCCGT
- 12. TTCCCACGGCTGAGAGGAGTTACCA
- 15 13. GGGAACCGCTGCAGCTGCATGTTGAC
- 14. GCTTTGTCAACATGCAGCTGCAGCGG
- 15. AAAGCAGTATCTGGCCTGAGATCTG
- 16. GATCCAGATCTCAGGCCAGATACT

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TABLE XI

ECEPO SECTION 2

~~ECORI Kpnl 1  
 A ATTCGGTACC AGACCCCAAG GTAACTTCT ACGTTGGAA ACGTATGGAA  
 GCCATGG TCTGGGTTC CATTGAAGA TCGGACCTT TGCATACCTT 4  
2  
5 GTTGGTCAAC AAGCAGTTGA AGTIGGCAG 7 GGTCTGGCAC TGCTGAGCCA  
 CACCAGTTG TTCGTCACT 6 TAACCTC CCAGACCGTG ACGACTCGCT 8  
9 GGCTGTACTG CGTGGCCAGG CACTGGTGGT AACTCTCTCT CAGCCGTGG  
 CCGACATGAC GCACCGGTCC GTGACCACCA TTTGAGGAA 10 GTCGGCACCC 12  
13 AACCGTGCA GCTGCATGTT GACAAAGCAG 15 BglI BamHI  
 TTGGCGACGT CGACGTACAA CTGTTTCTC ATAGACCGGA CTCTAGACCTAC 16~~

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TABLE XIII

ECEPO SECTION 3

- 1. GATCCAGATCTCTGACTACTCTGC
- 5 2. ACGCAGCAGAGTAGTCAGAGATCTG
- 3. TGCGTGCTCTGGGTGCACAGAAAGAGG
- 4. GATAGCCTCTTTCTGTGCACCCAGAGC
- 5. CTATCTCTCCGCCGGATGCTGCATCT
- 6. CAGCAGATGCAGCATCCGGCGGAGA
- 10 7. GCTGCACCGCTGCTTACCATCACTG
- 8. ATCAGCAGTGATGCTAGCCAGCCGTG
- 9. CTGATACCTTCCGCAAACTGTTTCG
- 10. ATACACGAAACAGTTTGCGGAAGGT
- 11. TGTATACTCTAACTTCCTGCGTGGTA
- 15 12. CAGTTTACCACGCAGGAAGTTAGAGT
- 13. AACTGAAACTGTATACTGGCGAAGC
- 14. GGCATGCTTCGCCAGTATACAGTTT
- 15. ATGCCGTA CTGGTGACCGCTAATAG
- 16. TCGAOTATTAGCGGTCACCCAGTAC

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TABLE XIII  
ECEPO SECTION 3

BamHI BglIII  
GA TCCAGATCTCTG  
GTCTAGAGAC

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ACTACTCTGC 1 TCGGTGCTCT 3 GCGTGCACAG AAAGAGGCTA 5 TCTCTCCGCC  
TGATGAGACG 2 ACGCACGAGA 4 CCCACGTGTC TTTCTCCGAT AGAGAGGCCG

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GGATGCTGCA TCTCTGAC 7 GCGTGCCTAC CATCACTGCT 9 GATACCTCC  
CCTACGACGT 6 AGACGAATG 8 GCGACGCATG GTAGTGACGA CTATGGAAGG

GCAAACGTGT TCGTGTATAC 11 TCTAACTTCC TCGTGGTAA 13 ACTGAAACTG  
CGTTTGACAA 10 AGCACATATG 12 AGATTGAAGG ACGCACCATT TGACTTTGAC

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TATACTGGCC 14 AAGCATGCCG 15 TACTGGTGAC CGCTAATAG SalI  
ATATGACCC 14 TTCGTACGGC 16 ATGACCACTG GCGATTATC AGCT

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TABLE XIV  
ECEPO GENE

	<u>XbaI</u>			<u>-1 1</u>	<u>MetAla</u>
	<u>CTAG</u>	AAACCATGAG	GGTAATAAAA	TAATGGCTCC	GCCGCGTCTG
5		TTTGGTACTC	CCATTATTTT	ATTACCGAGG	CGGCGCAGAC
		ATCTGCGACT	CGAGAGTTCT	GGAACGTTAC	CTGCTGGAAG
		TAGACGCTGA	GCTCTCAAGA	CCTTGCAATG	GACGACCTTC
					GATTTCTTCG
		TGAAAACATC	ACCACTGGTT	GTGCTGAACA	CTGTTCTTTG
		ACTTTTGTAG	TGGTGACCAA	CACGACTTGT	GACAAGAAAC
					TTGCTTTTGT
10		TTACGGTACC	AGACACCAAG	GTTAACTTCT	ACGCTTGGA
		AATGCCATGG	TCTGTGGTTC	CAATTGAAGA	TGCGAACCTT
					TGCATACCTT
		GTTGGTCAAC	AAGCAGTTGA	AGTTTGGCAG	GGTCTGGCAC
		CAACCAAGTTG	TTCGTCAACT	TCAAACCGTC	CCAGACCGTG
					ACGACTCGCT
		GGCTGTACTG	CGTGGCCAGG	CACTGCTGGT	AAACTCCTCT
15		CCGACATGAC	GCACCGGTCC	GTGACGACCA	TTTGAGGAGA
					GTCGGCACCC
		AACCGCTGCA	GCTGCATGTT	GACAAAGCAG	TATCTGGCCT
		TTGGCGACGT	CGACGTACAA	CTGTTTCGTC	ATAGACCGGA
					CTCTAGAGAC
		ACTACTCTGC	TGCGTGCTCT	GGGTGCACAG	AAAGAGGCTA
		TGATGAGACG	ACGCACGAGA	CCCACGTGTC	TTTCTCCGAT
					AGAGAGGCGG
20		GGATGCTGCA	TCTGCTGCAC	CGCTGCGTAC	CATCACTGCT
		CCTACGACGT	AGACGACGTG	GCGACGCATG	GTAGTGACGA
					CTATGGAAGG
		GCAAACCTGTT	TCGTGTATAC	TCTAACTTCC	TGCGTGGTAA
		CGTTTGACAA	AGCACATATG	AGATTGAAGG	ACGCACCATT
					TGACTTTGAC
		TATACTGGCG	AAGCATGCCG	TACTGGTGAC	CGCTAATAG
25		ATATGACCGC	TCGTACGGC	ATGACCACTG	GCGATTATCA
					GCT

SalI



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More particularly, <sup>Figure 10</sup> ~~Table VIII~~ illustrates oligo-  
 nucleotides employed to generate the Section 1 of the  
 ECEPO gene encoding amino terminal residues of the human  
 species polypeptide. Oligonucleotides were assembled  
 5 into duplexes (1 and 2, 3 and 4, etc.) and the duplexes  
 were then ligated to provide ECEPO Section 1 as in <sup>Figure 11</sup> ~~Table  
 IX~~. Note that the assembled section includes respective  
 terminal EcoRI and BamHI sticky ends, that "downstream"  
 10 of the EcoRI sticky end is a XbaI restriction enzyme  
 recognition site; and that "upstream" of the BamHI sticky  
 end is a KpnI recognition site. Section 1 could readily  
 be amplified using the M13 phage vector employed for  
 verification of sequence of the section. Some dif-  
 ficulties were encountered in isolating the section as an  
 15 XbaI/KpnI fragment from RF DNA generated in E.coli,  
 likely due to methylation of the KpnI recognition site  
 bases within the host. Single-stranded phage DNA was  
 therefore isolated and rendered into double-stranded form  
 in vitro by primer extension and the desired double-  
 20 stranded fragment was thereafter readily isolated.  
 ECEPO gene Sections 2 and 3 (<sup>Figures 13 and 15</sup> ~~Tables XI and XIII~~)  
 were constructed in a similar manner from the oligo-  
 nucleotides of <sup>Figures 12 and 14</sup> ~~Tables X and XII~~, respectively. Each  
 section was amplified in the M13 vector employed for  
 25 sequence verification and was isolated from phage DNA.  
 As is apparent from <sup>Figure 13</sup> ~~Table XI~~, ECEPO Section 2 was con-  
 structed with EcoRI and BamHI sticky ends and could be  
 isolated as a KpnI/BglII fragment. Similarly, ECEPO  
 Section 3 was prepared with BamHI and SalI sticky ends  
 30 and could be isolated from phage RF DNA as a BglII/SalI  
 fragment. The three sections thus prepared can readily  
 be assembled into a continuous DNA sequence (<sup>Figure 7</sup> ~~Table XIV~~)  
 encoding the entire human species EPO polypeptide with an  
 amino terminal methionine codon (ATG) for E.coli transla-  
 35 tion initiation. Note also that "upstream" of the ini-  
 tial ATG is a series of base pairs substantially

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duplicating the ribosome binding site sequence of the highly expressed OMP-f gene of E.coli.

Any suitable expression vector may be employed to carry the ECEPO. The particular vector chosen for expression of the ECEPO gene as the "temperature sensitive" plasmid pCFM536 -- a derivative of plasmid pCFM414 (A.T.C.C. 40076) -- as described in co-pending U.S. Patent Application Serial No. 636,727, filed August 6, 1984, <sup>(Published EPO Application No 136,440)</sup> by Charles F. Morris. More specifically, pCFM536 was digested with XbaI and HindIII; the large fragment was isolated and employed in a two-part ligation with the ECEPO gene. Sections 1 (XbaI/KpnI), 2 (KpnI/BglII) and 3 (BglII/SalI) had previously been assembled in the correct order in M13 and the EPO gene was isolated therefrom as a single XbaI/HindIII fragment. This fragment included a portion of the polylinker from M13 mp9 phage spanning the SalI to HindIII sites therein. Control of expression in the resulting expression plasmid, p536, was by means of a lambda P<sub>L</sub> promoter, which itself may be under control of the C<sub>1857</sub> repressor gene (such as provided in E.coli strain K12ΔHtrp).

The manufactured ECEPO gene above may be variously modified to encode erythropoietin analogs such as [Asn<sup>2</sup>, des-Pro<sup>2</sup> through Ile<sup>6</sup>]<sub>9</sub>hEPO and [His<sup>7</sup>]<sub>8</sub>hEPO, as described below.

A. [Asn<sup>2</sup>, des-Pro<sup>2</sup> through Ile<sup>6</sup>]<sub>9</sub>hEPO

Plasmid 536 carrying the ECEPO manufactured gene of ~~Table XIV~~ <sup>Figure 7</sup> as a XbaI to HindIII insert was digested with HindIII and XhoI. The latter endonuclease cuts the ECEPO gene at a unique, 6 base pair recognition site spanning the last base of the codon encoding Asp<sup>8</sup> through the second base of the Arg<sup>10</sup> codon. A XbaI/XhoI "linker" sequence was manufactured having the following sequence:

78

T0610X

	<u>XbaI</u>		+1	2	7	8	9		<u>XhoI</u>
		Met	Ala	Asn	Cys	Asp			
5'	-CTAG	ATG	GCT	AAT	TGC	GAC	-3'		
3'	-TAC	CGA	TTA	ACG	CTG	AGCT	-5'		

ρ The XbaI/XhoI linker and the XhoI/HindIII ECEPO gene sequence fragment were inserted into the large fragment resulting from XbaI and HindIII digestion of 13 plasmid pCFM526 -- a derivative of plasmid pCFM414 13 (A.T.C.C. 40076) -- as described in co-pending U.S. Patent Application Serial No. 636,727, filed August 10 6, 1984, by Charles F. Morris, to generate a plasmid-borne DNA sequence encoding E.coli expression of the 31 Met<sup>-1</sup> form of the desired analog.

ρ B. [His<sup>7</sup>]hEPO  
150 Plasmid 536 was digested with HindIII and XhoI as in part A above. A XbaI/XhoI linker was manufactured having the following sequence:

T0611X

	<u>XbaI</u>		+1	2	3	4	5	6	7	8	9		<u>XhoI</u>
		Met	Ala	Pro	Pro	Arg	Leu	Ile	His	Asp			
20	5'	-CTAG	ATG	GCT	CCG	CCA	CGT	CTG	ATC	CAT	GAC	-3'	
	3'	-TAC	CGA	GGC	GGT	GCA	GAC	TAG	GTA	CTG	AGCT	-5'	

ρ The linker and the XhoI/HindIII ECEPO sequence fragment were then inserted into pCFM526 to generate a plasmid-borne DNA sequence encoding E.coli expression of 3125 the Met<sup>-1</sup> form of the desired analog.

Construction of a manufactured gene ("SCEPO") incorporating yeast preference codons is as described in the following ~~Tables XV through XX~~ <sup>Figures 16 through 21 and 8</sup>. As was the case with the ECEPO gene, the entire construction involved 30 formation of three sets of oligonucleotides (~~Tables XV, XVII and XIX~~ <sup>Figures 16, 18 and 20</sup>) which were formed into duplexes and assembled into sections (~~Tables XVI, XVIII and XX~~ <sup>Figures 17, 19 and 31</sup>). Note that synthesis was facilitated in part by use of some sub-optimal codons in both the SCEPO and ECEPO construc-

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14 tions, i.e., oligonucleotides 7-12 of Section 1 of both  
14 genes were identical, as were oligonucleotides 1-6 of  
Section 2 in each gene.

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TABLE XV  
SCEPO SECTION 1 OLIGONUCLEOTIDES

- 1. AATTCAAGCTTGGATAAAAGAGCT
- 5 2. GTGGAGCTCTTTTATCCAAGCTTG
- 3. CCACCAAGATTGATCTGTGACTC
- 4. TCTCGAGTCACAGATCAATCTTG
- 5. GAGAGTTTTGGAAAGATACTTGTTG
- 6. CTTCCAACAAGTATCTTCCAAAAC
- 10 7. GAAGCTAAAGAAGCTGAAAACATC
- 8. GTGGTGATGTTTTCACTTCTTTAG
- 9. ACCACTGGTTGTGCTGAACACTGTTT
- 10. CAAAGAACAGTGTTCAGCACATCCA
- 11. TTTGAACGAAAACATTACGGTACCG
- 15 12. GATCCGGTACCGTAATGTTTTCGTT

TABLE XVI  
SCEPO SECTION 1

20 EcoRI HindIII 1  
AATTCA AGCTTGGATA  
GT TCGAACCTAT  
2

AAAGAGCTTC 3 ACCAAGATTG ATCTGTGACT CAGAGTTTT  
TTTCTCGAGG TGTTCYAAC TAGACACTGA GCTCTCAAAA  
4

25 5 GGAAAGATAC TTGTTGAAG 7 CTAAGAAGC TGAAAACATC ACCACTGGTT  
CCTTTCTATG ACAAACCTTC GATTTCTTCG ACTTTTGTAG TGGTACCAA  
6 8

9 GTGCTGAACA CTGTTCTTTG 11 AACGAAAACA TTACGGTACC KpnI BamHI  
CACGACTTGT GACAAGAAAC TTAGCTTTTGT AATGCCATGG CCTAG  
12

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TABLE XVII

SCEPO SECTION 2 OLIGONUCLEOTIDES

- 1. AATTCGGTACCAGACACCAAGGT
- 5 2. GTTAACCTTGGTGTCTGGTACCG
- 3. TAACTTCTACGCTTGGAAACGTAT
- 4. TTCCATACGTTTCCAAGCGTAGAA
- 5. GGAAGTTGGTCAACAAGCAGTTGAAGT
- 6. CCAAACCTCAACTGCTTGTGACCAAC
- 10 7. TTGGCAAGGTTTGGCCTTGTATCTG
- 8. GCTTCAGATAACAAGGCCAACCTTG
- 9. AAGCTGTTTTGAGAGGTCAAGCCT
- 10. AACAAGGCTGACCTCTCAAACA
- 11. TGTGGTTAACTCTTCTCAACCATGGG
- 15 12. TGGTTCCCATGGTTGAGAAGAGTTAACC
- 13. AACCATTGCAATTGCACGTCGAT
- 14. CTTTATCGACGTGCAATTGCAA
- 15. AAAGCCGTCTCTGGTTTGAGATCTG
- 16. GATCCAGATCTCAAACCAGAGACGG

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TABLE XVIII  
SCEPO SECTION 2

<sup>EcoRI</sup>  
<sup>KpnI</sup>  
<sup>1</sup>  
5 A ATTCGGTACC AGACACCAAG  
GCCATGG TCTGTGGTTC  
<sup>2</sup>

<sup>3</sup>  
GTTAACTTCT ACGCTTGGAA ACGATCGAA GTTGGTCAAC AAGCTGTTGA  
CAATTGAAGA TGCGAACCTT TGCATACCTT CAACCAGTTG TCGACAACT  
<sup>4</sup>

<sup>7</sup>  
10 AGTTGGCAA GGTTGGCT TGTTATCTGA AGCTGTTTGT AGAGGTCAAG  
TCAAACGTT CCAAACTGA ACAATAGACT TCGACAAAC TCTCCAGTTC  
<sup>8</sup>

<sup>11</sup>  
CCTTGTGGT TAACTCTTCT CAACCATGGG AACCATTGCA ATTGCACGTC  
GGAACACCA ATGAGAAGA GTTGGTACCC TTGGTAACGT TAACGTGCAG  
<sup>12</sup>

<sup>15</sup> <sup>BglII</sup> <sup>BamHI</sup>  
15 GATAAAGCCG TCTCTGGTTT GAGATCTG  
CTATTTCGGC AGAGACCAAA CTCTAGACCTA G  
<sup>16</sup>

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TABLE XIX

SCEPO SECTION 3 OLIGONUCLEOTIDES

1. GATCCAGATCTTTGACTACTTTGTT
- 5 2. TCTCAACAAAGTAGTCAAAGATCTG
3. GAGAGCTTTGGGTGCTCAAAGGAAG
4. ATGGCTTCCTTTTGAGCACCCAAAGC
5. CCATTTCCCCACCAGACGCTGCTT
6. GCAGAAGCAGCGTCTGGTGGGGAA
- 10 7. CTGCCGCTCCATTGAGAACCATC
8. CAGTGATGGTTCTCAATGGAGCG
9. ACTGCTGATACTTCAGAAAGTT
10. GAATAACTTTGAGGGTATCAG
11. ATTCAGAGTTACTCCAACTTCT
- 15 12. CTCAAGAAGTGGAGTAACTCT
13. TGAGAGGTAATTGAAGTTGTACAC
14. ACCGGTGTACAACCTTCAATTTACCT
15. CGGTGAAGCCTGTAGAACTGGT
16. CTGTCACCAGTTCTACAGGCTTC
- 20 17. GACAGATAAGCCCGACTGATAA
18. GTTGTATCAGTCGGGCTTAT
19. CAACAGTGTAGATGTAACAAAG
20. TCGACTTTGTTACATCTACACT

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TABLE XX  
SCEPO SECTION 3

BamHI BglII 1  
GATC CAGATCTTTG ACTACTTTGT TGAGAGCTTT  
GTCTAGAAAC TGATGAAACA ACTCTCGAAA

5

3 5  
GGGTGCTCAA AAGGAAGCCA TTCCCCACC AGACGCTGCT TCTGCCGCTC  
CCCACGAGTT TTCCTTCGGT AAGGGGTGG TCTGCGACGA AGACGGCGAG

4 6

7 9 11  
CATTGAGAAC CATCACTGCT GATACCTTCA GAAAGTTATT CAGAGTTTAC  
GTAACTCTTG GTAGTGACCA CTATCGAAGT CTTTCAATAA GTCTCAAATG

10

8 10 12

13 15  
TCCAACCTTCT TGAGAGGTAA ATTGAAGTTG TACACCGGTG AAGCCTGTAG  
AGTTTGAAGA ACTCTCCATT TAACTTCAAC ATGTGGCCAC TTCGGACATC

14 16

17 19  
AACTGGTAC AGATAAGCCC GACTGATAAC AACAGTGTAG  
TTGACCACTG TCTATTCCGGG CTGACTATTG TTGTCACATC

15

18

SalI  
ATGTAACAAA C  
TACATTGTTT CAGCT

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TABLE XXI  
SCEPO GENE

		-1 +1			
	<u>HindIII</u>	<u>ArgAla</u>			
	AGCTTGGATA	AAAGAGCTCC	ACCAAGATTG	ATCTGTGACT	CGAGAGTTT
5	ACCTAT	TTTCTCGAGG	TGTTCTAAC	TAGACACTGA	GCTCTCAAAA
	GGAAAGATAC	TTGTTGGAAG	CTAAAGAAGC	TGAAAACATC	ACCACTGGTT
	CCTTTCTATG	AACAACCTTC	GATTTCTTCG	ACTTTTGTAG	TGGTGACCAA
	GTGCTGAACA	CTGTTCTTTG	AACGAAAACA	TTACGGTACC	AGACACCAAG
	CACGACTTGT	GACAAGAAAC	TTGCTTTTGT	AATGCCATGG	TCTGTGGTTC
10	GTTAACTTCT	ACGCTTGGA	ACGTATGGAA	GTTGGTCAAC	AAGCTGTTGA
	CAATTGAAGA	TGCGAACCTT	TGCATACCTT	CAACCAGTTG	TTCGACAAC
	AGTTTGGCAA	GGTTTGGCCT	TGTTATCTGA	AGCTGTTTTG	AGAGGTCAAG
	TCAAACCGTT	CCAAACCGGA	ACAATAGACT	TCGACAAAAC	TCTCCAGTTC
	CCTTGTTGGT	TAACCTTCT	CAACCATGGG	AACCATTGCA	ATTGCACGTC
15	GGAACAACCA	ATTGAGAAGA	ATTGTTACCC	TTGGTAACGT	TAACGTGCAG
	GATAAAGCCG	TCTCTGGTTT	GAGATCTTTG	ACTACTTTGT	TGAGAGCTTT
	CTATTTCCGGC	AGAGACCAAA	CTCTAGAAAC	TGATGAAACA	ACTCTCGAAA
	GGGTGCTCAA	AAGGAAGCCA	TTTCCCCACC	AGACGCTGCT	TCTGCCGCTC
	CCCACGAGTT	TTCCTTCGGT	AAAGGGGTGG	TCTGCGACGA	AGACGGCGAG
20	CATTGAGAAC	CATCACTGCT	GATACCTTCA	GAAAGTTATT	CAGAGTTTAC
	GTAACCTTTG	GTAGTGACGA	CTATGGAAGT	CTTCAATAA	GTCTCAAATG
	TCCAACCTTCT	TGAGAGGTAA	ATTGAAGTTG	TACACCGGTG	AAGCCTGTAG
	AGGTTGAAGA	ACTCTCCATT	TAACCTCAAC	ATGTGGCCAC	TTCGGACATC
	AACTGGTGAC	AGATAAGCCC	GACTGATAAC	AACAGTGTAG	
25	TTGACCACTG	TCTATTCGGG	CTGACTATTG	TTGTACATC	
		<u>SaI I</u>			
	ATGTAACAAA	G			
	TACATTGTTT	CAGCT			

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The assembled SCEPO sections were sequenced in M13 and Sections 1, 2 and 3 were isolatable from the phage as HindIII/KpnI, KpnI/BglII, and BglII/SalI fragments.

5 The presently preferred expression system for SCEPO gene products is a secretion system based on  
 6) S.cerevisiae  $\alpha$ -factor secretion, as described in co-  
 pending U.S. Patent Application Serial No. 487,753, filed  
 April 22, 1983, by Grant A. Bitter, published October 31,  
 10 1984 as European Patent Application <sup>Nov</sup> 0 123,294. Briefly  
 put, the system involves constructions wherein DNA  
 60 encoding the leader sequence of the yeast  $\alpha$ -factor gene  
 40 product is positioned immediately 5' to the coding region  
 of the exogenous gene to be expressed. As a result, the  
 15 gene product translated includes a leader or signal  
 sequence which is "processed off" by an endogenous yeast  
 enzyme in the course of secretion of the remainder of the  
 60 product. Because the construction makes use of the  $\alpha$ -  
 factor translation initiation (ATG) codon, there was no  
 3/20 need to provide such a codon at the -1 position of the  
 SCEPO gene. As may be noted from <sup>Figure 8</sup> ~~Table XXX~~, the alanine  
 30 (+1) encoding sequence is preceded by a linker sequence  
 allowing for direct insertion into a plasmid including  
 60 the DNA for the first 80 residues of the  $\alpha$ -factor leader  
 6025 following the  $\alpha$ -factor promoter. The specific preferred  
 construction for SCEPO gene expression involved a four-  
 part ligation including the above-noted SCEPO section  
 fragments and the large fragment of HindIII/SalI  
 60 digestion of plasmid pC3. From the resulting plasmid  
 30 pC3/SCEPO, the  $\alpha$ -factor promoter and leader sequence and  
 SCEPO gene were isolated by digestion with BamHI and  
 ligated into BamHI digested plasmid pYE to form  
 expression plasmid pYE/SCEPO.

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CL EXAMPLE 12

P The present example relates to expression of

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recombinant products of the manufactured ECEPO and SCEPO genes within the expression systems of Example 11.

In use of the expression system designed for use of E.coli host cells, plasmid p536 of Example 11 was transformed into AM7 E.coli cells previously transformed with a suitable plasmid, pMW1, harboring a C<sub>1857</sub> gene.

32 Cultures of cells in LB broth (Ampicillin 50 µg/ml and  
82 kanamycin 5 µg/ml, preferably with 10 mM MgSO<sub>4</sub>) were  
20 maintained at 28°C and upon growth of cells in culture to  
32 10 O.D.<sub>600</sub> = 0.1, EPO expression was induced by raising the  
20 culture temperature to 42°C. Cells grown to about 40  
O.D. provided EPO production (as estimated by gel) of  
about 5 mg/OD liter.

Cells were harvested, lysed, broken with French  
15 Press (10,000 psi) and treated with lysozyme and NP-40  
detergent. The pellet resulting from 24,000 xg centrifuga-  
tion was solubilized with guanidine HCl and subjected  
to further purification in a single step by means of  
14 C<sub>4</sub> (Vydac) Reverse Phase HPLC (EtOH, 0-80%, 50 mM NH<sub>4</sub>Ac,  
20 pH 4.5). Protein sequencing revealed the product to be  
greater than 95% pure and the products obtained revealed  
two different amino terminals, A-P-P-R... and P-P-R... in  
a relative quantitative ratio of about 3 to 1. This  
8 9 latter observation of hEPO and [des Ala<sup>1</sup>]hEPO products  
25 indicates that amino terminal "processing" within the  
host cells serves to remove the terminal methionine and  
in some instances the initial alanine. Radioimmunoassay  
activity for the isolates was at a level of 150,000 to  
160,000 U/mg; in vitro assay activity was at a level of  
30 30,000 to 62,000 U/mg; and in vivo assay activity ranged  
from about 120 to 720 U/mg. (Cf., human urinary isolate  
standard of 70,000 U/mg in each assay.) The dose response  
curve for the recombinant product in the in vivo assay  
differed markedly from that of the human urinary EPO  
35 standard.

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The EPO analog plasmids formed in parts A and B of Example 11 were each transformed into pMW1-transformed AM7 E.coli cells and the cells were cultured as above. Purified isolates were tested in both RIA and in vitro assays. RIA and in vitro assay values for [Asn<sup>2</sup>, des-Pro<sup>2</sup> through Ile<sup>6</sup>]hEPO expression products were approximately 11,000 U/mg and 6,000 U/mg protein, respectively, while the assay values for [His<sup>7</sup>]hEPO were about 41,000 U/mg and 14,000 U/mg protein, respectively, indicating that the analog products were from one-fourth to one-tenth as "active" as the "parent" expression product in the assays.

In the expression system designed for use of S.cerevisiae host cells, plasmid pYE/SCEPO was transformed into two different strains, YSDP4 (genotype  $\alpha$  pep4-3 trp1) and RK81 (genotype  $\alpha$  pep4-3 trp1). Transformed YSDP4 hosts were grown in SD medium (Methods in Yeast Genetics, Cold Spring Harbor Laboratory, Cold Spring Harbor, N.Y., p. 62 (1983) supplemented with caseamino acids at 0.5%, pH 6.5 at 30°C. Media harvested when the cells had been grown to 36 O.D. contained EPO products at levels of about 244 U/ml (97  $\mu$ g/OD liter by RIA). Transformed RK81 cells grown to either 6.5 O.D. or 60 O.D. provided media with EPO concentrations of about 80-90 U/ml (34  $\mu$ g/OD liter by RIA). Preliminary analyses reveal significant heterogeneity in products produced by the expression system, likely to be due to variations in glycosylation of proteins expressed, and relatively high mannose content of the associated carbohydrate.

Plasmids PaC3 and pYE in HB101 E.coli cells were deposited in accordance with the Rules of Practice of the U.S. Patent Office on September 27, 1984, with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland, under deposit numbers A.T.C.C. 39881 and A.T.C.C. 39882, respectively. Plasmids pCFM526 in AM7 cells, pCFM536 in JM103 cells, and pMW1 in JM103

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a cells were likewise deposited on November 21, 1984 as A.T.C.C. ~~33932~~<sup>39132</sup>, ~~33934~~<sup>39134</sup>, and ~~33933~~<sup>39133</sup>, respectively.

Saccharomyces cerevisiae strains YSP04 and RK81 were deposited on November 21, 1984 as A.T.C.C. 20734 and  
5 20733, respectively.

It should be readily apparent from consideration of the above illustrative examples that numerous exceptionally valuable products and processes are provided by the present invention in its many aspects.

10 Polypeptides provided by the invention are conspicuously useful materials, whether they are microbially expressed products or synthetic products, the primary, secondary or tertiary structural conformation of which was first made known by the present invention.

15 As previously indicated, recombinant-produced and synthetic products of the invention share, to varying degrees, the in vitro biological activity of EPO isolates from natural sources and consequently are projected to have utility as substitutes for EPO isolates in culture  
20 media employed for growth of erythropoietic cells in culture. Similarly, to the extent that polypeptide products of the invention share the in vivo activity of natural EPO isolates they are conspicuously suitable for use in erythropoietin therapy procedures practiced on  
25 mammals, including humans, to develop any or all of the effects herefore attributed in vivo to EPO, e.g., stimulation of reticulocyte response, development of ferrokinetic effects (such as plasma iron turnover effects and marrow transit time effects), erythrocyte mass changes,  
30 stimulation of hemoglobin C synthesis (see, Eschbach, et al., supra) and, as indicated in Example 10, increasing hematocrit levels in mammals. Included within the class of humans treatable with products of the invention are patients generally requiring blood transfusions and  
35 including trauma victims, surgical patients, renal disease patients including dialysis patients, and

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patients with a variety of blood composition affecting disorders, such as hemophilia, sickle cell disease, physiologic anemias, and the like. The minimization of the need for transfusion therapy through use of EPO therapy  
 5 can be expected to result in reduced transmission of infectious agents. Products of the invention, by virtue of their production by recombinant methods, are expected to be free of pyrogens, natural inhibitory substances, and the like, and are thus likely to provide enhanced  
 10 overall effectiveness in therapeutic processes vis-a-vis naturally derived products. Erythropoietin therapy with products of the present invention is also expected to be useful in the enhancement of oxygen carrying capacity of individuals encountering hypoxic environmental conditions  
 15 and possibly in providing beneficial cardiovascular effects.

A preferred method for administration of polypeptide products of the invention is by parenteral (e.g., IV, IM, SC, or IP) routes and the compositions administered would ordinarily include therapeutically  
 20 effective amounts of product in combination with acceptable diluents, carriers and/or adjuvants. Preliminary pharmacokinetic studies indicate a longer half-life in vivo for monkey EPO products when administered IM rather  
 25 than IV. Effective dosages are expected to vary substantially depending upon the condition treated but therapeutic doses are presently expected to be in the range of  
 30 0.1 (~7U) to 100 (~7000U)  $\mu\text{g}/\text{kg}$  body weight of the active material. Standard diluents such as human serum albumin  
 30 are contemplated for pharmaceutical compositions of the invention, as are standard carriers such as saline.

Adjuvant materials suitable for use in compositions of the invention include compounds independently noted for erythropoietic stimulatory effects, such  
 35 as testosterone, progenitor cell stimulators, insulin-like growth factor, prostaglandins, serotonin,

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cyclic AMP, prolactin and triiodothyronine, as well as agents generally employed in treatment of aplastic anemia, such as methenolene, stanozolol and nandrolone [see, e.g., Resegotti, et al., Panminerva Medica, 23, 243-248 (1981); McGonigle, et al., Kidney Int., 25(2), 437-444 (1984); Pavlovic-Kantera, et al., Expt.Hematol., 8(Supp. 8), 283-291 (1980); and Kurtz, FEBS Letters, 14a(1), 105-108 (1982)]. Also contemplated as adjuvants are substances reported to enhance the effects of, or synergize, erythropoietin or asialo-EPO, such as the adrenergic agonists, thyroid hormones, androgens and BPA [see, Dunn, "Current Concepts in Erythropoiesis", John Wiley and Sons (Chichester, England, 1983); Weiland, et al., Blut, 44(3), 173-175 (1982); Kalmanti, Kidney Int., 22, 383-391 (1982); Shahidi, New.Eng.J.Med., 289, 72-80 (1973); Fisher, et al., Steroids, 30(6), 833-845 (1977); Urabe, et al., J.Exp.Med., 149, 1314-1325 (1979); and Billat, et al., Expt.Hematol., 10(1), 133-140 (1982)] as well as the classes of compounds designated "hepatic erythropoietic factors" [see, Naughton, et al., Acta.Haemat., 69, 171-179 (1983)] and "erythrotropins" [as described by Congote, et al. in Abstract 364, Proceedings 7th International Congress of Endocrinology (Quebec City, Quebec, July 1-7, 1984); Congote, Biochem.Biophys.Res.Comm., 115(2), 447-483 (1983) and Congote, Anal.Biochem., 140, 428-433 (1984)] and "erythrogenins" [as described in Rothman, et al., J.Surg.Oncol., 20, 105-108 (1982)]. Preliminary screenings designed to measure erythropoietic responses of ex-hypoxic polycythemic mice pre-treated with either 5- $\alpha$ -dihydrotestosterone or nandrolone and then given erythropoietin of the present invention have generated equivocal results.

Diagnostic uses of polypeptides of the invention are similarly extensive and include use in labelled and unlabelled forms in a variety of immunoassay techniques

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③ including RIA's, ELISA's and the like, as well as a variety of in vitro and in vivo activity assays. See, e.g., Dunn, et al., Expt.Hematol., 11(7), 590-600 (1983); Gibson, et al., Pathology, 16, 155-156 (1984); Krystal, Expt.Hematol., 11(7), 649-660 (1983); Saito, et al., Jap.J.Med., 23(1), 16-21 (1984); Nathan, et al., New Eng.J.Med., 308(9), 520-522 (1983); and various references pertaining to assays referred to therein. Polypeptides of the invention, including synthetic peptides comprising sequences of residues of EPO first revealed herein, also provide highly useful pure materials for generating polyclonal antibodies and "banks" of monoclonal antibodies specific for differing continuous and discontinuous epitopes of EPO. As one example, preliminary analysis of the amino acid sequences of <sup>Figure 6</sup> ~~Table VI~~ in the context of hydropathicity according to Hopp, et al., P.N.A.S. (U.S.A.), 78, pp. 3824-3828 (1981) and of secondary structures according to Chou, et al., Ann.Rev.Biochem., 47, p. 251 (1978) revealed that synthetic peptides duplicative of continuous sequences of residues spanning positions 41-57 inclusive, 116-<sup>128</sup>~~118~~ inclusive and 144-166 inclusive are likely to produce a highly antigenic response and generate useful monoclonal and polyclonal antibodies immunoreactive with both the synthetic peptide and the entire protein. Such antibodies are expected to be useful in the detection and affinity purification of EPO and EPO-related products.

Illustratively, the following three synthetic peptides were prepared:

- 30
- OJ (1) hEPO 41-57, V-P-D-T-K-V-N-F-Y-A-W-K~~E~~  
R-M-E-V-G;
  - PI (2) hEPO 116-128, K-E-A-I-S-P-P-D-A-A-S-A-A;
  - L (3) hEPO 144-166, V-Y-S-N-F-L-R-G-K-L-K-L-Y~~E~~  
T-G-E-A-C-R-T-G-D-R.
- 35

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14 Preliminary immunization studies employing the above-  
 14 noted polypeptides have revealed a relatively weak posi-  
 14 tive response to hEPO 41-57, no appreciable response to  
 14 hEPO 116-128, and a strong positive response to hEPO  
 14 144-166, as measured by capacity of rabbit serum antibo-  
 14 dies to immunoprecipitate <sup>125</sup>I-labelled human urinary EPO  
 isolates. Preliminary in vivo activity studies on the  
 three peptides revealed no significant activity either  
 alone or in combination.

10 While the deduced sequences of amino acid resi-  
 dues of mammalian EPO provided by the illustrative  
 examples essentially define the primary structural con-  
 formation of mature EPO, it will be understood that the  
 specific sequence of 165 amino acid residues of monkey  
 15 species EPO in Table V and the 166 residues of human spe-  
 cies EPO in ~~Table VI~~ <sup>Figure 5</sup> do not limit the scope of useful  
 polypeptides provided by the invention. Comprehended by  
 the present invention are those various naturally-  
 occurring allelic forms of EPO which past research into  
 20 biologically active mammalian polypeptides such as human  
 65  $\gamma$  interferon indicates are likely to exist. (Compare,  
 e.g., the human immune interferon species reported to  
 have an arginine residue at position No. 140 in EPO  
 published application <sup>NO. 1</sup> 0 077 670 and the species reported  
 25 to have glutamine at position No. 140 in Gray, et al.,  
 14 Nature, 295, pp. 503-508 (1982). Both species are  
 65 characterized as constituting "mature" human  $\gamma$  interferon  
 sequences.) Allelic forms of mature EPO polypeptides may  
 vary from each other and from the sequences of ~~Tables V~~ <sup>Figure 5 and 6</sup>  
 30 ~~and VI~~ in terms of length of sequence and/or in terms of  
 deletions, substitutions, insertions or additions of  
 amino acids in the sequence, with consequent potential  
 variations in the capacity for glycosylation. As noted  
 previously, one putative allelic form of human species  
 35 EPO is believed to include a methionine residue at posi-  
 tion 126. Expectedly, naturally-occurring allelic forms

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of EPO-encoding DNA genomic and cDNA sequences are also likely to occur which code for the above-noted types of allelic polypeptides or simply employ differing codons for designation of the same polypeptides as specified.

5 In addition to naturally-occurring allelic forms of mature EPO, the present invention also embraces other "EPO products" such as polypeptide analogs of EPO and fragments of "mature" EPO. Following the procedures of the above-noted published application by Alton, et al. 10 (WO/83/04053) one may readily design and manufacture genes coding for microbial expression of polypeptides having primary conformations which differ from that herein specified for mature EPO in terms of the identity or location of one or more residues (e.g., substitutions, 15 terminal and intermediate additions and deletions). Alternately, modifications of cDNA and genomic EPO genes may be readily accomplished by well-known site-directed mutagenesis techniques and employed to generate analogs and derivatives of EPO. Such EPO products would share at 20 least one of the biological properties of EPO but may differ in others. As examples, projected EPO products of the invention include those which are foreshortened by e.g., deletions [Asn<sup>2</sup>, des-Pro<sup>2</sup> through Ile<sup>6</sup>]hEPO, 8 9 66 [des-Thr<sup>163</sup> through Arg<sup>166</sup>]hEPO and "Δ27-55hEPO", the 25 latter having the residues coded for by an entire exon deleted; or which are more stable to hydrolysis (and, therefore, may have more pronounced or longer lasting effects than naturally-occurring EPO); or which have been altered to delete one or more a potential sites for gly- 30 cosylation (which may result in higher activities for yeast-produced products); or which have one or more cystein residues deleted or replaced by, e.g., histidine 5 9 or serine residues (such as the analog [His<sup>7</sup>]hEPO) and are potentially more easily isolated in active form from 35 microbial systems; or which have one or more tyrosine residues replaced by phenylalanine (such as the analogs

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[Phe<sup>15</sup>]hEPO, [Phe<sup>49</sup>]hEPO, and [Phe<sup>145</sup>]hEPO) and may bind more or less readily to EPO receptors on target cells.

Also comprehended are polypeptide fragments duplicating only a part of the continuous amino acid sequence or

5 secondary conformations within mature EPO, which fragments may possess one activity of EPO (e.g., receptor binding) and not others (e.g., erythropoietic activity).

Especially significant in this regard are those potential fragments of EPO which are elucidated upon consideration

10 of the human genomic DNA sequence of Table VI, i.e.,

"fragments" of the total continuous EPO sequence which are delineated by intron sequences and which may constitute distinct "domains" of biological activity. It is

noteworthy that the absence of in vivo activity for any

15 one or more of the "EPO products" of the invention is not wholly preclusive of therapeutic utility (see, Weiland,

et al., supra) or of utility in other contexts, such as in EPO assays or EPO antagonism. Antagonists of erythropoietin may be quite useful in treatment of polycythemia

20 or cases of overproduction of EPO [see, e.g., Adamson,

14 Hosp.Practice, 18(12), 49-57 (1983), and Hellmann, et

14 9 al., Clin.Lab.Haemat., 5, 335-342 (1983)].

According to another aspect of the present invention, the cloned DNA sequences described herein

25 which encode human and monkey EPO polypeptides are conspicuously valuable for the information which they provide concerning the amino acid sequence of mammalian erythropoietin which has heretofore been unavailable

30 despite decades of analytical processing of isolates of naturally-occurring products. The DNA sequences are also conspicuously valuable as products useful in effecting

the large scale microbial synthesis of erythropoietin by a variety of recombinant techniques. Put another way, DNA sequences provided by the invention are useful in

35 generating new and useful viral and circular plasmid DNA vectors, new and useful transformed and transfected

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microbial procaryotic and eucaryotic host cells (including bacterial and yeast cells and mammalian cells grown in culture), and new and useful methods for cultured growth of such microbial host cells capable of expression of EPO and EPO products. DNA sequences of the invention are also conspicuously suitable materials for use as labelled probes in isolating EPO and related protein encoding cDNA and genomic DNA sequences of mammalian species other than human and monkey species herein specifically illustrated. The extent to which DNA sequences of the invention will have use in various alternative methods of protein synthesis (e.g., in insect cells) or in genetic therapy in humans and other mammals cannot yet be calculated. DNA sequences of the invention are expected to be useful in developing transgenic mammalian species which may serve as eucaryotic "hosts" for production of erythropoietin and erythropoietin products in quantity. See, generally, Palmiter, et al., Science, 222(4625), 809-814 (1983).

Viewed in this light, therefore, the specific disclosures of the illustrative examples are clearly not intended to be limiting upon the scope of the present invention and numerous modifications and variations are expected to occur to those skilled in the art. As one example, while DNA sequences provided by the illustrative examples include cDNA and genomic DNA sequences, because this application provides amino acid sequence information essential to manufacture of DNA sequence, the invention also comprehends such manufactured DNA sequences as may be constructed based on knowledge of EPO amino acid sequences. These may code for EPO (as in Example 12) as well as for EPO fragments and EPO polypeptide analogs (i.e., "EPO Products") which may share one or more biological properties of naturally-occurring EPO but not share others (or possess others to different degrees).

DNA sequences provided by the present invention are thus seen to comprehend all DNA sequences suitable

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for use in securing expression in a procaryotic or eucaryotic host cell of a polypeptide product having at least a part of the primary structural conformation and one or more of the biological properties of erythropoietin, and selected from among: (a) the DNA sequences set out in <sup>Figures 5 and 6</sup> ~~Tables V and VI~~; (b) DNA sequences which hybridize to the DNA sequences defined in (a) or fragments thereof; and (c) DNA sequences which, but for the degeneracy of the genetic code, would hybridize to the DNA sequences defined in (a) and (b). It is noteworthy in this regard, for example, that existing allelic monkey and human EPO gene sequences and other mammalian species gene sequences are expected to hybridize to the sequences of <sup>Figures 5 and 6</sup> ~~Tables V and VI~~ or to fragments thereof. Further, but for the degeneracy of the genetic code, the SCEPO and ECEPO genes and the manufactured or mutagenized cDNA or genomic DNA sequences encoding various EPO fragments and analogs would also hybridize to the above-mentioned DNA sequences. Such hybridizations could readily be carried out under the hybridization conditions described herein with respect to the initial isolation of the monkey and human EPO-encoding DNA or more stringent conditions, if desired to reduce background hybridization.

In a like manner, while the above examples illustrate the invention of microbial expression of EPO products in the context of mammalian cell expression of DNA inserted in a hybrid vector of bacterial plasmid and viral genomic origins, a wide variety of expression systems are within the contemplation of the invention. Conspicuously comprehended are expression systems involving vectors of homogeneous origins applied to a variety of bacterial, yeast and <sup>mammalian</sup> ~~mammalian~~ cells in culture as well as to expression systems not involving vectors (such as calcium phosphate transfection of cells). In

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this regard, it will be understood that expression of, e.g., monkey origin DNA in monkey host cells in culture and human host cells in culture, actually constitute instances of "exogenous" DNA expression inasmuch as the EPO DNA whose high level expression is sought would not have its origins in the genome of the host. Expression systems of the invention further contemplate these practices resulting in cytoplasmic formation of EPO products and accumulation of glycosylated and non-glycosylated EPO products in host cell cytoplasm or membranes (e.g., accumulation in bacterial periplasmic spaces) or in culture medium supernatants as above illustrated, or in rather uncommon systems such as P.aeruginosa expression systems (described in Gray, et al., Biotechnology, 2, pp. 161-165 (1984)).

Improved hybridization methodologies of the invention, while illustratively applied above to DNA/DNA hybridization screenings are equally applicable to RNA/RNA and RNA/DNA screening. Mixed probe techniques as herein illustrated generally constitute a number of improvements in hybridization processes allowing for more rapid and reliable polynucleotide isolations. These many individual processing improvements include: improved colony transfer and maintenance procedures; use of nylon based filters such as GeneScreen and GeneScreen Plus to allow reprobing with same filters and repeated use of the filter, application of novel protease treatments [compared, e.g., to Taub, et al. Anal.Biochem., 126, pp. 222-230 (1982)]; use of very low individual concentrations (on the order of 0.025 picomole) of a large number of mixed probes (e.g., numbers in excess of 32); and, performing hybridization and post-hybridization steps under stringent temperatures closely approaching (i.e., within 4°C and preferably within 2°C away from) the lowest calculated dissociation temperature of any of the mixed probes employed. These improvements combine to

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provide results which could not be expected to attend  
their use. This is amply illustrated by the fact that  
mixed probe procedures involving 4 times the number of  
probes ever before reported to have been successfully  
5 used in even cDNA screens on messenger RNA species of  
relatively low abundance were successfully applied to the  
isolation of a unique sequence gene in a genomic library  
screening of 1,500,000 phage plaques. This feat was  
accomplished essentially concurrently with the publica-  
10 tion of the considered opinion of Anderson, et al.,  
supra, that mixed probe screening methods were  
"...impractical for isolation of mammalian protein genes  
3 when corresponding RNA's are unavailable.

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CM WHAT IS CLAIMED IS:

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1. A purified and isolated polypeptide having part or all of the primary structural conformation and one or more of the biological properties of naturally-occurring erythropoietin and characterized by being the product of procaryotic or eucaryotic expression of an exogenous DNA sequence.

10

2. A polypeptide according to claim 1 further characterized by being free of association with any mammalian protein.

15

3. A polypeptide according to claim 1 wherein the exogenous DNA sequence is a cDNA sequence.

20

4. A polypeptide according to claim 1 wherein the exogenous DNA sequence is a manufactured DNA sequence.

25

5. A polypeptide according to claim 1 wherein the exogenous DNA sequence is a genomic DNA sequence.

30

6. A polypeptide according to claim 1 wherein the exogenous DNA sequence is carried on an autonomously replicating circular DNA plasmid or viral vector.

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7. A polypeptide according to claim 1 possessing part or all of the primary structural conformation of human erythropoietin as set forth in Table VI or any naturally occurring allelic variant thereof.

8. A polypeptide according to claim 1 possessing part or all of the primary structural conformation of monkey erythropoietin as set forth in Table V or any naturally occurring allelic variant thereof.

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9. A polypeptide according to claim 1 which has the immunological properties of naturally-occurring erythropoietin.

5 10. A polypeptide according to claim 1 which has the in vivo biological activity of naturally-occurring erythropoietin.

10 11. A polypeptide according to claim 1 which has the in vitro biological activity of naturally-occurring erythropoietin.

15 12. A polypeptide according to claim 1 further characterized by being covalently associated with a detectable label substance.

13. A polypeptide according to claim 12 wherein said detectable label is a radiolabel.

20 *Sublet B5* 14. A DNA sequence for use in securing expression in a procaryotic or eucaryotic host cell of a polypeptide product having at least a part of the primary structural conformation and one or more of the biological properties of naturally-occurring erythropoietin, said DNA sequence selected from among:

(a) the DNA sequences set out in Tables V and VI or their complementary strands;

(b) DNA sequences which hybridize to the DNA sequences defined in (a) or fragments thereof; and

30 (c) DNA sequences which, but for the degeneracy of the genetic code, would hybridize to the DNA sequences defined in (a) and (b).

35 15. A procaryotic or eucaryotic host cell transformed or transfected with a DNA sequence according

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to claim 14 in a manner ~~allowing the host cell to express said polypeptide product.~~

16. A polypeptide product of the expression of a DNA sequence of claim 14 in a procaryotic or eucaryotic host.

17. A purified and isolated DNA sequence coding for procaryotic or eucaryotic host expression of a polypeptide having part or all of the primary structural conformation and one or more of the biological <sup>activities</sup> ~~properties~~ of erythropoietin.

18. A cDNA sequence according to claim 17.

19. A monkey species erythropoietin coding DNA sequence according to claim 18.

20. A DNA sequence according to claim 19 and including the protein coding region set forth in <sup>Figure 5</sup> ~~Table V~~.

21. A genomic DNA sequence according to claim 17.

22. A human species erythropoietin coding DNA sequence according to claim 21.

23. A DNA sequence according to claim 22 and including the protein coding region set forth in <sup>Figure 6</sup> ~~Table VI~~.

24. A manufactured DNA sequence according to claim 23.

25. A manufactured DNA sequence according to claim 24 and including one or more codons preferred for expression in E.coli cells.

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26. A manufactured DNA sequence according to claim 25, coding for expression of human species erythropoietin.

5 27. A manufactured DNA sequence according to claim 26 including the protein coding region set forth in Table <sup>Figure 7</sup> ~~XXIV~~.

8 28. A manufactured DNA sequence according to claim 24 and including one or more codons preferred for expression in yeast cells.

10 29. A manufactured DNA sequence according to claim 28, coding for expression of human species erythropoietin.

15 30. A manufactured DNA sequence according to claim 29 including the protein coding region set forth in Table <sup>Figure 8</sup> ~~XXI~~.

20 31. A DNA sequence according to claim 17 covalently associated with a detectable label substance.

25 32. A DNA sequence according to claim 31 wherein the detectable label is a radiolabel.

33. A single-strand DNA sequence according to claim 31.

30 34. A <sup>purified and isolated</sup> DNA sequence coding for a polypeptide fragment or polypeptide analog of naturally-occurring erythropoietin.

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35. A DNA sequence coding for [Phe<sup>15</sup>]hEPO, [Phe<sup>49</sup>]hEPO, [Phe<sup>145</sup>]hEPO, [His<sup>7</sup>]hEPO, [Asn<sup>2</sup> des-Pro<sup>2</sup> through Ile<sup>6</sup>]hEPO, [des-Thr<sup>163</sup> through Arg<sup>166</sup>]hEPO, or [ $\Delta$ 27-55]hEPO.

5

36. A DNA sequence according to claim 34 which is a manufactured sequence.

37. A biologically functional circular plasmid or viral DNA vector including a DNA sequence according to either of claims 14, 17, 34 or 35.

38. A procaryotic or eucaryotic host cell stably transformed or transfected with a DNA vector according to claim 37.

15

39. A polypeptide product of the expression in a procaryotic or eucaryotic host cell of a DNA sequence according to claims 17 or 34.

20

40. A glycoprotein product having a primary structural conformation sufficiently duplicative of that of a naturally-occurring erythropoietin to allow possession of one or more of the biological properties thereof and having an average carbohydrate composition which differs from that of naturally-occurring erythropoietin.

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41. A glycoprotein product having a primary structural conformation sufficiently duplicative of that of a naturally-occurring human erythropoietin to allow possession of one or more of the biological properties thereof and having an average carbohydrate composition which differs from that of naturally-occurring human erythropoietin.

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42. Vertebrate cells which can be propagated in vitro continuously and which upon growth in culture are capable of producing in the medium of their growth in excess of 100 U of erythropoietin per  $10^6$  cells in 48 hours as determined by radioimmunoassay.

43. Vertebrate cells according to claim 42 capable of producing in excess of 500 U erythropoietin per  $10^6$  cells in 48 hours.

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44. Vertebrate cells according to claim 42 capable of producing in excess of 1,000 U erythropoietin per  $10^6$  cells in 48 hours.

15

45. Vertebrate cells according to claim 42 which are mammalian or avian cells.

46. Vertebrate cells according to claim 45 which are COS-1 cells or CHO cells.

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47. A synthetic polypeptide having part or all of the amino acid sequence as set forth in Table V and having one or more of the in vivo or in vitro biological activities of naturally-occurring monkey erythropoietin.

25

48. A synthetic polypeptide having part or all of the amino acid sequence set forth in Table VI, other than a sequence of residues entirely within the sequence numbered 1 through 20, and having a biological property of naturally-occurring human erythropoietin.

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49. A synthetic polypeptide having part or all of the secondary conformation of part or all of the amino acid sequence set forth in Table VI, other than a sequence of residues entirely within the sequence numbered 1 through 20, and having a biological property of naturally-occurring human erythropoietin.

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50. A process for the production of a polypeptide having part or all of the primary structural conformation and one or more of the biological properties of naturally-occurring erythropoietin, said process comprising:

growing, under suitable nutrient conditions, procaryotic or eucaryotic host cells transformed or transfected with a DNA vector according to claim 37, and isolating desired polypeptide products of the expression of DNA sequences in said vector.

51. An antibody substance characterized by immunoreactivity with erythropoietin and with a synthetic polypeptide having a primary structural conformation substantially duplicative of a continuous sequence of amino acid residues extant in naturally-occurring erythropoietin except for any polypeptide comprising a sequence of amino acid residues entirely comprehended within sequence,

A-P-P-R-L-I-C-D-S-R-V-L-E-R-Y-L-L-E-A-K.

52. An antibody according to claim 51, which is a monoclonal antibody.

53. An antibody according to claim 51, which is a polyclonal antibody.

54. An antibody according to claim 51, which is immunoreactive with erythropoietin and a synthetic polypeptide having the sequence selected from the sequences:

V-P-D-T-K-V-N-F-Y-A-W-K-R-M-E-V-G,  
K-E-A-I-S-P-P-D-A-A-S-A-A, and  
V-Y-S-N-F-L-R-G-K-L-K-L-Y-T-G-E-A-C-R-T-G-D-R.

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55. A pharmaceutical composition comprising an effective amount of a polypeptide according to claims 1, 16, 39, 40 or 41 and a pharmaceutically acceptable diluent, adjuvant or carrier.

5

56. A method for providing erythropoietin therapy to a mammal comprising administering an effective amount of a polypeptide according to claims 1, 16, 39, 40 or 41.

10

57. A method according to claim 56 wherein the therapy comprises enhancing hematocrit levels.

15

58. A purified and isolated DNA sequence as set out in <sup>Figure 5 or 6</sup> Table V or VI or a fragment thereof or the complementary strand of such a sequence or fragment.

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59. A polypeptide product of the expression of a DNA sequence according to claim 58 in a procaryotic or eucaryotic host cell.

25

60. An improvement in the method for detection of a specific single stranded polynucleotide of unknown sequence in a heterogeneous cellular or viral sample including multiple single-stranded polynucleotides wherien:

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(a) a mixture of labelled single-stranded polynucleotide probes is prepared having uniformly varying sequences of bases, each of said probes being potentially specifically complementary to a sequence of bases which is putatively unique to the polynucleotide to be detected,

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(b) the sample is fixed to a solid substrate;  
(c) the substrate having the sample fixed thereto is treated to diminish further binding of polynucleotides thereto except by way of hybridization to polynucleotides in said sample,



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(d) the treated substrate having the sample fixed thereto is transitorily contacted with said mixture of labelled probes under conditions facilitative of hybridization only between totally complementary poly-  
5 nucleotides, and,

(e) the specific polynucleotide is detected by monitoring for the presence of a hybridization reaction between it and a totally complementary probe within said mixture of labelled probes, as evidenced by the presence  
10 of a higher density of labelled material on the substrate at the locus of the specific polynucleotide in comparison to a background density of labelled material resulting from non-specific binding of labelled probes to the substrate,

15 said improvement comprising using in excess of 32 mixed probes and performance of one or more of the following:

- (1) employing a nylon-based paper as said solid substrate;
- 20 (2) treating with a protease in step (c);
- (3) employing individual labelled probe concentrations of approximately 0.025 picomoles; and
- (4) employing as one of the hybridization con-  
25 ditions in step (d) stringent temperatures approaching to with 4°C away from the lowest calculated  $T_d$  of any of the probes employed.

*add C11*      *add 102*

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ABSTRACT

"PRODUCTION OF ERYTHROPOIETIN"

5 Disclosed are novel polypeptides possessing part  
or all of the primary structural conformation and one or  
more of the biological properties of mammalian erythro-  
poietin ("EPO") which are characterized in preferred  
forms by being the product of procaryotic or eucaryotic  
10 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 incor-  
porated into autonomously replicating plasmid or viral  
15 vectors employed to transform or transfect suitable pro-  
caryotic 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  
20 immunological properties and in vitro and in vivo biolo-  
gical activities of EPO of human or monkey species ori-  
gins. Disclosed also are chemically synthesized  
polypeptides sharing the biochemical and immunological  
properties of EPO. Also disclosed are improved methods  
25 for the detection of specific single stranded poly-  
nucleotides in a heterologous cellular or viral sample  
prepared from, e.g., DNA present in a plasmid or viral  
borne cDNA or genomic DNA "library".

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**DECLARATION FOR PATENT APPLICATION**

As a below named inventor, I hereby declare that my residence, post office address and citizenship are as stated below next to my name. I believe that I am the original, first and sole inventor (if only one name is listed below) or an original, first and joint inventor (if plural names are listed below) of the subject matter which is claimed and for which a patent is sought on the invention entitled "PRODUCTION OF ERYTHROPOIETIN"

the specification of which (check one):  is attached hereto;  was filed on \_\_\_\_\_ as Application Serial No. \_\_\_\_\_ and was amended on (or amended through) \_\_\_\_\_ (if applicable). I hereby state that I have reviewed and understand the contents of the above-identified specification, including the claims, as amended by any amendment(s) referred to above. I acknowledge the duty to disclose information which is material to the examination of this application in accordance with Title 37, Code of Federal Regulations, §1.56(a). I hereby claim foreign priority benefits under Title 35, United States Code, §119 of any foreign application(s) for patent or inventor's certificate listed below and have also identified below any foreign application for patent or inventor's certificate having a filing date before that of the application on which priority is claimed.

Prior Foreign Application(s)			Priority Claimed	
_____ (Number)	_____ (Country)	_____ (Day Month/Year Filed)	<input type="checkbox"/> Yes	<input type="checkbox"/> No
_____ (Number)	_____ (Country)	_____ (Day/Month/Year Filed)	<input type="checkbox"/> Yes	<input type="checkbox"/> No
_____ (Number)	_____ (Country)	_____ (Day/Month/Year Filed)	<input type="checkbox"/> Yes	<input type="checkbox"/> No

I hereby claim the benefit under Title 35, United States Code, §120 of any United States application(s) listed below and, insofar as the subject matter of each of the claims of this application is not disclosed in the prior United States application in the manner provided by the first paragraph of Title 35, United States Code, §112, I acknowledge the duty to disclose material information as defined in Title 37, Code of Federal Regulations, §1.56(a) which occurred between the filing date of the prior application and the national or PCT international filing date of this application:

<u>561,024</u> (Application Serial No.)	<u>December 13, 1983</u> (Filing Date)	<u>Pending</u> (Status - Patented, Pending or Abandoned)
<u>582,185</u>	<u>February 21, 1984</u>	<u>Pending</u>
<u>655,841</u>	<u>September 28, 1984</u>	<u>Pending</u>

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

**POWER OF ATTORNEY:** I (We) hereby appoint as my (our) attorneys, with full powers of substitution and revocation, to prosecute this application and transact all business in the Patent and Trademark Office connected therewith:

- |                              |                            |                             |
|------------------------------|----------------------------|-----------------------------|
| William E. Domruck (15,286)  | <del>John R. Brock</del>   | Nate F. Scarpelli (22,320)  |
| Albert W. Bicknell (15,389)  | Alvin D. Shulman (19,412)  | Edward M. O'Toole (22,477)  |
| William A. Marshall (17,053) | Donald J. Brot (19,490)    | Michael F. Borun (25,447)   |
| Jerome B. Klose (17,104)     | Owen J. Murray (22,111)    | Carl E. Moore, Jr. (26,487) |
| Basil P. Mann (18,464)       | Allen H. Gerstein (22,218) |                             |

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State or Country California	State or Country California
Date November 29, 1984	Signature Fu-Kuen Lin

See second page for additional joint inventors



Attorney's Docket No. 7183  
Applicant or Patentee: FU-KUEN LIN  
Serial or Patent No.: \_\_\_\_\_  
Filed or Issued: Herewith  
For: "PRODUCTION OF ERYTHROPOIETIN"

**VERIFIED STATEMENT (DECLARATION) CLAIMING SMALL ENTITY STATUS (37 CFR 1.9(f) and 1.27(c))—SMALL BUSINESS CONCERN**

I hereby declare that I am

- the owner of the small business concern identified below:
- an official of the small business concern empowered to act on behalf of the concern identified below:

NAME OF CONCERN Kirin-Amgen, Inc.  
ADDRESS OF CONCERN 1900 Oak Terrace Lane  
Thousand Oaks, California 91320

I hereby declare that the above identified small business concern qualifies as a small business concern as defined in 13 CFR 121.3-18, and reproduced in 37 CFR 1.9(d), for purposes of paying reduced fees under Section 41(a) and (b) of Title 35, United States Code, in that the number of employees of the concern, including those of its affiliates, does not exceed 500 persons. For purposes of this statement, (1) the number of employees of the business concern is the average over the previous fiscal year of the concern of the persons employed on a full-time, part-time or temporary basis during each of the pay periods of the fiscal year, and (2) concerns are affiliates of each other when either, directly or indirectly, one concern controls or has the power to control the other, or a third-party or parties controls or has the power to control both.

I hereby declare that rights under contract or law have been conveyed, to and remain with the small business concern identified above with regard to the invention, entitled "PRODUCTION OF ERYTHROPOIETIN" by inventor(s) Fu-Kuen Lin

described in

- the specification filed herewith
- application serial no. \_\_\_\_\_, filed \_\_\_\_\_
- patent no. \_\_\_\_\_, issued \_\_\_\_\_

If the rights held by the above identified small business concern are not exclusive, each individual, concern or organization having rights to the invention is listed below\* and no rights to the invention are held by any person, other than the inventor, who could not qualify as a small business concern under 37 CFR 1.9(d) or by any concern which would not qualify as a small business concern under 37 CFR 1.9(d) or a nonprofit organization under 37 CFR 1.9(e).

\*NOTE: Separate verified statements are required from each named person, concern or organization having rights to the invention availing to their status as small entities. (37 CFR 1.27).

(Small Entity-Small Business—page 1 of 2)

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NAME \_\_\_\_\_

- ADDRESS \_\_\_\_\_

INDIVIDUAL       SMALL BUSINESS CONCERN       NONPROFIT ORGANIZATION

NAME \_\_\_\_\_

ADDRESS \_\_\_\_\_

INDIVIDUAL       SMALL BUSINESS CONCERN       NONPROFIT ORGANIZATION

I acknowledge the duty to file, in this application or patent, notification of any change in status resulting in loss of entitlement to small entity status prior to paying, or at the time of paying, the earliest of the issue fee or any maintenance fee due after the date on which status as a small business entity is no longer appropriate. (37 CFR 1.28(b)).

I hereby declare that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code, and that such willful false statements may jeopardize the validity of the application, any patent issuing thereon, or any patent to which this verified statement is directed.

NAME OF PERSON SIGNING Robert D. Weist

TITLE OF PERSON OTHER THAN OWNER Secretary

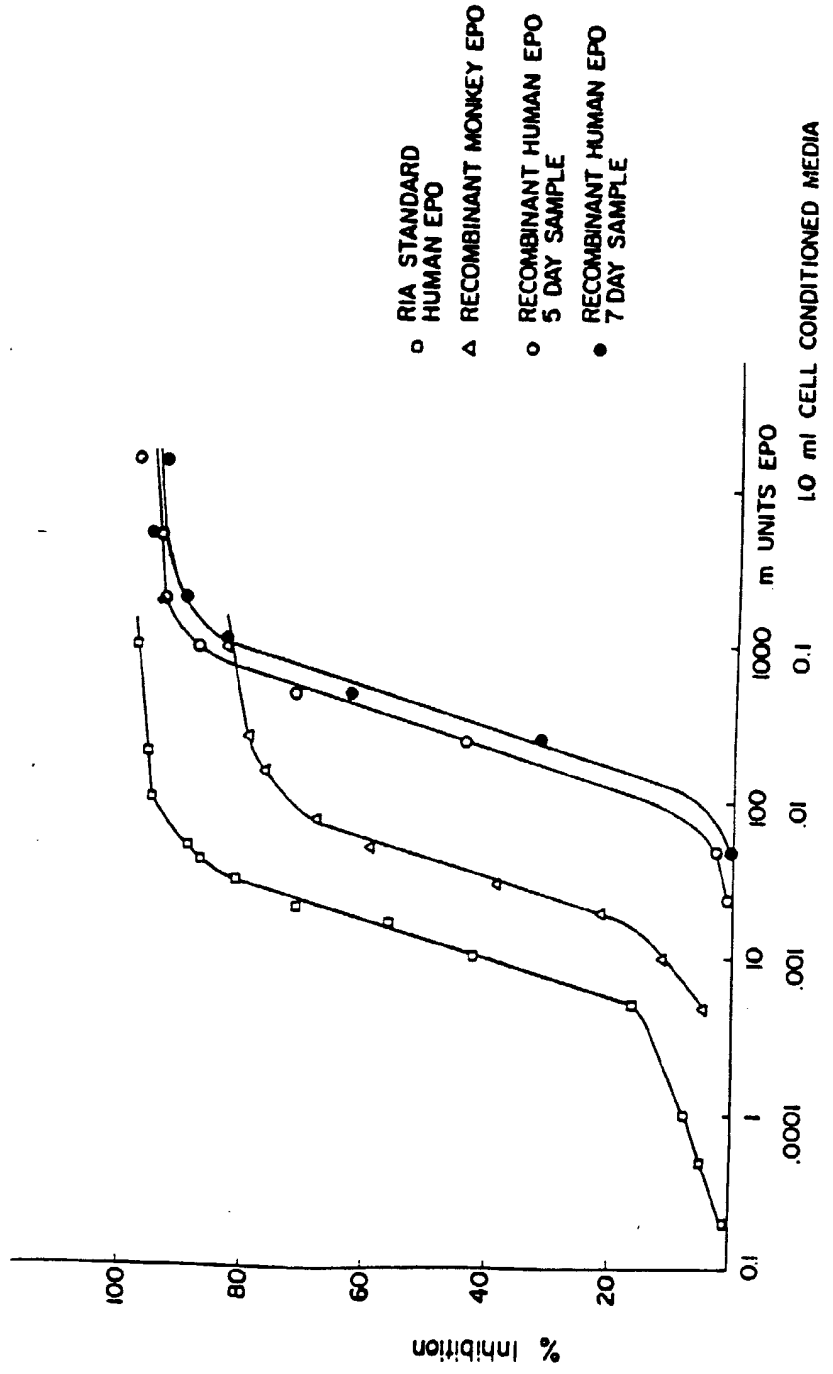
ADDRESS OF PERSON SIGNING 1900 Oak Terrace Lane  
Thousand Oaks, California 91320

SIGNATURE Robert D. Weist November 30, 1984  
Robert D. Weist      Date  
Secretary

(Small Entity-Small Business—page 2 of 2)

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FIG. 1 Comparison of Recombinant Human & Monkey EPO in Radioimmunoassay



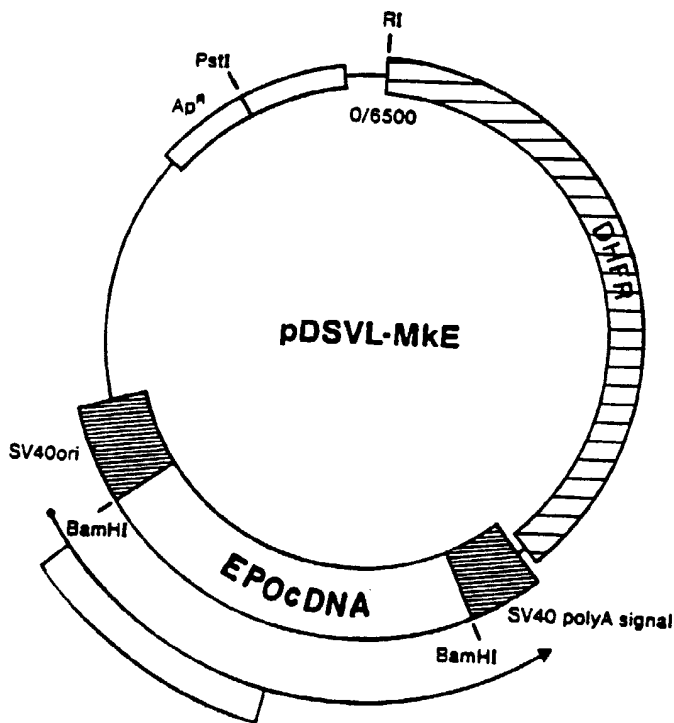
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FIG. 2



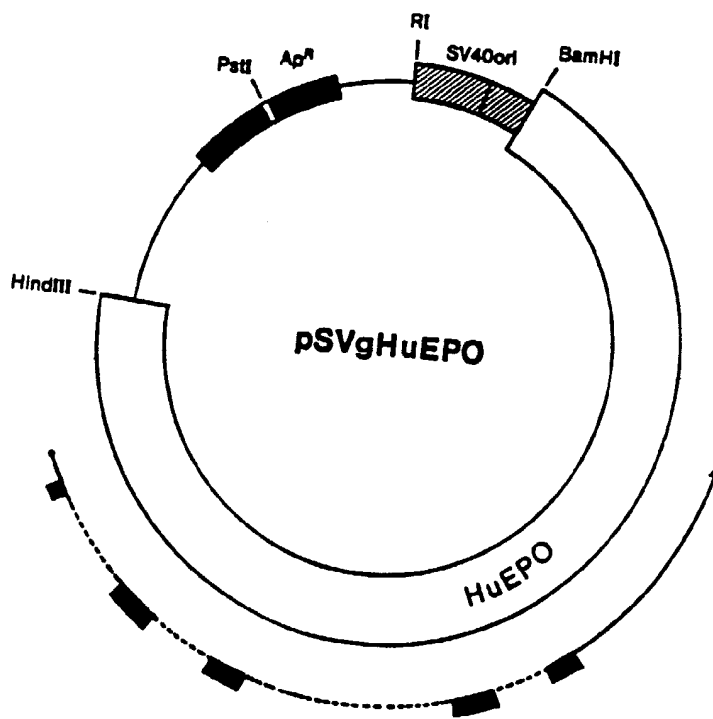
110

115

PRINTED DRAWING AS  
ORIGINALLY FILED.

675298

FIG. 3

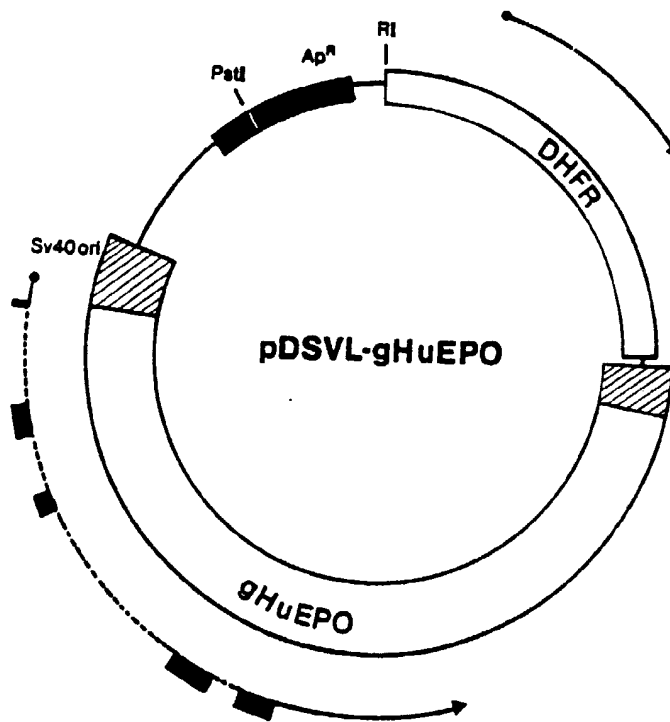




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ORIGINALLY FILED.

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FIG. 4



112

117

6 12242

FIG.5

150 Leu Arg Gly Lys Leu Lys Leu Tyr Thr Gly Glu Ala Cys Arg Arg  
CTC CGG GGA AAG CTG AAG CTG TAC ACG GGG GAG GCC TGC AGG AGA  
165 Gly Asp Arg DP  
GGG GAC AGA TGA CCAGTGGTCCAGCTGGGCACATCCACCACCTCCCTCACCACCA  
CTGCCTGTGCCACACCCCTCCCTCACCACCTCCCGAACCCCATCGAGGGCTCTCAGCTAAG  
CGCCAGCCTGTCCCATGGACACTCCAGTCCAGCAATGACATCTCAGGGGCCAGAGGAAC  
TGTCAGAGCACAACTCTGAGATCTAAGGATGTCCGAGGCCAACTTGAGGGCCAGAGC  
AGGAAGCATTGAGAGAGCAGCTTAAACTCAGGAGCAGAGACAATGCAGGGAACACCT  
GAGCTCACTCGGCCACCCTGC AAAATTTGATGCAGGACACGCTTGGAGGCAATTTACCCTG  
TTTTGCACCTACCAACAGGGACAGGATGACTGGAGAACTTAGGTGGCAAGCTGTGACTT  
CTCAAGGCCCTCAGGGCCACTCCCTTGGTGGCAGAGCCCCCTTGACACTGAGAGAAATTT  
TTGCAATCTGCAGCAGGAAAATTAACGACAGGTTTGGAGGTTGGAGGTTACTTGACAG  
GTGTGTGGGAAGCAGGGCGGTAGGGTGGAGCTGGGATGCCAGTGAAGACCTGAAGAC  
AGGATGGGGCTGGCCCTCGGTTCGGTGGGGTCCAGCTT  
HindIII

113

675 298

FIG.5

Translation of Monkey EPO cDNA

Sau3A  
GATCCGGGGCCCCCTGGACAGCCGCCCTCCCTCCAGGCCCGTGGGCTGGCCCTGGCC  
CGCTGAACCTCCCGGATGAGGACTCCCGGTGGTGGTACCCGCCGCCCTAGGTCGCTGAG

-27  
Met Gly Val His Glu Cys Pro Ala Trp  
GGACCCCGCCAGCCGGGAGATG GGG GTG CAC GAA TGT CCT GCC TGG

-20  
Leu Trp Leu Leu Ser Leu Val Ser Leu Pro Leu Gly Leu Pro  
CTG TGG CTT CTC CTC TCT CTC GTG TCG CTC CCT CTG GGC CTC CCA

-10  
Val Pro Gly Ala Pro Pro Arg Leu Ile Cys Asp Ser Arg Val Leu  
GTC CCG GGC GCC CCA CCA CGC CTC ATC TGT GAC AGC CGA GTC CTG

+1  
Glu Arg Tyr Leu Leu Glu Ala Lys Glu Ala Glu Asn Val Thr Met  
GAG AGG TAC CTC TTG GAG GCC AAG GAG GCC GAG AAT GTC ACG ATG

20  
Gly Cys Ser Glu Ser Cys Ser Leu Asn Glu Asn Ile Thr Val Pro  
GGC TGT TCC GAA AGC TGC AGC TTG AAT GAG AAT ATC ACC GTC CCA

30  
40

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FIG.5

50  
 Asp Thr Lys Val Asn Phe Tyr Ala Trp Lys Arg Met Glu Val Gly  
 GAC ACC AAA GTT AAC TTC TAT GCC TGG AAG AGG ATG GAG GTC GCG  
  
 60  
 Gln Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Leu Ser Glu  
 CAG CAG GCT GTA GAA GTC TGG CAG GGC CTG GCC CTG CTC TCA GAA  
  
 70  
  
 80  
 Ala Val Leu Arg Gly Gln Ala Val Leu Ala Asn Ser Ser Gln Pro  
 CCT GTC CTG CGG GGC CAG GCC GTG TTG GCC AAC TCT TCC CAG CCT  
  
 90  
 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 AGT GGC CTT  
  
 100  
  
 110  
 Arg Ser Ile Thr Thr Leu Leu Arg Ala Leu Gly Ala Gln Glu Ala  
 CGC AGC ATC ACC ACT CTG CTT CGG GCG CTG GGA GCC CAG GAA GCC  
  
 120  
 Ile Ser Leu Pro Asp Ala Ala Ser Ala Ala Pro Leu Arg Thr Ile  
 ATC TCC CTC CCA GAT GCG GCC TCG GCT GCT CCA CTC CGA ACC ATC  
  
 130  
  
 140  
 Thr Ala Asp Thr Phe Cys Lys Leu Phe Arg Val Tyr Ser Asn Phe  
 ACT GCT GAC ACT TTC TGC AAA CTC TTC CGA GTC IAC TCC AAT TTC

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FIG.6

AAGCTTC TGGGCTTCCAGACCAGGTACTTTGGGAAC TACGCAACCCAGGCACTCTGAGTCTCCGCCCA  
AGACCGGATGCCCCAGGGAGGTCCGGAGCCAGCCTTCCAGATAGCACGCTCCGCCAGTCCC  
AAGGTGCCCAACCGGCTGCATCCCTCCCGGACCCAGGCCCGGAGCAGCCCCCATGACCCACACGC  
ACGTC TGCAGCAGCCCGC TCAGCCCCCGGAGCCTCAACCCAGGGTCTCTGCCCTGCTGACCCCGG  
GTGGCCCC TACCCCTGGGAGCCCTCAGGCACACAGCCTCTCCCCCACCCACCCGCGCACACATG  
CAGATAACAGCCCGACCCCGCCAGACCGXAGAGTCCC TGGCCACCCCGGCCGTGCCCTGCCGCTG  
CGCCGACCGGCTGTCTCCCGGAGCCGGACCCGGCCACCGGCCXGCTCTGCTCCGACACCGGCC  
CTTGGACAGCCCTCTCTTAGGCCGTGGGGCTGGCCCTGCACCGCGACTTCCCGGATGAGGXX  
CCCGGTACCGGGCCGCCCAAGTGGTGGGGACCCCGCCAGCGCGGAG ATG GGG GTG CAC G  
GTGAGTACTCGCGGCTGGGGCTCCCGGCGCCGGGT TCC TGT TTAGCGGGGATTAGCCCCCGGCT

-27  
-24  
Met Gly Val His  
ATG GGG GTG CAC G

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FIG.6

YGGTGGCCCAACCATACCTGAACCTAGGCAAGGAGCAAGCCACAGATCCCTACGCCCTGGGCCAGGG

27 30  
Thr Gly Cys Ala Glu  
ACG GGC TGT GCT GAA

CCAGAGCCTTCAGGGACCCTTGACICCCCGGGCTGTGTGCATTTCAG

40  
His Cys Ser Leu Asn Glu Asn Ile Thr Val Pro Asp Thr Lys Val Asn Phe Tyr  
CAC TGC AGC TTG AAT GAG AAT ATC ACT GTC CCA GAC ACC AAA GTT AAT TTC TAT

50  
Ala Trp Lys Arg Met Glu  
GCC TGG AAG AGG ATG GAG GTCAGTCCCTTTTTTTTTTTTTTCTTTTGGAGAACTCAT

YGGAGCCCTGATTTGGATGAAGGGGAGAAATGATCGGGGAAAGGTAAAAATGGAGCAGCAGAGATGAGGCT

GCCTGGCCAGAGGCTCACGCTATAATCCAGGCTGAGATGGCCAGATGGGAGAAATGCTTGAGCCCT

GGAGTTTCAGACCAACC TAGGCAGCATAGTAGATECCCCCACTCTACAACATTTAAAAAAATTAAGTCAG

GTGAAGTGGTGCATGGTGTAGTCCCAGATATTTGGAGGCTGAGGGGGAGGATCGCTTGAGCCAGGAA

TTTGAGGCTCCAGTGAGCTGTGATCACACCACCTGCACTCCAGCTCAGTGACAGAGTGAGGCCCTGTCICA

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FIG. 6

AAAAGAAAAGAAAAGAAAATAATGAGGGCTGTAIGGAATACATTCATTATTCACCTCACTCACT  
 CACTCATTTCATTTCATTTCATTCACCAAGTCTTATTGCATACCTTCCTGTTGCCAGCTGGGCTTGG  
 GGGTGGCTGAGGGGAGGGGAGGGGACATGGGTCAGCTCGACTCCAGAGTCCACTCCCTGTAG  
 56 Val Gly Gln Ala Val Glu Val Trp Gln Gly Leu Ala Leu Ser Glu Ala  
 GTC GGG CAG CAG GCC GTA GAA GTC TGG CAG GCC CTG GCC CTG TCG GAA GCT  
 80 Val Leu Arg Gly Gln Ala Leu Val Asn Ser Ser Gln Pro Trp Glu Pro Leu  
 GTC CTG CCG GGC CAG GCC CTG TTG GTC AAC TCT TCC CAG CCG TGG GAG CCC CTG  
 \*  
 90  
 100 Gln Leu His Val Asp Lys Ala Val Ser Gly Leu Arg Ser Leu Thr Thr Leu Leu  
 CAG CTG CAT GTG GAT AAA GCC GTC AGT GGC CTT CGC ACC CTC ACC ACT CTG CTT  
 110 Arg Ala Leu Gly Ala Gln  
 CCG GCT CTG GGA GCC CAG GTGAGTAGGAGGGGACACTTCCTGCTGCCCTTCTGTAGAGGGGA  
 GAAGGGCTTGTCTAAGGAGTACAGGACIGTCCGTAITTCCTTCCCTTTCGTGGCAGTCCAGGACCTCCT  
 116 Lys Glu Ala Ile Ser Pro Pro Asp Ala Ala Ser Ala Ala  
 CTTTCTCCTTGGCAG AAG GAA GCC ATC TCC CCT CCA GAT GCG GCC TCA GCT GCT  
 120

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## FIG. 6

130 Pro Leu Arg Thr Ile Thr Ala Asp Thr Phe Arg Lys Leu Phe Arg Val Tyr Ser  
CCA CTC CGA ACA ATC ACT GCT GAC ACT TTC CGC AAA CTC TTC CGA GTC TAC TCC

140  
150 Asn Phe Leu Arg Gly Lys Leu Lys Leu Tyr Thr Gly Glu Ala Cys Arg Thr Gly  
AAT TTC CTC CGG GGA AAG CTG AAG CTG TAC ACA GGG GAG GCC TGC AGG ACA GGG

160  
166 Asp Arg OP  
GAC AGA TGA CCAGGIGTGTCCACCCTGGCCATATCCACCACCCTCCCTCACCACDATTGCTTGCCACA

CCC TCCCCCGCCACTCCTGAACCCCGTCCAGGGGCTCAGCTCAGCCCGAGCCGTGCCCATGGACACTCC

AGT GCCAGCAATGACATC TCAGGGGGCCAGAGGAAC TGCCAGAGAGCAACTCGAGATCTAAGGATGTCAC

AGGGCCCACTTGAAGGGCCAGAGCCAGGAGCATTCAGAGAGCAGCTTAAACTCAGGACAGAGCCCATGC

TGGGAGAGCGCCTGAGCTCAGCTGGCACCCCTGCCAAATTTGATGCCAGGACACGGCTTTGGAGCCGATTAC

CTGT T TCCGACCTACCATCAGGGACAGGATGACCTGGAGAACTTAGGTGGCAAGCTGTGACTTCTCCAGG

TCTCAGGGGCATGGGCATCCCTTGGTGGCAAGAGCCCCCTTGACACCCGGGTGGTGGCAACCATGAAGAC

AXCATXGGGGCTGGCCCTCGCCCTCATGTGGGTCCTCAAGTTTGGTATCTCAACCTATTGACAGACTGAA

ACACAATATGAC

FIG.7

ECEPO GENE

```

                -1 1
                MetAla
    xbaI
CTAG AAACCATGAG GGTAAATAAAA TAATGGCTCC GCCCGCTCTG
      TTTGGTACTC CCATTATTTT ATTACCGAGG CGGCCGAGAC

ATCTGGGACT CGAGAGTTCT GGAACGTTAC CTGCTGGAAG CTAAGAAGC
TAGACGCTGA GCTCTCAAGA CCTTGCAATG GACGACCTTC GATTTCTTCG

TGAAAACATC ACCACTGGTT GTGCTGAACA CTGTTCTTTG AACGAAAACA
ACTTTTGTAG TGGTGACCAA CACGACTTGT GACAAGAAAC TTGCTTTTGT

TTACGGTACC AGACACCAAG GTTAACTTCT ACGCTTGGAA ACGTATGGAA
AATGCCATGG TCTGTGGTTC CAATTGAAGA TGCGAACCTT TGCATACCTT

GTTGGTCAAC AAGCAGTTGA AGTTTGGCAG GGTCTGGCAC TGCTGAGCGA
CAACCAGTTG TTCGTCAACT TCAAACCGTC CCAGACCGTG ACGACTCGCT

GGCTGTACTG CGTGGCCAGG CACTGCTGGT AAACCTCTCT CAGCCGTGGG
CCGACATGAC GCACCGGTCC GTGACGACCA TTTGAGGAGA GTCGGCACCC

AACCGCTGCA GCTGCATGTT GACAAAGCAG TATCTGGCCT GAGATCTCTG
TTGGCGACGT CGACGTACAA CTGTTTCGTC ATAGACCGGA CTCTAGAGAC

ACTACTCTGC TCGTGTCTCT GGGTGCACAG AAAGAGGCTA TCTCTCCGC
TGATGAGACG ACGCACGAGA CCCACGTGTC TTTCTCCGAT AGAGAGGCGG

GGATGCTGCA TCTGCTGCAC CGCTGCGTAC CATCACTGCT GATACCTTCC
CCTACGACGT AGACGACGTG GCGACGCATG GTAGTGACGA CTATGGAAGG

GCAA~ACTGTT TCGTGTATAC TCTAACTTCC TCGTGGTAA ACTGAAACTG
CGTTTGACAA AGCACATATG AGATTGAAGG ACGCACCATT TGACTTTGAC

TATACTGGCG AAGCATGCCG TACTGGTGAC CGCTAATAG SalI
ATATSACCGC TTCGTACGGC ATGACCACTG GCGATTATCA GCT
    
```

FIG.8

SCEPC GENE

```

                -1 +1
HindIII      ArgAla
ACCTTGGATA AAAGAGCTCC ACCAAGATTG ATCTGTGACT CGAGAGTTTT
      ACCTAT TTTCTCGAGG TGGTTCTAAC TAGACACTGA GCTCTCAAAA

GGAAAGATAC TTGTTGGAAG CTAAAGAAGC TGAAAACATC ACCACTGGTT
CCTTCTATG AACAACTTC GATTTCTTCG ACTTTTGTAG TGGTGACCAA

GTGCTGAACA CTGTTCTTTG AACGAAAACA TTACGGTACC AGACACCAAG
CAGGACTTGT GACAAGAAAC TTGCTTTTGT AATGCCATGG TCTGTGGTTC

GTTAACTTCT ACGCTTGGAA ACGTATGGAA GTTGGTCAAC AAGCTGTTGA
CAATTGAAGA TCGGAACCTT TGCATACCTT CAACCAGTTG TTCGACAAC

AGTTTGGCAA GGTITGGCCT TGTTATCTGA AGCTGTTTTG AGAGGTCAAG
TCAAACCGTT CCAAACCGGA ACAATAGACT TCGACAAAAC TCTCCAGTTC

CCTTGTGGT TAACTCTTCT CAACCATGGG AACCATTGCA ATTGCACGTC
GGAACAACCA ATTGAGAAGA GTTGGTACCC TTGGTAACGT TAACGTGCAG

GATAAAGCCG TCTCTGGTTT GAGATCTTTG ACTACTTTGT TGAGAGCTTT
CTATTTCCGGC AGAGACCAA CTCTAGAAAC TGATGAAACA ACTCTCGAAA

GGGTGCTCAA AAGGAAGCCA TTTCCCCACC AGACGCTGCT TCTGCCGCTC
CCCACGAGTT TTCCTTCGGT AAAGGGGTGG TCTGCGACGA AGACGGCGAG

CATTGAGAAC CATCACTGCT GATACCTTCA GAAAGTTATT CAGAGTTTAC
GTAACCTCTG GTAGTGACGA CTATGGAAGT CTTTCAATAA GTCTCAAATG

TCCAACCTTCT TGAGAGGTAA ATTGAAGTTG TACACCGGTG AAGCCTGTAG
AGGTTGAAGA ACTCTCCATT TAACCTCAAC ATGTGGCCAC TTCGGACATC

AACTGGTGAC AGATAAGCCC GACTGATAAC AACAGTGTAG
TTGACCACTG TCTATTCGGG CTGACTATTG TTGTACATC

                SalI
ATGTAACAAA G
TACATTGTTT CAGCT
    
```

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Comparison of Human and Monkey EPD Polypeptides

		-10	+1	10	20	30	40	
Human	MGVHECPAWLWLLSLLSPLGLPVLGAPPRLICDSRVLERYLLEAKEAENITTCGCAEHC SLNENITVPDTK							
Monkey	MGVHECPAWLWLLSLSPLGLPVPAPPRLICDSRVLERYLLEAKEAENVTMCCSECSLNENITVPDTK							
		50	60	70	80	90	100	110
Human	VNFYAKRMEVGGQAVEVWQGLALLSEAVLRGQALLVNSSQPWEPLQLHVQKAVSGLRSLITLLRALGQAKE							
Monkey	VNFYAKRMEVGGQAVEVWQGLALLSEAVLRGQAVLANSSQPFEPQLHMDKAISGLRSITLLRALGQAE							
		120	130	140	150	160		
Human	AISPPDAASAAPLRTITADTFKRLFRVYSNFLRGKCLKLYTGEACRTGDR							
Monkey	AISLPDAASAAPLRTITADTFCKLFRVYSNFLRGKCLKLYTGEACRRGDR							

FIG. 9

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ECEPO SECTION 1 OLIGONUCLEOTIDES

1. AATTCTAGAAACCATGAGGGTAATAAAATA
2. CCATTATTTTATTACCCTCATGGTTTCTAG
3. ATGGCTCCGCCCGCTGATCTGCGAC
4. CTCGAGTCGCAGATCAGACGGCGGAG
5. TCGAGAGTTCTGGAACGTTACCTGCTG
6. CTTCCAGCAGGTAACGTTCCAGAACT
7. GAAGCTAAAGAAGCTGAAAACATC
8. GTGGTGATGTTTTTCAGCTTCTTTAG
9. ACCACTGGTTGTGCTGAACACTGTTC
10. CAAAGAACAGTGTTTCAGCACAACCA
11. TTTGAACGAAAACATTACGGTACCG
12. GATCCGGTACCGTAATGTTTTTCGTT

FIG. 10

ECEPO SECTION 1

```

      XbaI
EcoRI 1 3
AATTCTAG AAACCATGAG GGTAAATAAAA TATGGCTCC GCCGGTCTG
      GATC TTTGGTACTC CCATTATTTT ATTACGAGG CGGCGCAGAC
                2 4

ATCTGCGACT CGAGAGTTCT GGAACGTTAC CTGCTGAAG CTAAGAAGC
TAGACGCTGA GCTCAAGA CCTTGCAATG GACGACCTT GATTTCTTCG
                5 6

TGAAAACATC ACCACTGGTT GTGCTGAACA CTGTTCTTTG AACGAAAACA
ACTTTTGTAG TGGTGACCAA CACGACTTGT GACAAGAAAC TTGCTTTTGT
      7 8 9 10 11

      KpnI BamHI
TTACGGTACC G
AATGCCATGG CCTAG
      12
    
```

FIG. 11

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ECEPO SECTION 2 OLIGONUCLEOTIDES

1. AATTCGGTACCAGACACCAAGGT
2. GTTAACCTGGTGTCTGGTACCG
3. TAACTTCTACGCTTGGAAACGTAT
4. TTCCATACGTTTCCAAGCGTAGAA
5. GGAAGTTGGTCAACAAGCAGTTGAAGT
6. CCAAACCTCAACTGCTTGTGACCAAC
7. TTGGCAGGGTCTGGCACTGCTGAGCG
8. GCCTCGCTCAGCAGTGCCAGACCCTG
9. AGGCTGTACTGCGTGGCCAGGCA
10. GCAGTGCCTGGCCACGCAGTACA
11. CTGCTGGTAAACTCCTCTCAGCCGT
12. TTCCCACGGCTGAGAGGAGTTACCA
13. GGAACCGCTGCAGCTGCATGTTGAC
14. GCTTTGTCAACATGCAGCTGCAGCGG
15. AAAGCAGTATCTGGCCTGAGATCTG
16. GATCCAGATCTCAGGCCAGATACT

FIG. 12

ECEPO\_SECTION\_2

ECORI KpnI 1  
A ATTGGTACC AGAC<sup>1</sup>CCAAG GTTAACTTCT ACCT<sup>3</sup>TGGAA ACGTAT<sup>2</sup>CGAA  
GCCATGG TCTG<sup>2</sup>TGGTTC CAATT<sup>4</sup>GAAGA TGGCAACCTT TGCATACCTT<sup>4</sup>

<sup>5</sup> GTTGGTCAAC AAGCAGTTGA AGT<sup>7</sup>TGGCAG <sup>7</sup>GGTCTGGCAC TGC<sup>8</sup>TGAGCCG  
CAACCAGTTG TTCGTCAACT TCAAC<sup>6</sup>CTC CCAGACCGTG ACGACTCGCT<sup>8</sup>

GGCTGTACTG CGTGGCCAGG C<sup>9</sup>ATGGCTGGT <sup>11</sup>AACTCCTCT CAGCCGTGGG  
CCG<sup>10</sup>CATGAC GCACCGGTCC GTGAC<sup>12</sup>ACCA TTGAGGAGA GTCGGCACCC<sup>12</sup>

<sup>13</sup> AACCGTCCA GCTGCATGT GACAAGCCAG TATC<sup>15</sup>TGGCT GAGAT<sup>16</sup>CTG  
T<sup>14</sup>GGCGACGT CGACGTACAA CTGTT<sup>14</sup>CTG<sup>14</sup> ATAGACCGGA CTCTAGACCTAC

BglII

BamHI

FIG. 13



ECEPO SECTION 3

1. GATCCAGATCTCTGACTACTCTGC
2. ACGCAGCAGAGTAGTCAGAGATCTG
3. TGCGTGCCTCGGGTGCACAGAAAGAGG
4. GATAGCCTCTTTCTGTGCACCCAGAGC
5. CTATCTCTCCGCCGGATGCTGCATCT
6. CAGCAGATGCAGCATCCGGCGGAGA
7. GCTGCACCGCTGCGTACCATCACTG
8. ATCAGCAGTGATGGTACGCAGCGGTG
9. CTGATACCTCCGCAAACTGTTTCG
10. ATACACGAAACAGTTTGCGGAAGGT
11. TGTATACTCTAACTTCCTGCGTGGTA
12. CAGTTTACCACGCAGGAAGTTAGAGT
13. AACTGAACTGTATACTGGCGAAGC
14. GGCATGCTTCGCCAGTATACAGTTT
15. ATGCCGTA CTGGTGACCGCTAATAG
16. TCGACTATTAGCGGTCACCAAGTAC

FIG. 14

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ECEPO SECTION 3

BamHI BglII  
 GA TCCAGATCTCTG  
 GTCTAGAGAC

<sup>1</sup> ACTACTCTGC <sup>2</sup> TGGGTGCTCT <sup>3</sup> GGGTGCACAG <sup>4</sup> AAAGAGGCTA <sup>5</sup> TCTCTCCGCC  
 TGATGAGACG <sup>2</sup> ACGCAAGAGA <sup>4</sup> CCCACGTGTC <sup>5</sup> TTTCTCCGAT <sup>5</sup> AGAGAGGCGG

GGATGCTGCA <sup>6</sup> TCTGCTGCAC <sup>7</sup> CGCTGCGTAC <sup>8</sup> CATCACTCT <sup>9</sup> GATACCTCC  
 CCTACGACGT <sup>6</sup> AGACGACCTG <sup>8</sup> GCGACGCATG <sup>9</sup> GTAGTGACCA <sup>9</sup> CTATGGAAGG

GCAAACTGTT <sup>10</sup> TCGTGTATAC <sup>11</sup> TCTAACTTCC <sup>12</sup> TCGGTGGTAA <sup>13</sup> ACTGAAACTG  
 CGTTTGACAA <sup>10</sup> AGCACATAATG <sup>12</sup> AGATTGAAGG <sup>13</sup> ACGCACCATT <sup>13</sup> TGACTTTGAC

TATACTGGCG <sup>14</sup> AAGCATGCCG <sup>15</sup> TACTGGTGAC <sup>16</sup> CGCTAATAG SalI  
 ATATGACCGC <sup>14</sup> TTCGTACGGC <sup>16</sup> ATGACCACTG <sup>16</sup> GCGATTATC <sup>16</sup> AGCT

FIG. 15

SCEPO SECTION 1 OLIGONUCLEOTIDES

1. AATCAAGCTTGGATAAAAGAGCT
2. GTGGAGCTCTTTTATCCAAGCTTG
3. CCACCAAGATTGATCTGTGACTC
4. TCTCGAGTCACAGATCAATCTTG
5. GAGAGTTTTGAAAGATACTTGTG
6. CTTCCAACAAGTATCTTTCCAAAAC
7. GAAGCTAAAGAAGCTGAAAACATC
8. GTGGTGATGTTTTCAGCTTCTTTAG
9. ACCACTGGTTGTGCTGAACACTGTTC
10. CAAAGAACAGTGTTCAGCACAACCA
11. TTTGAACGAAAACATTACGGTACCG
12. GATCCGGTACCGTAATGTTTTCGTT

FIG. 16

SCEPO SECTION 1

EcoRI HindIII 1  
AATTCA AGCTTGGATA  
GT TCGAACCTAT  
2

AAAGAGCTCC ACCAAGATTG ATCTGTGACT CAGAGTTTT  
TTTCTCGAGG TGGTTCTAAC TAGACACTGA GCTCTCAAAA  
3  
4

GGAAAGATAC TTGTTGGAAG CTAAAGAAGC TGAAAACATC ACCACTGGTT  
CCTTTCTATG AACAACTTC GATTTCTTCG ACTTTTGTAG TGGTACCAA  
5 6 7 8

GTGCTGAACA CTGTTCTTIG AACGAAAACA TTACGGTACC G  
CACGACTTGT GACAAGAAAC TTGCTTTTGT AATGCCATGG CCTAG  
9 10 11 KpnI BamHI  
12

FIG. 17

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SCEPD SECTION 2 OLIGONUCLEOTIDES

1. AATTCGGTACCAGACACCAAGGT
2. GTTAACCTTGGTGTCTGGTACCG
3. TAACTTCTACGCTTGGAAACGTAT
4. TTCCATACGTTTCCAAGCGTAGAA
5. GGAAGTTGGTCAACAAGCAGTTGAAGT
6. CCAAACCTCAACTGCTTGTGACCAAC
7. TTGGCAAGGTTTGGCCTTGTTATCTG
8. GCTTCAGATAACAAGGCCAAACCTTG
9. AAGCTGTTTTGAGAGGTCAAGCCT
10. AACAAGGCTTGACCTCTCAAACA
11. TGTTGGTTAACTCTTCTCAACCATGGG
12. TGGTTCCCATGGTTGAGAAGAGTTAACC
13. AACCATTGCAATTGCACGTCGAT
14. CTTTATCGACGTGCAATTGCAA
15. AAAGCCGTCTCTGGTTTGAATCTG
16. GATCCAGATCTCAAACCAGAGACGG

FIG. 18

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SCEPO SECTION 2

```

      KpnI
EcoRI 1
A ATTCGGTACC AGACACCAAG
      GCCATGG TCTGTGGTTC
      2

GTAACTTCT3 ACGCTTGGAA ACGTATGGAA5 GTTGGTCAAC AAGCTGTTGA
CAATTG4AAGA TCGGAACCTT TGCATACCTT CAACCCAGTTG6 TTCGACAAC

AGTTGGCAA7 GGTTGGCCT TGTATCTGA9 AGCTGTTT9AG AGAGGTCAAG
TCAAAC8CTT CCAAACCGGA ACAATAGACT TCG10CAAAAAC TCTCCAGTTC10

CCTTGTGGT11 TAACTCTTCT CAACCATGGG13 ACCATTGCA13 ATTGCACGTC
GGAACA12CCA ATTGAGAAGA GTTGGTACCC TTGGT14ACGT TAACGTGCAG14

GATAAAGCCG15 TCTCTGGTTT GAGATCTG BamHI
CTATTT16CGC AGAGACCAAA CTCTAGACCTA G

```

FIG. 19

SCEPO SECTION 3 OLIGONUCLEOTIDES

1. GATCCAGATCTTTGACTACTTTGTT
2. TCTCAACAAAGTAGTCAAAGATCTG
3. GAGAGCTTTGGGTGCTCAAAGGAAG
4. ATGGCTTCCTTTTGAGCACCCAAAGC
5. CCATTTCCCCACCAGACGCTGCTT
6. GCAGAAGCAGCGTCTGGTGGGGAA
7. CTGCCGCTCCATTGAGAACCATC
8. CAGTGATGGTTCTCAATGGAGCG
9. ACTGCTGATACCTTCAGAAAGTT
10. GAATAACTTTCTGAAGGTATCAG
11. ATTCAGAGTTTACTCCAACTTCT
12. CTCAAGAAGTTGGAGTAAACTCT
13. TGAGAGGTA AATTGAAGTTGTACAC
14. ACCGGTGTACA ACTTCAATTTACCT
15. CGGTGAAGCCTGTAGA ACTGGT
16. CTGTCACCAGTTCTACAGGCTTC
17. GACAGATAAGCCCGACTGATAA
18. GTTGTATCAGTCGGGCTTAT
19. CAACAGTGTAGATGTAACAAAG
20. TCGACTTTGTTACATCTACACT

FIG. 20

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SCEPO SECTION 3

BamHI BglII 1  
 GATC CAGATCCTTG ACTACTTTGT TGAGAGCTTT  
 GTCTAGAAAC TGATGAAACA ACTCTCGAAA  
2

3 5  
 GGGTGCTCAA AAGGAAGCCA TTTCCCACC AGACGCTGCT TCTGCCGCTC  
 CCCACGAGTT TTCCTTCGGT AAAGGGGTGG TCTGCGACGA AGACGGCGAG  
4 6

7 9 11  
 CATTGAGAAC CATCACTGCT GATACCTTCA GAAAGTTATT CAGAGTTTAC  
 GTAACCTTTG GTAGTGACCA CTATGGAAGT CTTTCAATAA GTCCTCAAATG  
8 10 12

13 15  
 TCCAACCTTCT TGAGAGGTAA ATTGAAGTTG TACACCGGTG AAGCCTGTAG  
 AGGTTGAAGA ACTCTCCATT TAACTTCAAC ATGTGGCCAC TTCGGACATC  
14 16

17 19  
 AACTGGTAC AGATAAGCCC GACTGATAAC AACAGTGTAG  
 TTGACCACTG TCATTTCGGG CTGACTATTG TTGTCACATC  
18

SalI  
 ATGTAACAAA G  
 TACATTGTTT CAGCT  
20

FIG. 21

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