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(estimated M.W. = 18,399). Also revealed in the Fable is 5 the DNA sequence coding for a 27 residue leader sequence 4 along with 5' and 3' DNA sequences which may be significant to promoter/operator functions of the human gene 5 operon. Sites for potential glycosylation of the mature human EPO polypeptide are designated in the Table by 19 asterisks. It is worthy, of note that the specific amino acid sequence of table VI likely constitutes that of a سولا naturally occurring allelic form of human erythropoietin. 10 Support for this position is found in the results of continued efforts at sequencing of urinary isolates of human erythropoietin which provided the finding that a significant number of erythropoietin molecules therin have a methionine at residue 126 as opposed to a serine as shown 15 in the table. ρ Table VII below, illustrates the extent of B polypeptide sequence homology between human and monkey EPO. In the upper continuous line of the table, single K. letter designations are employed to represent the deduced 20 translated polypeptide sequences of human EPO commencing _ / with residua =2 and the lower continuous line shows the deduced polypeptide sequence of mankey EPO commencing at assigned residue number -27. Asterisks are employed to highlight the sequence homologies. It should be noted 25 that the deduced human and monkey EPO sequences reveal an "additional" lysine (K) residue at (human) position 115. Cross-reference to $\frac{Figure 6}{100 \log 100}$ indicates that this residue K is at the margin of a putative mRNA splice junction in the genomic sequence. Presence of the lysine residue in 30 the human polypeptide sequence was further verified by sequencing of a cDNA human sequence clone prepared from mRNA isolated from COS-1 cells transformed with the human genomic DNA in Example 7, infra.

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Comparison of Human and Monkey EPO Polypeptides

120 150 160 ASPPDAASAAPLRIIADTFRKLFRVYSNFLRGKLKLYTGEACRIGDR A I SLPDA A SA A PL R I I TAD I FCKL FRYY SNFLRGKLKL Y I GE ACRRODR

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Human

Monkey

Monkey

Human

Monkey

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EXAMPLE 6

The expression system selected for initial attempts at microbial synthesis of isolatable quantities 5 of EPO polypeptide material coded for by the monkey cDNA provided by the procedures of Example 3 was one involving mammalian host cells (i.e., COS-1 cells, A.T.C.C. No. CRL-1650). The cells were transfected with a "shuttle" vector capable of autonomous replication in E.coli host 10 (by virtue of the presence of pBR322-derived DNA) and the mammalian hosts (by virtue of the presence of 5V40 virusderived DNA).

More specifically, an expression vector was constructed according to the following procedures. The 15 plasmid clone 83 provided in Example 3 was amplified in E.coli and the approximately 1.4kb monkey EPO-encoding DNA was isolated by EcoRI and HindIII digestion. Separately isolated was an approximately 4.0 kb, HindIII/SalI fragment from pBR322. An approximately 30 20 bo, EcoRI/SalI "linker" fragment was obtained from Ml3mplO RF DNA (P and L Laboratories). This linker included, in series, an EcoRI sticky end, followed by SstI, Smal, BamHI and Xbal recognition sites and a Sall sticky end. The above three fragments were ligated to 25 provide an approximately 5.4 kb intermediate plasmid ("pERS") wherein the EPO DNA was flanked on one side by a "bank" of useful restriction endonuclease recognition sites. pERS was then digested with HindIII and SalI to yield the EPO DNA and the EcoRI to SalI (Ml3mplO) linker. 30 The 1.4 kb fragment was ligated with an approximately 4.0 kb BamHI/SalI of pBR322 and another Ml3mplO HindIII/BamHI RF fragment linker also having approximately 30 bp. The Ml3 linker fragment was characterized by a HindIII sticky end, followed by PstI, SalI, XbaI recognition sites and a 35 BamHI sticky end. The ligation product was, again, a useful intermediate plasmid ("pBR-EPO") including the EPO DNA flanked on both sides by banks of restriction site.

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The vector chosen for expression of the EPO DNA in COS-1 cells ("pDSvL1") had previously been constructed to allow for selection and autonomous replication in E.coli. These characteristics are provided by the origin 5 of replication and Ampicillin resistance gene DNA sequences present in the region spanning nucleotides 2448 through 4362 of pBR322. This sequence was structurally modified by the addition of a linker providing a $\underbrace{\text{Hind}}_{}$ III recognition immediately adjacent nucleotide 2448 prior to 10 incorporation into the vector. Among the selected vec-· tor's other useful properties was the capacity to autonomously replicate in COS-1 cells and the presence of a viral promoter sequence functional in mammalian cells. These characteristics are provided by the origin of 15 replication DNA sequence and "late gene" viral promoter DNA sequence present in tre 342 bp sequence spanning nucleatide numbers 5171 through 270 of the SV40 genome. A unique restriction site (BamHI) was provided in the vector and immediately adjacent the viral promoter 20 sequence through use of a commercially available linker sequence (Collaborative Research). Also incorporated in the vector was a 237 base pair sequence (derived as nucleotide numbers 2553 through 2770 of SV40) containing the "late gene" viral mRNA polyadenylation signal 25 (commonly referred to as a transcription terminator). This fragment was positioned in the vector in the proper orientation vis-a-vis the "late gene" viral promoter via the unique BamHI site. Also present in the vector was another mammalian gene at a location not material to 30 potential transcription of a gene inserted at the unique BamHI site, between the viral promoter and terminator sequences. [The mammalian gene comprised an approximately 2,500 bo mouse dinydrofolate reductase (DHER) minigene isolated from plasmid pMG-1 as in Gasser, et al., 35 P.N.A.S. (U.S.A.), 79, pp. 6522-6526, [1982].] Again,

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the major operative components of plasmid pDSVL1 comprise nucleotides 2448 through 4362 of pBR322 along with nucleotides 5171 through 270 (342bp) and 2553 through 2770 (237bp) of SV40 DNA.

Maniatis, et al., supra, the EPO-encoding DNA was isolated from plasmid pBR-EPO as a BamHI fragment and
ligated into plasmid pDSVL1 cut with BamHI. Restriction
enzyme analysis was employed to confirm insertion of the
EPO gene in the correct orientation in two of the
resulting cloned vectors (duplicate vectors H and L).
See Figure 2, illustrating plasmid pDSVL-MKE. Vectors
with EPO genes in the wrong orientation were saved for
use as negative controls in transfection experiments
designed to determine EPO expression levels in hosts
transformed with vectors having EPO DNA in the correct
orientation.

Vectors H, L, F, X and G were combined with carrier DNA (mouse liver and spleen DNA) were employed to transfect duplicate 60mm plates by calcium phosphate microprecipitate methods. Duplicate 60 mm plates were also transfected with carrier DNA as a "mock" transformation negative control. After five days all culture media were tested for the presence of polypeptides possessing the immunological properties of naturally-occurring EPO.

EXAMPLE 7

30 A. Initial EPO Expression System
Involving CCS-1 Cells

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The system selected for initial attempts at microbial synthesis of isolatable quantities of human EPO polypeptide material coded for by the human genomic DNA EPO clone, also involved expression in mammalian host cells (i.e., CCS-1 cells, A.T.C.C. No. CRL-1650). The

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human EPO gene was first sub-cloned into a ™shuttle™ vector which is capable of autonomous replication in both E.coli hosts (by virtue of the presence of pBR322 derived DNA) and in the mammalian cell line COS- (by virtue of 5 the presence of SV4O virus derived DNA). The shuttle vector, containing the EPO gene, was then transfected into COS-1 cells. EPO polypeptide material was produced in the transfected cells and secreted into the cell culture media.

More specifically, an expression vector was constructed according to the following procedures. isolated from lambda clone $\lambda h E 1$, containing the human genomic EPO gene, was digested with $\underline{\mathsf{Bam}}\mathsf{HI}$ and $\underline{\mathsf{Hind}}\mathsf{III}$ restriction endonucleases, and a 5.6 Kb DNA fragment 15 known to contain the entire EPO gene was isolated. This fragment was mixed and ligated with the bacterial plasmid pUC8 (Bethesda Research Laboratories, Inc.) which had been similarly digested, creating the intermediate plasmid "pUC8-HuE", plasmid convenient source of this 20 restriction fragment.

The vector chosen for expression of the EPO DNA in COS-1 cells (pSV452t) had previously been constructed. Plasmid p5V4SEt contained DNA sequences allowing selection and autonomous replication in E.coli. These charac-25 teristics are provided by the origin of replication and Ampicillin resistance gene DNA sequences present in the region spanning nucleotides 2448 through 4362 of the bacterial plasmid pBR322. This sequence was structurally modified by the addition of a linker providing a $\underbrace{\text{Hind}}_{\text{III}}$ 30 recognition site immediately adjacent to nucleotide 2448. Plasmid pSV4SEt was also capable of autonomous replication in COS-1 cells. This characteristic was provided by a 342 bp fragment containing the SV40 virus origin of replication (nucleotide numbers 5171 through 270). This 35 fragment had been modified by the addition of a linker providing an E coR1 recognition site adjacent to

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nucleotide 270 and a linker providing a Sall recognition site adjacent nucleotide 5171. A 1061 bp fragment of 5V40 was also present in this vector (nucleotide numbers 1711 through 2772 plus a linker providing a Sall recogni-5 tion site next to nucleotide number 2772). Within this fragment was an unique BamHI recognition sequence. In summary, plasmid pSV4SEt contained unique BamHI and HindIII recognition sites, allowing insertion of the human EPO gene, sequences allowing replication and selec-10 tion in E.coli, and sequences allowing replication in COS-1 cells.

In order to insert the EPO gene into pSV4SEt, plasmid puC8-HuE was digested with BamHl and HindIII restriction endonucleases and the 5.6 kb EPO encoding DNA 15 fragment isolated. pSV4SEt was also digested with BamHl and HindIII and the major 2513 bp fragment isolated (preserving all necessary functions). These fragments were mixed and ligated, creating the final vector "pSvgHuEPO". (See, Figure 3.) This vector was propa-20 gated in E.coli and vector DNA isolated. Restriction enzyme analysis was employed to confirm insertion of the EPO gene.

Plasmid pSVgHuEPO DNA was used to express human EPO polypeptide material in COS-1 cells. More specifi-25 cally, pSYgHuEPO DNA was combined with carrier DNA and transfected into triplicate 60 mm plates of COS-1 cells. As a control, carrier DNA alone was also transfected into COS-1 cells. Cell culture media were sampled five and seven days later and tested for the presence of polypep-30 tides possessing the immunological properties of naturally occurring human EPO.

B. Second EPO Expression System Involving COS-1 Cells

Still another system was designed to provide improved production of human EPO polypeptide material

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AM670088731 AM-ITC 00873398 coded by the human genomic DNA EPO clone in COS-1 cells (A.T.C.C. No. CRL-1650).

In the immediately preceding system, EPO was expressed in COS-1 cells using its own promoter which is 5 within the 5.6 Kb BamHI to HindIII restriction fragment. In the following construction, the EPO gene is altered so that it is expressed using the SV40 late promoter.

More specifically, the cloned 5.6 Kb BamHI to HindIII genomic human EPO restriction fragment was 10 modified by the following procedures. Plasmid pUC8-HuE, as described above, was cleaved with BamHI and with BstEII restriction endonucleases. BstEII cleaves within the 5.6 Kb EPO gene at a position which is 44 base pairs 5' to the initiating ATG coding for the pre-peptide and 1 15 approximately 680 base pairs 3' to the HindIII restriction site. The approximately 4900 base pair fragment was isolated. A synthetic linker DNA fragment, containing Sall and BstEII sticky ends and an internal BamHI recognition site was synthesized and purificu. The two 20 fragments were mixed and ligated with plasmid pBR322 which had been cut with Sal and BamHI to produce the intermediate plasmid pBRgHE. The genomic human EPO gene can be isolated therefrom as a 4900 base pair $\underline{\mathsf{Bam}}\mathsf{HI}$ digestion fragment carrying the complete structural gene - 25 with a single ATG 44 base pairs 3' to BamHI site adjacent the amino terminal coding region.

This fragment was isolated and inserted as a BamHI fragment into BamHI cleaved expression vector plasmid pDSVLl (described in Example 6). The resulting 30 plasmid, pSVLgHuEPO, as illustrated in Figure 4, was used to express EPO polypeptide material from COS-1 cells, as described in Examples 6 and 7A.

EXAMPLE B

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Culture media from growth of the six transfected COS-1 cultures of Example 6 were analyzed by radioim-

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munoassay according to the procedures set forth in Example 2, Part B. Each sample was assayed at 250, 125. 50, and 25 microliter aliquot levels. Supernatants from growth of cells mock transfected or transfected with vec-5 tors having incorrect EPO gene orientation were unambiguously negative for EPO immunoreactivity. For each sample of the two supernatants derived from growth of COS-1 cells transfected with vectors (H and L) having the EPO DNA in the correct orientation, the % inhibition of 10^{-125} I-EPO binding to antibody ranged from 72 to 88%, which places all values at the top of the standard curve. The exact concentration of EPO in the culture supernatant could not then reliably be estimated. A quite conservative estimate of 300 mU/ml was made, however, from the 15 value calculation of the largest aliquot size (250 microliter).

A representative culture fluid according to Example 6 and five and seven day culture fluids obtained according to Example 7A were tested in the RIA in order 20 to compare activity of recombinant monkey and human EPO materials to a naturally-occurring human EPO standard and the results are set out in graphic form in Figure 1. Briefly, the results expectedly revealed that the recombinant monkey EPO significantly competed for anti-human 25 EPO antibody although it was not able to completely inhibit binding under the test conditions. The maximum percent inhibition values for recombinant human EPO. however, closely approximated those of the human EPO standard. The parallel nature of the cose response 30 curves suggests immunological identity of the sequences (epitopes) in common. Prior estimates of monkey EPO in culture fluids were re-evaluated at these higher dilution levels and were found to range from 2.91 to 3.12 U/ml. Estimated human EPO production levels were correspon-35 dingly set at 392 mU/ml for the five-day growth sample

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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.

5 EXAMPLE 9

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., 1410 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 recom-15 binant 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, op. 1065-1067 (1961) and Hammonc, et . al., Ann.N.Y.Acad.Sci., 149, pp. 516-527 (1968) and acti-20 vity levels ranged from 0.94 to 1.24 U/ml.

EXAMPLE 10

In the previous examples, recombinant monkey or 25 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 30 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 35 ovary (CHO) DHFRT cells and the selectable marker, DHFR. [For discussion of related expression systems, see

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