## Properties of human testis-specific lactate dehydrogenase expressed from Escherichia coli

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The cDNA encoding the  $C_4$  isoenzyme of lactate dehydrogenase (LDH- $C_4$ ) was engineered for expression in *Escherichia* coli. The Ldh-c open reading frame was constructed as a cassette for production of the native protein. The modified Ldh-c cDNA was subcloned into the prokaryotic expression vector pKK223-3. Transformed E. coli cells were grown to mid-exponential phase, and induced with isopropyl  $\theta$  p thiogalactopyranoside for positive regulation of the tac promoter.  $\frac{1}{2}$  induced the 35 kDa subunity which spontaneously formed the positive 140 kDa tetramer. Induced cells expressed the 35 kDa subunit, which spontaneously formed the enzymically active 140 kDa tetramer. Human LDH-C<sub>4</sub> was purified over 200-fold from litre cultures of cells by AMP and oxamate affinity chromatography to a specific activity of 106 units/mg. The enzyme was inhibited by pyruvate concentrations above 0.3 mm, had a  $K_m$  for pyruvate of 0.03 mm, a turnover number (nmol of NADH oxidized/mol of  $LDH-C_4$  per min at 25 °C) of 14000 and was heat-stable.

#### INTRODUCTION

Testis-specific lactate dehydrogenase (LDH-C4) from the resus-specific factate denyarogenase  $(LDH-C_4)$  from the mouse has been purified, crystallized and characterized biochemically (reviewed in Goldberg, 1987a). Different kinetic properties of the testicular and somatic isoenzymes of LDH have been attributed to amino acid sequence differences, especially in the highly conserved coenzyme-binding domain (Musick  $\&$ Rossmann, 1979). The putative amino acid sequence of human LDH-C<sub>4</sub> deduced from a cDNA clone (Millan et al., 1987) was found to be dissimilar to active-centre residues of mouse LDH- $C<sub>4</sub>$  (Li et al., 1983), implying that generalization concerning the biochemical properties of the testis-specific isoenzymes might not be applicable. In order to obtain sufficient quantities of human LDH- $C_4$  for further investigation, and because of the impracticality of purifying the protein from healthy human testes, we have used recombinant DNA technologies to provide a source of human LDH- $C_4$ .

The present paper describes the engineering of the full-length human Ldh-c cDNA clone for over-expression of human LDH- $C_4$  from bacteria. The human protein was produced as a catalytically active tetramer of 140 kDa and purified to homogeneity by two-step affinity chromatography. The kinetics of pyruvate reduction by human LDH-C<sub>4</sub> from E. coli were compared with those for mouse LDH- $C_4$  as well as with those for the somatic isoenzymes of LDH. In addition, we have demonstrated that human LDH- $C_4$  has heat-stability properties similar to those of the mouse isoenzyme (Hawtrey & Goldberg, 1970).

#### MATERIALS AND METHODS

#### Engineering of human  $Ldh-c$  cDNA for expression

The human Ldh-c cDNA was cloned by screening a  $\lambda$ gtll human testis cDNA expression library with rabbit antisera and monoclonal antibodies against mouse LDH-C<sub>4</sub> (Millan et al., 1987). The entire 1523 bp human  $Ldh-c$  cDNA fragment was isolated from the  $\lambda$  clone by a partial *EcoRI* digestion, separated by agarose-gel electrophoresis, transferred to a DEAE-cellulose

membrane (Schleicher and Schuell), recovered by high-salt elution and purification and purification (Dretzen et al., 1991) and precipitation (Dretzen et al., 1991) and purification (Dretzen et al., 1991) and purification (Dretzen et al., 1991) and purification (Dretzen et al., 19 elution and purified by ethanol precipitation (Dretzen et al., 1981). A 1250 ng portion of the 1.5 kbp fragment was ligated into 500 ng of a 2.9 kbp vector (a 5-fold molar excess of fragment over vector), an EcoRI digested and phosphatase-treated pGEM-3 plasmid (Promega). The HB101 strain of  $E$ . coli was transformed with 100 ng of the ligation mixture, and colonies containing recombinant plasmids were selected on Luria-Bertani (LB) medium plates with 50  $\mu$ g of ampicillin/ml. Plasmid DNA from the transformants was isolated by the small-scale alkalinelysis method, digested with restriction enzymes, electrophoresed through a  $1\%$  agarose gel containing 0.5  $\mu$ g of ethidium bromide/ml and detected by photographing the gel under u.v. light. Three out of nine transformants contained the insert in the correct orientation. One was prepared by the large-scale alkalinelysis method, purified by centrifugation through a CsCl equilibrium gradient, desalted by ethanol precipitation and designated pG3HC 1.5. Molecular-biology procedures were adapted from standard protocols (Maniatis et al., 1982), and enzymes were purchased from New England Biolabs.

Most of the  $Ldh-c$  open reading frame (ORF) is contained within a 1390 bp HincII fragment. Digestion with HincII removed the 5' untranslated sequences and the first 5 bp of the ORF. A  $5'$ linker was constructed with a *Smal* site adjacent to the first 5 bp of the ORF by synthesizing two oligomers with the following  $C<sub>1</sub>$   $C<sub>2</sub>$   $C<sub>3</sub>$   $C<sub>4</sub>$   $C<sub>5</sub>$   $C<sub>6</sub>$ 

# $CCGGGGATGTC$

Oligonucleotides were synthesized by the phosphoramidite method on an Applied Biosystems synthesizer, and purified on a Waters Delta-Pak  $C_{18}$  (30 nm-pore-size; 5  $\mu$ m-particle-size) column by reverse-phase h.p.l.c. The latter oligomer only was kinased to increase the chances of only one linker attaching and in the proper orientation. The two oligomers were boiled for 5 min, annealed by slow cooling to room temperature, and ligated with a 100:1 linker/Ldh-c fragment ratio. The regenerated

Abbreviations used: LDH, lactate dehydrogenase; ORF, open reading frame. Abbreviations used: LDH, lactate dehydrogenase; ORF, open reading frame.

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ORF was digested with DraI, which removed the AA of the stop codon TAA, and all of the 3' untranslated sequences. An oligomer was synthesized with the palindromic sequence:

#### AAACCCGGGTTT

treated with kinase, annealed to itself and ligated. Tandem repeats of the <sup>5</sup>' and <sup>3</sup>' linkers were removed by digestion with XmaI. The Ldh-c ORF engineered with XmaI (SmaI) linkers was subcloned by the above methods into the *XmaI* site of pGEM-4 (Promega) to construct pHumLdh-c ORF (Fig. 1).



#### Fig. 1. Genetic engineering scheme for construction of pHumLdh-c ORF and pKKHC4

The black bar denotes the human Ldh-c cDNA contained within the Agtll vector (Millan et al., 1987). Both the pGEM-4 and pKK223-3 vectors are derived from pBR332, a multiple-copy-number plasmid<br>ectors are derived from pBR332, a multiple-copy-number plasmid  $\mu$ K233-3 allows expression of human LDH-C-by induction of the  $\frac{1}{2}$  by multipliers and  $\frac{1}{2}$  by multipliers of the  $\frac{1}{2}$  conserved hybrid *tac* promoter. The *tac* promoter consists of the  $-35$  conserved TTGACA sequence from the *trp* promoter, and the  $-10$  conserved TOACA sequence from the *trp* promoter, and the  $-10$  conserved<br> $T_{A}T_{A}A$ . The sequence, the operator and the Shine-Dalgarno sequence comprising the ribosome-binding site from the lac UV-5 promoter comprising the ribosome-binding site from the lac UV-5 promoter (de Boer et al., 1983; Amann et al., 1983). The human  $Ldh-c$  ORF was inserted asymmetrically into the EcoRI and Small sites of the<br>M13mp8 polylinker (Vicire & Messing, 1982), which is followed by M13mp8 polylinker (Vieira & Messing, 1982), which is followed by the strong  $rrnB$  transcription terminators (Brosius *et al.*, 1981*a*,*b*). See the Materials and methods section for cloning details.

The Ldh-c ORF was modified for subcloning into the plasmid expression vector pKK223-3 (Pharmacia). The HinclI fragment containing all but the first <sup>5</sup> bp of the Ldh-c ORF was isolated from pHumLdh-c ORF. However, the <sup>5</sup>' and <sup>3</sup>' untranslated sequences were already removed, and the stop codon and <sup>3</sup>' SmaI remained intact since HincII cuts downstream of SmaI in the multiple cloning site of pGEM-4. The  $Ldh-c$  natural  $EcoRI$ site was methylated with EcoRI methylase. Two oligomers were synthesized with the complementary sequences:

GAATTCATGTC CTTAAGTACAG

and purified. Again, the latter oligomer only was treated with kinase, and the two oligomers were annealed and ligated with a 100: <sup>1</sup> linker/Ldh-c fragment ratio. Tandem repeats of <sup>5</sup>' and <sup>3</sup>' linkers were removed by digestion with EcoRI and XmaI. The Ldh-c ORF fragment was subcloned by the above methods into an EcoRI- and XmaI-digested pKK223-3 vector and designated pKKHC4 (Fig. 1).

#### Over-expression and purification of human  $LDH-C<sub>4</sub>$

Cell growth and induction. The W3110  $lac I<sup>q</sup>$  strain of E. coli cells transformed with pKKHC4 were grown in 1-litre volumes of Luria-Bertani medium with ampicillin (50  $\mu$ g/ml) at 37 °C. Cells were grown to mid-exponential phase  $(A_{600} = 0.5)$  and induced with 1 mm-isopropyl  $\beta$ -D-thiogalactopyranoside. For negative controls, parallel cultures were grown of transformed cells that were not induced and of non-transformed host cells that were not induced. All cultures were incubated for an additional 2 h beyond mid-exponential phase. Cells were harvested at 5000 g for 30 min at 4  $^{\circ}$ C, and pellets were frozen at  $-20$  °C.

Cell lysis and protein fractionation. Frozen pellets were combined from 20-litre cultures (60 g wet wt. of cells), thawed quickly at 37 °C, resuspended in 200 ml of 0.1 M-sodium phosphate buffer, pH 7.0 containing 1 mm-EDTA and 1 mm-dithiothreitol and sonicated at 4 'C until the cells were lysed. The crude extract was adjusted to 33% saturated with solid  $(NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>$ (196 g/l), and after stirring for 1 h at  $4^{\circ}$ C the precipitated protein was removed by centrifugation at  $12100 g$  for 30 min at  $4^{\circ}$ C. The supernatant was adjusted to 55% saturation with  $(NH_4)_2SO_4$  (351 g/l) and stirred overnight at 4 °C. The precipitate was collected by centrifugation at  $12100 g$  for 30 min at 4 °C, and resuspended in a minimum volume of 0.1 M-sodium phosphate buffer, pH 7.0, containing <sup>1</sup> mM-EDTA and <sup>1</sup> mM-dithiothreitol. The protein was then heated for 15 min at 55  $\degree$ C, cooled and centrifuged at  $12100 g$  for 30 min, and dialysed against 0.1 M-sodium phosphate buffer, pH 7.0, containing 1 mM-EDTA and <sup>1</sup> mM-dithiothreitol at 4 °C.

AMP-agarose chromatography. The dialysed cell extract was loaded at a flow rate of 1 ml/min on to a 40 ml volume column of 5'-AMP-agarose (Pharmacia) equilibrated with 0.1 M-sodium phosphate buffer, pH 7.0, containing <sup>1</sup> mM-EDTA and <sup>1</sup> mmdithiothreitol, and washed through with the same buffer until the  $A_{280}$  of the eluate fell to baseline (Lee et al., 1982). LDH was  $\frac{1280}{2}$  of the cluster for the baseline (EQ et al., 1702). LDIT was pecifically eluted with a reduced NAD<sup>-1</sup>-pyruvate adduct (50 mg)<br>of NAD+ and 77 mg of sodium pyruvate in 100 ml of 0.1 M- $\mu$  ivel and  $\mu$  ing or sourcin pyravate in 100 in 01 0.1 M<sup>2</sup> odium phosphate builet, pri .o. Equate fractions with peak<br>DH activity were pooled, concentrated by ultrafiltration and LDH activity were pooled, concentrated by ultrafiltration and dialysed against 0.02 M-sodium phosphate buffer, pH 6.8, containing <sup>1</sup> mM-EDTA and <sup>1</sup> mM-dithiothreitol.

Oxamate-agarose chromatography. The dialysed protein was

loaded on to a 14 ml-volume oxamate-agarose (Sigma Chemical Co.) column equilibrated with 0.02 M-sodium phosphate buffer, pH 6.8, containing 0.2 M-NaCl and 200  $\mu$ M-NADH (O'Carra & Barry, 1972; Spielmann et al., 1973; O'Carra et al., 1974; Kolk et al., 1978) and washed with the same buffer until no more protein was detected by the Bio-Rad micro-assay.  $LDH-C<sub>a</sub>$  was  $\frac{1}{2}$  must use the contract of the  $\frac{1}{2}$  must be contracted with  $\frac{1}{2}$  must be contracted buffer, photo with 10 mm-pyruvale in 0.02 m-sourching phosphate building  $\frac{1}{2}$ . pH 6.8, containing 0.2 M-NaCl. The 10 ml fraction with the highest LDH activity was concentrated by ultrafiltration and dialysed against 0.1 M-sodium phosphate buffer, pH 7.0, containing 1 mM-EDTA and 1 mM-dithiothreitol.

Protein analysis. Each step in the protein purification was **a rotein anarysis.** Each step in the protein purineation was assayed for total protein (Bio-Rad standard or micro-assay) and LDH activity (see below). Catalytic activity of the over-expressed human LDH- $C_A$  was also demonstrated by PAGE on native 7.5% polyacrylamide gels stained for LDH activity (Goldberg, 1964). Degree of purity of human LDH-C subunits was shown by SDS/12%-PAGE gels stained with Coomassie Brilliant Blue (Laemmli, 1970).

## $\mathbf{C}$  studies routinely measured spectro- $\mathbf{C}$  measured spectro- $\mathbf{C}$

LDH enzyme activity was routinely measured spectrophotometrically by the decrease in absorbance at 340 nm due to NADH oxidation (Hawtrey & Goldberg, 1970) in 3 ml of 0.5 msodium phosphate buffer, pH 7.0, containing 0.15 mm-NADH and 0.25 mM-pyruvate at 25 °C. To detect specifically human  $L$ -LDH activity and not  $E$ . coli D-LDH activity, lactate oxidation was measured spectrophotometrically by the increase in absorbance at 340 nm due to  $NAD<sup>+</sup>$  reduction (Neilands, 1955) in 0.1 M-Tris/HCl buffer, pH 8.4, containing  $1 \text{ mM-}NAD^+$  and 24 mm-DL-lactate. One unit of enzyme activity is defined as the amount of LDH that oxidizes 1  $\mu$ mol of NADH/min. The  $K_m$ for pyruvate was calculated from a Lineweaver-Burk plot.

Heat-stability assays were performed by diluting the pure enzyme in  $0.1$  M-sodium phosphate buffer, pH 7.0, containing  $f(0.1 \%)$  BSA, incubating for 10 min at temperatures increasing from  $0^{\circ}$ C, to  $85^{\circ}$ C, cooling and measuring activity spectrophotometrically.

The turnover number is defined as the number of mol of NADH oxidized/mol of human LDH-C<sub>4</sub> per min at 25 °C. The activity was measured spectrophotometrically as above, and converted by using a molar absorption coefficient of  $6.2 \times 10^3$  M<sup>-1</sup>·cm<sup>-1</sup> for NADH (Horecker & Kornberg, 1948) and a molecular mass of 140000 Da for LDH- $C_4$ .

### $\text{SULIS}$

The human Ldh-c cDNA (1523 bp) contains 66 bp of 5' untranslated sequences, a 999 bp ORF including the stop codon and 130 bp of 3' untranslated sequences (Millan et al., 1987) followed by 328 bp of unidentified DNA, probably an artifact of the cDNA cloning. The cDNA was modified by removing the 5' untranslated sequences by digestion with *HincII*, and the  $3'$ untranslated sequences by digestion with DraI (Fig. 1). The 5' linker added an Smal site and replaced the start codon and half of the HincII site. Regeneration of the HincII site implied that the linker and start codon were engineered in the correct orientation and reading frame. The 3' linker replaced the stop codon, the DraI site and added an SmaI site. Regeneration of the DraI site implied that the linker was ligated on properly and the stop codon was reconstructed in the correct reading frame. Two of 18 clones contained the engineered Ldh-c cDNA ORF bounded by Smal linkers in pGEM-4. This construction was designated pHumLdh-c ORF.



Fig. 2. Demonstration of  $LDH-C<sub>4</sub>$  enzymic activity in E. coli

rallel cultures of W3110 lacT<sup>a</sup> cells transformed with pKKHC4 were grown to mid-exponential phase, and induced with isopropyl  $\beta$ -D-thiogalactopyranoside for expression of human LDH-C<sub>4</sub> (lane 1), or not induced (lane 2), or induced and blocked with antiserum to mouse LDH-C<sub>4</sub> (lane 3). Non-transformed W3110 lac $I<sup>n</sup>$  host cells were grown and not induced (lane 4). Cell lysates were analysed by PAGE on a native 7.5% polyacrylamide gel, stained for LDH activity with lactate as the substrate. Purified human  $LDH-C<sub>4</sub>$  (lane 5) was electrophoresed along with human testis extract (lane  $6$ ) and human testis extract blocked with antiserum to mouse  $LDH-C_4$ (lane 7) for controls. The testis extract (lanes  $6$  and  $7$ ) also contains LDH isoenzymes consisting of the A- and B-subunit-containing heterotetramers from the anode (bottom of gel) to the cathode as follows: LDH-B<sub>4</sub>, LDH-B<sub>3</sub>A, LDH-B<sub>2</sub>A<sub>2</sub>, LDH-C<sub>4</sub> (lane 6 only) and LDH-BA<sub>3</sub>. In addition, faintly staining sub-bands can be seen in these extracts.

The cDNA was then engineered for expression of human The cDNA was then engineered for expression of human LDH-C<sub>a</sub> from bacteria. The *HincII* fragment was isolated from pHumLdh-c ORF and a 5' linker was attached to add an  $EcoRI$ site, the start codon and half the HincII site (Fig. 1). Again, regeneration of the *HincII* site implied correct construction. Five out of 18 clones contained the Ldh-c ORF bounded by  $5'$  EcoRI and 3' Smal linkers in  $pKK223-3$ , and were designated  $pKKHCA$ .

E. coli cells transformed with pKKHC4 were grown to midexponential phase, induced with isopropyl  $\beta$ -D-thiogalactopyranoside and lysed by sonication. Synthesis of human  $LDH-C<sub>4</sub>$ in these cells was shown by non-denaturing PAGE. Native gels stained for LDH activity demonstrated that induced cells express LDH as an enzymically active tetramer (Fig. 2). The crude LDH- $C_4$  migrated as a diffuse zone of activity near the LDH- $C_4$  band in the human testis control. Non-induced cells and host cells containing no plasmid had no LDH activity, as expected, since E. coli do not contain L-LDH (Tarmy & Kaplan, 1968b; Everse & Kaplan, 1973). Therefore only cells transformed with the human Ldh-c sequence under control of the tac promoter, and induced with isopropyl  $\beta$ -D-galactopyranoside, expressed LDH-C<sub>4</sub>. When the cell lysate was incubated with rabbit anti-(mouse LDH-C<sub>4</sub>) serum before electrophoresis, migration of the LDH- $C_4$  was specifically blocked. The remaining activity was due to heterotetramers of LDH-A and -B subunits. The purified human LDH-C<sub>4</sub> co-migrated with the LDH-C<sub>4</sub> band from human testis.

Human LDH-C<sub>4</sub> was purified over 200-fold from bacteria with a final specific activity of 106 units/mg of protein, and a yield of 13 $\%$  (Table 1). Lactate oxidation, which was measured to detect specifically L-LDH activity, was consistently about one-quarter the pyruvate reduction activity, with a final specific activity of 26 units/mg (LeVan, 1989).<br>The degree of purity was analysed initially by SDS/PAGE

#### Table 1. Summary of human  $LDH-C<sub>4</sub>$  purification from E. coli

For experimental details see the Materials and methods section.



(Fig. 3). The crude lysate contained no predominant band around 35 kDa, which indicated that the subunit was not overexpressed as <sup>a</sup> major cellular protein. Thus excess LDH-C subunits had not denatured to form inclusion bodies. A band seen in the total cell lysate was absent from the run-through of the AMP-agarose column. The eluate from the AMP-agarose column contained a predominant band of the same mobility and a few contaminants. The eluate from the oxamate-agarose column demonstrated that the protein was purified to homogeneity, and this band migrated at 35 kDa, as expected for the LDH subunit. Subsequently, the first <sup>20</sup> N-terminal residues of the pure protein were sequenced. Serine, as the first amino acid residue, indicated that the initiation methionine residue was cleaved off.



#### Fig. 3. Purification of human  $LDH-C_4$  from E. coli

 $\frac{1}{\sqrt{1-\frac{1$ Litre cultures of world *incluent* cents, transformed with  $\mu$ **A-KIIC-thiography** were grown and induced with isopropyl  $\beta$ -D-thiogalactopyranoside for expression of human LDH-C<sub>4</sub>. Cells were harvested and lysed, protein was precipitated between 33% and 55% saturation with  $(NH_4)_2SO_4$ , heated at 55 °C for 15 min and dialysed. LDH-C<sub>4</sub> was purified by AMP-agarose and then oxamate-agarose affinity chromatography. Fractions of crude cell lysate (lane 1), AMP wash (lane 2), AMP-agarose eluate (lane 3) and oxamate-agarose eluate (lane 4) with LDH activity were electrophoresed on a SDS  $12\%$ polyacrylamide gel and proteins were detected by staining with Coomassie Blue. The arrow at the left denotes the position of the human LDH-C subunit, and the molecular masses of the protein markers (lane 5) are indicated on the right.



Fig. 4. Pyruvate inhibition of human  $LDH-C<sub>4</sub>$  enzymic activity

LDH enzyme activity was measured spectrophotometrically by the decrease in absorbance at <sup>340</sup> nm due to NADH oxidation in 0.05 M-sodium phosphate buffer, pH 7.0, containing 0.15 mm-NADH and increasing concentrations of pyruvate from 0.1 to 5.0 mm at 25 °C.

Maximal LDH-C<sub>4</sub> activity was achieved between 0.25 mm- and 0.30 mm-pyruvate (Fig. 4). LDH- $C_4$  showed substrate inhibition with increasing concentrations of pyruvate. At 5.0 mm-pyruvate, activity was 50% maximal. The  $K<sub>m</sub>$  for pyruvate was 0.03 mm. These optimal pyruvate concentration and  $K<sub>m</sub>$  values are comparable with the respective values of 0.2 mm and 0.053 mm for LDH-C<sub>4</sub> derived from human testes (Svasti & Viriyachai, 1975b), 0.62 mm and 0.05 mm for LDH- $C_4$  derived from human spermatozoa (Wilkinson & Withycombe, 1965), and 0.5 mm and 0.026 mm for LDH- $C_4$  from mouse testes (Goldberg, 1972). At 1 mm-pyruvate mouse LDH-C<sub>4</sub> activity was 58 $\%$  maximal (Hawtrey & Goldberg, 1970).

Human LDH-C<sub>4</sub> was stable at temperatures up to 55 °C, and activity subsequently decreased at higher temperatures (Fig. 5). LDH-C<sub>4</sub> maintained at 65 °C was stable for 10 min, with 70 % of the activity remaining after 1 h at 65  $\rm{°C}$  (results not shown).

The turnover number with pyruvate was calculated to be 14000 mol of NADH oxidized per mol of human LDH- $C_4$  per min. This is higher than the value (2980) for mouse  $LDH-C_4$ (Goldberg, 1972), yet lower than the values (ranging from 41 500 to 160000) for the somatic LDH isoenzymes (Pesce et al., 1967).



Fig. 5. Heat-stability of purified human LDH- $C_4$ 

Heat-stability assays were carried out by diluting the pure enzyme in 0.1 M-sodium phosphate buffer, pH 7.0, containing  $0.1 \%$  BSA, incubating for 10 min at temperatures increasing from  $0^{\circ}$ C to 85 °C, cooling, and measuring LDH activity spectrophotometrically in 0.05 M-sodium phosphate buffer, pH 7.0, containing 0.15 mM-NADH and 0.25 mM-pyruvate at 25 °C.

#### **DISCUSSION**

The human Ldh-c cDNA was engineered for expression of the catalytically active tetramer in  $E$ . *coli* in milligram quantities. Human LDH-C<sub>4</sub> is a 140 kDa protein consisting of four identical subunits of <sup>331</sup> amino acids residues each. A cDNA cassette containing only the LDH-C subunit coding sequences was subcloned into pKK223-3 because this vector allows expression of the native protein. To maximize expression, all the <sup>5</sup>' and <sup>3</sup>' non-coding sequences were removed, because they have the potential to interfere with the regulatory mechanisms of the host  $\epsilon$  including the Ldh-c coding sequence including the pression system. The  $Ldd - c$  coding sequence including the initiation codon ATG was inserted into a unique cloning site 10 bp past the Shine-Dalgarno sequence (Shine  $\&$  Dalgarno, 1975) for efficient ribosome-binding (Gold et al., 1981; Stormo<br>
1975) for efficient ribosome-binding (Gold et al., 1981; Stormo<br>
1981; Stormon from *et al.*, 1982; Kozak, 1983). Analysis of the mRNA sequence from the Shine-Dalgarno sequence through the ATG and the cDNA encoding the first 33 amino acid residues indicated no potential. for significant mRNA secondary structure (Tinoco et al., 1973). Although the human  $Ldh-c$  codon usage was not optimal for all amino acids in  $E.$  coli, the protein was synthesized.

pKK223-3 has been used with some success for over-production of prokaryotic and eukaryotic proteins in E. coli. However, replacement of the replication origin of  $pKK223-3$ derived from pBR322 with that of pUC9 produced chicken lysozyme at about 25% of total cellular protein, which was 10-fold more than achieved with the analogous construction in unmodified  $pKK233-3$  (Miki et al., 1987). Also, replacement of the ribosome-binding site of the tac promoter with a synthetic sequence containing greater complementarity to the 3' end of 16 S rRNA produced a fusion protein of  $\beta$ -galactosidase and polyoma T antigen 5-10-fold more than in control cells (Pallas et al., 1986). Thus it may be possible to increase the production of LDH-C<sub>4</sub> further with an improved expression vector.

Purification of human LDH-C<sub>4</sub> from E. coli is not complicated by the presence of an endogenous  $E.$  coli L-LDH. The primitive labile D-LDH contributes to a portion of the initial LDH activity measured spectrophotometrically. This is not detected on the native gel stained for oxidation of lactate because  $D-LDH$  is functionally a unidirectional enzyme that acts as a pyruvate reductase (Tarmy & Kaplan, 1968 $a,b$ ). Other prokaryotic dehydrogenases with the conserved subunit size of 35 kDa (Kaplan, 1965) would co-migrate with the LDH-C band on the SDS/ polyacrylamide denaturing gel, but with their different charges and stabilities are lost during the purification procedure. Also, isolation of  $LDH-C_4$  from bacteria is not complicated by the presence of the eukaryotic LDH- $A_4$  and LDH- $B_4$  somatic isoenzymes, as occurs with isolation from human testes (Svasti  $\&$ Viriyachai, 1975a). The yield of human LDH-C<sub>4</sub> purified from E. coli is 5 mg per 20 litres of cells, as compared with 4 mg of LDH-C<sub>4</sub> from 225 g of human testes (Svasti & Viriyachai, 1975a).

Human LDH- $C_4$  is heat-stable relative to the somatic LDH isoenzymes (Wilkinson & Withycombe, 1965; McKee et al., 1972), as is mouse LDH-C<sub>4</sub> (Hawtrey & Goldberg, 1970). Similarly to the LDH isoenzymes containing B subunits, human  $LDH-C<sub>4</sub>$  is inhibited by higher concentrations of pyruvate (Plagemann et al., 1960; Wilkinson & Withycombe, 1965; Latner et al., 1966). A turnover number intermediate between those of mouse  $LDH-C<sub>a</sub>$  and the somatic isoenzymes may be indicative of the greater similarity of the sequence of the coenzyme-binding site of human LDH-C (Millan et al., 1987) to the somatic isoenzymes than to the sequence of mouse LDH-C (Pan et al., 1983). Although the kinetic properties of  $LDH-C<sub>4</sub>$  do not suggest a function for the testis-specific isoenzyme, its expression and purification from bacteria has provided us with an abundant source of a human sperm antigen for testing as an immunocontraceptive vaccine. The testis isozyme is predominantly cytosolic; however, a portion of the enzyme is adsorbed on the sperm cell surface (Erickson et al., 1975; Storey & Kayne, 1977; Alvarez & Storey, 1984). Antibodies to  $LDH-C<sub>4</sub>$  do not crossreact with the somatic isoenzymes LDH- $A_4$  and LDH- $B_4$ (Goldberg, 1971; Liang et al., 1986). Because LDH-C, is restricted to the germinal epithelium of testes and to spermatozoa (Blanco & Zinkham, 1963; Goldberg, 1963; Goldberg & Hawtrey, 1967), it would be seen as a foreign protein in the female reproductive tract. These cell-specific properties have allowed  $LDH-C<sub>4</sub>$  to be used as a model for development of a contraceptive vaccine by eliciting an immune response directed against sperm (Goldberg, 1987b). The antigenic determinants can be mapped, and correlated with the three-dimensional structure derived from X-ray crystallography, as was done with<br>notice derived from X-ray crystallography, as was done with mouse LDH-C<sub>4</sub> (Wheat & Goldberg, 1985a,b; Wheat et al., 1985; Hogrefe et al., 1987; Hogrefe et al., 1989). Ultimately,  $h_{\text{tot}}$ , hogicic et al., 1707, hogicic et al., 1707). Ottiniately, antigens may be developed into an effective multivalent into an effective multivalent in  $\frac{1}{n}$ antigens may be developed into an effective multivalent immuno-<br>contraceptive vaccine.

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