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RESEARCH ARTICLE

An alternative purification method for human serum paraoxonase 1 and its interactions with anabolic compounds

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Abstract

In this study, an alternative purification method for human paraoxonase 1 (hPON1) enzyme was developed using two-step procedures, namely, ammonium sulfate precipitation and Sepharose-4B-L-tyrosine-3-aminophenantrene hydrophobic interaction chromatography. SDS-polyacrylamide gel electrophoresis of the enzyme indicates a single band with an apparent $M_{\rm W}$ of 43 kDa. The enzyme was purified 219-fold with a final specific activity of 4408 400 U/mg and a yield of 10%. Furthermore, we examined the *in vitro* effects of some anabolic compounds, such as zeranol, 17 β -estradiol, diethylstilbestrol, oxytocin, and trenbolone on the enzyme activity to understand the better inhibitory properties of these molecules. The five anabolic compounds dose dependently decreased the activity of hPON1 with inhibition constants in the millimolar–micromolar range. The results show that these compounds exhibit inhibitory effects on hPON1 at low concentrations with IC₅₀ values ranging from 0.064 to 16.900 μ M.

Keywords

Anabolic compounds, hydrophobic interaction chromatography, inhibition, purification

History

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Introduction

Paraoxonase 1 (PON1: EC 3.1.8.1) is a calcium-dependent serum esterase that is synthesized by the liver. In serum, it is closely associated with high-density lipoproteins^{1,2}. Paraoxonase hydrolyze organophosphate compounds which are widely used as insecticides and nerve gases. Therefore, it plays a major role in the detoxification of these compounds and other artificial substrates, so that it may alter significantly an individual's susceptibility to the toxicity of these chemicals. In addition, paraoxonase is involved in lipid metabolism, since this enzyme probably hydrolyzes multiple oxygenated forms of polyunsaturated fatty acids of low-density lipoproteins associated phospholipids. For this reason, paraoxonase can be defined as an antioxidant enzyme^{3,4}.

The use of anabolic steroids for growth promotion purposes in meat producing animals results in an improvement in muscle growth, more lean meat, and a higher feed efficiency. However, toxicological/epidemiological studies show that there are harmful effects to consumers; as a result, the public health is placed in risk. As a consequence, the use of anabolic steroids for fattening purposes has been banned in the European Union since 1986⁵. These anabolic agents are used for increasing the rate of weight gain, improving the feed efficiency, storing protein, and decreasing fatness^{6–8}. But, depending on the use of anabolic in animal feed, anabolic residues that may occur in meat and meat products present risks to human health⁹.

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Zeranol is a resorcyclic acid lactone and a synthetic oestrogenic derivative of the mycotoxin zearalenone, which is produced by Fusarium moulds. It is a weak estrogen and is currently used to improve feed conversion efficiency and promote growth rates in livestock production. It has been widely used since 1969 as a growth promoter in the USA to improve the fattening rates of cattle¹⁰.

Trenbolone acetate (TBA), a kind of 19-nortestesteron, is a synthetic steroid with anabolic properties^{11–13}. TBA decreases the rate of both protein synthesis and degradation, and when the rate of degradation is less than the rate of synthesis, muscle protein rate increases¹⁴.

Diethylstilbestrol (DES) is a synthetic estrogenic compound with carcinogenic and anabolic effects. Its most important effect is to improve the growth rate by increasing the quantity of digestible feed in livestock. As diethylstilbestrol is a carcinogenic compound, its use has been banned in animal production in European Union countries¹¹.

Estradiol (ES) is a sort of natural anabolic steroid hormone. It exists as 17α - and 17β -isomers. Estradiol- 17β has been widely applied in clinics for its most potent biological effects of the endogenous estrogens⁵, and has also been used to promote unisexualization, improve feed conversion efficiency, and increase the rate of weight gain in aquaculture because of its protein synthesis stimulation and sex reversal effects. However, chronic exposure of humans to ES- 17β through food chain can cause toxic effects as caused by other steroid hormones on public health, such as children precocious puberty, teratogenicity, and carcinogenesis. Ingestion of ES- 17β residues in treated aquatic animal tissues may be potentially hazardous to consumers. Therefore, monitoring the residual content of this compound in

fishery products is necessary for controlling the illegal use of such substances to ensure public health and trade contacts¹⁵.

In this study, we developed an alternative purification method for the purification of the hPON1 enzyme. Specifically, human serum PON1 was purified by two-step procedures using ammonium sulfate precipitation and Sepharose-4B-L-tyrosine-3-aminophenantrene hydrophobic interaction chromatography which was specifically designed to the retained N-terminal hydrophobic signal peptide for PON1 enzyme.

In recent years, anabolisants are used for increasing the rate of weight gain, improving the feed efficiency, storing protein, and decreasing fatness. But, depending on the use of anabolic in animal feed, residues which may occur in meat and meat products present human health risks. Thus, the determination of the effects of these compounds on human PON1 activity is vital. However, to our knowledge, no study is available on the *in vitro* effects of anabolic compounds on PON1 activity. In this study, we aimed to determine any possible effect of some anabolic compounds on pure hPON1 activity.

Materials and methods

The materials used include Sepharose 4B, L-tyrosine, 3-aminophenantrene, paraoxon, protein assay reagents, and chemicals for electrophoresis were obtained from Sigma Chem. Co. (St. Louis, MO). All other chemicals used were of analytical grade. The anabolic compounds were provided by the local pharmacy.

Ammonium sulfate precipitation

Human serum was isolated from fresh human blood taken to dry tube. The blood samples were centrifuged at $1500 \,\mathrm{rpm}$ for $15 \,\mathrm{min}$. and the serum was removed. First, serum paraoxonase was isolated by ammonium sulfate precipitation $(60-80\%)^{16}$. The precipitate was collected by centrifugation at $15\,000 \,\mathrm{rpm}$ for $20 \,\mathrm{min}$, and redissolved in $6.5 \,\mathrm{mL}\,100 \,\mathrm{mM}$ Tris–HCl buffer (pH 8.0).

Hydrophobic interaction chromatography

The pooled precipitate obtained from human serum by using ammonium sulfate precipitation was subjected to hydrophobic interaction chromatography. The final saline concentration of precipitate was adjusted to 1 M ammonium sulfate, prior to that it was loaded onto the hydrophobic column prepared from Sepharose 4B-L-tyrosine-3-aminophenantrene. The preparation of hydrophobic column is as follows. About 10% CNBr was

Figure 1. Schematic representation of the Sepharose-4B-L-tyrosine-3-aminophenantrene hydrophobic gel. L-Tyrosine by using saturated L-tyrosine solution in the same buffer was coupled to Sepharose-4B-L-tyrosine activated with CNBr. The functional group of L-tyrosine (-NH2) was covalently bound with Sepharose 4B by means of an amide bond. After that, L-tyrosine was attached to the activated gel as a spacer arm, and finally diazotized 3-aminophenantrene was clamped to the meta position of L-tyrosine molecule as ligand. In this way, Sepharose-4B-L-tyrosine-3-aminophenantrene hydrophobic interaction gel was obtained. The hydrophobic interaction chromatography column was equilibrated with 0.1 M Na₂HPO₄ buffer pH 8.00 including 1 M (NH₄)₂SO₄.

prepared in 1:1 dilution of Sepharose 4B and water. The mixture was titrated to pH 11 in an ice bath and maintained at that pH for 8-10 min. The reaction was stopped by filtering the gel on a Buchner funnel and washing with cold 0.1 M NaHCO₃ buffer pH 10. L-Tyrosine was coupled to Sepharose-4B-L-tyrosine which was activated with CNBr by using saturated L-tyrosine solution in the same buffer. The reaction was completed by stirring with a magnet for 90 min. In order to remove excess of L-tyrosine from the Sepharose-4B-L-tyrosine gel, the mixture was washed with distilled water. The hydrophobic gel was obtained by diazotization of 3-aminophenantrene and coupling of this compound to the Sepharose-4B-L-tyrosine. The pH was adjusted to 9.5 with 1 M NaOH and, after gentle stirring for 3 h at room temperature; the coupled red Sepharose derivative was washed with 1L of water and then 200 mL of 0.05 M Tris-sulfate buffer pH 7.5. The column was equilibrated with 0.1 M Na₂HPO₄ buffer pH 8.00 including 1 M ammonium sulfate. The paraoxonase was eluted with ammonium sulfate gradient using 0.1 M Na₂HPO₄ including 1 M ammonium sulfate buffer with and without ammonium sulfate pH 8.00. The purified hPON1 enzyme was stored in the presence of 2 mM CaCl₂ at +4 °C, in order to maintain activity.

Total protein determination

The absorbance at 280 nm was used to monitor the protein in the column effluents and ammonium sulfate precipitation. Quantitative protein determination was achieved by absorbance measurements at 595 nm according to Bradford¹⁷, with bovine serum albumin standard.

SDS polyacrylamide gel electrophoresis

SDS polyacrylamide gel electrophoresis was performed in order to verify the purified enzyme. It was carried out in 12 and 3% acrylamide concentrations, containing 0.1% SDS, for the running and stacking gel, respectively, according to Laemmli¹⁸.

Paraoxonase enzyme assay

Paraoxonase enzyme activity towards paraoxon was quantified spectrophotometrically by the method described by Gan et al. ¹⁹. The reaction was followed for 2 min at 37 °C by monitoring the appearance of *p*-nitrophenol at 412 nm in Biotek automated recording spectrophotometer (Biotek, Winooski, VT). A molar extinction coefficient (ε) of *p*-nitrophenol at pH 8.0 in 100 mM Tris-base buffer of 17 100 M⁻¹ cm⁻¹ was used for the calculation. PON1 activity (1 U/L) was defined as 1 µmol of *p*-nitrophenol formed per minute.

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Sepharose-4B L-tyrosine 3-aminophenantrene

In vitro inhibition kinetic studies

For the inhibition studies, in the presence of purified hPON1, different concentrations of anabolic compounds were added to the cuvette. Paraoxonase activity with anabolic compounds was assayed by following the hydration of paraoxon. Activity % values of paraoxonase for five different concentrations for each of the anabolic compounds were determined by regression analysis using the Microsoft Office 2000 Excel. Paraoxonase activity without an anabolic compound was accepted as 100% activity. The inhibitor concentration causing up to 50% inhibition (IC $_{50}$ values) on the hPON1 enzyme activity were determined from the graphs.

Result and discussion

Paraoxonase has been purified so far from different sources with different yields and purification folds^{19–24}. However, most of these previous studies either included many steps or had low purification yields. For these reasons, we focused in this study to develop a new and simpler chromatographic method for the purification of hPON1. In this study, a new strategy for the purification of the PON1 enzyme was developed. Human serum paraoxonase was purified by two sequential procedures, ammonium sulfate precipitation followed by hydrophobic interaction chromatography specifically designed for PON1 enzyme.

Subsequently, prior to loading onto hydrophobic interaction column; the precipitate was saturated with 1 M ammonium sulfate in order to improve its efficiency for binding to hydrophobic gel of the column. A new hydrophobic gel has been synthesized in order to reduce the number of the purification steps of paraoxonase enzyme. The hydrophobic gel was designated based

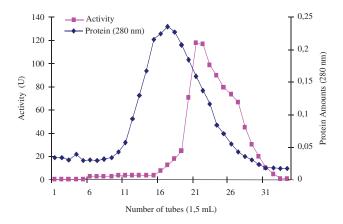


Figure 2. Elution graphic of PON1 with hydrophobic interaction chromatography. Purification of human serum PON1 by Sepharose 4B-L-tyrosine-3-aminophenantrene hydrophobic interaction chromatography with ammonium sulfate gradient. Fractions from the ammonium sulfate extraction were pooled as described in Methods section. This material was eluted by increasing the ammonium sulfate concentration. Protein concentration was determined by measuring an absorbance of 280 nm and PON1 activities of fractions were assayed activity using paraoxon substrate. 1 unit = 1 μ mol min⁻¹ per ml. U, units.

on the retained N-terminal hydrophobic signal peptide for PON1 enzyme. 3-Aminophenantrene, which is a hydrophobic group, was added to Sepharose-4B gel matrix with the extension of L-tyrosine arm (Figure 1).

Figure 2 shows the typical elution pattern of the enzyme activity on hydrophobic column. The enzyme activity and the total protein concentration were determined from all fractions collected from each of the purification steps. The fractions with the highest paraoxonase activity and the lowest protein contents, i.e., 21, 22m and 23 tubes, were pooled. Finally, PON1 was purified 219.14-fold. In another study, paraoxonase activity from pooled plasma of Q and R phenotypes shows quite variation 122.7 and 737 units, respectively 19. As seen in Table 1, each purification step yielded excellent results compared with the final specific activity and purification values reported for other purification procedures 20,24.

Different purification protocols have been used for PON enzyme from different sources. Furlong et al. ²⁰ reported 62.1-fold PON purification from human serum using four-step purification protocols, namely, Agarose Blue, Sephadex G-200, and DEAE—Trisacryl M Sephadex G-75. Sheep serum paraoxonase was purified in 330–385-fold using ethanol, pH, and ionic strength fractionation²⁵. Rodrigo et al. purified the liver paraoxonase in 415-fold by hydroxyapatite adsorption, chromatography on DEAE—Sepharose CL-6B, non-specific affinity chromatography

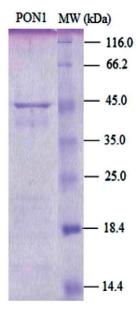


Figure 3. SDS-PAGE of human serum paraoxonase. The pooled fractions from ammonium sulfate precipitation and hydrophobic interaction chromatography were analyzed by SDS-PAGE (12 and 3%) and revealed by Coomassie Blue staining. Experimental conditions were as described in the method. Lane 2 contained 3 μ g of various molecular mass standards: β -galactosidase (116,0), bovine serum albumin (66.0), ovalbumin (45.0), carbonic anhydrase, (33,0), ∞ -lactoglobulin (25.0), lysozyme (19.5). Thirty microgram of purified hPON (lane 1) migrated with a mobility corresponding to an apparent M_r 43.0 kDa.

Table 1. Purification of human serum paraoxonase 1.

Purification step	Volume (mL)	Activity (U/mL.dak)	Total activity	Total protein (mg)	Specific activity (U/mg protein)	Purification fold	Yield %
Extract	18	512.98	9233.64	0.4590	20 116.86	_	100
Ammonium sulfate precipitation	6.5	633.21	4115.87	0.2990	13 765.45	0.68	45
Hydrophobic interaction chromatography	2	440.84	881.68	0.0002	44 08400	219.14	10

on Cibacron Blue 3 GA, and anion exchange on Mono Q HR 5/5. In addition, liver PON3 has been purified in 177-fold using a protocol consisting of seven steps²⁶. Colak and Gencer purified human PON1 enzyme using two-step procedures, namely ammonium sulfate precipitation and Sepharose-4B-L-tyrosine-9-aminophenantrene hydrophobic interaction chromatography. Overall purification rate was found 526-fold²⁷. In another work, the effect

Table 2. The IC_{50} values of anabolisant compounds.

Compounds	IC ₅₀ (μM)			
Oxytocin	16.900			
DES	0.066			
Trenbolone	0.127			
Estradiol-17β	0.064			
Zeranol	6.860			

of ammonium sulfate on the activities of Paraoxonase isoenzymes Q and R was researched. For this purpose, ammonium sulfate precipitation was performed before the Q and R isoenzymes. After ammonium sulfate precipitation, the specific activity of R isoform is 20.7 mU/mg. However, after ammonium sulfate precipitation, the specific activity of Q isoform is 6.6 mU/mg²⁸. Paraoxonase was purified shark Scyliorhinus canicula serum by Sayın et al. Purification of shark serum Paraoxonase was performed using the following methods: ammonium sulfate fractionation (60-80%) and Sepharose 4B-L-yrosine-1-naphtylamine hydrophobic interaction chromatography. A 37-fold purification with a yield of 1.127% was found²⁹. PON was purified and characterized from the Merino and Kivircik sheep's blood serums by a two-step procedure using ammonium sulfate precipitation and Sepharose-4B-L-tyrosine-1-napthylamine hydrophobic interaction chromatography for the first time by Erol et al. On SDS polyacyrilamide gel electrophoresis, purified human serum paraoxonase yielded a single band of 66 kDa on SDS-PAGE³⁰.

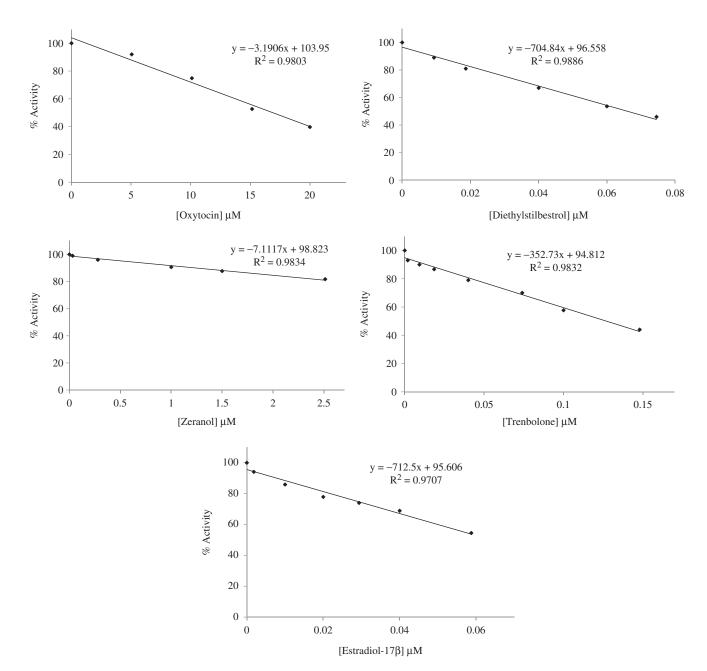


Figure 4. Inhibition graphics of anabolizan compounds.

Figure 3 illustrates the final purification patterns determined by SDS gel electrophoresis. The purified human serum A-esterase gives a single band on SDS-PAGE with a weight of 43 kDa. This corresponds to the previous studies ^{19,20,31}. Some purification studies indicate the differences of the migration of PON bands in SDS gel electrophoresis ¹⁹. Furlong et al. ²⁵ demonstrated two PON bands purified from rabbit serum. Gan et al. ¹⁹ report that human paraoxonase contains 15.8% carbohydrate. Sequence analysis ³² indicates five potential N-glycosylation sites in rabbit paraoxonase and four in humans. Therefore, a minimum molecular weight of human PON1 was 43 kDa ³². Moreover, a molecular weight of PON enzyme may increase up to 47–54 kDa in case of the contamination of albumin and ApoA1 ³³. However, our purification protocol yielded a single 43 kDa band suggesting the purification free from contaminants.

The kinetic parameters for the various anabolic compounds are presented in Table 2. The IC $_{50}$ values obtained for each of the anabolic compounds are significantly different (Figure 4). We have determined the IC $_{50}$ values of 0.064– $16.900\,\mu M$ for the inhibition of hPON1 activity. As it understood, these anabolic compounds were effective inhibitors on purified human serum PON1 activity. Paraoxon was used as a substrate at this work. Relatively studies have not reported on investigations of the inhibition of paraoxonase as substrate using paraoxon.

Many chemical species influence metabolism at low concentrations by decreasing or increasing the normal enzyme activity, especially by inhibiting enzymes with critical function³⁴, being thus drug targets³⁵. PON is important in the metabolism as prganophosphates (OP) hydrolyzer. OP are pesticides that inhibit cholinesterase. They cause poisonings and deaths^{36–38}. Paraoxonase, a member of the A-oxonase family, breaks down acetylcholinesterase inhibitors before they bind to the cholinesterases, and thus protects people from harmful effects caused by exposure to low doses of OP pesticides^{35,39}. Yet, it is estimated that worldwide 220 000 people are killed each year from such exposures. This is one reason why inhibitors of paraoxonase must be well investigated. PON is also a drug target 40-43. We have performed a number of studies regarding the interactions of different inhibitors with several such enzymes, including PON1^{44–50}. Sinan et al. showed that gentamycin sulfate and cefazolin sodium salt inhibited human serum PON1 dose and time dependently, with IC50 values of 0.887 and 0.0084 mM, respectively, but did not affect liver PON1 activity in human hepatoma HepG2 cells¹⁶. In another work, human serum paraoxonase (hPON1) was purified and the *in vitro* effects of commonly used antibiotics, namely clarithromycin and chloram phenicol, on purified human serum, paraoxonase enzyme activity (serum hPON1) and human hepatoma (HepG2) cellparaoxonase enzyme activity (liver hPON1) were determined. And they were determined to inhibit serum hPON1 and liver hPON1⁵¹. It was determined that commonly used antibiotics, namely sodium ampicillin, ciprofloxacin, and clindamycin phosphate, were effective inhibitors on human serum PON1⁵². Kıranoglu et al. showed that while mouse liver PON activity showed a statistically significant decrease for ethinyl estradiol in combination with desogestrel and levonorgestrel all three drugs, serum PON activity increased⁵³. In another work, in vitro inhibitory effects of oxytocin, dexamethasone, atropine sulfate, gentamicin sulfate, sulfadoxine-trimethoprim, furosemid, metamizole sodium, and toldimfos sodium were investigated. The IC50 values obtained varied markedly from 0.014 to 507.72 mg/mL. According to these findings, most potent and significant inhibition was displayed by dexamethasone, atropine sulfate, and furosemid⁵⁴.

However, it is thought that more extensive inhibition studies are necessary for a better understanding of the protective role of PONs against the toxic effects of xenobiotics, including environmental heavy metals and oxidative stress by-products^{16,55}. But, there are only few studies regarding effects of drugs on PON1 activity in the literature. Considering these, we report in the present study, the effects of some anabolic compounds against purified hPON1.

Declaration of interest

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