

A validated method to assess glutathione peroxidase enzyme activity

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Research Article

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A validated method to assess glutathione peroxidase enzyme activity

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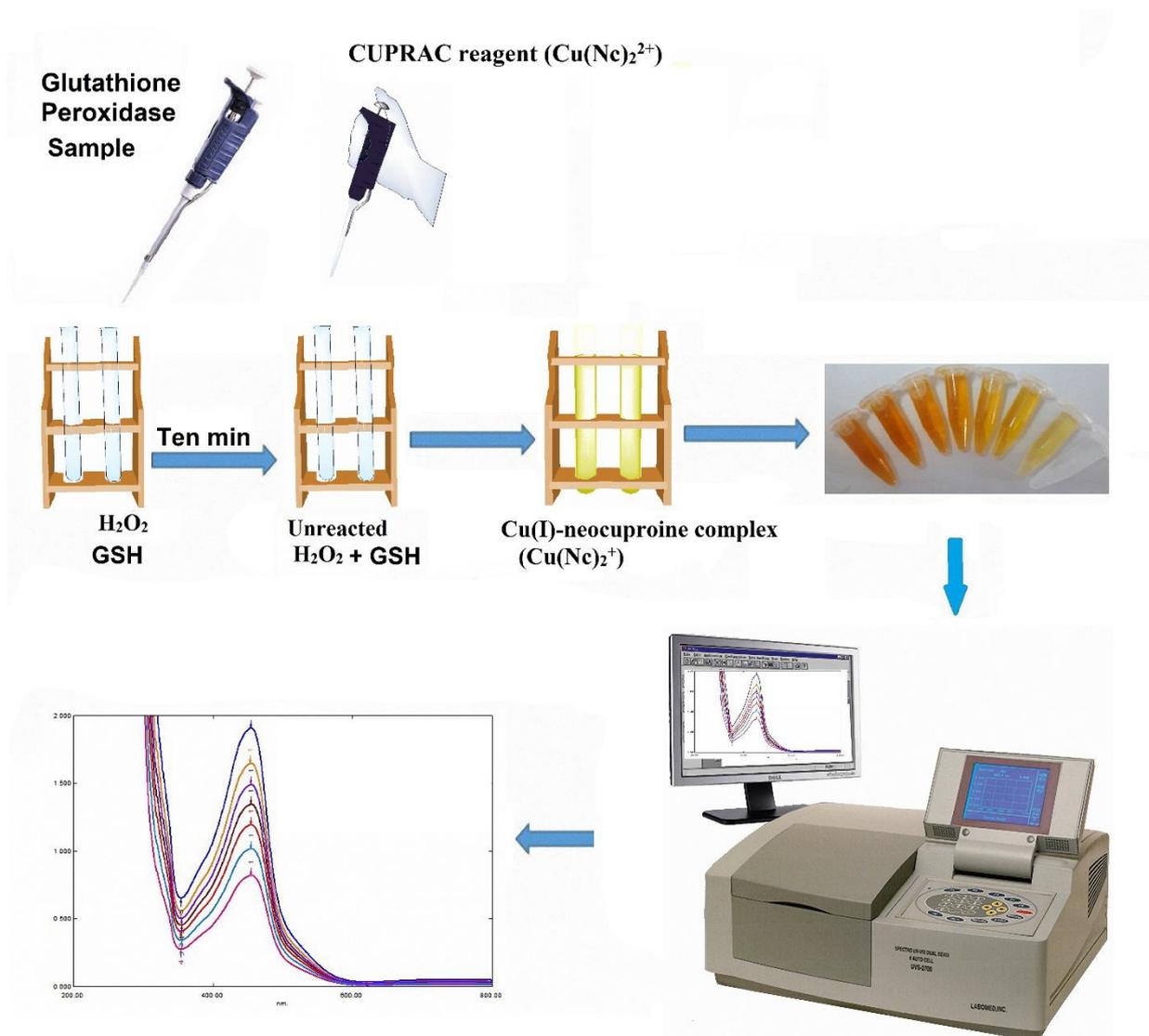
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Graphical abstract:



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Abstract:

This essay presents a reliable, effective and easy procedure for measuring glutathione peroxidase activity (Gpx). The enzyme samples were incubated with phosphate buffer, which included appropriate concentrations of glutathione and peroxide as substrates, to determine the Gpx activity. After a sufficient incubation time, the CUPRAC reagent ($\text{Cu}(\text{Nc})_2^{2+}$) was added to stop the enzyme's reaction. The unreacted substrates act to reduce $\text{Cu}(\text{II})$ -neocuproine complex ($\text{Cu}(\text{Nc})_2^{2+}$) to strongly coloured $\text{Cu}(\text{I})$ -neocuproine complex ($\text{Cu}(\text{Nc})_2^+$) that was measured spectrophotometrically at 450 nm (CUPRAC method). The glutathione peroxidase activity was linked to a decrease in the absorbance of the coloured $\text{Cu}(\text{I})$ -neocuproine complex ($\text{Cu}(\text{Nc})_2^+$).

The procedure uses the Box–Behnken design (BBD) to optimise the formation of the $\text{Cu}(\text{I})$ -neocuproine complex ($\text{Cu}(\text{Nc})_2^+$). The response surface methodology (RSM) is used to determine the accuracy of the method. This new protocol was confirmed by applying a Bland–Altman plot analysis of Gpx activity in matched samples using the Gpx-DTNB assay. The correlation coefficient between the two protocols was 0.9967. This means that the new protocol was very accurate and on par with the comparison method.

Keywords: Bland–Altman plot, Box–Behnken design, CUPRAC method, glutathione, response surface methodology.

Introduction

Glutathione peroxidases (Gpx) act to reduce hydroperoxides (ROOH) by glutathione (GSH):



R may be an aliphatic, aromatic, or hydrogen-containing organic group. H₂O, alcohol (ROH) (or a second H₂O when H₂O₂ is the substrate), and glutathione disulphide (GSSG) are the products. The enzyme glutathione reductase is responsible for regenerating GSH from GSSG in the cell [1, 2]. The glutathione peroxidase family (GPx1–8) catalyse the reduction of organic and inorganic peroxides by using reduced GSH. Proteins that contain selenocysteine make up five of the eight glutathione peroxidases (GPx1–4 and GPx6) [3]. The tendency of various GPxs to catalyse the degradation of hydroperoxides by thiols is their common denominator [4]. Gpx is a significant system to protect against endogenously and exogenously mediated lipid peroxidation that is present in many animal tissues. The enzyme is stoichiometric in selenium, and it reacts with several organic hydroperoxides as well as hydrogen peroxide [5].

Even though several protocols for assessing glutathione peroxidase activity have been established, only a few are still useful. To assess glutathione peroxidase activity in biological tissues, only two different test systems have been used. The first system [1, 6] was based on measuring ROOH or GSH consumption at regular interval. The second system monitors GSSG production by coupling to the glutathione reductase-catalysed reaction [1, 7]. The decrease in NADPH concentration is continuously measured spectrophotometrically or fluorometrically [6, 7].

Ellman's reagent (DTNB) is most commonly used in the first system to colourimetrically evaluate glutathione consumption as a function of glutathione peroxidase activity [6]. Compared to other tests, the GPx-DTNB assay is insensitive [8] and Ellman's reagent is relatively unstable [9]. Moreover, the process is time-consuming [10]. Flohé and Günzler [11], on the other hand, showed another polarographic method. It employed strong acid to stop enzyme-catalysed or spontaneous GSH-hydrogen peroxide reactions at a specified time (t). Polarography is then used to determine the GSH content.

Ugar et al. [12] recommended a microplate-based method that reduced the Cu(II)-neocuproine complex to highly coloured Cu(I)-neocuproine complex by using unreacted GSH. Catalase enzyme with high activity was used to stop the Gpx reaction. The absorbance decrement was correlated with Gpx activity. The method was suitable to assess Gpx activity in pure samples but not for assessing its activity in biological tissues because this does not take into account the interference arising from the presence of the catalase enzyme. Glutathione peroxidase and catalase act on hydrogen peroxide as a common substrate. All previous methods work to block interference with the catalase enzyme by adding sodium azide, which inhibits the catalase enzyme selectively.

Fluorescent methods occupy an important part of the second system of methods used to estimate the Gpx activity. Weiss et al. [13] documented a fluorometric method to measure Gpx activity in less than 100 µg of tissue. That assay depends on the fluorometric behaviour of NADP⁺ that arises from the oxidised glutathione created by the Gpx reaction. Martinez et al. [14] developed a fluorometric procedure with high sensitivity that used the assay of oxidised glutathione with o-phthalaldehyde. Kamata et al. [15] developed a sensitive method to assess Gpx activity using the fluorometric reaction of oxidised glutathione with N-(9-acridinyl) maleimide. The procedure was used to evaluate human plasma samples and liver homogenates.

Paglia and Valentino [7] were developed the most widely used protocol for measuring Gpx activity. It was based on the change of absorbance at 340 nm when NADPH is consumed by oxidised glutathione (GSSG). The protocol was modified by Lawrence and Burk [16] to study the activity of the Gpx enzyme in the liver supernatant of rats fed with a Se-deficient diet. The protocol was simple and selective, but the sensitivity was poor, and the enzyme and NADPH are costly [12]. Furthermore, NADPH + H⁺ seems to be a potent Gpx inhibitor [10]. Since proteins and DNA absorb UV light, the method cannot have precise results when calculating Gpx activity in biological tissues.

A simple method for determining Gpx enzyme activity is identified in this paper. Phosphate buffer was used to incubate the enzyme samples, which had appropriate concentrations of glutathione and peroxide as substrates. After appropriate incubation, the CUPRAC reagent (Cu(Nc)₂²⁺) was added to stop the enzyme's reaction. Unreacted substrates reduced the Cu(II)-neocuproine (Cu(Nc)₂²⁺) complex to highly coloured

Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex, which has a maximum absorbance at 450 nm (CUPRAC method). The Gpx activity was correlated inversely with the decrease of absorbance of coloured Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex.

The current protocol is precise, efficient, and trustworthy. The method is interference-free, simple to implement in laboratory experiments, and appropriate for clinical diagnosis.

Materials and Methods

Chemicals and materials

Ammonium acetate, calcium chloride, copper(II) chloride, dipotassium phosphate (K_2HPO_4), hydrochloric acid, sodium dihydrogen phosphate dihydrate ($\text{NaH}_2\text{PO}_4 \cdot 2\text{H}_2\text{O}$), hydrogen peroxide (30%), di-sodium hydrogen phosphate dihydrate ($\text{Na}_2\text{HPO}_4 \cdot 2\text{H}_2\text{O}$), sodium azide, sodium hydroxide, sodium nitrate, and trichloroacetic acid were purchased from BDH.

Bovine serum albumin, glutathione, homocysteine, neocuproine (2,9-dimethyl-1,10-phenanthroline) and tert-butyl hydroperoxide were purchased from Sigma-Aldrich.

Animals: The albino rats were obtained from Animal House, University of Babylon, Babylon governorate, Iraq. They were kept in well-ventilated cages with monitored light and humidity, as well as free access to regular food and water. The current study was conducted in accordance with the WSAVA Animal Welfare Recommendations [17].

Reagents

1. The phosphate buffer solution (100 mM M, pH 7.0) consisted of 1:1.5 volumes of A and B solutions. Solution A contained 13.62 g of KH_2PO_4 dissolved in 1 L of distilled water (D.W.). Solution B contained 17.8 g of $\text{Na}_2\text{HPO}_4 \cdot 2\text{H}_2\text{O}$ dissolved in 1 L of D.W. A 0.372 g EDTA and 0.6501 g of NaN_3 was added to the final solution and mixed gently.
2. Reduced glutathione (4 mM) was prepared by dissolving 0.1228 g of reduced glutathione in 100 ml of 100 mM phosphate buffer solution (pH 7.0).
3. Hydrogen peroxide H_2O_2 (2 mM) was prepared daily, in 100 mM phosphate buffer solution (pH 7.0). The final concentration was prepared by using a molar extinction coefficient of $43.6 \text{ M}^{-1} \text{ cm}^{-1}$ at 240 nm.

4. Copper(II) chloride (10^{-2} M) comprised 0.4262 g of $\text{CuCl}_2 \cdot 2\text{H}_2\text{O}$ dissolved in 250 ml of D.W.
5. Ammonium acetate buffer (NH_4Ac) (1.816M, pH 7.0) was composed of 35 g of NH_4Ac dissolved in 250 ml D.W.
6. Neocuproine (7.5×10^{-3} M OF 2,9-dimethyl-1,10-phenanthroline) (Nc) was composed 0.039 g Nc that was dissolved in 25 ml of 96% ethanol.
7. Fresh working reagent (CUPRAC reagent) was used for the experiment. The reagent was composed of Cu(II) :Nc: NH_4Ac at a ratio of 1:1:1 (v/v/v).

Instrument

A Shimadzu 1800 spectrophotometer was used to take the measurements in the current study.

Ethical agreement

Ethics Committee (University of Babylon/ College of Science/ Iraq), Ref. no.: 5114
Date: 1/10/ 2020.

Glutathione peroxidase purification

Rat's liver Gpx was purified, as described by Chafik et al. [18]. Gpx activity was measured using the GPx-DTNB assay [6]. The specific activity of Gpx was estimated to be about $1.2 \text{ U} \cdot \text{mg}^{-1}$ protein.

Tissue preparation

After the animals were sacrificed, the livers were washed thoroughly and rinsed with ice. They were weighted in an objective equilibrium after being carefully blotted between folds of filter paper. A polytron homogeniser was used to prepare 10% of the homogenate in 0.1 M phosphate buffer (pH 7.0). Unbroken cells, cell debris, nuclei, mitochondria, and erythrocytes were removed from the homogenate by centrifugation at 10,000 rpm for 20 minutes. The Gpx level was measured in the supernatant.

Detailed of procedure

The details of the procedure are shown in table 1.

Table 1. The details of the protocol used to measure glutathione peroxidase activity.

Reagents	Test	Control	STD	Blank
Sodium phosphate buffer	1500µL	1900µL	1600µL	2000µL
Reduced glutathione	200 µL	----	200 µL	-----
Sample	100 µL	100 µL	-----	-----
The reaction was initiated by adding peroxide:				
Peroxide	200 µL	-----	200 µL	-----
Mix by vortex and incubate for 10 minutes at 37°C, after that, the reaction was terminated with 0.5 ml of 8% TCA				
Mix well and centrifuge for 15 minutes at 3000 xg, then remove 1 ml of supernatant in a clean tube , and add:				
Working reagent	3ml	3ml	3ml	3ml

Absorbance was read against blank at 450 nm after 30 min.

Calculation

Unit definition: one unit of GPx was defined as the amount of enzyme capable of oxidising 1.0 µmole GSH to GSSG per minute at 25 °C, pH 7.0 [1, 6].

The residual glutathione concentration in test tube = $(\frac{A_{Test} - A_{Control}}{A_{STD}}) \times \text{Conc. of STD}$

The concentration of standard glutathione was 400 µM.

Also, the residual glutathione concentration in the test tube could be calculated from the glutathione standard curve (fig. 1). Gpx activity equals the number of micromoles of consumed glutathione.

Glutathione peroxidase activity (U/L) =

$$\left(\frac{\text{Conc. of GSH in STD} - \text{Conc. of residual GSH in test tube}}{\text{time (10min)}} \right) \times \left(\frac{\text{Total volume (ml)}}{\text{Volume of the sample (ml)}} \right) \times \text{D.f.}$$

Standard curve preparation

To create a standard curve for the assay, the stock solution of the standards (glutathione and peroxide) were diluted with the phosphate buffer (0.1 M, pH 7.0) according to the layout in table 2. Once each standard tube was created and mixed, 1000 µL of each was added to the three ml of working solution according to the protocol listed in table 1. Absorbance was read against blank at 450 nm after 30 min (as shown in fig. 1).

Table 2: Standard Curve Preparation

Standard	Volume of Glutathione 4mM	Final concentration of glutathione	Volume of peroxide 2mM	Final concentration of peroxide	Volume of phosphate buffer solution
S1	400 μ l	800 μ M	400 μ l	400 μ M	1200 μ l
S2	300 μ l	600 μ M	300 μ l	300 μ M	1400 μ l
S3	200 μ l	400 μ M	200 μ l	200 μ M	1600 μ l
S4	150 μ l	300 μ M	150 μ l	150 μ M	1700 μ l
S5	100 μ l	200 μ M	100 μ l	100 μ M	1800 μ l
S6	50 μ l	100 μ M	50 μ l	50 μ M	1900 μ l
S7	25 μ l	50 μ M	25 μ l	25 μ M	1950 μ l
S8	5 μ l	10 μ M	5 μ l	5 μ M	1990 μ l
B	-----	-----	-----	-----	2000 μ l

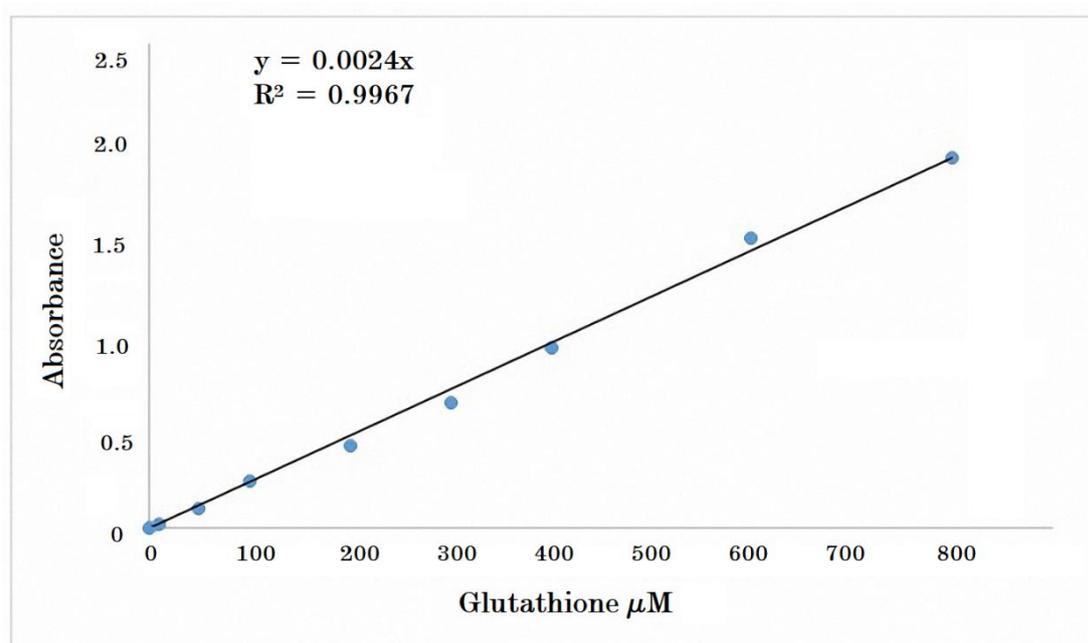


Fig. 1. The standard curve of glutathione obtained by using the current protocol.

Performance of the method

Performance of the current method was achieved according to the guideline on bioanalytical method validation from the Committee for Medicinal Products for Human Use [19].

Optimising the protocol

To optimise the Gpx-CUPRAC protocol, the BBD was used to apply RSM. To design the Gpx assay experiment and optimise the method, the statistical parameters were calculated using Chemoface software, Version 1.5 [20]. The enzymatic reaction was

performed with Gpx solution (500 U.L⁻¹), which was prepared fresh before the experiment by dissolving 417 mg of Gpx standard in 100 ml of 100 mM phosphate buffer solution (pH 7.0). The Gpx-DTNB procedure [6, 11] was used to adjust the final Gpx activity to 500 U.L⁻¹. The independent variables were glutathione, peroxide, and neocuproine concentrations (Table 3), and the dependent variable was the obtained Gpx activity compared to the current spectrophotometric protocol.

The mathematical modelling of a second-order polynomial equation was used to determine the interaction between the dependent and independent variables.

$$Y = \beta_0 + \sum \beta_i X_i + \sum \beta_{ii} X_i^2 + \sum \beta_{ij} X_i X_j + \varepsilon \quad (1)$$

where Y refers to the response variable, $X_i X_j$ refers to the independent variables, and β_0 , β_i , β_{ii} , and β_{ij} refer to the intercept, linear, quadratic and interaction coefficients, respectively. Random error is represented by the symbol ε .

Table 3. shows the results of using the Box–Behnken design to optimise the glutathione peroxidase activity assay. The independent variables were glutathione, peroxide, and neocuproine concentrations, and the dependent variable was glutathione peroxidase activity based on current spectrophotometric sensitivity.

Run	Glutathione Concentration mM	Peroxide Concentration mM	Neocuproine Concentration mM	Glutathione Peroxidase Activity (U/L)
1	2	1	5	401
2	2	1	10	473
3	2	3	5	482
4	2	3	10	482
5	6	1	5	429
6	6	1	10	464
7	6	3	5	491
8	6	3	10	508
9	0.636	2	7.5	446
10	7.364	2	7.5	508
11	4	0.318	7.5	410
12	4	3.682	7.5	517
13	4	2	3.296	446
14	4	2	11.705	500
15	4	2	7.5	500
16	4	2	7.5	497
17	4	2	7.5	502

Accuracy, selectivity, and reproducibility

Three kinds of interfering biochemicals were dissolved in four separate flasks to test the accuracy of the new Gpx procedure. The first included 100 mM phosphate buffer solution (pH 7.4), the second 5 mM sucrose, glucose, mannose, galactose, and ribose, the third 5 mM isoleucine, leucine, aspartic acid, methionine, and valine, and the fourth 3% casein and 3% bovine albumin. The phosphate buffer was used to dissolve all the interfering biomolecules. One ml of 3000 U/l Gpx enzyme was combined with 9 ml of the solutions containing possibly interfering biochemicals in an enzymatic reaction. Using the Gpx-DTNB method, the activity of Gpx was calibrated to 300 U/l. The Gpx-CUPRAC protocol recovery was calculated for each possible interfering biomolecule. The association between relative percentage errors and interfering biological substances is shown in Table 4. Several biological samples were used to test the current method's inter-and intra-day reproducibility, and the results were shown using RSD.

Signal stability

A Gpx solution (150 U.L⁻¹) was used to study the stabilisation of the coloured chelate complex. After adding the working solution, the absorbance reading at 450 nm was measured after 15 minutes, 30 minutes, 45 minutes, 60 minutes, 5 hours, one day, three days and one week.

Sensitivity and linearity

The linearity and sensitivity of the protocol were tested using a range of Gpx activities (0, 5, 10, 25, 50, 100, 200, 400, 600, 700 and 1000 U.L⁻¹). The linearity was assessed by comparing it to the Gpx-DTNB method [6, 11] using a web-based program for bias assessment and comparison of analytical methods [21]. Limits of quantitation (LOW) and detection (LOD) were used to calculate the sensitivity of the Gpx-CUPRAC assay [22].

Validation

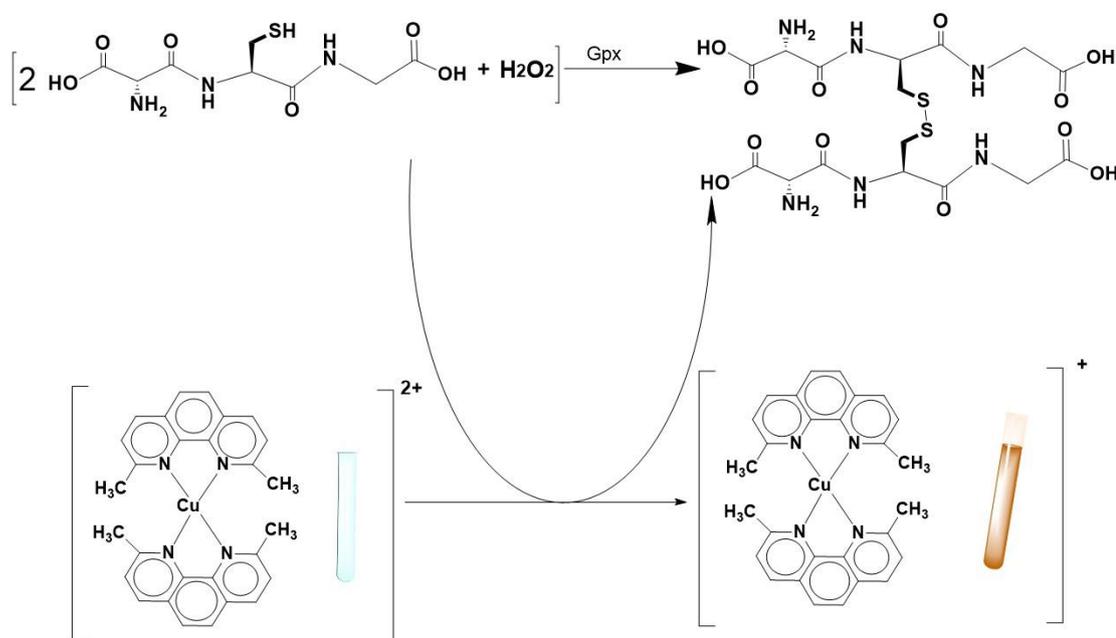
The Passing–Bablok regression [23] and Bland–Altman analysis [24] were used to compare the Gpx-CUPRAC method to the Gpx-DTNB method. The QiMacros program linked to Microsoft Excel 2016 was used for mathematical analyses (QiMacros, Know Ware International, Denver, USA).

Results and Discussion

CUPRAC reagent as a suitable probe to measure Gpx activity [the Gpx-CUPRAC method]

By using the cupric neocuproine complex ($\text{Cu}(\text{Nc})_2^{2+}$) as a suitable chromogenic oxidising probe, the present work explains a basic procedure to assay Gpx activity in biological samples (CUPRAC method). Apak et al. [25] introduced the CUPRAC method to calculate antioxidant capacity. To establish Gpx activity, the enzyme samples were incubated with phosphate buffer, which contains suitable concentrations of the glutathione and peroxide as substrates. To stop the enzyme's reaction, the CUPRAC reagent ($\text{Cu}(\text{Nc})_2^{2+}$) was added after a suitable incubation time.

Reducing the Cu(II)-neocuproine ($\text{Cu}(\text{Nc})_2^{2+}$) complex to coloured Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex by the unreacted substrates was quantified spectrophotometrically at 450 nm (CUPRAC method), as shown in Scheme 1. The decrease of absorbance of coloured Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex correlated with the Gpx activity. The Gpx enzyme reaction formed the Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex, which had a single peak at 450 nm. The absorbance was specifically associated with the unreacted substrates (glutathione and peroxide), as shown in Figure 2.



Scheme 1: Glutathione peroxidase uses peroxide to convert reduced glutathione (GSH) to oxidised glutathione (GSSG). Cu(II)-neocuproine complex reacts with unreacted substrates (glutathione and peroxide) to produce yellow-orange Cu(I)-neocuproine stable complex.

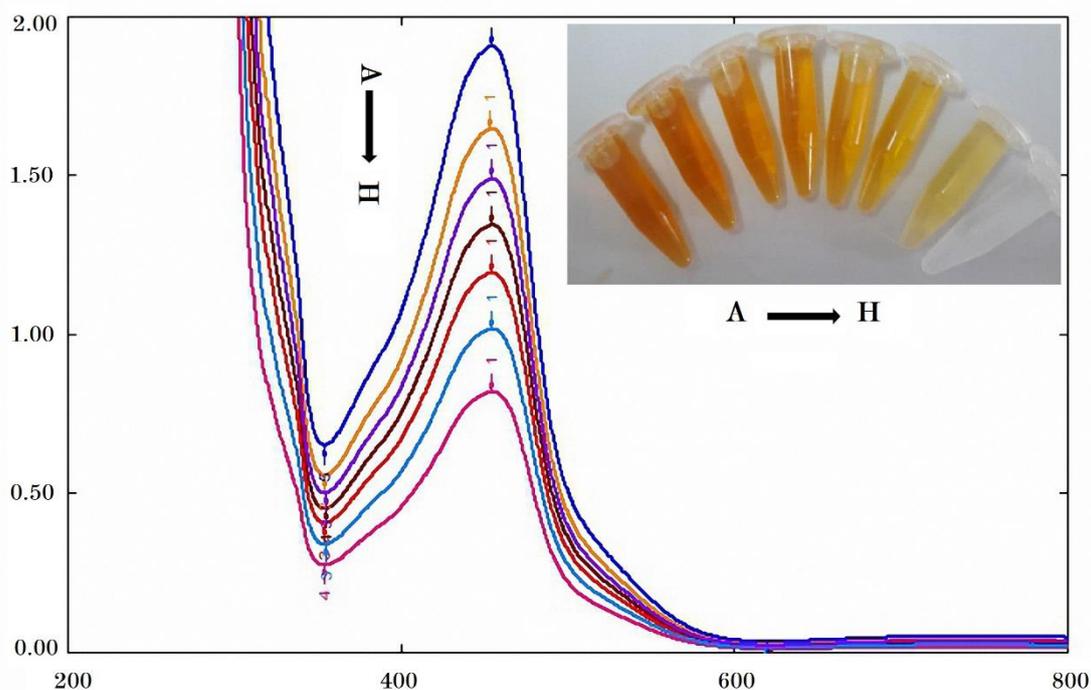


Fig. 2. Glutathione enzyme activity correlated inversely with the intensity of the formed Cu(I)-neocuproine ($\text{Cu}(\text{Nc})_2^+$) complex. Absorption spectra was achieved by reducing ($\text{Cu}(\text{Nc})_2^{2+}$) to coloured Cu(I)-neocuproine complex ($\text{Cu}(\text{Nc})_2^+$) as a result of adding 1 ml of the solution prepared by mixing a suitable concentration of glutathione (GSH) with peroxide (H_2O_2). (a) to (h) represent $\{(800/400), (700/350), (600/300), (550/275), (500/250), (450/225), (400/200)\}$ μM of (GSH/ H_2O_2).

Optimising the Gpx-CUPRAC assay

To achieve the optimum conditions, statistical methods were adapted with the Box–Behnken design (BBD) [26]. To optimise the Gpx-CUPRAC assay, BBD is an important measuring method with three central points to optimise glutathione, peroxide, and neocuproine concentrations to achieve optimal Gpx activity (see Table 3). The regression model for the Gpx-CUPRAC assay was determined using the analysis of variance (ANOVA) of the response surface methodology (RSM), as shown in Table 4. The model's F-value (12.82) showed that it was significant, while the lack-of-fitness F-value (2.8405) showed that it was not significant as compared to the related p-value. The significance of model terms was proved by obtained p-value ($p = 0.0019$).

The adjusted response (Adjusted $R^2 = 0.9918$) was in acceptable agreement with the coefficient predicted response (Predicted $R^2 = 0.9482$). As a result, the Gpx-CUPRAC assay's ANOVA showed that the specific correlation between the independent variables of the proposed model was appropriate for description and highly significant.

To investigate the graphical results of the independent variables, contour diagrams and three dimensional (3D) of the BBD were used. When the third factor was constant, the creation of 2D and 3D graphs at the midpoint stage was based on a combination of two variables. In the response plot in Fig. 3a–f, the relationships between the variables (glutathione, peroxide, and neocuproine concentrations) are depicted. All the figures exhibited good significant curvature. Gpx activity was optimum at 4 mmol L⁻¹ glutathione, 2 mmol L⁻¹ peroxide, and 7.5 mmol L⁻¹ neocuproine concentrations. The actual activity level was 500 U/L. The actual value was consistent with the predicted value, indicating that the RSM investigation was reliable and well-suited to experimental conditions.

Table 4. ANOVA values for the Gpx-CUPRAC assay's experimental variables.

	Sum of squares	Degree of freedom	Mean square	F-value	p-value
Regression	1.9322e+03	9	2.1469e+03	12.8227	0.0019
Residual	1.1720e+03	7	167.4265		
Lack-of-fit	1.0273e+03	5	205.4637	2.8405	0.2806
Pure error	144.6667	2	72.3333		
Total	2.0494e+04	16			
R ²	0.9428				
Explainable R ²	0.9918				

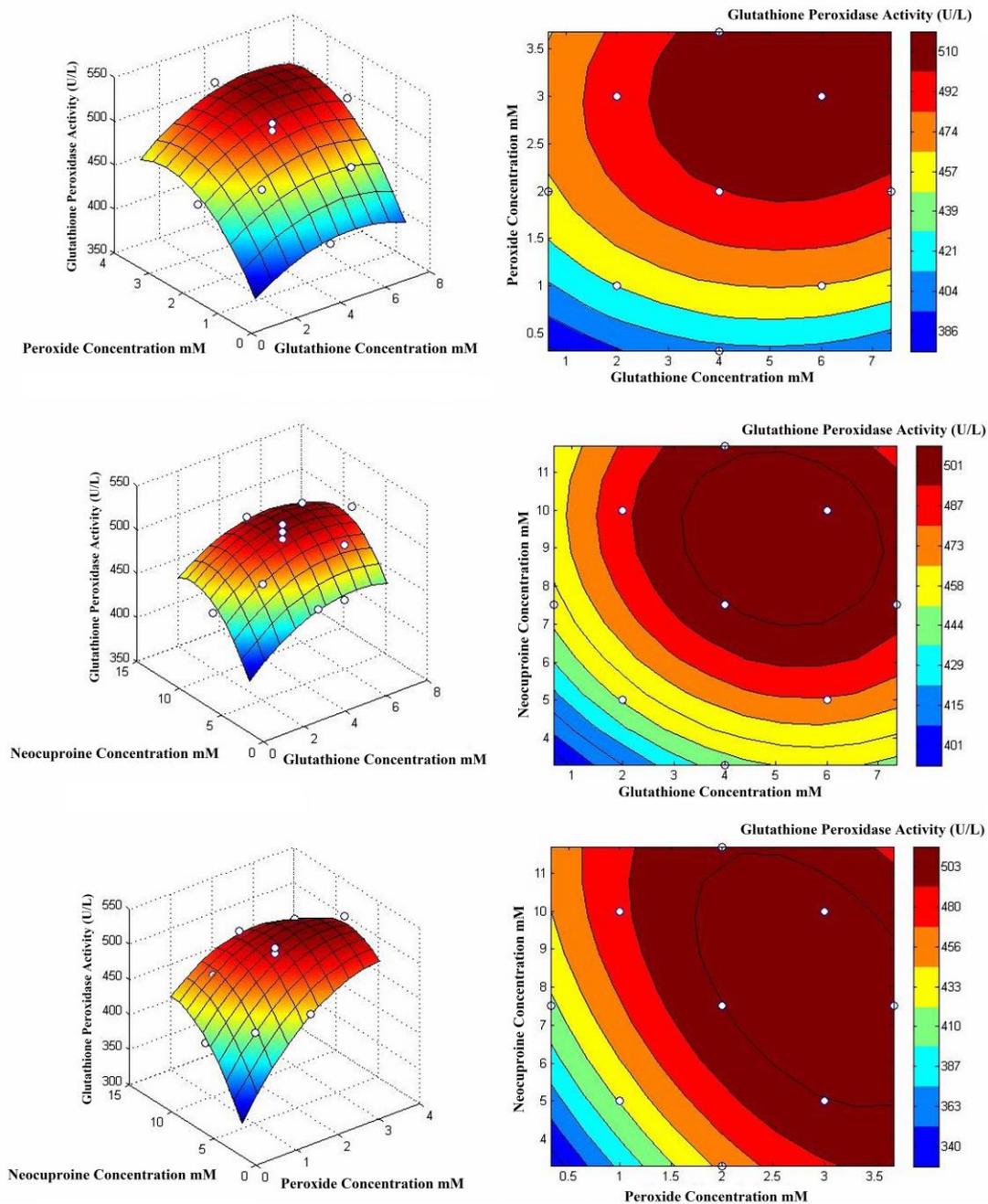


Fig. 3. 3D surface plot graphs and contours, demonstrating the interactions between glutathione concentration, peroxide concentration, and neocuproine concentration. The relationships between the variables (glutathione, peroxide, and neocuproine concentrations) are depicted in (a) through (f).

Signal stability

The coloured complex in the current study was remarkably stable at room temperature. At 25 °C, the CUPRAC complex's 450 nm absorbance remained remarkably stable for more than a week.

Linearity and sensitivity

According to the results shown in Fig 4, the Gpx-CUPRAC method was linear ($y = 1.0006 x$) within the range of 2–1000 U.L⁻¹ of Gpx activity (Pearson's $r = 0.999$). The LOQ (1 U.L⁻¹) and LOD (3 U.L⁻¹) values demonstrated the high sensitivity of the Gpx-CUPRAC assay. The linearity of the new method was like that of the Gpx-DTNB assay.

Figure 4 The Gpx activity of diluted tissue homogenates that were obtained by using the Gpx-CUPRAC method compared to the values achieved using the GPx-DTNB method.

Selectivity, reproducibility, and accuracy of the Gpx-CUPRAC method

The findings in Table 5 showed that the analysed biomolecules cannot interact with the Gpx-CUPRAC assay. The current method differs from previous methods for determining Gpx activity because it does not interfere with biomolecules. The current assay employs a control test tube to eliminate interference caused by organic biomolecules in the sample containing Gpx enzyme activity. In the current procedure, the absorbance of the test tube belongs to two kinds of compounds: unreacted substrates and sample interferences. The absorbance of the control test tube corresponds to interference compounds only. We excluded the interference of any compound that could reduce Cu(II)-neocuproine complex ($\text{Cu}(\text{Nc})_2^{2+}$) to coloured Cu(I)-neocuproine complex ($\text{Cu}(\text{Nc})_2^+$) by subtracting the absorbance of the control test tube from the absorbance of the test tube. That means the remaining absorbance was exclusively for unreacted substrates.

Table 5. By using the Gpx-CUPRAC assay to assess Gpx activity, there is a relationship between relative percentage error and interfering biological interferences.

	Added Gpx U/l	Found Gpx U/l	Relative error (%)
Volumetric flask 1	300	300	0.00
Volumetric flask 2	300	302	0.66
Volumetric flask 3	300	303	1.0
Volumetric flask 4	300	305	1.66

The Gpx-CUPRAC assay was used to determine the Gpx activity of homogenates of liver tissue. The findings revealed that Gpx activity was elevated as predicted in liver tissue homogenates (Fig. 5). The Gpx-CUPRAC method demonstrated reasonable inter-day ($\text{RSD}\% = 2.2.8\%–3.2\%$) and intra-day ($\text{RSD}\% = 2.7\%–3.8\%$).

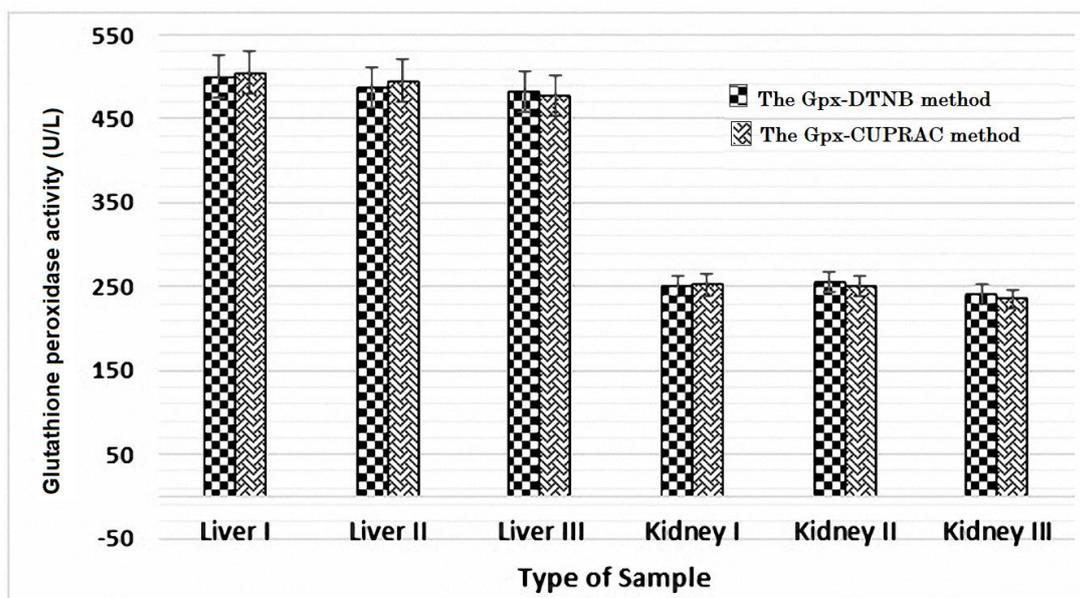


Fig. 5. Comparison of Gpx activity of diluted tissue homogenates that were obtained by the Gpx-CUPRAC method and the GPx-DTNB method.

Gpx activity assessment is a useful parameter for evaluating the liver's ability to reduce the susceptibility to oxidative stress. Furthermore, several scientific experiments have focused on Gpx activity in the livers of some types of lab animals to assess the oxidative stress inclination [27, 28].

Validation

Using matched enzymatic samples, Bland–Altman plot analyses (QI Macros, 2016) were used to compare the Gpx activity assessed by the present method with the Gpx activity assessed by the GPx-DTNB process [6]. Bland–Altman plot shows the relative differences between Gpx-CUPRAC and GPx-DTNB methods, as well as the mean relative bias (Fig. 6). The correlation coefficient between the two protocols was 0.9994. This means that the new protocol is almost as accurate as the reference protocol. The comparison between the Gpx-CUPRAC method and the GPx-DTNB method using the Passing–Bablok similarity analysis showed a good agreement correlation (Fig. 7).

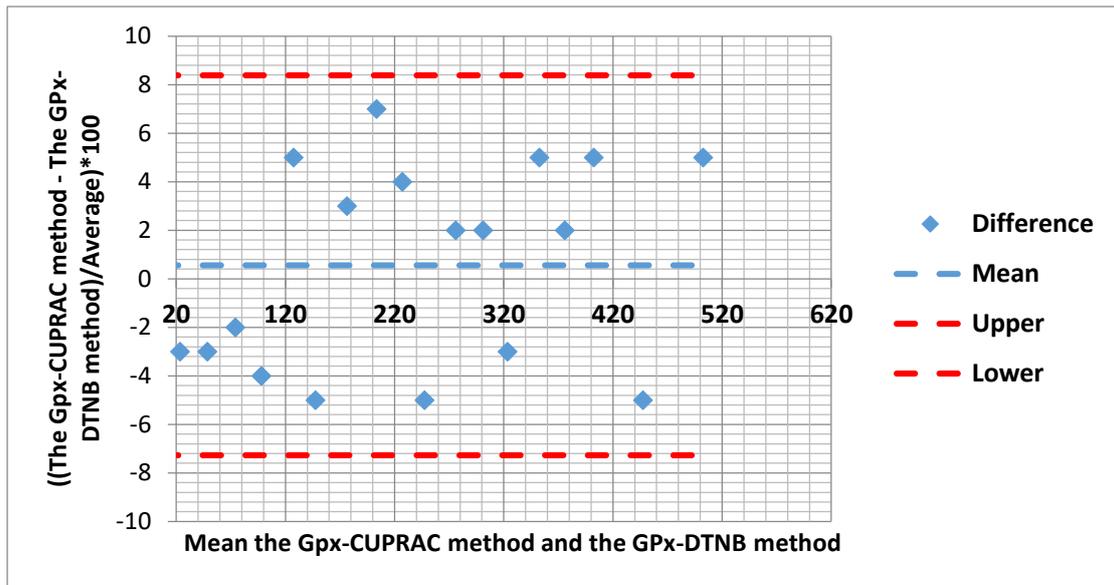


Fig. 6. Bland–Altman plot shows the relative differences between the Gpx-CUPRAC and GPx-DTNB protocols, as well as the mean relative bias.

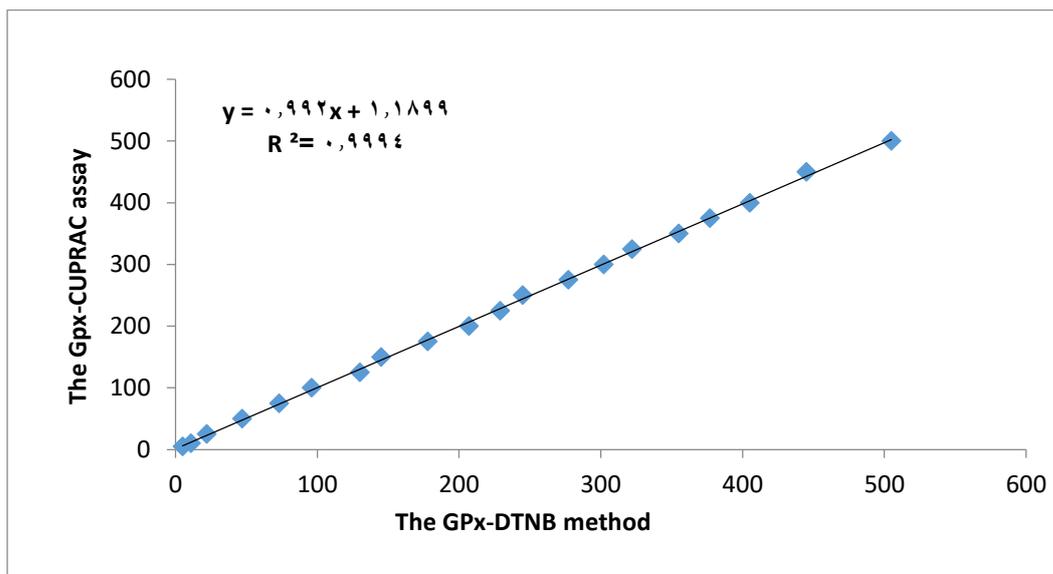


Fig. 7. Gpx activities were measured using the Gpx-CUPRAC and GPx-DTNB methods over a series of Gpx dilutions.

Conclusion

In this research, a simple and accurate protocol for calculating Gpx activity with a single reagent solution was developed. The BBD calculated the optimum glutathione, peroxide, and neocuproine concentrations. The new method is free of the interference that can arise when proteins, amino acids, and sugars are present. The CUPRAC assay's working solution allows for calculating Gpx activity at low concentrations of substrate.

Competing financial interests

The author declares no competing financial interests.

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