

Square Wave Voltammetric and Computational Study of the Thyroxine-Uracil Interaction

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(Received 19 /9/ 2018 ; Accepted 25 / 10 / 2018)

ABSTRACT

The voltammetric behavior of Thyroxine (T_4) was studied using square wave voltammetry in phosphate buffer solution at (pH 7.0) as supporting electrolyte. Thyroxine gives two well-defined reduction peaks at Ep_1 (-0.359) volt and Ep_2 (-1.01) volt verses (Ag/AgCl/Sat.KCl) as reference electrode. The Gibb's free energy (ΔG), enthalpy (ΔH) and entropy (ΔS) changes of temperature dependent on (K) were calculated using Van't Hoff equation for Thyroxine and Uracil binding. The molecular docking between Thyroxine and Uracil has been studied, and the results indicate that the interaction between T_4 and Ur was mainly hydrogen bonding and van der Waals interaction.

Keywords: Thyroxine, Uracil, Interaction, Molecular docking.

الملخص

7.0
(Ag/AgCl/Sat.KCl) (-1.01) (-0.359)

($\Delta G, \Delta S, \Delta H$)

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INTRODUCTION

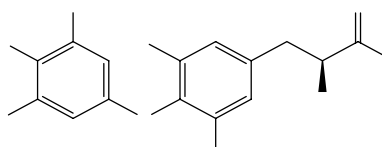
L-Thyroxine (L- T_4) (+)-3,5,3',5'-tetraiodo-L-thyronine (1) is an important biological compound derived from tyrosine and produced in the thyroid gland (Voet *et al.*, 2002). Also T_4 is the main hormone secreted into the bloodstream by the thyroid gland. It is inactive and most of it is converted to an active form called triiodothyronine (T_3) by organs such as the liver and kidneys. Thyroid hormones play vital roles in regulating the body's metabolic rate, heart, digestive functions, muscle control, brain development and maintenance of bones, among many other effects not fully studied. The thyroid hormones T_3 and T_4 are unique in that iodine (as iodide) as an essential component of both (Abdul-Fattah *et al.*, 2018).

The usual methods for the determination of T_4 were UV- absorption (Gregorini *et al.*, 2013), Time resolved fluorescence (Wu *et al.*, 1999), Enzyme immunoassays (Tsoncheva, 1988), HPLC (Sawabe *et al.*, 2011), Radioimmunoassay (RIA) (Ping-Jun, 1983), and Chemiluminescence (CL) (Gok and Ates, 2004).

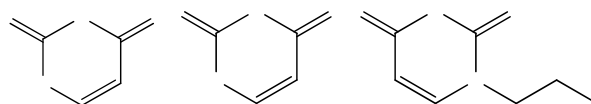
However, these methods have some disadvantages such as expensive instrumentation, time consuming and complicated operations. Cathodic reduction of T_4 on silver electrode was studied by Iwamoto & Co-workers (Iwamoto *et al.*, 1984) in comparison with its multi-step reduction at HMDE. Cathodic stripping square wave voltammetry was applied to determine T_4 in urine, showed that T_4 in Britton Robinson buffer has two reduction peaks in the pH range(2-9) and which involve two steps reduction, at pH 10 has one reduction peak, refer to C-I bonds which is reduced in a single step (Hernandez *et al.*, 1994). Chemically modified carbon paste electrodes are used by Hu's group in the presence of CTAB (Hu *et al.*, 2004) (Chitravathi *et al.*, 2009) and Chitravathi used phenyl hydrazine as mediator to determine T_4 and the methods are applied for the determination of T_4 in commercial tablets (Chitravathi *et al.*, 2010).

Aboul-Enein and Stefan construct an amperometric biosensor to determine thyroxine based on the immobilization of L-amino acid oxidase (LAAO) on carbon paste electrodes and the two methods were applied to determine thyroxine tablets (Stefan and Aboul-Enein, 2002) (Aboul-Enein *et al.*, 2002). And for the determination of thyroxine by potentiometric sensor and applied method to determine T_4 in levothyroxine tablets and whole blood (Alimadadi *et al.*, 2014) (Moldoveanu *et al.*, 2014).

In the present study, the electrochemical behavior of T_4 and its interaction with Uracil (Ur) were studied as related simple compound to the antithyroid drugs (2). In addition, the binding constant and thermodynamic parameters were evaluated.



(1) Structure of L-thyroxine



(2) antithyroid drugs

EXPERIMENTAL

Reagents and Chemicals :

A stock solution (10^{-3} M) of L- T_4 was prepared by dissolving T_4 (obtained from Alfa company, Germany) in (0.1 M NaOH in 70% ethanol solution); they were kept in darkness at 4°C , 0.2M K_2HPO_4 & 0.2M KH_2PO_4 (obtained from Alfa company, Germany) to prepare 0.1M phosphate buffer solution (PBS) at pH 7.0. The buffer was adjusted to the required pH with the same solutions. Uracil was obtained from BDH laboratory reagent, and all solutions were prepared using deionized water and with no further purification.

Apparatus:

All voltammetric measurements were performed using 797- VA Computrace stand (Metrohm AG, CH-9101 Herisau, Switzerland). Reference electrode (RE) was Ag/AgCl/ Sat.KCl and Pt wire was used as auxiliary electrode (AE) and Hanging Mercury Drop Electrode (HMDE) was used as working electrode (WE). pH measurements were performed by using a digital pH meter (HAYANA) calibrated with standard buffers; for temperature control, a HAAKE G water bath was used.

Computational study:

The Molecular Operating Environment MOE v.(2009) software developed by (Chemical Computing Group, Montreal, Canada) was used for the graphical illustrations and molecular interaction study.

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Molecular mechanics and quantum chemical calculations were performed to study the geometries, electronic structures. The 3D structures were drawn and used as the starting point for energy minimization. The energy minimizations were performed until the gradient was below (Minimum RMS Gradient 0.0001 Kcal/mol/Å³). Initial geometry optimization of molecule was carried out using molecular mechanics by the force field method (MMFF94x).

RESULT AND DISCUSSION

Electrochemical behavior of L-T₄ :

Preliminary measurements of T₄ using SWV and the three-electrode system with HMDE as working electrode in PBS at pH 7.0 as supporting electrolyte gives two well-defined peaks at (-0.359 and -1.01) V versus Ag/AgCl/Sat.KCl. The Fig. (1) using optimum instrument conditions.

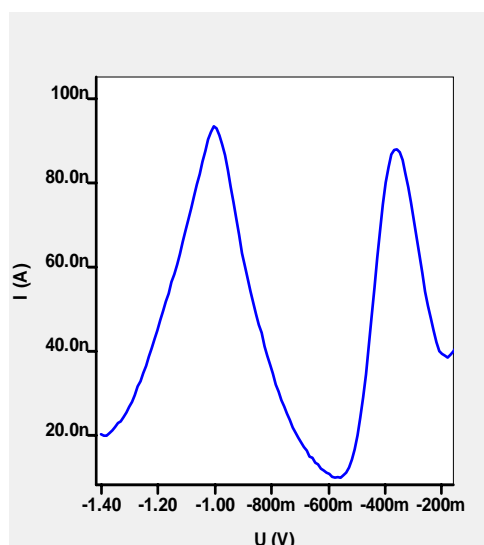


Fig.1: The S.W.Voltammogram of (4.98×10^{-6} M) T₄ in PBS at pH 7.0

Also the optimum condition has been studied and the results obtained are shown in (Table 1) and all subsequent experiments used these conditions (Abdul-Fattah *et al.*, 2018).

Table 1: The optimum condition values of thyroxine by using SWV technique

Conditions	Optimum Condition Values
Deposition Potential (V)	-0.4
Deposition Time (Sec.)	70
Equilibrium Time (Sec)	5.0
Voltage Step (V)	0.010
Amplitude (V)	0.04
Frequency (Hz)	50
Drop size (mm)	7
pH	7.0

The calibration curve of T₄ was constructed using SWV under the optimum conditions (Table 1) and potential between (-1.4 -0.1)V and gives a two straight lines; the first, at (1.996×10^{-7} - 19.61×10^{-7})M with the R² equal to (0.999) and (0.9963) for Ep₁ and Ep₂ respectively, the second at (0.996×10^{-6} - 11.857×10^{-6})M range, with the R² equal to (0.9819) and (0.9848) for Ep₁ and Ep₂ respectively (Abdul-Fattah *et al.*, 2018).

Effect of Temperature on T_4 :

The S.W.Voltammogram of ($9.9 \times 10^{-6} M$) (L-Thyroxine) in phosphate buffer solution at (pH=7), using the optimal conditions, were recorded at different temperatures (288,293,298,303,308) K. The peak potential E_p and the diffusion current (I_p) for the reduction of L-Thyroxine were measured and the results are shown in (Table 2) :

Table 2: The value of I_p of T_4 at different temperature

Temp. (K)	288	293	298	303	308
E_{p1} (V)	-0.339	-0.329	-0.319	-0.309	-0.299
$I_{p^0_1}$ (nA)	167	188	197	209	237

The result shows that the diffusion current (I_{p^0}) was found here to be increased with increasing temperature. This in fact is due to that the diffusion rate increased with temperature .

Voltammetric study of T_4 -Ur Interaction:

To study the interaction between Thyroxine and Uracil, a successive amount of Uracil ($1 \times 10^{-4} M$ as a stock solution) was added to voltammetric cell containing ($9.9 \times 10^{-6} M$) (L-Thyroxine) in phosphate buffer solution at (pH 7.0) at different temperatures (288,293,295,303,308) °K and the voltammogram was recorded for each addition. The peak current was measured at $E_{p1} = (-0.365 V)$ because it was more sensitive than E_{p2} , which belongs to reduction peak of L- T_4 ; denoted as I_{p^0} Fig (2 A). It is very clear from Fig. (2), the peak current I_p decreased gradually with the sequence additions of Uracil until reaches constant value (saturation).

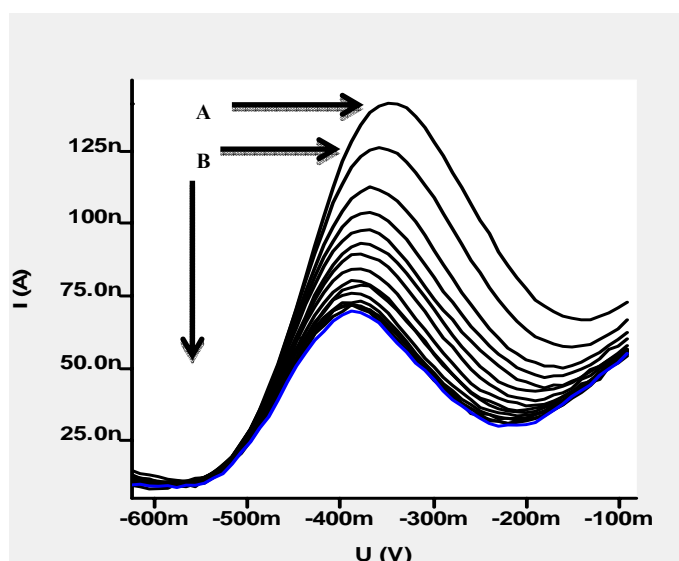
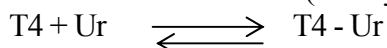


Fig.2: SW Voltammogram [9.9×10^{-6}] molar (L-Thyroxine) in the A)absence of UracilB) with the successive additionalof Uracil

Determination of Binding Constant (K) for (L-Thyroxine – Uracil):-

The interaction of (L-Thyroxine) withUracil can be described using the following equation :-



An equation for voltammetric determination can be deduced according to(Jalali and Dorraji, 2012) the current diffusion equation was described as follows :-

$$\ln (I_p / (I_{p^0} - I_p)) = \ln (1 / [Conc.(M)]) - \ln (K) \dots\dots\dots [1]$$

Where K is apparent binding constant, I_{p^0} and I_p the peak current of the free (T_4) and the complex (T_4 -Ur), respectively. then the plot of $\ln (1 / [Conc.Uracil (M)])$ versus $\ln (I_p / (I_{p^0} - I_p))$ give linear relation with intercept of $\ln (K)$ Equation (1), the results shown in Fig. (3) and (Table 3):

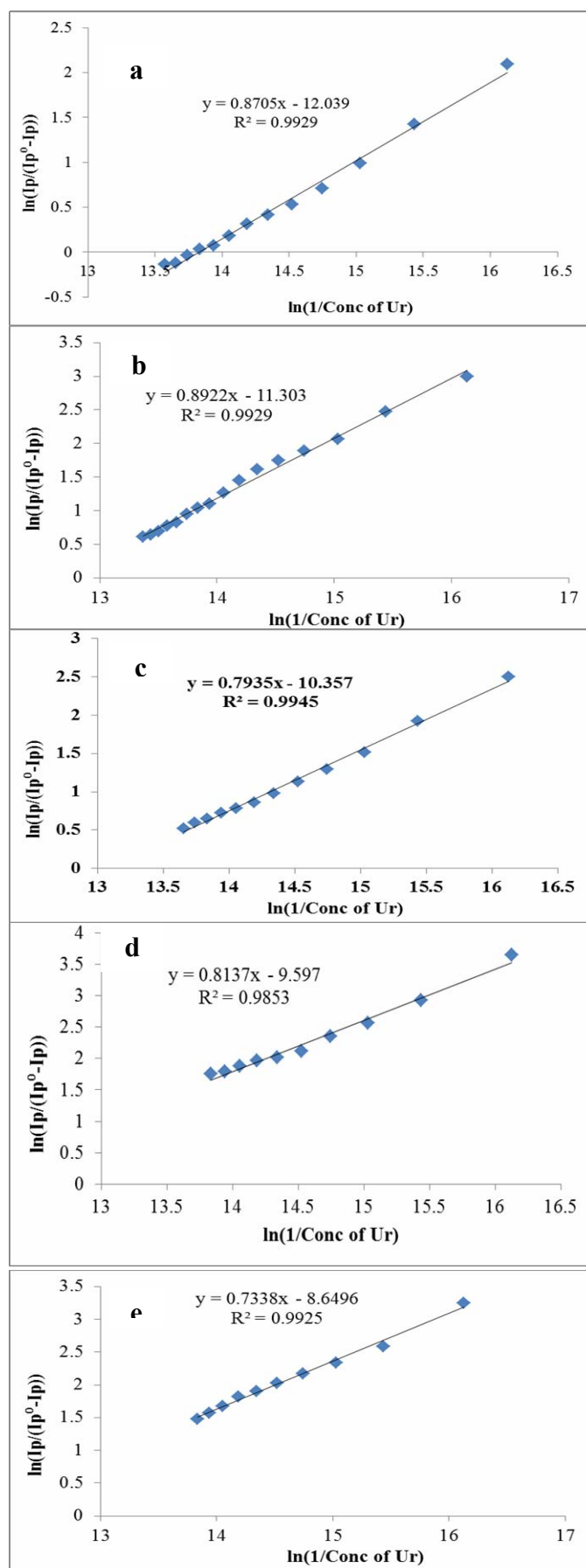


Fig. 3: (a-e) plot $\ln(1/[\text{Conc. of Uracil (M)}])$ versus $\ln(I_p/(I_p^\circ - I_p))$ of Thyroxine and Uracil interaction at (a=288, b=293, c=295, d=303, e=308) °K

Table 3: The binding constant at different temperatures (288,293,298,303,308)°K

Temp. °K	288	293	298	303	308
ln K (Ep ₁)	12.039	11.303	10.357	9.597	8.6496
Kx10 ⁴ , molar ⁻¹	16.92	8.11	3.15	1.47	0.571

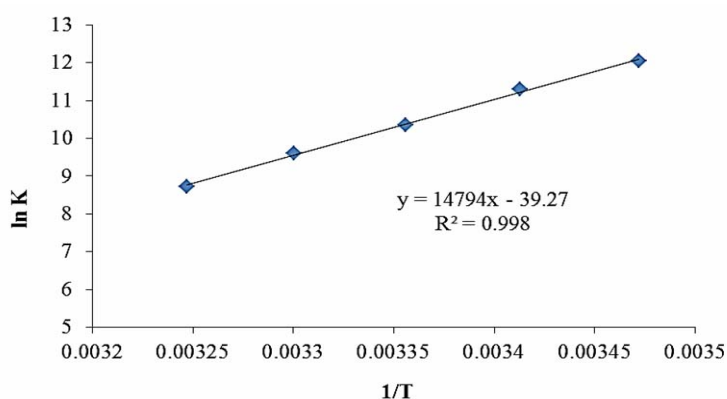
The result indicates that the value of K was found decreasing with increasing temperature in Ep₁.

Calculation of Thermodynamic Parameters :-

The plotting of ln K against 1/T using Van't Hoff equation (equation 2), gives a linear relationship Fig. (4). The enthalpy change (ΔH) was obtained from the slope, ΔS from intercept and Gibb's free energy (ΔG) was calculated from Equation (3) :

$$\ln K = -\Delta H / RT + \Delta S / R \quad \text{..... [2]}$$

$$\Delta G = -RT \ln K \quad \text{..... [3]}$$

**Fig.4: The relation between ln K and 1/T K⁻¹ for interaction between L-Thyroxine and Uracil****Table 4: The thermodynamic parameters at different temperatures (288,293,298,303,308)°K**

Temp °K	1/T	ln K	K x10 ⁴ molar ⁻¹	ΔH (KJ/mole)	ΔG (KJ/mole)	ΔS (J/mole.K)
288	0.003472	12.039	16.92	-122.997	-28.83	-326.491
293	0.003413	11.303	8.11		-27.53	
298	0.003356	10.357	3.15		-25.66	
303	0.003300	9.597	1.47		-24.18	
308	0.0032468	8.6496	0.571		-22.33	

From (Table 4), one can see that the negative value of ΔH indicates the exothermicity of the binding interaction while a negative Gibb's free energy change (ΔG) represents a spontaneous occurrence of the interaction and the negative energy change (ΔS) shows that the system becomes more order.

From thermodynamics parameters ($\Delta H < 0$, $\Delta S < 0$), it is clear that the van der Waals and hydrogen bonding is the main force in the interaction (Zhao, 2010).

Molecular Docking:

To predict the structure of molecular complex between two or more molecules (Ferreira *et al.*, 2015), the molecular docking technique was performed to get the best orientation and conformation of complex. As shown in Fig. (5a, 5b, 5c).

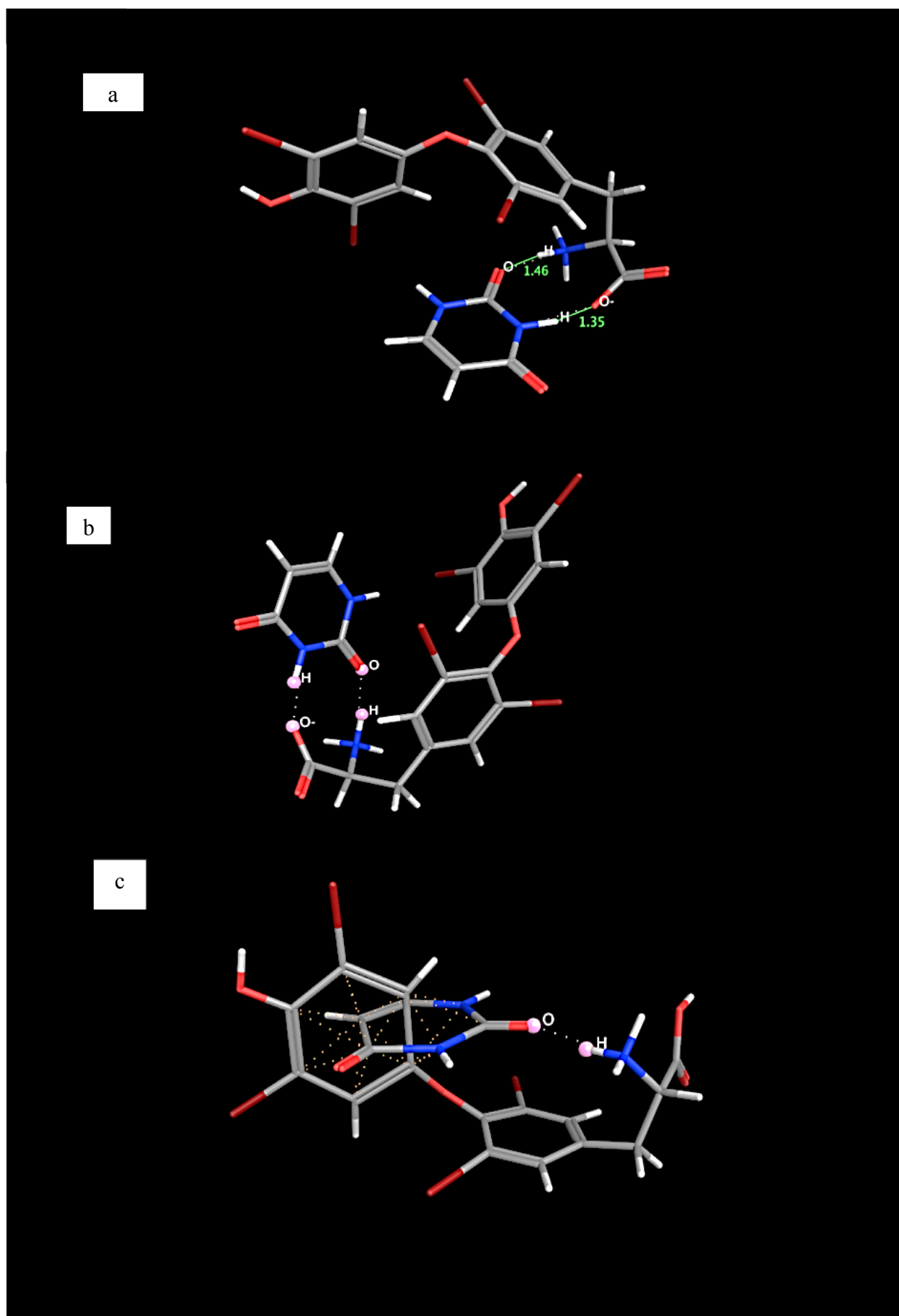


Fig. 5 : (a,b,c) Molecular Docking between Thyroxine and Uracil

From Fig. (5a,5b, 5c), we observe that Thyroxine interacts with Uracil by H-bonding and electrostatic forces.

The oxygen of carboxylic group of T₄ was very closed with hydrogen of Ur with distance (1.35°A) and oxygen of carbonyl group (C=O) of Ur also was very closed with hydrogen of amine group in T₄ with distance (1.64°A) as hydrogen bonding between them (as shown in figure 7a and 7b with white dashed line) and that agrees with thermodynamic result about Ep₁ ($\Delta H < 0$ and $\Delta S < 0$).

On the other hand, the phenolic ring (π electrone) of T₄ was also interacted with two nitrogen's atoms of Urring making a cation– π interaction and phenolic ring (π electrone) of T₄ with Ur aromatic ring's as π - π stacking (as shown in figure5c with a yellow dashed line) that suggestsan electrostatic forces, as shown in Fig. (5c). The result of molecular docking between T4 and Ur is shown in (Table 5):

Table 5: Molecular Docking result between T₄ and Ur

E_{min} Of ligand(Ur) Kcal/mol	T₄, Kcal/mol	T₄-Lig aft. Docking, Kcal/mol
-29.1091	89.1110	8.7169

CONCLUSION

In this paper, the interaction of thyroxine hormone with uracil has been studied by electrochemical method. The experimental results indicate that uracil can interact with thyroxine through hydrogen bond and van der Waals forces. The binding constant (K) between thyroxine and uracil was determined ($16.92 \times 10^4 - 0.571 \times 10^4$) at temperature range (288-308)°K, thermodynamic parameters also were calculated. The molecular docking also has been studied between thyroxine and uracil.

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