

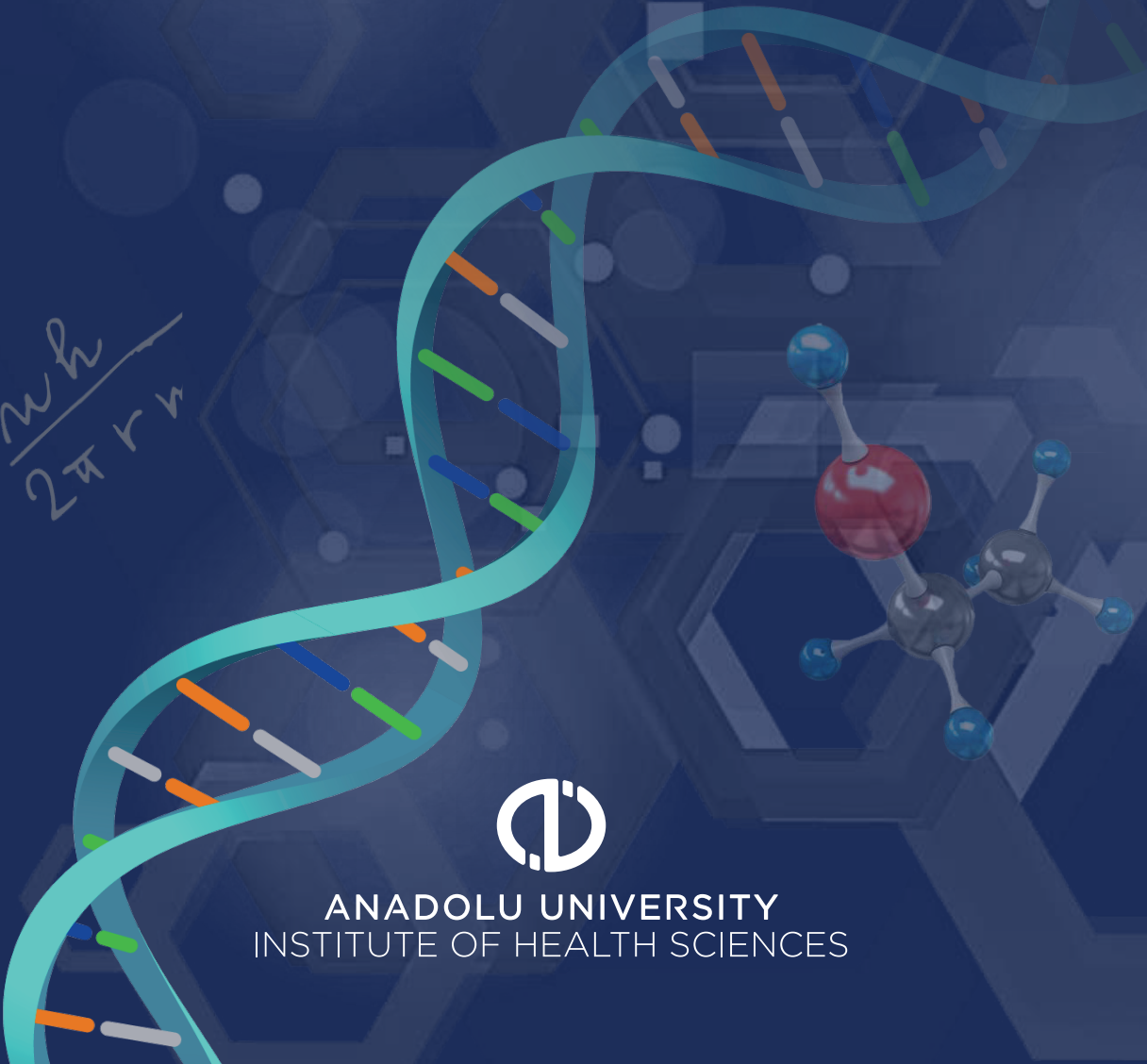
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Synthesis and characterization of new piperazine-dithiocarbamate compounds as potent MAO-A inhibitors

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ABSTRACT

Monoamine oxidase (MAO) enzymes have an important place in neurodegenerative diseases. While MAO-A inhibitors are used in depression; MAO-B enzyme has an important place in Parkinson's disease. Within the scope of this study, 7 new piperazine-dithiocarbamate derivative compounds were synthesized. Structure determinations of the compounds were performed using ¹H-NMR, ¹³C-NMR and HRMS spectroscopic methods. The MAO inhibitory activities of the compounds were determined by *in vitro* fluorometric methods. According to the obtained results, compound **2d** with IC₅₀=0.108±0.004 μM; compound **2e** exhibited inhibitory activity with an IC₅₀=0.074±0.003 μM.

Keywords: Dithiocarbamate, HRMS, Monoamine oxidase, NMR, Piperazine

1. INTRODUCTION

Monoamine oxidase (MAO), an intracellular flavoenzyme in the outer mitochondrial membrane, is responsible for the oxidative deamination of dietary amines, monoamine neurotransmitters and hormones [1,2]. Inhibitors of monoamine oxidases (MAOIs), a member of a protein family of flavin-containing amine oxidoreductases that play an important role in the regulation and metabolism of various neurotransmitters, may be useful in the treatment of psychiatric and neurological diseases [3]. Two MAO isoforms encoded by separate genes and localized to the outer membranes of the mitochondria are termed MAO-A and MAO-B [4]. These isoforms have highly conserved structures. However, the two isozymes have different substrate and inhibitor specificities due to the difference in inhibitor specificities of a pair of gating residues due to a difference in a pair of gating residues [5]. The two isoforms of MAO are dependent on substrate specificity and inhibitor selectivity based on their

amino acid sequences, their three-dimensional structure. MAO-A and MAO-B have similar amino acid residues and orientations in their active sites, with only six of the sixteen active site residues differing between the two isozymes. In reality, structural analyses have revealed that MAO-A has a single Å3 550 cavity, but MAO-B has a tighter and longer dipartite cleft known as the entrance and substrate cavities. These two pockets can Å3 unite into a single ~700 cavity [6-8]. Both the MAO-A and -B are present in the majority areas of brain [9]. The MAO-A isoform's substrate specificity includes larger endogenous amines such as norepinephrine, serotonin, and epinephrine, whereas the MAO-B isoform's substrate specificity includes small amines such as β-phenyl ethylamine and benzylamine [10].

There are two types of MAOIs that are used for medicinal purposes nowadays. MAO-A inhibitors are mostly used to treat mental illnesses such as depression and anxiety, whereas MAO-B inhibitors are used to treat Parkinson's and Alzheimer's disease

[11]. Monoaminoxidase inhibitors can reduce peroxide production, which increases neuronal damage and causes nervous system diseases such as Alzheimer's and Parkinson's disease. Monoaminoxidase inhibitors have been used in the treatment of nervous disease [12]. Because central serotonin deficiency and to a lesser extent norepinephrine play a role in depressive illness, MAO-A inhibitors are used as antidepressant agents [13]. WHO predicts that by 2020, depression will be the second most disabling condition in the world, after stress and cardiovascular system [9,14].

Although some MAO-A inhibitors, including phenelzine, isocarboxazid, tranylcypromine, iproniazid, moclobemide and toloxatone, tranylcypromine, iproniazid, moclobemide and toloxatone have made significant contributions to the treatment of depression, their clinical use has been limited due to potential adverse effects, food and medication combinations, and the emergence of other pharmacological classes [15]. In reality, MAO-A inhibitors, especially irreversible inhibitors, are used with caution in the clinic due to their potential to produce a possibly fatal hypertensive crisis when taken with tyramine-containing foods [16]. The development of reversible MAO-A inhibitors, on the other hand, eliminates this difficulty because the reversible inhibitor is displaced from MAO-A when substrate concentrations rise, enabling metabolism to occur [17]. Due to a lack of affinity and selectivity for one of the isoforms, the majority of currently used MAO inhibitors cause adverse effects. As a result, more powerful, reversible, and selective MAO-A and MAO-B inhibitors must be developed [18].

In this study, we aim to present the new piperazine derivatives including dithiocarbamate moiety as selective MAO-A inhibitors.

2. MATERIALS AND METHODS

2.1. Chemistry

All of the chemicals were obtained from commercial sources and utilized without further purification. The uncorrected melting points (M.p.) were determined using the Mettler Toledo-MP90 Melting Point

System. ¹H-NMR spectroscopy (nuclear magnetic resonance) FT-NMR spectrometer Bruker DPX 300; ¹³C-NMR spectrometer Bruker DPX 75 MHz (Bruker Bioscience, Billerica, MA, USA). ESI was used to record mass spectra on an LCMS-IT-TOF (Shimadzu, Kyoto, Japan).

2.1.1. Synthesis of Sodium 4-(4-fluorobenzyl)piperazine-1-carbodithioate (1)

1-(4-Fluorobenzyl)piperazine (0.05 mol) was dissolved in EtOH (absolute) in presence of NaOH. Carbon disulfide was dissolved in absolute EtOH, and this solution placed in dropping funnel. It was added dropwise to the reaction medium in an ice bath. The precipitated product was filtered, washed with diethyl ether, and dried at the end of the process.

2.1.2. Synthesis of the target compounds (2a-2c)

Sodium 4-(4-fluorobenzyl)piperazine-1-carbodithioate (0.001 mol) and appropriate 2-bromoacetophenone (0.001 mol) was reacted in 20 ml acetone in room temperature. At the end of the reaction, acetone was evaporated under reduced pressure. The precipitated product was filtered, washed with water, dried, and crystallized from ethanol.

2-oxo-2-phenylethyl 4-(4-fluorobenzyl)piperazine-1-carbodithioate (2a)

Yield: 81 %, ¹H-NMR (300 MHz, DMSO-*d*₆): δ = 2.47 (4H, br.s., piperazine), 3.54 (2H, s, -CH₂-), 4.00 (2H, br.s., piperazine), 4.19 (2H, br.s., piperazine), 4.96 (2H, s, -CH₂-), 7.16 (2H, t, *J*=8.9 Hz), 7.36 (2H, dd, *J*₁=5.7 Hz, *J*₂=8.6 Hz), 7.55 (2H, t, *J*=7.7 Hz), 7.64-7.67 (1H, m), 8.04 (2H, d, *J*=7.1 Hz). ¹³C-NMR (75 MHz, DMSO-*d*₆): δ = 44.17, 50.30, 51.76, 52.28, 60.72, 115.48 (*J*=21.01 Hz), 128.67, 128.68, 129.26, 131.36 (*J*=7.9 Hz), 133.88, 136.72, 161.87 (*J*=242.6 Hz), 193.25, 194.83. HRMS (m/z): [M+H]⁺ calcd for C₂₀H₂₁N₂OFS₂: 389.1152; found 389.1134.

2-oxo-2-(*p*-tolyl)ethyl 4-(4-fluorobenzyl)piperazine-1-carbodithioate (2b)

Yield: 85 %, ¹H-NMR (300 MHz, DMSO-*d*₆): δ = 2.39 (3H, s, -CH₃), 2.47 (4H, br.s., piperazine), 3.53 (2H, s, -CH₂-), 3.99 (2H, br.s., piperazine), 4.19 (2H, br.s., piperazine), 4.93 (2H, s, -CH₂-), 7.16

(2H, t, $J=8.9$ Hz), 7.34-7.39 (4H, m), 7.94 (2H, d, $J=8.2$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 21.66, 44.11, 50.36, 51.70, 52.28, 60.74, 115.47 ($J=21.1$ Hz), 128.79, 129.78, 129.79, 131.34 ($J=7.8$ Hz), 134.19, 144.33, 161.86 ($J=242.7$ Hz), 192.72, 194.87. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{21}\text{H}_{23}\text{N}_2\text{OFS}_2$: 403.1309; found 403.1284.

2-(4-methoxyphenyl)-2-oxoethyl 4-(4-fluorobenzyl) piperazine-1-carbodithioate (2c)

Yield: 75 %, $^1\text{H-NMR}$ (300 MHz, DMSO- d_6): δ = 2.48 (4H, br.s., piperazine), 3.54 (2H, s, $-\text{CH}_2-$), 3.85 (3H, s, $-\text{OCH}_3$), 3.99 (2H, br.s., piperazine), 4.20 (2H, br.s., piperazine), 4.91 (2H, s, $-\text{CH}_2-$), 7.07 (2H, d, $J=8.9$ Hz), 7.16 (2H, t, $J=8.9$ Hz), 7.37 (2H, dd, $J_1=5.8$ Hz, $J_2=8.4$ Hz), 8.02 (2H, d, $J=8.9$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 43.98, 50.22, 51.78, 52.25, 56.08, 60.71, 114.46, 115.49 ($J=21.1$ Hz), 129.50, 131.06, 131.07, 131.39 ($J=6.7$ Hz), 161.88 ($J=242.9$ Hz), 163.78, 191.53, 194.96. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{21}\text{H}_{23}\text{N}_2\text{O}_2\text{FS}_2$: 419.1258; found 419.1238.

2-(4-cyanophenyl)-2-oxoethyl 4-(4-fluorobenzyl) piperazine-1-carbodithioate (2d)

Yield: 89 %, $^1\text{H-NMR}$ (300 MHz, DMSO- d_6): δ = 2.46 (4H, br.s., piperazine), 3.52 (2H, s, $-\text{CH}_2-$), 3.98 (2H, br.s., piperazine), 4.18 (2H, br.s., piperazine), 4.93 (2H, s, $-\text{CH}_2-$), 7.16 (2H, t, $J=8.9$ Hz), 7.36 (2H, dd, $J_1=5.7$ Hz, $J_2=8.6$ Hz), 7.62 (2H, d, $J=8.6$ Hz), 8.65 (2H, d, $J=8.6$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 44.02, 50.33, 51.70, 52.18, 60.65, 115.50 ($J=21.1$ Hz), 115.69, 118.62, 129.24, 129.56, 131.43 ($J=5.3$ Hz), 133.31, 140.11, 161.91 ($J=241.5$ Hz), 163.78, 193.05, 194.59. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{21}\text{H}_{20}\text{N}_3\text{OFS}_2$: 414.1105; found 414.1119.

2-(4-nitrophenyl)-2-oxoethyl 4-(4-fluorobenzyl) piperazine-1-carbodithioate (2e)

Yield: 80 %, $^1\text{H-NMR}$ (300 MHz, DMSO- d_6): δ = 2.46 (4H, br.s., piperazine), 3.52 (2H, s, $-\text{CH}_2-$), 3.98 (2H, br.s., piperazine), 4.17 (2H, br.s., piperazine), 4.97 (2H, s, $-\text{CH}_2-$), 7.16 (2H, T, $J=8.8$ Hz), 7.35 (2H, dd, $J_1=6.0$ Hz, $J_2=8.1$ Hz), 8.26 (2H, d, $J=8.9$ Hz), 8.37 (2H, d, $J=8.9$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 44.13, 51.57, 51.91, 52.30, 60.77, 115.45 ($J=21.1$ Hz), 124.37, 130.02, 131.28 ($J=7.9$

Hz), 134.15, 134.20, 141.62, 150.37, 161.83 ($J=242.7$ Hz), 192.93, 194.47. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{20}\text{H}_{20}\text{N}_3\text{O}_3\text{FS}_2$: 434.1003; found 434.1006.

2-(4-fluorophenyl)-2-oxoethyl 4-(4-fluorobenzyl) piperazine-1-carbodithioate (2f)

Yield: 70 %, $^1\text{H-NMR}$ (300 MHz, DMSO- d_6): δ = 2.48 (4H, br.s., piperazine), 3.54 (2H, s, $-\text{CH}_2-$), 3.99 (2H, br.s., piperazine), 4.19 (2H, br.s., piperazine), 4.94 (2H, s, $-\text{CH}_2-$), 7.16 (2H, t, $J=8.9$ Hz), 7.34-7.42 (4H, m), 8.13 (2H, dd, $J_1=5.5$ Hz, $J_2=8.9$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 44.03, 50.34, 51.77, 52.26, 60.71, 115.48 ($J=21.1$ Hz), 116.29 ($J=21.9$ Hz), 131.37 ($J=7.7$ Hz), 131.69 ($J=9.5$ Hz), 133.47, 133.50, 161.87 ($J=242.78$ Hz), 165.55 ($J=251.9$ Hz), 191.94, 194.77. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{20}\text{H}_{20}\text{N}_2\text{OF}_2\text{S}_2$: 407.1058; found 407.1039.

2-(4-chlorophenyl)-2-oxoethyl 4-(4-fluorobenzyl) piperazine-1-carbodithioate (2g)

Yield: 80 %, $^1\text{H-NMR}$ (300 MHz, DMSO- d_6): δ = 2.46 (4H, br.s., piperazine), 3.52 (2H, s, $-\text{CH}_2-$), 3.98 (2H, br.s., piperazine), 4.18 (2H, br.s., piperazine), 4.93 (2H, s, $-\text{CH}_2-$), 7.16 (2H, t, $J=8.9$ Hz), 7.33-7.38 (2H, m), 7.63 (2H, d, $J=8.6$ Hz), 8.06 (2H, d, $J=8.6$ Hz). $^{13}\text{C-NMR}$ (75 MHz, DMSO- d_6): δ = 43.98, 50.49, 51.88, 52.31, 60.78, 115.46 ($J=21.1$ Hz), 129.37, 130.58, 131.29 ($J=8.1$ Hz), 134.17, 134.21, 135.47, 138.75, 161.83 ($J=242.7$ Hz), 192.46, 194.65. HRMS (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{20}\text{H}_{20}\text{N}_2\text{OFS}_2\text{Cl}$: 423.0762; found 423.0746.

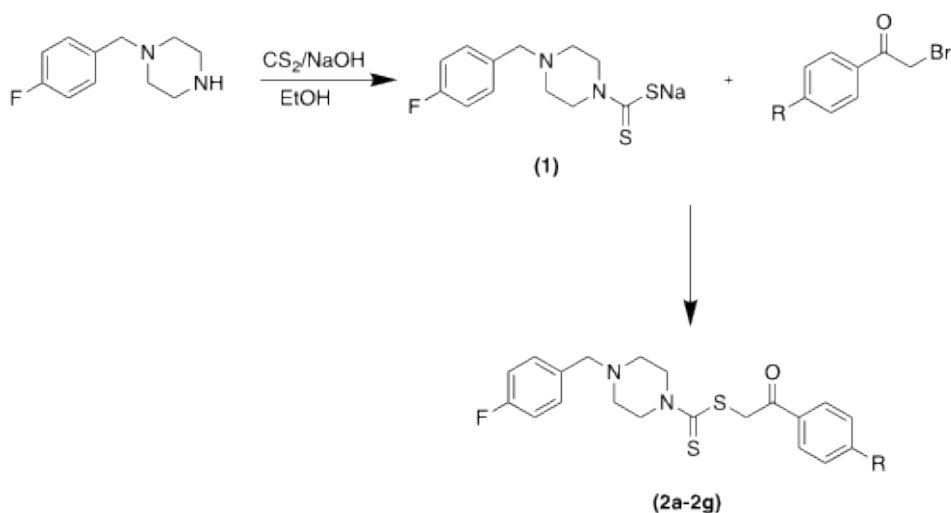
2.2. MAO Enzymes Inhibition Assay

The in vitro MAO inhibition test was carried out using the standard fluorometric method, and the percentages and IC_{50} values of the compounds obtained were reported as described by our research group already [19-25].

3. RESULTS AND DISCUSSION

3.1. Chemistry

The compounds **2a-2g** were obtained as presented in **Scheme 1**. Initially, a sodium 4-(4-fluorobenzyl) piperazine-1-carbodithioate (**1**) was obtained by means of the reaction between 1-(4-fluorobenzyl)



Compounds	R
2a	-H
2b	-CH ₃
2c	-OCH ₃
2d	-CN
2e	-NO ₂
2f	-F
2g	-Cl

Scheme 1. Synthesis pathway for obtained compounds (2a-2g)

piperazine and carbon disulfide using NaOH. Secondly, the resulting products were obtained by reacting the sodium 4-(4-fluorobenzyl)piperazine-1-carbodithioate with appropriate 2-bromoacetophenone derivatives. The structures of the compounds obtained were established by

spectroscopic methods, namely ¹H-NMR, ¹³C-NMR, and HRMS.

3.2. MAO Enzymes Inhibition Assay

The in vitro fluorometric approach published previously by our research group was used to

Table 1. % Inhibition and IC₅₀ values of the synthesized compounds, moclobemide and selegiline against MAO-A and MAO-B.

Compounds	MAO-A % Inhibition		MAO-A IC ₅₀ (μM)	MAO-B % Inhibition		MAO-B IC ₅₀ (μM)
	10-3 M	10-4 M		10-3 M	10-4 M	
2a	65.152±1.234	42.861±0.948	>100	45.451±0.763	24.451±0.632	>1000
2b	72.647±1.052	40.542±0.891	>100	41.894±0.612	20.132±0.741	>1000
2c	68.496±1.319	46.668±0.961	>100	36.135±0.754	28.648±0.665	>1000
2d	90.497±1.316	84.433±0.868	0.108±0.004	35.532±0.817	27.745±0.662	>1000
2e	93.134±1.034	90.225±0.921	0.074±0.003	33.177±0.793	20.647±0.765	>1000
2f	63.320±1.263	41.150±0.918	>100	38.736±0.806	20.349±0.621	>1000
2g	69.033±0.958	44.962±0.845	>100	43.940±0.730	31.033±0.733	>1000
Moclobemid	94.121±2.760	82.143±2.691	6.061±0.262	-	-	-
Selegiline	-	-	-	98.258±1.052	96.107±1.165	0.037±0.001

assess the inhibitory capability of the synthesized compounds on MAO isoenzymes [19-25]. % Inhibition and IC_{50} values of the synthesized compounds, moclobemide and selegiline against MAO-A and MAO-B were presented in **Table-1**. As seen in **Table-1**, none of the compounds showed more than 50% inhibition against MAO-B enzyme at 10^{-4} M concentration. In contrast, compounds **2d** and **2e** showed greater than 50% inhibitory activity against the MAO-A enzyme. Therefore, serial dilutions of these compounds at 10^{-3} - 10^{-9} M concentrations were prepared and their inhibitory activities on MAO-A enzyme were investigated. According to the results obtained, compound **2d** with $IC_{50}=0.108\pm 0.004$ μ M; compound **2e** exhibited inhibitory activity with an $IC_{50}=0.074\pm 0.003$ μ M.

3.2.1. Statistical analysis

Percentage of % inhibitions compared to the control group and values of IC_{50} were determined. All descriptive data were expressed as the mean \pm standart error of mean (SEM) from 5 times repetition within the experiments. Data were calculated on Microsoft Excel software with the sigmoidal dose-response curves by using Hill equation.

4. CONCLUSION

MAO-A inhibitors are used in the treatment of depression. However, this group of drugs is not preferred because of their side-effect profiles. Therefore, there is a need for new inhibitors with a reduced side-effect profile. Within the scope of this study, 7 new piperazine-dithiocarbamate derivative compounds were synthesized. Structure determinations of the compounds were performed using 1 H-NMR, 13 C-NMR and HRMS spectroscopic methods. The MAO inhibitory activities of the compounds were determined by *in vitro* fluorimetric methods. None of the compounds showed inhibitory activity against the MAO-B enzyme. In contrast, compounds **2d** and **2e** showed greater than 50% inhibitory activity against the MAO-A enzyme. According to the obtained results, compound **2d** with $IC_{50}=0.108\pm 0.004$ μ M; compound **2e** exhibited inhibitory activity with an $IC_{50}=0.074\pm 0.003$ μ M. When the structures of the compounds are examined,

compound **2d** contains CN group and compound **2e** contains NO_2 group. Both groups are active groups that can withdraw electrons. Therefore, it is thought that such substituents contribute positively to the activity.

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Ethical approval

Not applicable, because this article does not contain any studies with human or animal subjects.

Author contribution

Concept: ZAK, YO; Design: ZAK, YO; Supervision: ZAK; Materials: DO, BNS, SL; Data Collection and/or Processing: DO, BNS, SL; Analysis and/or Interpretation: DO, BNS, SL; Literature Search: DO, ZAK; Writing: DO, ZAK; Critical Reviews: YO, ZAK.

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Conflict of interest

The authors declared that there is no conflict of interest.

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Synthesis and *in silico* evaluation of some new 2,4-disubstituted thiazole derivatives

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ABSTRACT

The synthesis and antiproliferative activity investigation of methyl 3/4-[[4-(2-substituted thiazol-4-yl)phenyl]amino]-3-oxopropanoate/(4-oxobutanoate) (**3a-h**) derivatives were aimed in this work. The synthesis of the new compounds was carried out by a simple, multiple-step synthetic procedure. The physicochemical properties of the compounds were determined using SwissADME and QikProp software systems. Additionally, virtual target and toxicity predictions were carried out for all final compounds. The pharmacokinetics/physicochemical, druglikeness properties and biological target and toxicity predictions of the compounds were determined to possess satisfying findings. Since the target determination of the compounds according to literature is point out their cytotoxic properties, the DNA gyrase enzyme was chosen as common enzymatic pathway, and evaluated via docking studies. Compound **3b**, namely methyl 4-[[4-(2-methylthiazol-4-yl)phenyl]amino]-4-oxobutanoate was detected to possess a potential to inhibit DNA gyrase-ATPase activity.

Keywords: DNA gyrase inhibition, druglikeness, molecular docking, pharmacokinetic properties, thiazole

1. INTRODUCTION

The thiazole ring is an aromatic heterocyclic ring commonly found in natural and synthetic molecules. It is existing in the structure of alkaloids, steroids, vitamins (Thiamine-vit B1), flavones, pigments and secondary metabolites [1-3]. Secondary metabolites are organic intermediates and products produced by bacteria, fungi, or plants metabolism which of those constitute important drugs in treatment [4]. Thiazole including seconder metabolites are quite large scale that some of them are penicillins, epothilones, ascidians, tubulysins, cystothiazoles, scleritodermin, leinamycin, lyngbyabellin thiostrepton and micrococcin P1 known with various biological activity potential [5].

In 1887, Hantzsch and Weber described thiazole ring, synthesis and properties [6]. In this method, known today as the Hantzsch method, α -halo ketones and thioamides are reacted to form thiazoles [7]. In later years, Prop and coworkers synthesized new thiazole derivatives, in particular 2-aminothiazoles. Gabriel acquired various thiazoles by using a different implementation of Robinson-Gabriel oxazole synthesis. Tcherina obtained thiazole derivatives by cyclization reaction of α -thiocyanoketones whereas Erlenmeyer used the reaction of mercapto ketones and nitriles. Lastly, Cook and Heilborn were developed synthetic method via α -aminonitriles, α -aminoamides and carbon disulfide. In following years, thiazole synthesis was carried out using these methods and continued in this context [8,9]. In this direction, the

synthetic suitability of the thiazole ring leads to new applications in industrial and medicinal chemistry [10]. Mono/di/polysubstituted aromatic/nonaromatic and condensed thiazoles were widely studied and it was observed that the variety of biological activities increased with changes in substitution [3,11,12].

Thiazole derivatives were reported with a broad range of biological activities such as antifungal, antibacterial, antiviral, analgesic, anticancer, anticholinesterase, antihypertensive, and antiproliferative [13-17]. Besides, in present day, thiazole core is existing in many clinically used drugs; abafungin and ravuconazole (antifungal), ritonavir (anti-HIV), febuxostat (antigut), nizatidine (antiulcer), imidacloprid (insecticide), myxothiazols and melithiazols (fungicide), fatostatin (sterol regulatory element-binding proteins (SREBPs) inhibitor), tiazofurin, bleomycin and dasatinib (anticancer), fanetizole, meloxicam and fentiazac (anti-inflammatory) sulfathiazole (antimicrobial), and nitazoxanide (antiparasitic agent) and penicillin (antibiotic) [18-20].

According to previous studies, DNA gyrase inhibition by thiazole derivatives was investigated and obtained remarkable enzyme inhibition results, even against drug-resistance pathogens [21-23]. The action mechanism of the DNA gyrase enzyme is need ATP energy. The role of the gyrase subunit B (GyrB) is to provide the energy necessary for the DNA ligation process through ATP hydrolysis, hence, the inhibition of this subunit cause stopping ATP hydrolysis, so bacteria cells lose their functions. According to literature, thiazole ring is generally linked to GyrB subunit [24,25]. Therefore, we determined the GyrB as a target.

Furthermore, in many of our studies, we conducted biological activity studies on similar [26-28] and different thiazole derivatives [29,30]. Likewise, we synthesized eight novel thiazole derivatives and evaluated their *in silico* physicochemical properties and evaluated their anti-DNA-gyrase activity, in this study.

2. MATERIALS AND METHODS

2.1. Chemistry

All chemicals were purchased from Sigma-Aldrich Chemical Co., (Sigma-Aldrich Corp., St. Louis, MO, USA) and Merck (Merck, Darmstadt, Germany). All melting points (m.p.) were determined by Electrothermal 9300 digital melting point apparatus (Electrothermal, Essex, UK) and are uncorrected. All the reactions were monitored by thin-layer chromatography (TLC) using Silica gel 60 F254 TLC plates (Merck, Darmstadt, Germany). Spectroscopic data were recorded with the following instruments: IR, Shimadzu 8400S spectrophotometer (Shimadzu, Tokyo, Japan); ¹H-NMR, Bruker 500 MHz spectrometer (Bruker Bioscience, Billerica, MA, USA); ¹³C-NMR, Bruker 100 FT spectrometer (Bruker Bioscience, Billerica, MA, USA); MS, MS-FAB, VG Quattro Mass spectrometer (Fisons Instruments GmbH, Mainz, Germany) and elemental analyses were performed on a Perkin Elmer EAL 240 elemental analyser (Perkin Elmer, Norwalk, CT, USA).

2.1.1. General procedure for the synthesis of methyl 3/4-[[4-(2-substituted thiazol-4-yl)phenyl]amino]-3-oxopropanoate/(4-oxobutanoate) derivatives (3a-h)

4-(2-Substituted-4-thiazolyl)aniline derivatives (0.05 mol) (**2a-d**) were solved in tetrahydrofuran (75 mL) and triethylamine (0.06 mol). Methyl malonyl chloride/methyl succinyl chloride (0.06 mol) was added dropwise over 15 min to a magnetically stirred solution at 0-5°C. After the reaction was completed, the solvent was evaporated under reduced pressure, the residue was reacted with water and filtered. The obtained resulting solid was crystallized from ethanol after dryness to gain pure final products.

Methyl 3-[[4-(2-methylthiazol-4-yl)phenyl]amino]-3-oxopropanoate (**3a**)

80-82 % yield; mp 175-179 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 2.76 (s, 3H, CH₃), 3.49 (s, 2H, CH₂),

3.66 (s, 3H, OCH₃), 7.63 (d, 2H, *J*=8.5 Hz, Ar-H), 7.81 (s, 1H, thiazole C₅-H), 7.88 (d, 2H, *J*=8.5 Hz, Ar-H), 10.29 (s, 1H, N-H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ 19.39, 43.97, 53.45, 113.07, 119.66, 126.93, 130.09, 138.91, 153.98, 164.48, 165.87, 168.57. For C₁₄H₁₄N₂O₃S calculated: 57.92 % C, 4.86 % H, 9.65 % N; found: 57.81 % C, 4.94 % H, 9.55 % N. MS [M+1]⁺: *m/z* 291.

Methyl 4-[[4-(2-methylthiazol-4-yl)phenyl]amino]-4-oxobutanoate (**3b**)

76-80 % yield; mp 162-165 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 2.61-2.65 (m, 4H, CH₂), 2.70 (s, 3H, CH₃), 3.37 (s, 3H, OCH₃), 7.63 (d, *J*=8.5 Hz, 2H, Ar-H), 7.78 (s, 1H, thiazole C₅-H), 7.85 (d, 2H, *J*=8.5 Hz, Ar-H), 10.09 (s, 1H, N-H). For C₁₅H₁₆N₂O₃S calculated: 59.19 % C, 5.30 % H, 9.20 % N; found: 59.11 % C, 5.15 % H, 9.35 % N. MS [M+1]⁺: *m/z* 305.

Methyl 3-[[4-(2-phenylthiazol-4-yl)phenyl]amino]-3-oxopropanoate (**3c**)

76-78 % yield; mp 147-150 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 3.51 (s, 2H, CH₂), 3.67 (s, 3H, OCH₃), 7.51-7.56 (m, 3H, Ar-H), 7.69 (d, 2H, *J*=8.5 Hz, Ar-H), 7.99-8.03 (m, 5H, Ar-H), 8.08 (s, 1H, thiazole C₅-H), 10.36 (s, 1H, N-H). For C₁₉H₁₆N₂O₃S calculated: 64.76 % C, 4.85 % H, 7.95 % N; found: 64.61 % C, 4.96 % H, 7.85 % N. MS [M+1]⁺: *m/z* 353.

Methyl 4-[[4-(2-phenylthiazol-4-yl)phenyl]amino]-4-oxobutanoate (**3d**)

71-74 % yield; mp 203-206 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 2.62-2.64 (m, 4H, CH₂), 3.61 (s, 3H, OCH₃), 7.51-7.54 (m, 3H, Ar-H), 7.69 (d, 2H, *J*=8.5 Hz), 7.98 (d, 2H, *J*=8.5 Hz, Ar-H), 8.02 (d, 2H, *J*=8.5 Hz, Ar-H), 8.06 (s, 1H, thiazole C₅-H), 10.14 (s, 1H, N-H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ 28.92, 31.36, 51.85, 113.77, 119.49, 126.66, 127.08, 129.31, 129.74, 130.81, 133.53, 139.71, 155.51, 167.27, 170.38, 173.34. For C₂₀H₁₈N₂O₃S calculated: 65.55 % C, 4.95 % H, 7.64 % N; found: 65.64 % C, 4.99 % H, 7.75 % N. MS [M+1]⁺: *m/z* 367.

Methyl 3-[[4-(2-(4-methoxyphenyl)thiazol-4-yl)phenyl]amino]-3-oxopropanoate (**3e**)

65-68 % yield; mp 153-158 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 3.51 (s, 2H, CH₂), 3.67 (s, 3H, OCH₃), 3.84 (s, 3H, OCH₃), 7.08 (d, 2H, *J*=9 Hz, Ar-H), 7.67 (d, 2H, *J*=9.0 Hz, Ar-H), 7.95-8.0 (m, 5H, Ar-H), 10.34 (s, 1H, N-H). For C₂₀H₁₈N₂O₄S calculated: 62.81 % C, 4.74 % H, 7.33 % N; found: 62.88 % C, 4.82 % H, 7.36 % N. MS [M+1]⁺: *m/z* 383.

Methyl 4-[[4-(2-(4-methoxyphenyl)thiazol-4-yl)phenyl]amino]-4-oxobutanoate (**3f**)

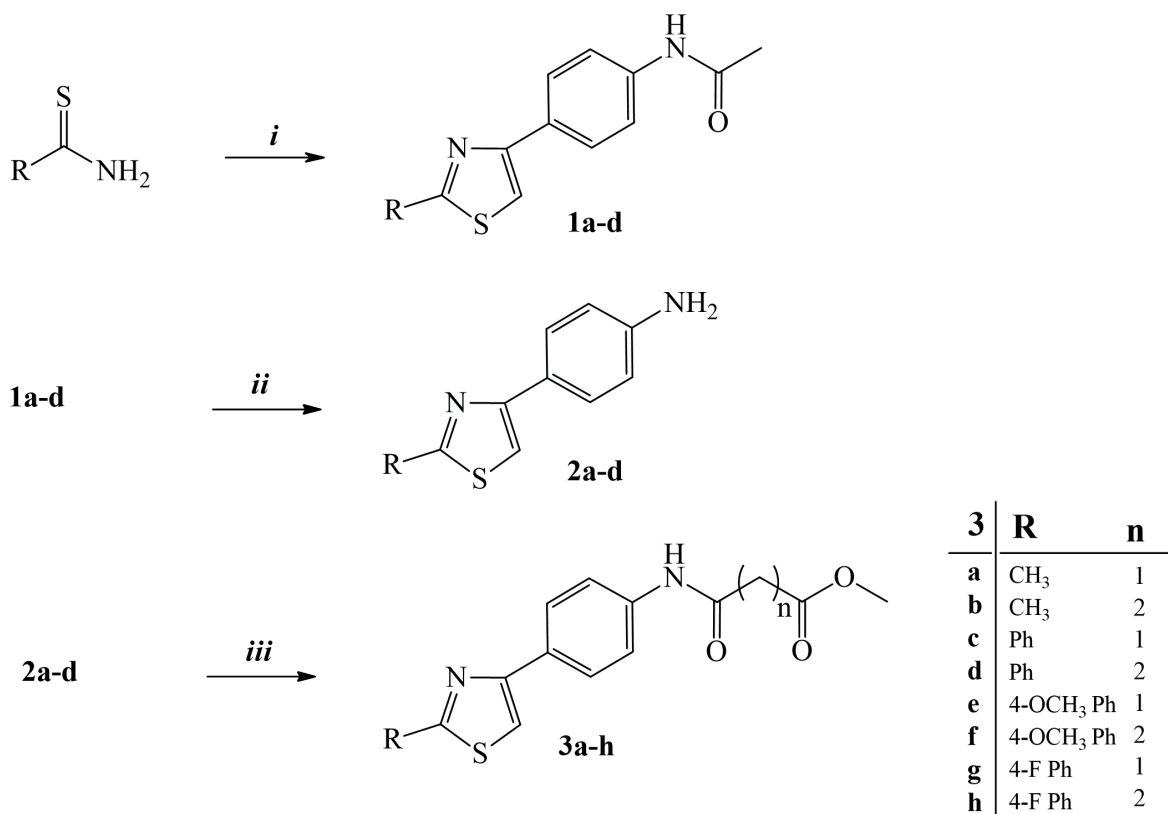
71-74 % yield; mp 203-206 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 2.61-2.65 (m, 4H, CH₂), 3.61 (s, 3H, OCH₃), 3.84 (s, 3H, OCH₃), 7.08 (d, 2H, *J*=8.5 Hz, Ar-H), 7.66 (d, 2H, *J*=8.5 Hz, Ar-H), 7.94-7.97 (m, 5H, Ar-H), 10.12 (s, 1H, N-H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ 43.51, 52.03, 55.40, 112.54, 114.64, 119.23, 125.98, 126.54, 127.43, 129.52, 138.71, 154.62, 160.90, 164.11, 166.87, 168.12. For C₂₁H₂₀N₂O₄S calculated: 63.62 % C, 5.08 % H, 7.07 % N; found: 63.66 % C, 4.98 % H, 7.15 % N. MS [M+1]⁺: *m/z* 397.

Methyl 3-[[4-(2-(4-fluorophenyl)thiazol-4-yl)phenyl]amino]-3-oxopropanoate (**3g**)

63-68 % yield; mp 184-186 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 3.67 (s, 2H, CH₂), 3.83 (s, 3H, OCH₃), 7.37 (brs, 2H, Ar-H), 7.69 (brs, 2H, Ar-H), 7.99-8.21 (m, 5H, Ar-H), 10.33 (s, 1H, N-H). For C₁₉H₁₅FN₂O₃S calculated: 61.61 % C, 4.08 % H, 7.56 % N; found: 61.69 % C, 4.19 % H, 7.65 % N. MS [M+1]⁺: *m/z* 370.

Methyl 4-[[4-(2-(4-fluorophenyl)thiazol-4-yl)phenyl]amino]-4-oxobutanoate (**3h**)

65-69 % yield; mp 200-203 °C. ¹H NMR (500 MHz, DMSO-*d*₆) δ 2.62-2.64 (m, 4H, CH₂), 3.80 (s, 3H, OCH₃), 7.38 (brs, 2H, Ar-H), 7.72 (brs, 2H, Ar-H), 7.98-8.20 (m, 5H, Ar-H), 10.35 (s, 1H, N-H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ 28.42, 30.93, 51.42, 113.34, 116.25, 116.38, 126.68, 128.42,



Scheme 1. The synthetic protocol of the compounds **3a-h**. Reactants and conditions: *i*: 4-(2-bromoacetyl)acetanilide, EtOH, r.t., 8h; *ii*: 10 % HCl, EtOH, reflux; *iii*: methyl malonyl chloride/methyl succinyl chloride, TEA, THF, r.t.

125.53, 128.79, 129.76, 139.25, 155.0, 162.25, 164.23, 165.65, 169.80, 172.83. For C₂₀H₁₇FN₂O₃S calculated: 62.49 % C, 4.46 % H, 7.29 % N; found: 62.59 % C, 4.35 % H, 7.25 % N. MS [M+1]⁺: *m/z* 384.

2.2. In Silico Prediction

The physicochemical properties of the compounds were calculated using SwissADME [31] and Qikprop [32] software programs. Potential activity and toxicity profiles were predicted using SwissTargetPrediction [33] and pkCSM–pharmacokinetics [34] software programs. The DNA-gyrase crystal was retrieved from RCSB Protein Data Bank (PDBID: 4DUH), and the protein and final compounds prepared using protein preparation and ligand preparation wizards at 7.4±1.0 pH. Their binding modes were investigated using Maestro software program.

3. RESULTS AND DISCUSSION

Present study was undertaken to synthesize methyl 3/4-[[4-(2-substituted thiazol-4-yl)phenyl]amino]-3-oxopropanoate/(4-oxobutanoate) (**3a-h**) derivatives and to evaluate their *in silico* physicochemical properties. The synthetic protocol was realized according to previously reported studies [26] and it was depicted in **Scheme 1**. Initially, thiazole derivatives were synthesized according to Hantzsch thiazole synthesis with the reaction of alkyl/aryl thioamides and α -bromo-(4'-acetylamino)acetophenone. Then the obtained thiazole intermediates (**1a-d**) were hydrolyzed to convert acetanilide to corresponding aniline. Lastly, the acquired 4-(2-substituted-4-thiazolyl)aniline derivatives (**2a-d**) were reacted with methyl malonyl chloride or methyl succinyl chloride at acetylation conditions to reach ester compounds (**3a-h**). The

Table 1. Some properties of synthesized compounds from SwissADME

	HBA	HBD	TPSA	Log P _{ow}	SC	Log K _p	GIA	RoF (V)	Ghose	Leadlikeness (V)
3a	4	1	96.53	2.38	Moderately	6.11	High	0	Yes	Yes
3b	4	1	96.53	2.48	Moderately	6.77	High	0	Yes	Yes
3c	4	1	96.53	3.50	Poorly	5.59	High	0	Yes	No (2)
3d	4	1	96.53	3.62	Poorly	6.26	High	0	Yes	No (2)
3e	5	1	105.76	3.48	Poorly	5.79	High	0	Yes	No (3)
3f	5	1	105.76	3.60	Poorly	6.46	High	0	Yes	No (2)
3g	5	1	96.53	3.77	Poorly	5.63	High	0	Yes	No (2)
3h	5	1	96.53	3.93	Poorly	6.29	High	0	Yes	No (2)

MW: Molecular Weight, **HBA:** H-bond acceptor, **HBD:** H-bond acceptor, **TPSA:** Topologic polar surface area (Å²) **Log P_{ow}:** *Consensus* Log P_{ow} (Average of all five predictions), **SC:** Solubility Class (Water), **GIA:** Gastrointestinal absorption, **Log K_p:** skin permeation (-cm/s), **RoF (V):** Rule of Five (violation number), **Ghose:** Ghose Filter, **Leadlikeness (V):** Suitability score (violation number).

Table 2. Some properties of synthesized compounds from Schrodinger (QikProp)

	HBA	HBD	PSA	Log P _{ow}	AS	-Log K _p	%HOA	RoF (V)	RoT (V)	Similar Molecules
3a	5	0	81.10	2.74	-4.20	-2.56	96	0	0	Azapropazone
3b	6	1	82.09	2.75	-4.76	-2.53	96	0	0	Diloxanide
3c	5	0	79.52	4.08	-5.80	-1.83	100	0	1	Phenylbutazone
3d	6	1	80.51	3.98	-6.20	-1.80	100	0	1	Oxyphenbutazone
3e	5.75	0	88.0	4.09	-5.84	-1.94	100	0	1	Bisacodyl
3f	6.75	1	88.99	4.09	-6.35	-1.91	100	0	1	Famprofazone
3g	6	1	80.51	4.28	-6.60	-1.94	100	0	1	Bisacodyl
3h	5	0	79.5	4.32	-6.17	-1.96	100	0	1	Tribuzone

HBA: H-bond acceptor (2 / 20), **HBD:** H-bond acceptor (0 / 6), **PSA:** vdW Polar SA (7.0 / 200.0), **%HOA:** % Human Oral Absorption in GI (poor<%25), **Log P_{ow}:** Log P_{ow} (-6.5 / 0.5), **AS:** log S for aqueous solubility (-6.5 / 0.5), **Log K_p:** skin permeability (cm/hr), **RoF (V):** Rule of Five (violation number), **RoT (V):** Jorgensen Rule of 3 Violations, **Similar Molecules:** At least 90% similarity and above were considered, but if there is nothing, at this time the closest one indicated.

structure of final compounds was assigned on the basis of spectroscopic and analytical data. In the ¹H-NMR spectra of the compounds, singlet peaks at about 3.37-3.83, 7.78-8.08 and 10.09-10.36 ppm were assigned to methoxy proton (OCH₃), thiazole C₅ proton and NH protons, respectively. The other alkylic and aromatic protons were determined at about 2.61-3.51 ppm and 7.08-8.21 ppm. In the ¹³C-NMR spectra of the compounds, peaks at about 51.85-53.45 ppm indicated the methoxy carbon existed in ester structure, the signals observed in the down field over 160 ppm were defined for carbonyl carbons and vicinal aromatic carbons to heteroatoms as expected. M+1 peaks were determined for all compounds. Elemental analyses results for C, H, and N elements were satisfactory within calculated values of the compounds.

The physicochemical/pharmacokinetic and druglikeness properties of the final compounds (**3a-h**) were predicted using SwissADME (**Table 1**) and QikProp (**Table 2**) softwares. ADME parameters (for Absorption, Distribution, Metabolism and Excretion) including H-bond acceptor (HBA), H-bond acceptor (HBD), polar surface area (PSA/TPSA), partition coefficient (Log Po/w), skin permeability (Log K_p), Rule of five (violation number-RoF) and Gastrointestinal absorption (GIA) were calculated virtually in both programs. Additionally, water solubility (Log S), Jorgensen Rule of 3 Violations (RoT(V), Molecular Weight (MW) were identified and evaluated [35]. According to the Lipinski Rule of Five, all compounds were detected in the compliant with the rule to be an oral bioavailable drug. The rule suggests that a high absorption is available for an

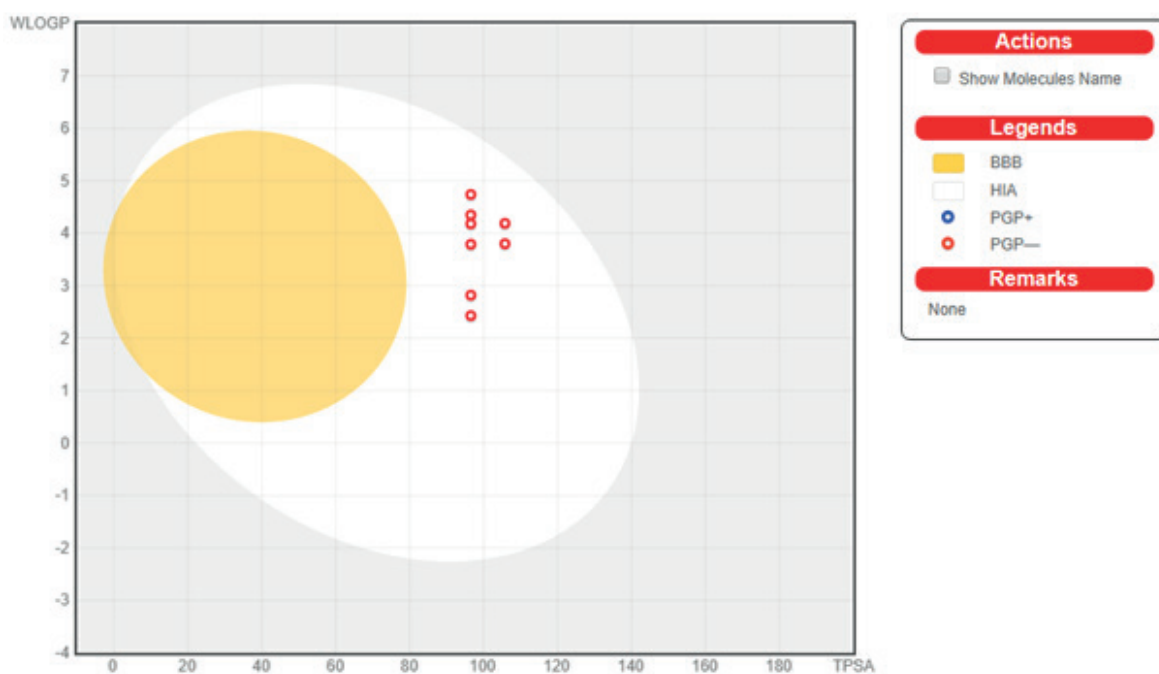


Figure 1. Boiled-egg plot of the compounds **3a-h**

oral drug when it possesses HBD less than 5, HBA less than 10, MW less than 500, and log P less than 5 [36]. All compounds passed Ghose filter developed by him [37] for drug discovery. Leadlikeness is the filter identified by Teague and co-workers which only compounds **3a** and **3b** passed through this filter [38]. Also, similar molecules (anti-inflammatory, antiprotozoal and laxative drugs) to these compounds were detected by QikProp software. The boiled-egg

Table 3. SwissTarget prediction of the compounds **3a-h** in percentage (%)

	Kinase	Enzyme	Family A G protein-coupled receptor
3a	33.3	13.3	13.3
3b	26.7	13.3	13.3
3c	60	-	-
3d	53.3	-	13.3
3e	40	-	13.3
3f	40	20	-
3g	53.3	13.3	-
3h	33.3	20	20

plot between WLOGP and TPSA was also studied in SwissADME to predict gastrointestinal absorption and brain penetration of the selected molecules. According to **Figure 1**, none of the compounds have BBB permeability but they possess gastrointestinal absorption. Besides, red dots show molecules predicted not to be effluxed from central nervous system by P-glycoprotein [39].

Target prediction of the final compounds was identified using SwissTarget prediction software. Top fifteen targets were estimated; frequently predicted target regions were given in **Table 3** and over 10 % were in explanation. Compounds **3a-h** were determined to target on kinase, enzyme and Family A G protein-coupled receptor with percentages of 26.7-60, 13.3-20 and 13.3-20 %, respectively. Additionally, compound **3a** and **3c** exhibited phosphodiesterase target with 13.3 and 26.7 %. Compound **3b** showed Family B G protein-coupled receptor (13.3%) whereas **3d** showed nuclear receptor (26.7%). Besides, compound **3e**, **3f** and **3g** represented targets phosphatase, electrochemical

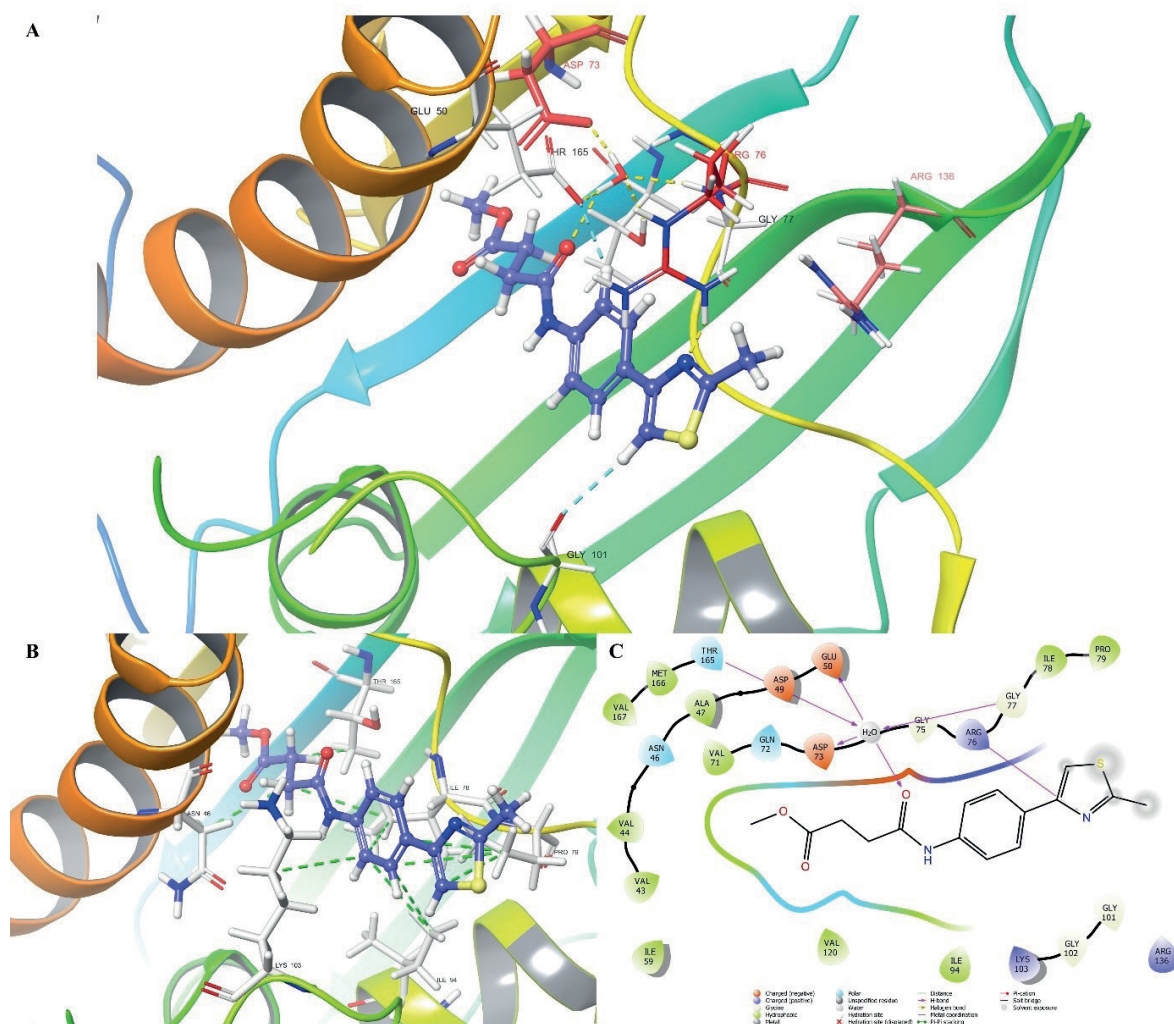


Figure 2. The best 2D and 3D poses of compound **3b**. **A)** The carbon atoms of key amino acids were represented as red, the carbon atoms interacted residues were represented as white. **B)** Hydrophobic interactions were represented as green dashes. **C)** The ligand, the residues around its 5 Å, and the important water molecule were displayed.

transporter and protease with percentages 13.3, 13.3 and 20 %, respectively. The possible sites of target which the compound may bind to are mostly the targets which are predicted by the software and the probability score are very less that is between 0.153 to 0.097. This makes an inference that the small compound may have high target attraction towards the specific binding site it is directed to [40].

In addition to the above information, we investigated the probable antimicrobial action mechanism of

the final compounds through the docking study. The docking study point out that compound **3b** may have potent antimicrobial activity via ATP-competitive DNA gyrase B inhibition. Obtained 2D and 3D pose showed that compound **3b** interacted with Asp49, Glu50, Asp73 and Gly77 amino acids via water-mediated H-bonds. On the other hand, there were one direct H-bond with Arg76 and also 2 aromatic H-bonds with Gly101 and Thr165 residues. Moreover, compound **3b** have showed van der Waals

interactions with Asn46, Ile78, Pro79, Ile94, Lys103 and Thr165 amino acids. All those connections were envisaged as binding action (**Figure 2**). The role of the GyrB subunit is to provide the energy necessary for the DNA ligation process through ATP hydrolysis, hence, the inhibition of this subunit cause stopping ATP hydrolysis, so bacteria cells lose their functions. In previous studies, Asp73 and Gly101 residues have been described as key amino acids for inhibition of the GyrB subunit [41,42]. However, Gly101 amino acid and the other loop amino acids (Gly102, Lys103, and Phe104) could stabilize the loop region, which reduces the flexibility of the enzyme. Therefore, it can be concluded that be a direct proportion between the strong inhibition potential and those loop amino acids interactions. Moreover, final compounds except **3b**, could not properly enter the cavity, thus, not dock with important amino acids well at GyrB

active pocket (**Figure 3**). For compounds **3c-3h**, It is probably caused by their bulky structures due to phenyl substitutions at the 4th position of the thiazole ring. On the other hand, compound **3a** has a small moiety (methyl) at thiazole ring like **3b**, but its localization was observed like other derivatives. It may be related to the absence of one carbon at the side chain. This absence may have led to unfavorable localization between ligand and protein. Indeed, there weren't any connections between **3a** and Gly101. It's clear that only compound **3b** was found as an appropriate analog to inhibit DNA gyrase via blocking ATP hydrolysis. In further studies, the advanced experimental studies should be improved. Although this potential hit was determined, this compound may also be useful for developing more powerful molecules.

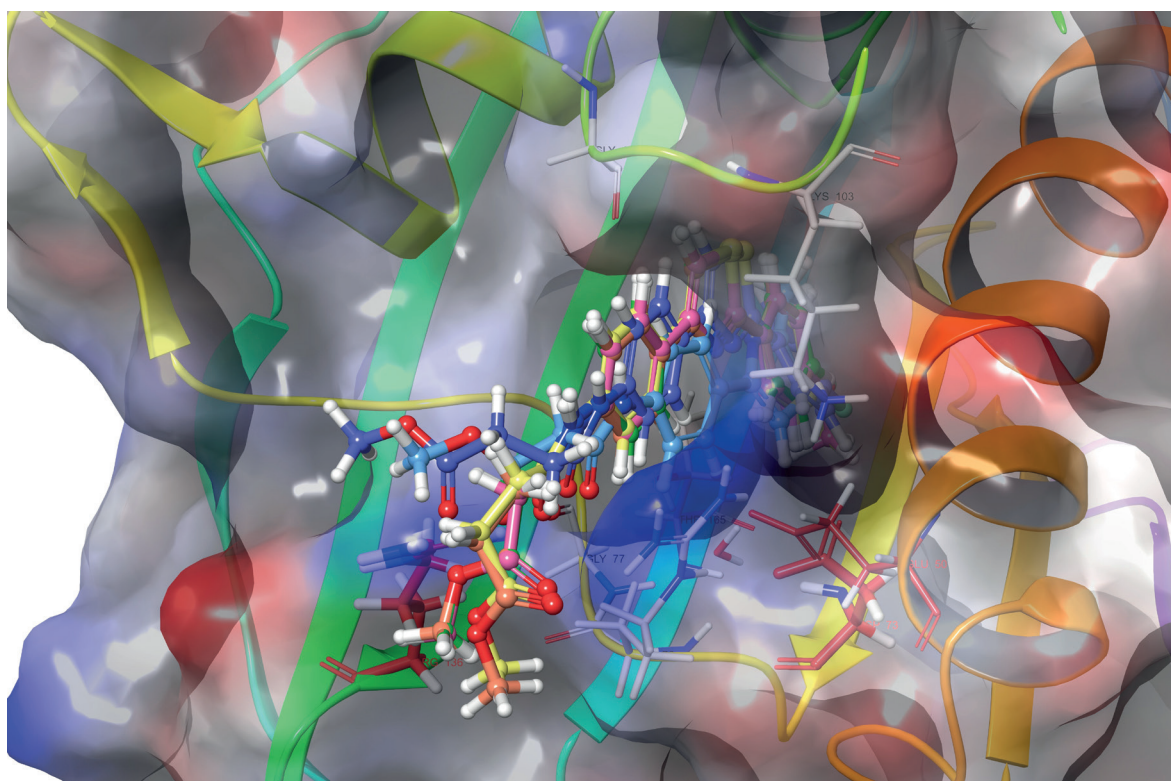


Figure 3. The final compounds at DNA gyrase active region

Table 4. Toxic profiles of the compounds (**3a-h**) predicted using pkCSM software

	3a	3b	3c	3d	3e	3f	3g	3h
Ames toxicity	No	No	No	No	No	No	No	No
Max. tolerated dose (human, log mg/kg/day)	-0.156	-0.198	-0.054	-0.025	-0.118	-0.107	-0.102	-0.089
hERG I inhibitor	No	No	No	No	No	No	No	No
hERG II inhibitor	No	No	No	No	No	No	No	No
Oral Rat Acute Toxicity (LD ₅₀ , mol/kg)	2.686	2.643	2.243	2.147	2.447	2.37	2.38	2.304
Oral Rat Chronic Toxicity (LOAEL, log mg/kg_bw/day)	1.178	1.169	0.911	0.909	1.01	0.936	0.974	0.901
Hepatotoxicity	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes
Skin sensitisation	No	No	No	No	No	No	No	No
<i>T. pyriformis</i> toxicity (log µg/L)	0.887	1.127	0.567	0.562	0.46	0.451	0.462	0.455
Minnow toxicity (log mM)	0.881	0.719	-0.382	-0.511	-0.837	-0.966	-0.631	-0.76

Toxic profiles of the compounds were also predicted using pkCSM software and the results were displayed in **Table 4**. The website can provide details of toxicology effects in the fields of AMES toxicity, human maximum tolerance dose, hERG-I inhibitor, hERG-II (human Ether-à-go-go-Related Gene) inhibitor, LD₅₀ (lethal dose), chronic oral rat toxicity, hepatotoxicity, skin toxicity, *T. pyriformis* toxicity and minnow toxicity. None of the compounds possessed Ames toxicity which mean they have no mutagenic potential and also skin sensitization, however they all cause hepatotoxicity. They don't inhibit hERG-I and hERG-II which is a satisfying finding that provides non-cardiotoxic profile to the compounds [43]. Acute oral rat toxicity (LD₅₀) of the compounds were found in between 2.147-2.686 mol/kg, whereas chronic oral rat toxicity (LOAEL) were found 0.901-1.178 log mg/kg_bw/day. Toxicity to *T. pyriformis* which is a ciliate protozoan was found between the range of 0.451-1.127 log µg/L.

Compounds **3a** and **3b** including alkyl function (methyl) which differ from others showed the highest toxicity to the protozoan and minnow (fish species), too.

4. CONCLUSION

New eight thiazole compounds (**3a-h**) were synthesized containing ester and amide function using known synthetic procedures and starting from Hantzsch thiazole ring cyclization reaction. Final ester compounds were evaluated for their physicochemical/pharmacokinetic and druglikeness properties. All compounds were found to be in compliance with such rules, Lipinski and Ghose. Kinases were *in silico* determined as the main biological target for these compounds and toxicity profile of them was identified at the acceptable ranges, virtually. Compound **3b** was found a hit molecule to inhibit DNA gyrase B subunit enzyme.

Ethical approval

Not applicable, because this article does not contain any studies with human or animal subjects.

Author contribution

Concept: LY, YÖ; Design: LY, YÖ; Supervision: YÖ; Materials: LY; Data Collection and/or Processing: LY, AEE; Analysis and/or Interpretation: LY, YÖ, AEE; Literature Search: LY, AEE; Writing: LY, AEE; Critical Reviews: LY, AEE.

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Conflict of interest

The authors declare that there is no conflict of interest.

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Design, synthesis and *in vitro* COX inhibitory profiles of a new series of tetrazole-based hydrazones

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ABSTRACT

Inhibition of cyclooxygenases (COXs) by selective and nonselective inhibitors is a favorable approach for pharmacologic intervention in a variety of disorders such as cancer. For this purpose, a new class of tetrazole-hydrazone hybrids (**1-12**) was designed. A facile and efficient three-step procedure was applied for the preparation of compounds **1-12**, which were tested for their inhibitory activities towards cyclooxygenases (COXs) by means of an *in vitro* colorimetric inhibitor screening method. The most potent and selective COX-1 inhibitors were determined as 2-[(1-methyl-1*H*-tetrazol-5-yl)thio]-*N*'-(4-(piperidin-1-yl)benzylidene)acetohydrazide (**1**) (40.88±2.79%) and 2-[(1-methyl-1*H*-tetrazol-5-yl)thio]-*N*'-(4-(morpholin-4-yl)benzylidene)acetohydrazide (**2**) (39.80±2.78%), whereas the most potent and selective COX-2 inhibitor was found as 2-[(1-phenyl-1*H*-tetrazol-5-yl)thio]-*N*'-(4-(pyrrolidin-1-yl)benzylidene)acetohydrazide (**10**) (42.38±1.16%). In general, 1-methyl-1*H*-tetrazole moiety resulted in selective COX-1 inhibition, whereas 1-phenyl-1*H*-tetrazole moiety gave rise to preferential COX-2 inhibition.

Keywords: Cyclooxygenase, hydrazone, synthesis, tetrazole

1. INTRODUCTION

Prostaglandins (PGs) and thromboxane A₂ (TXA₂) belong to a class of lipid signaling molecules derived from arachidonic acid (AA) [1]. Biosynthesis of PGs and TXA₂ is catalyzed by two cyclooxygenase (COX) isoforms, COX-1 and COX-2 [1,2], which have similar structural and catalytic properties [3]. However, both enzymes differ in terms of their regulation of expression, tissue distribution, and pharmacological functions [3,4]. COX-1 is constitutively expressed in many tissues (e.g., thrombocytes, gastrointestinal tract, lung, kidney, liver, brain, spleen and prostate), where COX-1-derived prostanoids participate in homeostatic functions. Under normal physiological conditions,

COX-1-derived prostanoids play a fundamental role in the maintenance of platelet function, renal perfusion, and cytoprotective levels of PGs in the gastric mucosa [3-6]. On the contrary, COX-2 is frequently referred to as the inducible isoform, responsible for enhanced PG production in response to inflammatory and mitogenic stimuli [4-7]. The up-regulation and overexpression of COX-2 lead to inflammation, loss of apoptosis, uncontrolled cell proliferation, metastasis, and angiogenesis finally resulting in cancer [8].

Today, the concept of “constitutive” and “inducible” isoforms has been challenged [4] by mounting evidence pointing out that COX-2 is also constitutively expressed in some tissues [9]

and COX-1 is up-regulated in various pathological conditions including thrombosis, atherosclerosis, and tumorigenesis [10].

Inhibition of COXs by a variety of selective and nonselective inhibitors is a promising approach for pharmacologic intervention in many pathological conditions [11,12].

Hydrazides-hydrazones are not only useful intermediates for the synthesis of various heterocyclic compounds but also frequently occurring motifs in drug molecules [13-16] due to their diverse pharmacological activities such as anti-inflammatory, analgesic, antidepressant, anticonvulsant, antitumor, antimycobacterial, and antimalarial activities [13-21]. In particular, the anti-inflammatory potency of hydrazones is well-documented and some of them exert their action through the inhibition of COXs [17-21].

Tetrazole scaffold is reported to be among the top 25 most commonly utilized nitrogen heterocycles in pharmaceutical agents [22]. The tetrazole ring is a well-known bioisostere of the carboxylic acid group [23] and therefore this core is frequently used for the design of promising COX inhibitors for the management of inflammation [24,25].

Prompted by aforementioned data [17-25], the facile and efficient synthesis of a novel series of tetrazole-based hydrazones (**1-12**) was performed and *in vitro* studies were conducted to assess their inhibitory effects on COXs.

2. MATERIALS AND METHODS

2.1. Chemicals and Equipments

All chemicals were purchased from commercial suppliers and were used without further purification. The melting points (M.p.) of the compounds were detected on a MP90 digital melting point apparatus (Mettler Toledo, Ohio, USA) and are uncorrected. Rotavapor® R-100 (Büchi, Switzerland) was used for the evaporation of the solvents. Nuclear Magnetic Resonance (NMR, ¹H and ¹³C) spectra were recorded on a Bruker spectrometer (Bruker, Billerica, MA, USA). Chemical shifts were reported in parts per

million (ppm) and the coupling constants (*J*) were expressed in Hertz (Hz). and High Resolution Mass Spectrometry (HRMS) spectra were obtained from a Shimadzu LCMS-IT-TOF system (Shimadzu, Kyoto, Japan).

2.2. General procedure for the preparation of ethyl 2-[(1-methyl/phenyl-1*H*-tetrazol-5-yl)thio]acetate

A mixture of 5-mercapto-1-methyl/phenyl-1*H*-tetrazole (0.05 mol) and ethyl chloroacetate (0.05 mol) in the presence of potassium carbonate (0.05 mol) in acetone was refluxed for 10 h. Upon completion of the reaction, the solvent was evaporated. The crude product was solved in water and then extracted with diethyl ether. After extraction, the solvent was evaporated [26,27].

2.3. General procedure for the preparation of 2-[(1-methyl/phenyl-1*H*-tetrazol-5-yl)thio]acetohydrazide

A mixture of the appropriate ester (0.05 mol) and hydrazine hydrate (0.1 mol) in ethanol was stirred at room temperature for 3 h [26,27]. Upon completion of the reaction, the precipitate was collected by filtration and dried.

2.4. General procedure for the preparation of 2-[(1-methyl/phenyl-1*H*-tetrazol-5-yl)thio]-*N'*-(4-substituted benzylidene)acetohydrazides (**1-12**)

A mixture of the appropriate hydrazide (0.01 mol) and aromatic aldehydes (0.01 mol) was refluxed in ethanol for 7 h. Upon completion of the reaction, the precipitate was collected by filtration and dried. The product was crystallized from ethanol.

2.4.1. 2-[(1-Methyl-1*H*-tetrazol-5-yl)thio]-*N'*-(4-(piperidin-1-yl)benzylidene)acetohydrazide (**1**)

M.p.: 138.4 °C. Yield: 84%.

¹H NMR (300 MHz, DMSO-*d*₆): 1.57 (brs, 6H), 3.25 (brs, 4H), 3.98 (s, 3H), 4.14 and 4.57 (2s, 2H), 6.95 (d, *J*= 8.91 Hz, 2H), 7.49 (dd, *J*= 2.37 Hz, 2.49 Hz, 8.94 Hz, 2H), 7.89 and 8.03 (2s, 1H), 11.51 (s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 24.39 (CH₂), 25.42 (2CH₂), 34.13 (CH₃), 36.37 (CH₂), 48.81 (CH₂),

48.87 (CH₂), 114.97 (CH), 115.09 (CH), 123.40 (C), 128.64 (CH), 128.93 (CH), 145.05 (CH), 148.20 (C), 152.84 (C), 167.96 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₆H₂₁N₇OS: 360.1601, found: 360.1596.

2.4.2. 2-[(1-Methyl-1H-tetrazol-5-yl)thio]-N'-(4-(morpholin-4-yl)benzylidene)acetohydrazide (2)

M.p.: 199.2 °C. Yield: 90%.

¹H NMR (300 MHz, DMSO-*d*₆): 3.19-3.21 (m, 4H), 3.72-3.75 (m, 4H), 3.98 (s, 3H), 4.15 and 4.58 (2s, 2H), 6.98 (d, *J* = 8.79 Hz, 2H), 7.54 (dd, *J* = 2.43 Hz, 2.58 Hz, 8.90 Hz, 2H), 7.92 and 8.06 (2s, 1H), 11.54 (s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 34.11 (CH₃), 36.34 (CH₂), 47.86 (CH₂), 47.92 (CH₂), 66.40 (2CH₂), 114.74 (CH), 114.83 (CH), 124.57 (C), 128.56 (CH), 128.84 (CH), 144.88 (CH), 148.04 (C), 152.64 (C), 168.05 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₅H₁₉N₇O₂S: 362.1394, found: 362.1387.

2.4.3. 2-[(1-Methyl-1H-tetrazol-5-yl)thio]-N'-(4-(4-methylpiperazin-1-yl)benzylidene)acetohydrazide (3)

M.p.: 139.2 °C. Yield: 88%.

¹H NMR (300 MHz, DMSO-*d*₆): 2.21 (s, 3H), 2.41-2.43 (m, 4H), 3.21-3.24 (m, 4H), 3.98 (s, 3H), 4.15 and 4.57 (2s, 2H), 6.97 (d, *J* = 8.76 Hz, 2H), 7.52 (dd, *J* = 3.12 Hz, 3.15 Hz, 8.84 Hz, 2H), 7.90 and 8.04 (2s, 1H), 11.52 (s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 34.13 (CH₃), 36.35 (CH₂), 46.21 (CH₃), 47.51 (CH₂), 47.57 (CH₂), 54.87 (2CH₂), 114.89 (CH), 114.98 (CH), 124.08 (C), 128.57 (CH), 128.85 (CH), 144.96 (CH), 148.10 (C), 152.53 (C), 168.01 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₆H₂₂N₈OS: 375.1710, found: 375.1712.

2.4.4. 2-[(1-Methyl-1H-tetrazol-5-yl)thio]-N'-(4-(pyrrolidin-1-yl)benzylidene)acetohydrazide (4)

M.p.: 203.4 °C. Yield: 82%.

¹H NMR (300 MHz, DMSO-*d*₆): 1.93-1.97 (m, 4H), 3.25-3.29 (m, 4H), 3.98 (s, 3H), 4.13 and 4.56 (2s, 2H), 6.56 (d, *J* = 8.76 Hz, 2H), 7.48 (dd, *J* = 3.09 Hz, 3.12 Hz, 8.75 Hz, 2H), 7.87 and 8.01 (2s, 1H), 11.42 (s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 25.41 (2CH₂), 34.10 (CH₃), 36.40 (CH₂), 47.69 (2CH₂), 112.00 (2CH), 120.94 (C), 128.84 (CH), 129.13 (CH), 145.63 (CH), 148.81 (C), 149.32 (C), 167.77 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₅H₁₉N₇OS: 346.1445, found: 346.1428.

2.4.5. 2-[(1-Methyl-1H-tetrazol-5-yl)thio]-N'-(4-(1H-imidazol-1-yl)benzylidene)acetohydrazide (5)

M.p.: 191.1 °C. Yield: 88%.

¹H NMR (300 MHz, DMSO-*d*₆): 3.99 (s, 3H), 4.20 and 4.63 (2s, 2H), 7.14 (brs, 1H), 7.76 (d, *J* = 8.76 Hz, 2H), 7.82-7.85 (m, 3H), 8.06 and 8.23 (2s, 1H), 8.35 (brs, 1H), 11.81 and 11.85 (2s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 34.13 (CH₃), 36.19 (CH₂), 118.26 (CH), 120.79 (2CH), 128.86 (CH), 129.08 (CH), 130.63 (CH), 132.76 (C), 136.02 (CH), 138.29 (C), 143.45 (CH), 146.72 (C), 168.60 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₄H₁₄N₈OS: 343.1084, found: 343.1074.

2.4.6. 2-[(1-Methyl-1H-tetrazol-5-yl)thio]-N'-(4-(1H-1,2,4-triazol-1-yl)benzylidene)acetohydrazide (6)

M.p.: 257.9 °C. Yield: 86%.

¹H NMR (300 MHz, DMSO-*d*₆): 3.99 (s, 3H), 4.20 and 4.64 (2s, 2H), 7.88 (d, *J* = 8.64 Hz, 2H), 7.96 (d, *J* = 8.73 Hz, 2H), 8.07 and 8.24 (2s, 1H), 8.27 (s, 1H), 9.37 (s, 1H), 11.84 (s, 1H).

¹³C NMR (75 MHz, DMSO-*d*₆): 34.14 (CH₃), 36.17 (CH₂), 120.00 (2CH), 128.78 (CH), 129.00 (CH), 133.67 (C), 137.99 (C), 142.96 (CH), 143.36 (CH), 146.54 (CH), 153.07 (C), 168.62 (C).

HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₃H₁₃N₉OS: 344.1037, found: 344.1028.

2.4.7. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(piperidin-1-yl)benzylidene)acetohydrazide (7)

M.p.: 189.4 °C. Yield: 83%.

¹H NMR (300 MHz, DMSO-*d*₆): 1.56 (brs, 6H), 3.24 (brs, 4H), 4.28 and 4.69 (2s, 2H), 6.94 (dd, *J* = 2.37 Hz, 2.64 Hz, 8.99 Hz, 2H), 7.50 (dd, *J* = 3.63 Hz, 3.66 Hz, 8.75 Hz, 2H), 7.65-7.68 (m, 5H), 7.91 and 8.06 (2s, 1H), 11.55 and 11.59 (2s, 1H).¹³C NMR (75 MHz, DMSO-*d*₆): 24.38 (CH₂), 25.42 (2CH₂), 36.72 (CH₂), 48.87 (2CH₂), 114.96 (CH), 115.08 (CH), 123.39 (C), 124.91 (2CH), 128.63 (CH), 128.94 (CH), 130.53 (2CH), 131.09 (CH), 133.57 (C), 145.16 (CH), 148.24 (C), 152.84 (C), 167.71 (C).HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₂₁H₂₃N₇OS: 422.1758, found: 422.1756.**2.4.8. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(morpholin-4-yl)benzylidene)acetohydrazide (8)**

M.p.: 251.6 °C. Yield: 89%.

¹H NMR (300 MHz, DMSO-*d*₆): 3.20 (brs, 4H), 3.74 (brs, 4H), 4.27 and 4.69 (2s, 2H), 6.99 (d, *J* = 8.25 Hz, 2H), 7.54 (d, *J* = 6.90 Hz, 2H), 7.69 (brs, 5H), 7.92 and 8.07 (2s, 1H), 11.58 and 11.62 (2s, 1H).¹³C NMR (75 MHz, DMSO-*d*₆): 36.68 (CH₂), 47.91 (2CH₂), 66.40 (2CH₂), 114.75 (CH), 114.84 (CH), 124.55 (C), 124.95 (2CH), 128.56 (CH), 128.84 (CH), 130.54 (2CH), 131.12 (CH), 133.57 (C), 144.99 (CH), 148.24 (C), 152.84 (C), 167.71 (C).HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₂₀H₂₁N₇O₂S: 424.1550, found: 424.1551.**2.4.9. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(4-methylpiperazin-1-yl)benzylidene)acetohydrazide (9)**

M.p.: 209.2 °C. Yield: 87%.

¹H NMR (300 MHz, DMSO-*d*₆): 2.21 (s, 3H), 2.43 (brs, 4H), 3.21-3.23 (m, 4H), 4.27 and 4.69 (2s, 2H), 6.97 (dd, *J* = 2.34 Hz, 2.43 Hz, 8.87 Hz, 2H), 7.52 (dd, *J* = 2.97 Hz, 8.76 Hz, 2H), 7.69-7.71 (m, 5H), 7.91 and 8.06 (2s, 1H), 11.56 and 11.61 (2s, 1H).¹³C NMR (75 MHz, DMSO-*d*₆): 36.67 (CH₂), 46.20(CH₃), 47.55 (2CH₂), 54.85 (2CH₂), 114.89 (CH), 114.99 (CH), 124.07 (C), 124.94 (2CH), 128.57 (CH), 128.86 (CH), 130.54 (2CH), 131.12 (CH), 133.56 (C), 145.06 (CH), 148.12 (C), 152.54 (C), 167.76 (C).HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₂₁H₂₄N₈OS: 437.1867, found: 437.1884.**2.4.10. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(pyrrolidin-1-yl)benzylidene)acetohydrazide (10)**

M.p.: 215.1 °C. Yield: 81%.

¹H NMR (300 MHz, DMSO-*d*₆): 1.95 (brs, 4H), 3.27 (brs, 4H), 4.26 and 4.68 (2s, 2H), 6.57 (d, *J* = 8.67 Hz, 2H), 7.48 (d, *J* = 8.67 Hz, 2H), 7.69 (brs, 5H), 7.88 and 8.03 (2s, 1H), 11.47 and 11.50 (2s, 1H).¹³C NMR (75 MHz, DMSO-*d*₆): 25.41 (2CH₂), 36.73 (CH₂), 47.68 (2CH₂), 112.01 (2CH), 120.92 (C), 124.94 (2CH), 128.83 (CH), 129.14 (CH), 130.53 (2CH), 131.10 (CH), 133.57 (C), 145.72 (CH), 148.83 (C), 149.33 (C), 167.53 (C).HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₂₀H₂₁N₇OS: 408.1601, found: 408.1604.**2.4.11. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(1H-imidazol-1-yl)benzylidene)acetohydrazide (11)**

M.P.: 236.4 °C. Yield: 86%.

¹H NMR (300 MHz, DMSO-*d*₆): 4.32 and 4.75 (2s, 2H), 7.14 (brs, 1H), 7.65-7.71 (m, 5H), 7.76 (d, *J* = 8.70 Hz, 2H), 7.84 (d, *J* = 8.70 Hz, 3H), 8.07 and 8.25 (2s, 1H), 8.36 (brs, 1H), 11.86 and 11.92 (2s, 1H).¹³C NMR (75 MHz, DMSO-*d*₆): 36.54 (CH₂), 118.29 (CH), 120.80 (2CH), 124.92 (2CH), 128.84 (CH), 129.08 (CH), 130.55 (2CH), 131.13 (CH), 132.75 (CH), 132.83 (C), 133.55 (C), 136.02 (CH), 138.30 (C), 143.56 (CH), 146.75 (C), 168.35 (C).HRMS (ESI) (*m/z*): [M+H]⁺ calcd. for C₁₉H₁₆N₈OS: 405.1241, found: 405.1250.**2.4.12. 2-[(1-Phenyl-1H-tetrazol-5-yl)thio]-N'-(4-(1H-1,2,4-triazol-1-yl)benzylidene)acetohydrazide (12)**

M.p.: 240.6 °C. Yield: 84%.

^1H NMR (300 MHz, $\text{DMSO-}d_6$): 4.32 and 4.75 (2s, 2H), 7.67-7.70 (m, 5H), 7.89 (d, $J = 8.73$ Hz, 2H), 7.96 (d, $J = 8.49$ Hz, 2H), 8.09 and 8.26 (2s, 1H), 8.28 (s, 1H), 9.38 (s, 1H), 11.88 (s, 1H).

^{13}C NMR (75 MHz, $\text{DMSO-}d_6$): 36.48 (CH_2), 120.02 (2CH), 124.94 (2CH), 128.77 (CH), 129.01 (CH), 130.56 (2CH), 131.14 (CH), 133.55 (C), 133.66 (C), 138.01 (C), 142.96 (CH), 143.46 (CH), 146.58 (CH), 153.07 (C), 168.36 (C).

HRMS (ESI) (m/z): $[\text{M}+\text{H}]^+$ calcd. for $\text{C}_{18}\text{H}_{15}\text{N}_9\text{OS}$: 406.1193, found: 406.11957

2.5. Biochemistry

2.5.1. Determination of COX inhibitory potency

COX colorimetric inhibitor screening assay kit (item no: 701050), which includes both ovine COX-1 and human recombinant COX-2, is used to measure the peroxidase component of COXs. In this method, the peroxidase activity is detected colorimetrically *via* monitoring the appearance of oxidized *N,N,N',N'*-tetramethyl-*p*-phenylenediamine (TMPD) at 590 nm.

In the current work, this kit was applied to determine the inhibitory activities of compounds **1-12** (at 100 μM) towards both COXs according to the

manufacturer's instructions (Cayman, Ann Arbor, MI, USA). All measurements were performed in triplicate and the data were expressed as mean \pm SD. SC-560 (at 1 μM) was used as a selective COX-1 inhibitor, whilst rofecoxib (at 10 μM) was used as a selective COX-2 inhibitor.

3. RESULTS AND DISCUSSION

3.1. Chemistry

The synthesis of the hitherto unreported compounds (**1-12**) was performed as depicted in Figure 1. Ethyl 2-[(1-methyl/phenyl-1*H*-tetrazol-5-yl)thio]acetate was prepared *via* the treatment of 5-mercapto-1-methyl/phenyl-1*H*-tetrazole with ethyl chloroacetate in the presence of potassium carbonate. The reaction of this ester with hydrazine hydrate yielded 2-[(1-methyl/phenyl-1*H*-tetrazol-5-yl)thio]acetohydrazide, which underwent a subsequent nucleophilic addition-elimination reaction with aromatic aldehydes affording compounds **1-12**. Their structures were confirmed by ^1H and ^{13}C NMR, HRMS data. In the ^1H NMR spectra of compounds **1-12**, the signal due to $\text{CH}=\text{N}$ proton was detected in the region 7.87-8.26 ppm as two singlets. The signal due to N-H proton appeared in the region 11.42-11.92 ppm as a singlet or two singlets. The

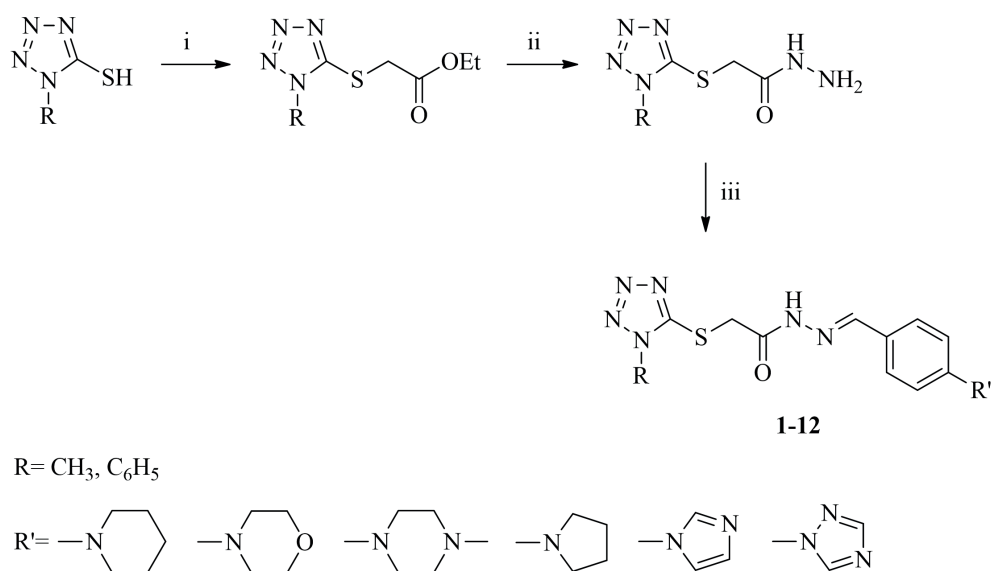


Figure 1. The synthetic route for the preparation of compounds **1-12**. Reagents and conditions: (i) $\text{ClCH}_2\text{COOEt}$, K_2CO_3 , acetone, reflux, 10h; (ii) $\text{NH}_2\text{NH}_2 \cdot \text{H}_2\text{O}$, ethanol, rt, 3h; (iii) ArCHO , ethanol, reflux, 7h.

signal due to the S-CH₂ protons was observed as two singlets in the region 4.13-4.75 ppm. In the ¹H NMR spectra of compounds **1-6**, the protons of the methyl group attached to the N¹ atom of the tetrazole ring gave rise to a singlet peak at 3.98-3.99 ppm. In the ¹³C NMR spectra of compounds **1-12**, the signal due to the CH=N carbon was observed at 143.45-146.58 ppm, whereas the signal due to the C=O carbon was detected in the region 167.53-168.62 ppm. The S-CH₂ carbon gave rise to a singlet peak at 36.17-36.73 ppm. In the ¹³C NMR spectra of compounds **1-6**, the methyl carbon attached to the N¹ atom of the tetrazole ring gave rise to a singlet peak at 34.10-34.14 ppm. The HRMS data were also in agreement with the proposed structures of compounds **1-12**.

3.2. In vitro COX inhibition

A colorimetric assay was employed to assess the COX inhibitory profiles of compounds **1-12** (Table 1). Compounds **7** and **11** did not show any inhibitory activity towards COXs. Compounds **3**, **8**, **9** and **12**

showed inhibitory activity towards both COXs. Among them, compound **3** preferentially inhibited COX-1, whilst compounds **8** and **9** preferentially inhibited COX-2. Compound **12** showed nonselective COX inhibitory activity and its inhibitory effects on COX-1 and COX-2 were found to be insignificant. The 1,2,4-triazole ring at the 4th position of the benzylidene motif caused a diminution in COX inhibitory potency.

The most potent and selective COX-1 inhibitors were found as compounds **1** (40.88±2.79%) and **2** (39.80±2.78%) carrying 1-methyl-1H-tetrazole moiety as compared to SC-560 (97.36±2.62%). These compounds did not show any inhibitory activity towards COX-2. The replacement of the methyl group of compound **1** with the phenyl moiety of compound **7** resulted in the loss of COX-1 inhibitory activity as well as COX-2 inhibitory potency. The replacement of the methyl group of compound **2** with the phenyl moiety of compound **8** led to a decrease in COX-1 inhibitory potency

Table 1. COX inhibition (%) caused by compounds **1-12** and reference agents

Compound (100 μM)	R	R'	Inhibition%	
			COX-1	COX-2
1	CH ₃	Piperidin-1-yl	40.88±2.79	----
2	CH ₃	Morpholin-4-yl	39.80±2.78	----
3	CH ₃	4-Methylpiperazin-1-yl	27.97±2.69	12.45±1.22
4	CH ₃	Pyrrolidin-1-yl	35.36±3.93	----
5	CH ₃	1H-Imidazol-1-yl	33.41±3.09	----
6	CH ₃	1H-1,2,4-Triazol-1-yl	19.55±1.96	----
7	C ₆ H ₅	Piperidin-1-yl	----	----
8	C ₆ H ₅	Morpholin-4-yl	14.37±2.31	30.26±0.76
9	C ₆ H ₅	4-Methylpiperazin-1-yl	21.77±1.58	38.82±0.24
10	C ₆ H ₅	Pyrrolidin-1-yl	----	42.38±1.16
11	C ₆ H ₅	1H-Imidazol-1-yl	----	----
12	C ₆ H ₅	1H-1,2,4-Triazol-1-yl	9.86±3.06	12.46±1.46
SC-560 (1 μM)	-	-	97.36±2.62	ND
Rofecoxib (10 μM)	-	-	ND	98.36±1.86

ND: Not determined.

(14.37±2.31%) and a substantial increase in COX-2 inhibitory activity (30.26±0.76%).

The most potent COX-2 inhibitors were found as compounds **10** (42.38±1.16%) and **9** (38.82±0.24%) carrying 1-phenyl-1*H*-tetrazole moiety as compared to rofecoxib (98.36±1.86%). Compound **9** also inhibited COX-1 (21.77±1.58%), whilst compound **10** did not show any inhibitory potency on COX-1.

In general, 1-methyl-1*H*-tetrazole moiety gave rise to selective COX-1 inhibition, whereas 1-phenyl-1*H*-tetrazole moiety led to preferential COX-2 inhibition. It can be concluded that the substituent at the 1st position of the tetrazole ring is of great importance for selectivity. The replacement of the alkyl (methyl) group with the bulky aryl moiety shifted the selective inhibition from COX-1 to COX-2.

4. CONCLUSION

In this paper, we conducted a simple and efficient protocol for the preparation of a series of tetrazole-based hydrazones (**1-12**), which were investigated for their inhibitory effects on COXs at 100 µM using an *in vitro* colorimetric assay. Based on *in vitro* experimental data, compounds **1** (40.88±2.79%) and **2** (39.80±2.78%) carrying 1-methyl-1*H*-tetrazole moiety were found as the most potent and selective COX-1 inhibitors. On the other hand, compound **10** (42.38±1.16%) carrying 1-phenyl-1*H*-tetrazole moiety was determined as the most potent and selective COX-2 inhibitor in this series. This work could represent a rational guideline for further structural modifications to generate a new class of tetrazole-based hydrazones endowed with selective COX-1 or COX-2 inhibitory activity.

Ethical approval

Not applicable, because this article does not contain any studies with human or animal subjects.

Author contribution

Concept: MDA, ZAK and AÖ; Design: MDA and ZAK; Materials: MDA, BS, HET, ZAK and AÖ; Data Collection and/or Processing: MDA, BS and HET; Analysis and/or Interpretation: MDA and HET; Literature Search: MDA; Writing: MDA; Critical Reviews: MDA, BS, HET, ZAK and AÖ.

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Conflict of interest

The authors declare that there is no conflict of interest.

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Chaerophyllum libanoticum Boiss. Et Kotschy: The fruit essential oil, composition, skin-whitening and antioxidant activities

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ABSTRACT

This study was aimed to evaluate the essential oil composition of *Chaerophyllum libanoticum* fruits and its potential uses in the cosmetic industry. The essential oil was analyzed by Gas Chromatography and Gas Chromatography-Mass Spectrometry (GC-GC/MS) systems, simultaneously. The yield of essential oil was calculated as 0.22 % (v/w). Major components of the oil were characterized as limonene (26.7%), *p*-cymene (25.5%), and β -phellandrene (7.0%). In addition, antioxidant and antityrosinase activities of the essential oil were evaluated. The oil exhibited moderate antioxidant activity (TEAC). In the DPPH[•] assay, the oil was tested at 5 mg/mL concentration, and the inhibition ratio was calculated as 31.3 \pm 1.1%. At 1 mg/mL of concentration, TEAC (mmol/L) value was determined as 0.027 \pm 0.008. As evidence to its skin whitening properties, the oil inhibited the tyrosinase 17.7 \pm 1.6 % at 1 mg/mL..

Keywords: *Chaerophyllum libanoticum*, Apiaceae, Essential oil, Antioxidant, Antityrosinase

1. INTRODUCTION

Apiaceae is considered as one of the most important families due to its economic value in the field of pharmaceutical, cosmetic, flavor, and fragrance industries and is represented by about 3780 species in 434 genera [1,2]. The genus *Chaerophyllum* L. is represented in the Flora of Turkey by 15 species [3], and the members of the genus have a distinctive aromatic character [4]. *Chaerophyllum* Boiss. et Kotschy. is known as “Mentik” in Southern parts of Turkey, and is consumed as food like some other *Chaerophyllum* species [5].

Essential oil components are responsible for the characteristic fragrance of *Chaerophyllum* species,

and various studies have been reported on the volatile composition of different parts and origins [6-11]. Previous investigations on biological properties of *Chaerophyllum* species revealed that they possess various biological activities such as antimicrobial [7,10], antioxidant [7,10], anti-inflammatory [7], angiotensin-converting enzyme (ACE) inhibition [12], glutathione-S-transferase inhibition [13], cytotoxic properties [14].

Essential oils have been used in cosmetic preparations due to their health benefit effects in skin disorders related to their antioxidant, antimicrobial and anti-inflammatory activities. Recently, tyrosinase, a key enzyme in the production of melanin, has become

one of the most important targets in the cosmetic industry [15]. Overproduction of melanin leads to hyperpigmentation, age spots, melasma, and this process might be related to increased oxidative stress on the skin. Essential oils, their components and essential oil-bearing plants are considered natural cosmetic ingredients. They may also possess antioxidant and antityrosinase properties.

This present study was aimed to characterize volatile constituents of *Chaerophyllum libanoticum* Boiss. Et Kotschy fruit essential oil collected from Osmaniye. The essential oil was analyzed by GC-FID and GC/MS system, simultaneously. Furthermore, antioxidant and antityrosinase activities were determined by *in vitro* methods.

2. MATERIALS AND METHODS

2.1. Plant Material

Chaerophyllum libanoticum was collected from Osmaniye; Zorkun, Mitisin Plateau at an altitude 1340 m in Turkey, on October 14, 2018 (Ahmet Duran, 10733).

2.2. Chemicals and Reagents

All chemicals and solvents were of high purity and at least of analytical grade. 1,1-diphenyl-2-picrylhydrazyl (DPPH•) (Aldrich), 2,2'-azinobis-(3-ethylbenzothiazoline)-6-sulfonic acid (Sigma-Aldrich), Trolox ((*S*)-6-hydroxy-2,5,7,8-tetramethylchromane-2-carboxylic acid, Sigma-Aldrich), gallic acid (Sigma), ascorbic acid (Sigma-Aldrich), L-DOPA (Sigma), kojic acid (Sigma-Aldrich), tyrosinase from mushroom (Sigma-Aldrich), dipotassium hydrogen phosphate (Merck), sodium dihydrogen phosphate dihydrate (Merck), sodium sulfate anhydrous (Sigma-Aldrich), potassium persulfate (Sigma-Aldrich), methanol (Merck), ethanol (Sigma-Aldrich), and *n*-hexane (Sigma-Aldrich) were purchased.

2.3. Isolation of Essential Oil

The air-dried fruits of *Chaerophyllum libanoticum* were subjected to hydrodistillation, using a Clevenger-type apparatus for 3h. The obtained oils were kept at +4°C in the dark until the experiments.

2.4. GC-FID Analysis

GC analyses were performed using an Agilent 6890N GC system. FID temperature was set to 300°C and the same operational conditions were applied to a triplicate of the same column used in GC/MS analyses. Simultaneous auto injection was employed to obtain equivalent retention times. Relative percentages of the separated compounds were calculated from integration of the peak areas in the GC-FID chromatograms.

2.5. GC/MS Analysis

The GC/MS analysis was carried out with an Agilent 5975 GC-MSD system (Agilent, USA; SEM Ltd., Istanbul, Turkey). Innnowax FSC column (60m x 0.25mm, 0.25µm film thickness) was used with helium as carrier gas (0.8 mL/min.). GC oven temperature was kept at 60°C for 10 min and programmed to 220°C at a rate of 4°C/min, and kept constant at 220°C for 10 min and then programmed to 240°C at a rate of 1°C/min. Split ratio was adjusted 40:1. The injector temperature was at 250°C. The interphase temperature was at 280°C. MS were taken at 70 eV. Mass range was from m/z 35 to 450.

2.6. Identification of Compounds

The components of essential oils were identified by comparison of their mass spectra with those in the in-house Baser Library of Essential Oil Constituents, Adams Library [16], MassFinder Library [17], Wiley GC/MS Library [18], and confirmed by comparison of their retention indices. These identifications were accomplished by comparison of retention times with authentic samples or by comparison of their relative retention index (RRI) to a series of *n*-alkanes. Alkanes were used as reference points in the calculation of relative retention indices (RRI) [19]. Relative percentage amounts of the separated compounds were calculated from FID chromatograms.

2.7. DPPH Radical Scavenging Activity

DPPH• scavenging activity was determined by the method of Ağalar and Temiz, 2021 [20]. 0.2 mM DPPH solution was prepared in a dark environment with methanol. The experiment was carried out in 96-well microplates. Incubation was carried out for

30 minutes at room temperature and in the dark (A). For the sample control, 100 µL of methanol was added instead of DPPH (B). 0.1 mM DPPH as a blank control (C), methanol (D) as solvent control, and Vitamin C and gallic acid as a positive control were used. Absorbances were measured at 517 nm, and inhibition percentages were calculated according to the formula below (Eq.1).

$$\% \text{ inhibition: } [(C-D) - (A-B) / (C-D)] \times 100 \quad (\text{Eq.1})$$

2.8. Trolox Equivalent Antioxidant Capacity (TEAC)

The trolox equivalent antioxidant capacity of the essential oils was performed *via* the method described by Re et al. (1999) using ABTS^{•+} radical [21]. 7 mM ABTS^{•+} radical and 2.5 mM potassium persulfate were dissolved in water and kept in the dark for 16h at room temperature. Essential oils were dissolved in ethanol at 1 mg/mL concentration. Trolox were prepared in absolute ethanol with 2.5, 2, 1.5, 1, 0.5, 0.25, 0.125 and 0.0625 mM concentrations. 990 µL ABTS^{•+} solution and 10 µL samples were mixed. Absorbances were measured at 734 nm for 30 min, and the results were expressed as Trolox equivalent antioxidant capacity (mmol/L Trolox).

2.9. Tyrosinase Inhibition

The tyrosinase inhibitory activity of the oil was evaluated by using L-DOPA [20]. Tyrosinase and L-DOPA were prepared in the phosphate buffer (100 mM, pH 6.8). Essential oils were dissolved in buffer containing 5% DMSO. For each sample solution, four wells designated as A, B, C and D each contained a reaction mixture (40 µL) as follows: A, 20 µL buffer (pH 6.8) and 20 µL tyrosinase (200 U/mL); B, 40 µL buffer; C, 20 µL tyrosinase (200 U/mL) and 20 µL sample; D, 20 µL sample and 20 µL buffer. The contents of each well were mixed and incubated at 37 °C for 10 min. Then, 5 mM of L-DOPA (160 µL) was added. After second incubation at 37 °C for 10 min, the absorbance at 475 nm of each well was measured. The percentage inhibition of the tyrosinase activity was calculated by the following equation 2.

$$\% \text{ inhibition: } [(A-B) - (C-D) / (A-B)] \times 100 \quad (\text{Eq. 2})$$

2.10. Statistical Analysis

All the experiments were carried out in triplicate, and data were expressed as means ± standard deviation (SD). Statistical analysis were performed using Sigmaplot 14.0 software (Systat Software, Inc., San Jose, CA, USA). IC₅₀ values were calculated by regression analysis.

3. RESULTS AND DISCUSSION

3.1. Chemical Composition of *C. libanoticum* Oil

The essential oil obtained from the fruits of *C. libanoticum* was analyzed by GC-GC/MS systems, simultaneously. The essential oil yield was calculated as 0.22% (v/w).

Sixty-four constituents were characterized as representing 95.9% of the essential oil. Limonene (26.7%), *p*-cymene (25.5%), and β-phellandrene (7.0%) were found to be the most abundant compounds.

β-Pinene (5.1%), cryptone (4.1%), hexadecanoic acid (3.4%), α-pinene (2.7%), octanal (2.0%), γ-terpinene (1.4%), *trans*-pinocarveol (1.3%), and myrtenol (1.0%) were present in relatively high amounts (Table 1).

In the essential oil, monoterpene hydrocarbons (69.9%) were the most abundant group. Also, the percentage of oxygenated monoterpenes (12.7%) and other compound groups (11.5%) were considerable. In addition, sesquiterpene hydrocarbons (0.7%) and oxygenated sesquiterpenes (1.0%) were present in relatively low amounts. (Figure 1).

Several studies identified essential oil compositions of *Chaerophyllum* species. Previous reports on volatile constituents of *Chaerophyllum* species growing wild in Turkey are given in Table 2. Studies were mainly focused on the aerial parts and fruits of these species. Essential oils obtained from different species were characterized with various major components. Demirci et al. (2007) detected that the fruit essential oil was rich in limonene (15.9 %) and β-phellandrene (17.6 %) [10]. Our results revealed

Table 1. Chemical composition of the fruit essential oil of *C. libanoticum*

RRI	Compounds	%	IM
1032	α -Pinene	2.7	t _R , MS
1118	β -Pinene	5.1	t _R , MS
1132	Sabinene	0.3	t _R , MS
1174	Myrcene	0.1	t _R , MS
1183	Pseudolimonene	0.3	MS
1194	Heptanal	0.2	MS
1203	Limone	26.7	t _R , MS
1218	β-Phellandrene	7.0	t _R , MS
1246	(Z)- β -Ocimene	0.2	t _R , MS
1255	γ -Terpinene	1.4	t _R , MS
1280	p-Cymene	25.5	t _R , MS
1290	Terpinolene	0.4	t _R , MS
1296	Octanal	2.0	t _R , MS
1400	Nonanal	0.1	MS
1435	γ -Campholene aldehyde	0.1	MS
1452	p-Cymenene	0.2	MS
1458	cis-1,2-Limonene epoxide	0.1	MS
1463	Heptanol	0.1	t _R , MS
1468	trans-1,2-Limonene epoxide	0.4	MS
1474	trans-Sabinene hydrate	0.1	t _R , MS
1476	4,8-Epoxyterpinolene	0.7	MS
1499	α -Campholene aldehyde	0.5	MS
1535	β -Bourbonene	0.2	t _R , MS
1548	(E)-2-Nonenal	0.8	MS
1553	Linalool	0.5	t _R , MS
1565	8,9-Limonene epoxide-I	0.3	MS
1570	8,9-Limonene epoxide-II	0.1	MS
1571	trans-p-Menth-2-en-1-ol	0.7	MS
1586	Pinocarvone	0.2	MS
1607	Thymol methyl ether	0.1	t _R , MS
1611	Terpinen-4-ol	0.3	t _R , MS
1617	trans-Dihydrocarvone	tr	t _R , MS
1638	trans-p-Menth-2,8-dien-1-ol / cis-p-Menth-2-en-1-ol	0.7	MS
1648	Myrtenal	0.4	MS
1655	(E)-2-Decenal	0.3	MS
1670	trans-Pinocarveol	1.3	t _R , MS
1678	cis-p-Menth-2,8-dien-1-ol	0.5	MS
1690	Cryptone	4.1	MS
1706	α -Terpineol	0.4	t _R , MS
1744	Phellandral	0.7	MS
1751	Carvone	0.8	t _R , MS
1783	β -Sesquiphellandrene	0.2	MS
1786	ar-Curcumene	0.3	MS

Table 1. Continued

RRI	Compounds	%	IM
1804-	Myrtenol	1.0	MS
1811	trans-p-Mentha-1(7),8-dien-2-ol	0.3	MS
1815	p-Mentha-1,3-dien-7-al	0.1	MS
1845	trans-Carveol	0.6	t _R , MS
1864	p-Cymen-8-ol	0.9	t _R , MS
1867	cis-Carveol	0.3	t _R , MS
1896	cis-p-Mentha-1(7),8-dien-2-ol	0.1	MS
2029	Perilla alcohol	0.2	MS
2113	Cumin alcohol	0.3	t _R , MS
2131	Hexahydrofarnesyl acetone	0.2	t _R , MS
2144	Spathulenol	0.2	t _R , MS
2198	Thymol	0.1	t _R , MS
2239	Carvacrol	0.2	t _R , MS
2241	ar-Turmerol	0.2	MS
2278	Torilenol	0.1	MS
2296	Myristicine	0.1	MS
2369	Eudesma-4(15), 7-dien-1 β -ol	0.3	MS
2392	Caryophyllenol II	0.2	MS
2931	Hexadecanoic acid	3.4	MS
<i>Grouped compounds (%)</i>			
<i>Monoterpene hydrocarbons</i>			69.9
<i>Oxygenated monoterpenes</i>			12.8
<i>Sesquiterpene hydrocarbons</i>			0.7
<i>Oxygenated sesquiterpenes</i>			1.0
<i>Others</i>			11.5
TOTAL %			95.9

RRI: Relative retention indices calculated against n-alkanes; %: calculated from the FID chromatograms; tr: Trace (<0.1 %). Identification method (IM): t_R, identification based on the retention times (t_R) of genuine compounds on the HP Innowax column; MS, identified on the basis of computer matching of the mass spectra with those of the in-house Baser Library of Essential Oil Constituents, Adams, MassFinder and Wiley libraries and comparison with literature data.

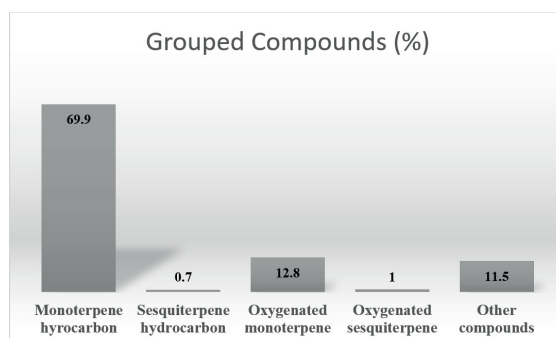


Figure 1. Compound group distribution of the essential oil

Table 2. Main constituents of the essential oils of *Chaerophyllum* species growing wild in Turkey

Plant	Collection place	Parts	Main constituents	Reference
<i>C. aksekiense</i>	Antalya	Fruits	Heptacosane (10.1%), humulene epoxide II (7.8%), (<i>E</i>)- β -farnesene (6.2%), caryophyllene oxide (6.0%),	[6]
<i>C. aromaticum</i>	İstanbul	Aerial parts	Sabinene (28.1%), terpinolene (16.7%), γ -terpinene (16.1%)	[7]
<i>C. byzantinum</i>	Bursa	Aerial parts	Sabinene (30.0%), <i>p</i> -cymen-8-ol (16.0%), terpinolene (11.5%)	[8]
<i>C. crinitum</i>	Bitlis	Aerial parts	(<i>E</i>)- β -Ocimene (38.1%), terpinolene (12.7%)	[9]
<i>C. libanoticum</i>	Osmaniye	Fruits	Limonene (15.9 %), β -phellandrene (17.6 %)	[10]
<i>C. macrospermum</i>	Hakkari	Aerial parts	Terpinolene (21.4%), myristicin (18.9%), <i>p</i> -cymen-8-ol (11.9%)	[9]

that the *p*-cymene ratio was higher than that of β -phellandrene. Moreover, Gnannadi et al. (2011) reported that the main constituents of the essential oil obtained from flowering aerial parts of *C. macropodium* were *trans*- β -ocimene (34.5%), *trans*- β -farnesene (11.8%), *cis*- β -ocimene (10.4%), and *p*-cymene [11]. A study on *C. aromaticum* showed that γ -terpinene (21.0-33.8%), β -phellandrene (14.3-30.0%) and β -pinene (14.3-17.8%) were the main compounds in the fruit essential oil [22].

3.2. Antioxidant Properties of *C. libanoticum* Oil

The antioxidant activity of the essential oil was investigated by two assays, namely DPPH radical scavenging activity and Trolox equivalent antioxidant capacity (TEAC), and the results are given in Table 3. The essential oil of *C. libanoticum* at a 5 mg/mL concentration inhibited DPPH radical by 31.3 ± 1.1 %. The effects were relatively weak when compared with positive controls (vitamin C, IC_{50} value of 9.3 ± 0.01 μ g/mL, gallic acid, IC_{50} value of 1.93 ± 0.02 μ g/mL). Furthermore, the essential oil presented 0.027 ± 0.008 Trolox equivalent antioxidant capacity at 1 mg/mL concentration.

Demirci et al. (2007) evaluated the DPPH• scavenging activity of *C. libanoticum* fruit essential oil, and they reported the IC_{50} value with >30 mg/mL [10]. Kurkcuoglu et al. (2018) found that the essential oil obtained from aerial parts of *C. aromaticum* inhibited DPPH radical by 2.06% at 20 mg/mL [7]. Ebrahimabadi et al. (2010) evaluated the DPPH

radical scavenging activity of *C. macropodium* leaf and flower's essential oils and extracts. The results revealed that the extracts had higher activity than the essential oils [23].

3.3. Tyrosinase Inhibition of *C. libanoticum* Oil

Tyrosinase is a rate-limiting enzyme for the biosynthesis of melanocytes. Thus, it is considered the primary target for skin hyperpigmentation disorders. In the present study, tyrosinase inhibition of the essential oil was tested at 1 mg/mL of concentration, and it exhibited considerable effect (Table 3). However, the results were found to be weak when compared to standard, kojic acid. To the best of our knowledge, tyrosinase inhibition of the essential oil of *Chaerophyllum* species was evaluated for the first time.

4. CONCLUSION

According to our findings, the essential oil isolated from the fruits of *C. libanoticum* was rich in monoterpene hydrocarbons and oxygenated monoterpenes. The essential oil possesses moderate antioxidant and antityrosinase activity. The effects were weak when compared with positive controls. activity.

Ethical approval

Not applicable, because this article does not contain any studies with human or animal subjects.

Table 3. Activity results of the essential oil of *C. libanoticum*

	DPPH ^c scavenging activity % ^a	TEAC ^b (mmol/L)	Tyrosinase inhibition % ^b
<i>C. libanoticum</i> EO	31.3 ± 1.1	0.027 ± 0.008	17.7 ± 1.6
Gallic acid ^c	1.93 ± 0.02	-	-
Ascorbic acid ^c	9.3 ± 0.01	-	-
Kojic acid ^c	-	-	3.6 ± 0.01

Data was given as mean ± SD ($n = 3$) ^a ; Samples tested at 5 mg/mL concentration, ^b ; Samples tested at 1 mg/mL concentration;

^c ; Positive controls, values represented IC₅₀^c (µg/mL), EO: essential oil

Author contribution

Concept: HGA, BT, MK; Design: HGA, BT, MK; Supervision: KHCB; Materials: AD; Data Collection and/or Processing: MK, BT; Analysis and/or Interpretation: MK, BT; Literature Search: MK, HGA, BT; Writing: BT, MK, HGA, KHCB; Critical Reviews: KHCB.

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Conflict of interest

The authors declared that there is no conflict of interest.

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High-density lipoprotein: Quality is more important than quantity!

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ABSTRACT

High density lipoproteins (HDLs) are complex lipid carriers which replace their own structures and functions according to different conditions including healthy or pathological status. HDL has substantial roles in reverse cholesterol transport system for regulation of lipid metabolism. Also HDL has many functions such as antiinflammation, antioxidation, antithrombotic and vasodilator actions. However, these functions may be compromised under pathological conditions. Nowadays, it's known that the function of HDL is more important than it's dose for preventing against cardiovascular diseases. This review mostly summarizes available information of HDL structure, metabolism and the real information related to the physicochemical changes of HDLs causing their different functions. Also HDLs, that lost their protective functions have been associated with oxidative stress and systemic inflammation diseases.

Keywords: Antiinflammatory, antioxidant, dysfunctional HDL, high-density lipoproteins

1. INTRODUCTION

Lipoproteins carry apolar lipids in the center of their structures including cholesterol esters and triglycerides. Also, they have hydrophilic membrane carrying phospholipids, cholesterol, and apolipoproteins. These lipid carriers are grouped according to their size, apolipoproteins and lipid composition. Lipoproteins are grouped as chylomicrons, chylomicron remnants, very low density lipoproteins (VLDL), intermediate density lipoproteins (IDL), low density lipoproteins (LDL), high density lipoproteins (HDL) and Lp(a). Table 1 shows the classification of lipoproteins and the major lipids and major apoproteins that these lipoproteins contain (Table 1) [1,2].

HDLs, which will be discussed, have different roles such as antiinflammatory, antioxidant and

reverse cholesterol transport. These roles of HDLs can be observed under both physiological and pathological situations. Such as, HDLs undertake antiinflammatory roles in the immune system when inflammatory conditions occurs and change their structures for carrying proinflammatory molecules [3]. This review reports the novel knowledges linking up the relationships between physicochemical characteristics of HDLs and those changes found in the pathological circumstances such as oxidative stress and systemic inflammation.

HDL Structure:

HDLs have complex and heterogeneous structures among other class of lipoproteins by carrying important differences in composition, shape, size and function [4]. HDLs have small (5-17 nm diameter) and most dense (1,063-1,25 g/mL) structures in the

Table 1. Lipoprotein classes

Lipoprotein	Density (g/ml)	Major Lipids	Major Apoproteins
Chylomicrons	<0.930	Triglycerides	Apo B-48, Apo C, Apo E, Apo A-I, A-II, A-IV
Chylomicron Remnants	0.930- 1.006	Triglycerides Cholesterol	Apo B-48, Apo E
VLDL	0.930- 1.006	Triglycerides	Apo B-100, Apo E, Apo C
IDL	1.006- 1.019	Triglycerides Cholesterol	Apo B-100, Apo E, Apo C
LDL	1.019- 1.063	Cholesterol	Apo B-100
HDL	1.063- 1.210	Cholesterol Phospholipids	Apo A-I, Apo A-II, Apo C, Apo E
Lp (a)	1.055- 1.085	Cholesterol	Apo B-100, Apo A

group of lipoproteins. HDLs have different shaped forms such as discoidal (nascent) and spherical (mature).

Nascent HDL consists of 100 to 150 phospholipid molecules in phospholipid bilayer as found in plasma membrane and 2 to 5 Apo A-I molecules creating a band around the lipid structure. Nascent HDL contains approximately 10% of overall circulating HDL. Apo A-I and Apo A-II are the most existing protein contents of the nascent HDL. Numerous phospholipids and few triglycerides, free cholesterol and cholesteryl esters are also found in this unique structure. Nascent HDL is converted into mature HDL, that is the main type of HDL in the blood stream [5]. The structure of these mature HDL particle contains of 3–5% cholesterol, approximately 5% triglycerides, 15–20% esterified cholesterol, 26–32% phospholipids and 45–55% apolipoprotein molecules. The lipid core of HDL is consist of cholesterol esters (CE). Apo A-I and Apo A-II are main apoproteins of this mature HDL and few Apo E and Apo C are also found around of lipid structure. HDL is divided into three classes considering their densities: HDL1, HDL2 and HDL3 which have densities of 1,050-1,063 g/ml, 1,063-1,12 g/ml and 1,12-1,21 g/ml respectively.

Moreover, these classes of HDLs are also consists of five different subclasses such as HDL2 and 2b, HDL3a, 3b, and 3c, according to their size. HDL3 which is the first small and mature form of HDL is the best acceptor of free cholesterol. When the amount of esterified free cholesterol increases, the particle size increases and HDL2 occurs. HDL2 is further enhanced by cholesterol esters and also obtain apoE. This apo E-containing particle (HDL1)

actually constitute a small fraction of HDL but is known as metabolically active subclass [6,7].

Besides, HDL particles comprise some other components with different biological activity such as vitamins, hormones and microRNAs. HDL successfully takes these functional miRs to their target cells. Hence, microRNAs consolidate structure of HDL particles [8].

Apo A-I is an effective molecule that contributes in nearly all known functions of HDL particle. Apo A-I provides the coaction of HDL particle with ATP-binding cassette transporters A1 (ABCA1), G1, scavenger receptor B1 and lecithin cholesterol acyltransferase (LCAT) [9]. These cholesterol-phospholipid transporters expressed in hepatocytes, enterocytes, macrophages and other tissues [10]. The main structure of high-density lipoprotein is shown in figure 1.

HDL Metabolism and Function:

Firstly, Apo A-I is synthesized in the metabolism of HDL particles. Nearly 80% of these Apo A-I molecules are obtained by de novo synthesis in hepatocytes and others secreted by intestinal mucosa (20%). The name of this firstly synthesized nascent protein and phospholipid structure is the form of pre- β 1 HDL. In blood stream, coaction of Apo A-I or pre- β 1 HDL with ABCA1 transporters is an important process for beginning of HDL functions. The Apo A-I also starts activation of LCAT enzyme. These mentioned receptors cause transportation of phospholipids and free cholesterol from peripheral tissues to HDL particle and transformation of initial particles into nascent HDL (α 4 HDL). Cholesteryl esters obtained by LCAT enzyme located with

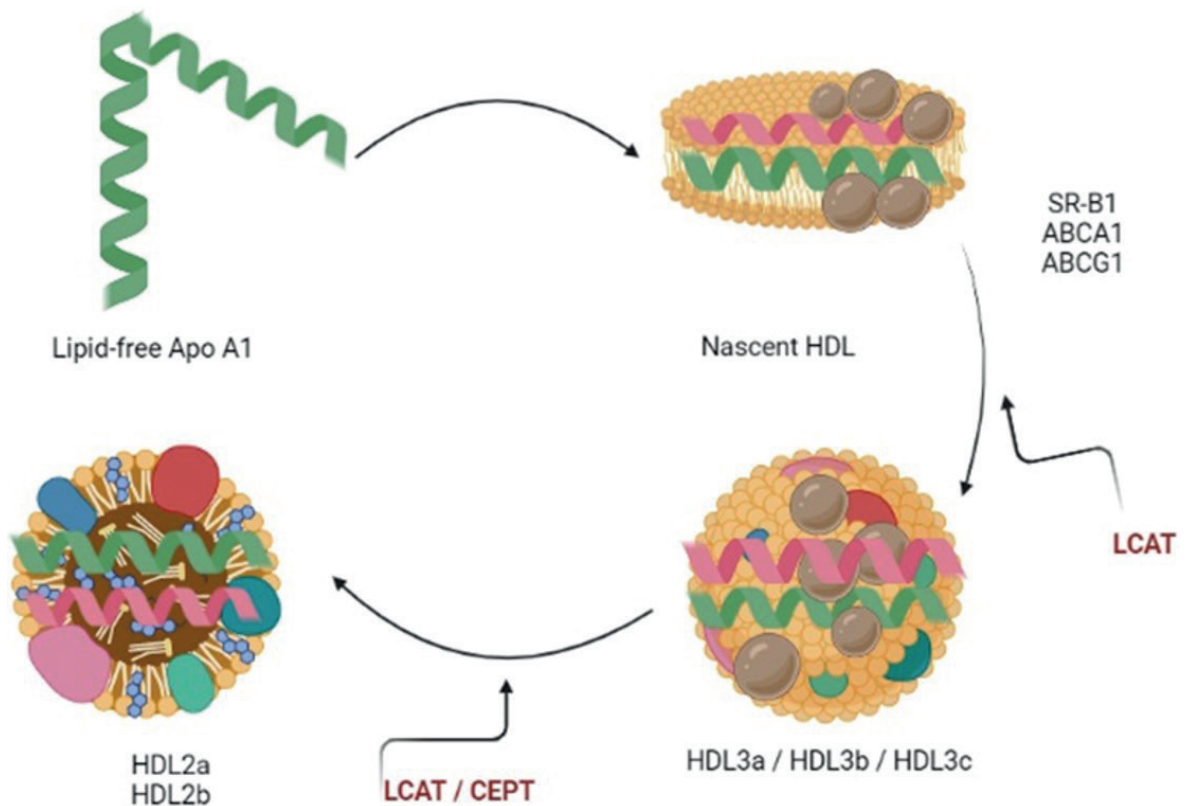


Figure 1. The main structure of high-density lipoprotein

triglycerides in the hydrophobic core of discoidal HDL. Formation of mature HDL structure begins with accumulation of triglycerides and cholesteryl ester. This spherical HDL form is the main particles in circulation. Conformational changes in this Apo A-I are critical for formation of spherical HDL and functions for transport of cholesterol from peripheral tissues to HDL. The spherical HDL is developed by continuous of the LCAT enzyme effect and the activities of lipid transfer proteins. Mature HDL formation is further continued by different proteins such as cholesteryl ester transfer protein (CETP) and the phospholipid transfer protein (PLTP). HDL concentrations and structure are controlled by these proteins in circulation [5]. Another important feature of HDL is that it contains paraoxonase 1 (PON-1) enzyme in its structure [10].

Apo A-I has the HDL associated enzymes PON-1 and platelet activating factor acetylhydrolase (PAF-AH) which suppress LDL oxidation by hydrolyzing oxidized lipids [11]. HDL performs multiple physiological functions. Especially HDL

is a substantial molecule for continuity of whole lipid metabolism [12]. The most important role of HDL is the continuous activity of RCT pathway which removes cholesterol from peripheral tissues to the liver exerting antiatherosclerotic character of this lipoprotein. It is therefore considered an antiatherosclerotic lipoprotein. This process is called the HDL cycle and protects endothelial cells. Actually, HDLs prevent toxic accumulation of cholesterol and cause transformation of cholesterol into bile acids [13]. HDL has roles to interfere with nitric oxide (NO) metabolism because of its preventive effects on endothelial cells and vascular function. Except that Apo A-I, HDL related enzymes and apolipoproteins such as PON-1 and Apo E stimulate eNOS and increase NO production [14]. Antiinflammatory action of HDL is associated with Apo A-I, PON-1 and sphingosine-1-phosphate contents. HDL prevents migration and infiltration of monocytes and macrophages to the arterial wall by inhibiting monocyte chemoattractant protein 1 (MCP 1) which induced by LDL [15].

The other indispensable role of HDL is its antioxidant action. HDL success this action by preventing of lipid oxidation and removing oxidized products. Because of interaction of inflammation and oxidative stress, HDL's antioxidant characteristic contributes to it's antiinflammatory behaviors [6]. Furthermore, HDL has antithrombotic effects because of it's roles on the prostacyclin signaling pathway. It's known that prostacyclins interact with NO to protect against platelet activation and aggregation. HDL also induce the expression of cyclooxygenase 2, that induce prostacyclin synthesis in endothelial cells. Moreover, HDLs protect against apoptosis of endothelial cells by reduction of CPP32-like protease activity, resulting in a decrease in the effect of tumor necrosis factor (TNF- α) [16]. HDL-C physiological functions are shown in figure 2.

To summarize the functions of HDL; HDL transfers apoproteins to other lipoproteins. It takes lipids from other lipoproteins and esterifies cholesterol by the LCAT reaction. By transferring appropriate esters to other lipoproteins, it ensures the transport of cholesterol to the liver. Also, antithrombotic

property of HDL is it's ability to induce the generation of prostacyclin. HDL particles are highly antiatherogenic and inhibit endothelial cell apoptosis and prevents against LDL oxidation.

What conditions make HDL go bad?

HDLs obtained from healthy people have frequently antiinflammatory characteristics. On the other hand, when systemic inflammation occurs in organism, HDL can turn into proinflammatory form as part of an acute phase response. Coronary atherosclerosis, surgery, diabetes mellitus, influenza, sepsis, chronic systemic inflammation are a good examples of this effect [17].

Proinflammatory properties of HDL

HDL has an antiinflammatory effect through two mechanisms. 1) Normal HDLs protect LDL-C against oxidation. Then it decrease attraction of monocytes into arterial tissue and occurrence of foam cells that caused by oxidized LDL. 2) Normal HDLs carry out reverse cholesterol transport by removing cholesterol from peripheral tissues.

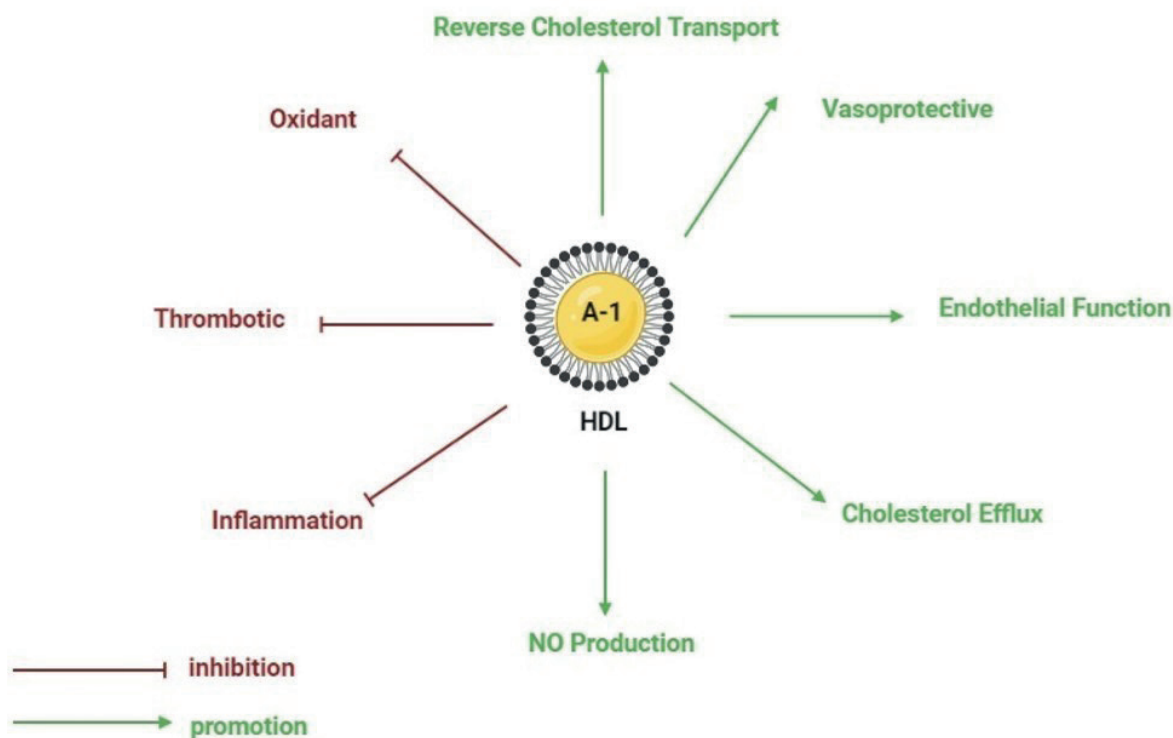


Figure 2. Physiological functions of HDL-C

HDLs have some antioxidant enzymes to work the continuity of an antiinflammatory statement. More than twenty years ago, it was shown that HDL can be in proinflammatory state during an acute phase response in humans and animal after a surgery [18].

During systemic inflammatory state antioxidants can decrease and HDL can increase oxidized lipids and proteins that enhance its proinflammatory state. Dysfunctional HDL causes releasing of proinflammatory cytokines that evoke transportation of monocytes into arterial walls, which eventually causes to occurrence of macrophage accumulation [19]. Atherosclerosis, in its simplest form, is accumulation of oxide LDL-C and triglycerides in peripheral vascular cells. ABCA1 and ABCG1 have roles in prevention and regression of atherosclerosis by removing of excess oxidized cholesterol from macrophages and forming of normal high density lipoprotein [20].

Dysfunctional HDL is also created by myeloperoxidase (MPO). MPO, released from macrophages in atherosclerotic lesions, causes oxidative damage to Apo A-I. It's known that HDLs isolated from the atherosclerotic lesions contain in large quantities of MPO modified proteins such as nitrated and chlorinated Apo A-I [21].

Pro-oxidant properties of HDL

HDL may incorporate lipid peroxidation products and phospholipid containing hydroperoxides that given by oxidized LDL. Furthermore, HDL has roles in hydrolyzing oxidized phospholipids, such as F2-isoprostanes, formed during the oxidative modification of LDL [22]. On the other hand, oxidized HDL may induce the production of reactive oxygen species (ROS) and enhance the risk of cardiovascular diseases by upregulating expression of some proinflammatory genes such as tumor necrosis factor alfa, cyclooxygenase 2 and plasminogen activator inhibitor-1 (PAI-1) [23]. In addition to all this, the enrichment of HDL with triglycerides, MPO, phospholipase A2, ceruloplasmin and serum amyloid A decrease its antioxidant activity and cause the formation of prooxidant HDL [24]. It was shown that oxidized HDLs can not carry out cholesterol efflux from foam cells [25-28].

Methods of HDL Measurement in the Laboratory

Some *in vitro* and *ex vivo* reproducible methods were developed to determine HDL's heterogeneity and various functions. Therewithal detection of the presence of dysfunctional HDL in the patient will help to change the treatment process positively [24]. Standard method for measuring HDL-C; It is the precipitation of lipoproteins using a polyanion such as heparin $MnCl_2$, phosphotungstate $MnCl_2$, dextran sulfate $MnCl_2$ or polyethylene glycol, and then determining the cholesterol value by an enzymatic colorimetric test. These methods have limitations regarding insufficient precipitation of Apo B lipoproteins and assay environment that influence HDL-C measurement. These insufficiencies could produce inaccurate HDL-C measurement for the correct define of cardiovascular risk. On the other hand, electrophoresis, ultracentrifugation, chromatography and nuclear magnetic resonance can be seen more purified methods to separate HDL-C in plasma [29]. But this methods have also disadvantages because of requirement of specialized equipment or performing and learning of the methodology is difficult. Hence, the colorimetric methods are used to measure of HDL composition [30].

The increases or decreases in the change in fluorescence intensity (FI) caused by dichlorofluorescein diacetate (DCFHDA) oxidation can be measured to define the functional characteristics of HDL rather than quantity.

This method measure the capability of various doses of HDL to reduce the generation of intracellular H_2O_2 as reflected by the decrease in the generation of FI by DCF [31] (Figure 3).

2. CONCLUSION

The heterogeneity in the structure and content of HDL is the source of its functional diversity and importance. HDL components and their activity especially in the disease state undergoing significant modifications dynamically change. Such dysfunctional HDL particles appear as both a cause and a symptom of many diseases, especially atherosclerosis. Nowadays, routine lipid measurements in laboratories have

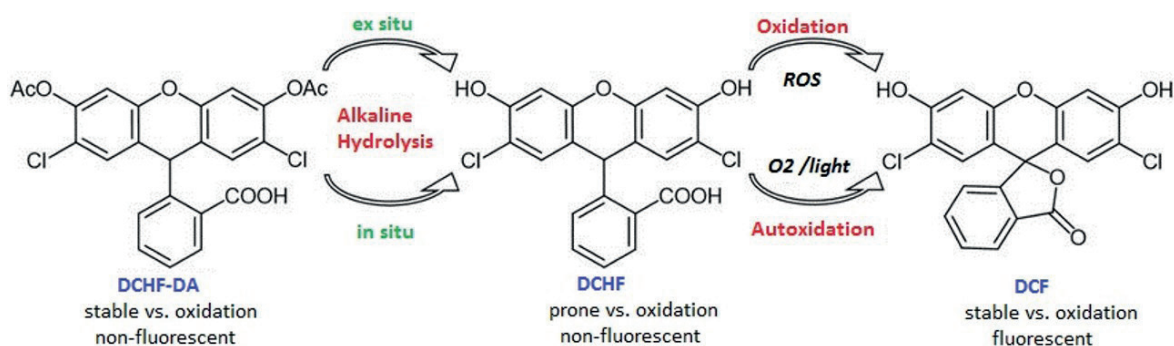


Figure 3. Oxidation of dichlorofluorescein diacetate

become a misleading indicator. In the light of current information, when assessing HDL in the body, it is clear that it is essential to make a qualitative analysis rather than a quantitative one.

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Author contribution

Concept: IE, GAC, OA; Design: IE and GAC; Supervision: OA; Data Collection and/or Processing: IE and GAC; Literature Search: IE; Writing: IE and GAC; Critical Reviews: IE, GAC and OA.

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Conflict of interest

The authors declared that there is no conflict of interest.

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