

Article Synthesis and Biological Evaluation of Thiazole-Based Derivatives with Potential against Breast Cancer and Antimicrobial Agents

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1. Introduction

New drug design is the perspective process from both aspects of discovery and synthesis. The main scope of the design of new compounds or complexes is based on the knowledge of the biological target. Over the past ten years, attention toward antifungal, antimicrobial, and anticancer properties have risen. Cancer ranks as a leading cause of death in every country of the world. Moreover, based on the statistics from the World Health Organization (WHO), there were an estimated 19.3 million new cancer cases in 2020. Among these, 2.26 million (11.7%) cases refer to female breast cancer [1].

A significant increase in disease cases is mainly associated with fungal infections. Invasive candidiasis is the most common fungal disease among hospitalized patients in developed countries [2,3]. Besides interest in antifungal activity, the development of new antimicrobial agents with novel structures remains the primary goal for finding a solution to increasing bacterial resistance [4–7]. *Escherichia coli* and *Staphylococcus aureus* are the two main causes of healthcare-associated and hospital-acquired invasive infections in adults [8]. The majority of *E. coli* and *S. aureus* bacteremia occur in individuals aged 65 years and older, with the highest rates in those aged 85 years and older [9,10].

A potentially commercial drug is mostly represented as the organic ligand that inhibits or activates the properties of a biomolecule. The affinity of organic ligands to coordinate



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Copyright: © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). transition metal ions has increased in the research field of coordination chemistry. Therefore, this study is focused on thiazole-based derivatives. Nitrogen-containing heterocycles with a sulfur nucleus are an important class of compounds [11,12]. Recent clinical trials have increased the significance of thiazole derivatives due to their application as antiinflammatory, anticancer, antifungal, antihypertensive, anticonvulsant, and antidiabetic compounds [13–15]. The selection of the appropriate metal salts is the second essential step in the architecture of coordination complexes. Metal-containing compounds are broadly used in a wide range of biological systems. The class of complexes on the basis of platinum derivatives is considered to be one of the most effective in the struggle against cancer tumors today, and cisplatin is the most widespread of them in clinical practice. However, its poor solubility in water and high general toxicity [16] serve as the reason for the development of new antitumoral medications on the basis of other metals compounds. Copper has attracted our scientific attention due to its essential role as the structural component of several enzymes and proteins: laccase, ascorbate oxidase, ceruloplasmin, amine oxidases, superoxide dismutase, and cytochrome C oxidase [17–20]. Endogenous metals are believed to be less toxic to healthy cells; therefore, the antitumor activity of a number of coppercontaining coordination compounds has been tested and encouraging results have been obtained [21].

This work is aimed at broadening research on the inhibition and binding affinity of thiazole-based derivatives complexes. To achieve our goal, three organic ligands, 1-[2-thiazol-4-yl-(4-substitutedphenyl)]-4-*n*-propylpiperazines, and three corresponding Cu(II) complexes were investigated. Moreover, the standard procedures for the characterization of synthesized compounds (F-AAS, FTIR, NMR, and TGA), ADME analysis, MTT assay, and microbiology tests were performed.

2. Results and Discussion

2.1. Crystal Structure Analysis

The crystal structure was determined for ligand L2 (Figure 1). In this case, a goodquality crystal structure (suitable for X-ray measurements) was obtained. The thiazole and phenyl rings are almost co-planar. The dihedral angle between their planes is 4.3 Å. The bond lengths in the thiazole ring are typical for its 2,4-substituted derivatives (Table 1) [22].



Figure 1. Molecular structure of ligand L2.

The piperazine possesses a chair conformation with the thiazole ring and propyl group bonded in the equatorial positions (trans configuration). All CN bonds of the piperazine ring have similar lengths. The piperazine CNC angle from the thiazole side is significantly larger than that from the propyl side (Table 1). It is a consequence of the partially aromatic character of the C9-N12 bond, which affects the nitrogen hybridization. Its hybridization is between SP3 and SP2 and that enlarges the CNC angle. In turn, the N15-C19 bond is a classical single bond, therefore the CNC angle from the propyl side is a typical tetrahedral angle.

i—j	d _{ij} (Å)	i—j—k	α _{ijk} (°)
C4—C7	1.4746(15)	C7—N8—C9	110.41(9)
C7—N8	1.3925(14)	N8-C9-S10	114.69(8)
N8-C9	1.3055(14)	C9—S10—C11	88.72(5)
C9—S10	1.7585(11)	S10-C11-C7	110.53(9)
S10-C11	1.7252(12)	C11-C7-N8	115.65(10)
C11—C7	1.3582(16)	N12-C13-C14	109.97(9)
C9—N12	1.3643(14)	C13-C14-N15	110.99(9)
N12-C13	1.4589(14)	C14—N15—C16	109.18(9)
C13—C14	1.5176(16)	N15-C16-C17	110.82(9)
C14—N15	1.4598(14)	C16-C17-N12	109.73(9)
N15-C16	1.4634(14)	C17—N12—C13	114.17(9)
C16—C17	1.5208(15)		
C17—N12	1.4643(14)		
N15—C19	1.4674(14)		

Table 1. Selected structural data of ligand L2.

The L2 molecules do not contain the donors of the classical hydrogen bonds. Consequently, the supramolecular structure of L2 is stabilized via non-classical hydrogen bonds, CH•••N, CH•••S, and CH•••Cl, as well as CH••• π interactions. Furthermore, the molecules do not form π - π stacking interactions despite possessing aromatic rings.

The Hirshfeld surface and the 2D fingerprint plot reflect the above-mentioned intermolecular interactions that exist in the structure of L2 (Figure 2). The domination of dispersive interactions is revealed in large contribution of $H \bullet \bullet \bullet H$ and $H \bullet \bullet \bullet C$ contacts in comparison to the rest (Table 2).



Figure 2. Hirshfeld surface (plotted over d_{norm}) and 2D fingerprint plot of L2.

Tabl	e 2.	Percentage	distribut	ion (%) of t	the intermo	lecular	contacts	in th	ne structure	of l	ligand	L2
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H●●●H	49.6	Cl●●●N	0.7
Н●●●С	21.1	Cl●●S	0.9
H●●●N	5.4	C•••C	0.0
H●●●S	7.0	C●●●N	0.7
H●●●Cl	13.2	C●●●S	1.2
Cl●●●Cl	0.0	N●●●S	0.1
Cl•••C	0.0	Sees	0.2

2.2. FTIR Spectra

Figure 3 represents the FTIR spectra of pure ligands and synthesized coordination compounds. The spectra of pure ligands do not have any peaks in the region of $3400-3300 \text{ cm}^{-1}$. The opposite situation occurs with the spectra of all copper(II) complexes, which contain peaks in this region. Peaks are assigned to v(NH) vibrations and correspond to the amino group formed during the complexation process due to hydrogen movement [23].



Figure 3. FTIR spectra of pure ligands and obtained complexes: (**a**) L1 and Cu(L1)₂Cl₂; (**b**) L2 and Cu(L2)Cl₂; (**c**) L3 and Cu(L3)Cl₂.

All obtained spectra include peaks in the range of $2700-3000 \text{ cm}^{-1}$, which correspond to the C-H stretching vibrations. In the case of pure ligands, the strong absorption peaks of v(C=N) in the region of $1600-1620 \text{ cm}^{-1}$ and $1200-1230 \text{ cm}^{-1}$ are observed. After the complexation processes, this vibration mode in the region of $1600-1620 \text{ cm}^{-1}$ is shifted to the higher wavenumbers. They are at 1627, 1620, and 1609 cm⁻¹ for free ligands L1, L2, and L3, respectively. In the complexes, these bands appear at 1599 cm⁻¹ for Cu(L1)₂Cl₂, at 1616 cm⁻¹ for Cu(L2)Cl₂, and at 1606 cm⁻¹ in the case of Cu(L3)Cl₂. These modes confirm the involvement of nitrogen atoms in the complexation processes. The absorption peaks of v(C-S) are observed in the range of $910-925 \text{ cm}^{-1}$. They are at 920, 923, and 919 cm^{-1} for free L1, L2, and L3, respectively. Due to the coordination process, the modes of complexes are shifted to 966 cm^{-1} (Cu(L1)₂Cl₂), 967 cm^{-1} (Cu(L2)Cl₂), and 964 cm^{-1} (Cu(L3)Cl₂) due to the coordination process.

Uncoordinated ligands include modes of β (CH) and γ (CH) vibrations between 1250–1030 and 840–690 cm⁻¹. For obtained complexes, they appear in similar ranges.

Thus, it is clear that all pure ligands act as N-donor and S-donor ligands. In the case of L2 and L3, the coordination appears through the sulfur atom from the thiazole ring and the nitrogen atom from the piperazine ring [24]. L2 and L3 ligands act as bidentate. From the FTIR spectra of the L1 ligand and complex based on it, it is clear that nitrogen and sulfur are also involved in the coordination process.

2.3. Thermogravimetric Study

Thermogravimetric analysis (TGA) is an alternative, fast, and useful technique for determining the composition of coordination compounds. Therefore, the thermogravimetric curves of three ligands and complexes are presented in Figure 4.



Figure 4. TG-DTG traces of the pure organic ligands and complexes: (**a**) L1; (**b**) Cu(L1)₂Cl₂; (**c**) L2; (**d**) Cu(L2)Cl₂; (**e**) L3; (**f**) Cu(L3)Cl₂.

Table 3 demonstrates the decomposition stages of the three pure ligands and complexes with the identified temperature ranges and mass losses.

Commlay	Channe	Temperature Range, °C			Mas	s Loss	Final Solid Product of the
Complex	Stages	Ι	II	III	Found	Calculated	Thermal Decomposition
L1	2	160-365	440-620	-	100.00	100.00	-
L2	2	160-340	410-620	-	100.00	100.00	-
L3	2	190-380	400-660	-	100.00	100.00	-
Cu(L1) ₂ Cl ₂	3	70–175	280-420	480-720	87.81	88.78	CuO
Cu(L2)Cl ₂	3	50-250	305-415	500-750	86.74	82.57	CuO
Cu(L3)Cl ₂	3	50-250	290–420	480–770	84.87	82.49	CuO

Table 3. Thermal decomposition data of L1, L2, L3, Cu(L1)₂Cl₂, Cu(L2)Cl₂, and Cu(L3)Cl₂.

Based on the obtained results, the thermolysis of all organic ligands is a two-stage process, which starts between 160–190 °C. The most stable is ligand L3 because it decomposes at 190 °C. The first and the biggest mass losses on the TG curves correspond to the destruction of organic ligands. In the range of 400–660 °C, the burning of organic fragments takes place. All organic ligands correspond to the thermal decomposition stability.

The results demonstrate the thermal decomposition of three copper(II) complexes. The most stable is compound $Cu(L1)_2Cl_2$; it starts to decompose at 70 °C. The other coordination compounds are stable up to 50 °C. The first small mass losses overlapping each other correspond to the initial fragmentation and degradation of the organic ligand. Above 175 °C in the case of $Cu(L1)_2Cl_2$, and 250 °C in the case of $Cu(L2)Cl_2$ and $Cu(L3)Cl_2$, the biggest mass losses on TG curves are observed; they are associated with the further degradation and decomposition of organic ligands. When the temperature rises, decomposition of the residual part of the ligand, loss of chloride anions, and formation of solid-state metal oxide take place. Above 700 °C, pure CuO appears. Higher than expected final mass losses indicate the formation of volatile polymeric Cu_xCl_x [25].

2.4. Biological Activity Predictions

The activity of the synthesized ligands was investigated. The obtained results are shown in Table 4. According to the results, all three ligands have demonstrated a wide range of biological activity.

Table 4. Activity prediction results for three ligands: L1, L2, L3, where Pa is the probability to be active; Pi is the probability to be inactive.

	Pa	Pi	Activity
	0.767	0.003	Anti-anorexic
	0.741	0.007	Phosphatase inhibitor
	0.634	0.012	Neurodegenerative diseases treatment
т 1	0.598	0.007	Transcription factor STAT inhibitor
LI	0.621	0.058	Glycosylphosphatidylinositol phospholipase D inhibitor
	0.568	0.010	Transcription factor STAT3 inhibitor
	0.635	0.086	Mucomembranous protector
	0.543	0.020	Anxiolytic
	0.797	0.003	Anti-anorexic
	0.740	0.007	Phosphatase inhibitor
	0.714	0.032	Glycosylphosphatidylinositol phospholipase D inhibitor
10	0.590	0.007	Transcription factor STAT inhibitor
LZ	0.596	0.015	Insulin promoter
	0.669	0.091	Phobic disorders treatment
	0.594	0.017	Neurodegenerative diseases treatment
	0.586	0.021	Cholesterol antagonist
	0.705	0.012	Phosphatase inhibitor
	0.694	0.004	Anti-anorexic
	0.607	0.006	Transcription factor STAT inhibitor
1.2	0.592	0.008	Transcription factor STAT3 inhibitor
L3	0.569	0.019	Neurodegenerative diseases treatment
	0.612	0.067	Antineurotic
	0.528	0.019	Anti-ulcerative
	0.526	0.032	Cholesterol antagonist

2.5. ADME Analysis

An ADME analysis is used to investigate the properties and calculate the pharmacokinetic parameters of novel compounds. Drug-likeness descriptors selected using the Lipinski and Veber rules were calculated with SwissADME. The Lipinski's rule of five by notes that good absorption is more likely when the molecular weight (MW) < 500 Da, number of hydrogen bond donors (HBDs) < 5—as shown in molecular docking studies—LogP < 5, and number of hydrogen bond acceptors (HBAs) < 10. Veber rules identified two other relevant descriptors: the number of rotatable bonds (NBR) < 10 and polar surface area (PSA) < 140 Å² [26].

The calculations and analysis of the newly-synthesized organic ligands and coordination compounds indicated that all compounds except $Cu(L1)_2Cl_2$ have no violations of these rules (Figure 5). The red-colored zone corresponds to the oral bioavailability [27].



Figure 5. Oral bioavailability graph generated using the SwissADME service: (**a**) L1, (**b**) L2, (**c**) L3, (**d**) Cu(L1)₂Cl₂, (**e**) Cu(L2)Cl₂, and (**f**) Cu(L3)Cl₂.

Additionally, the ADME analysis confirmed the effectiveness of the tested compounds crossing the blood-brain barrier, except for $Cu(L1)_2Cl_2$ (Figure 6) [28].



Figure 6. BOILED-Egg diagram for the synthesized ligands and complexes.

A ProTox II toolkit was used to classify the ligands and complexes into toxicity classes, which are shown in Table 5.

Table 5. Toxicity prediction result	s.
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Name	Toxicity Class	Predicted LD50
 L1	4	1000 mg/kg
L2	4	1000 mg/kg
L3	4	1000 mg/kg
$Cu(L1)_2Cl_2$	4	540 mg/kg
$Cu(L2)Cl_2$	4	540 mg/kg
$Cu(L3)Cl_2$	4	540 mg/kg

2.6. Antimicrobial Activity

Assays were performed using reference strains of bacteria: *Staphylococcus aureus* ATCC 33591, *Escherichia coli* ATCC 25922, and fungus *Candida glabrata* ATCC 15126 (Table 6). The tested compounds were dissolved in DMSO solution, which was recommended because of their chemical structures. This solvent has proven antimicrobial activity and the control with a serial dilution of DMSO was also included in the tests [29]. Moreover, nystatin, a conventional antifungal antibiotic, was examined and used as reference material (MIC = 4 µg/co). The activity of synthesized complexes was considered only if the DMSO impact was excluded. Bacterial strains were not sensitive to the tested substances and the MIC values were mainly > 1024 µg/mL, except L1 (128 µg/mL). More promising results were obtained for the *C. glabrata* strain where the MIC values were in the range of 32–128 µg/mL, except for Cu(L1)₂Cl₂, where MIC > 1024 µg/mL. The lowest value was obtained for L1 (32 µg/mL). In the cases of L3, Cu(L2)Cl₂, and Cu(L3)Cl₂, the MIC was equal to 64 µg/mL. The obtained results from the minimal inhibitory concentration assays indicate that the tested complexes have antifungal activity, but more work in this field is needed to increase and examine their potential.

Table 6. Results of the minimal inhibitory concentration assays established for microbial strains (μ g/mL).

Complex	S. aureus	E. coli	C. glabrata
L1	>1000	128	32
L2	>1000	>1000	128
L3	>1000	>1000	64
$Cu(L1)_2Cl_2$	>1000	>1000	>1000
$Cu(L2)Cl_2$	>1000	>1000	64
$Cu(L3)Cl_2$	>1000	>1000	64
Nystatin	-	-	4

2.7. MTT Analysis

Cell viability was determined through the MTT assay in L929 and MCF7 cells for all compounds (Figure 7). Only two copper complexes, i.e., $Cu(L1)_2Cl_2$ and $Cu(L3)Cl_2$, presented cytotoxic effects. The other compounds, i.e., compounds without copper and $Cu(L2)Cl_2$, were not cytotoxic in the concentration used. The IC50 values were calculated for $Cu(L1)_2Cl_2$ and $Cu(L3)Cl_2$. A differential effect of copper compounds $Cu(L1)_2Cl_2$ and $Cu(L3)Cl_2$ were demonstrated in two cell types. The MCF7 cells were more sensitive to both compounds compared to L929 cells. The calculated IC50 value (that is, the concentration of compound which exhibited 50% cell viability) for MCF7 cells was much lower (105.6 μ M and 82.64 μ M, respectively) than for L929 cells (185.56 μ M and 185.86 μ M). According to the IC50 results (experimental proof of the safety of the compounds), the compounds do not cause high toxicity in the non-tumorigenic cell line, which indicates a certain selectivity towards tumorigenic cells.



Figure 7. Cytotoxic activity of the investigated complexes (red line) and the ligands (blue line) against breast cancer cells MCF7 vs. mouse fibroblast cell line L929 (in corresponding concentrations): (a) mouse fibroblast cell line L929 for L1 and $Cu(L1)_2Cl_2$; (b) mouse fibroblast cell line L929 for L2 and $Cu(L2)Cl_2$; (c) mouse fibroblast cell line L929 for L3 and $Cu(L3)Cl_2$; (d) breast cancer cells MCF7 for L1 and $Cu(L1)_2Cl_2$; (e) breast cancer cells MCF7 for L2 and $Cu(L2)Cl_2$; (f) breast cancer cells MCF7 for L3 and $Cu(L3)Cl_2$; (f) breast cancer cells MCF7 for L3 and $Cu(L3)Cl_2$.

3. Materials and Methods

3.1. Chemistry

All chemicals used for synthesis were purchased from Sigma-Aldrich (Saint Louis, MO, USA) and Alfa Aesar (Haverhill, MA, USA). All melting points (mp) were measured in open capillaries in an electrothermal apparatus and are uncorrected.

3.1.1. ¹H, and ¹³C NMR

The ¹H spectra ligands (L1-L3) were recorded in CDCl₃ as a solvent in a 600 MHz spectrometer, which is a Bruker Avance III spectrometer at ambient temperature (Supplementary Materials Figures S1, S3 and S5). The chemical shifts are reported in ppm on a scale downfield from tetramethylsilane (TMS) as the internal standard, and the signal patterns are indicated as follows: s = singlet, d = doublet, t = triplet, m = multiplet, br = broad; a number of protons, and J approximates the coupling constant in Hertz. The ¹³C NMR spectra were recorded on a Bruker Avance III in a 600 MHz spectrometer (Supplementary Materials Figures S2, S4 and S6). The TLC data were obtained with Merck silica gel 60F254 aluminum sheets.

3.1.2. Flame Atomic Absorption Spectroscopy

The content of Cu(II) in the solid complex was determined by the F-AAS spectrometer with a continuum source of light and using air/acetylene flame (Analytik Jena, contraAA 300, Jena, Germany). The absorbance was measured at analytical spectral line 324.7 nm. The limit of quantification was 0.04 mg/L. The solid sample was decomposed using the Anton Paar Multiwave 3000 closed system instrument. Mineralization was carried out for 50 min at 240 °C under a pressure of 60 bar. The contents of carbon, hydrogen, and nitrogen were determined by a Vario-micro company Elementar Analysensysteme GmbH.

3.1.3. Fourier-Transform Infrared Spectroscopy

FTIR spectra were recorded with an IRTracer-100 Shimadzu Spectrometer (4000–600 cm⁻¹) with the accuracy of recording set as 1 cm⁻¹ using KBr pellets.

3.1.4. Thermogravimetric Analysis

The thermolysis of compounds in the air atmosphere was studied by TG-DTG techniques in the range of 25–800 °C at a heating rate of 10 °C min⁻¹; TG and DTG curves were recorded on Netzsch TG 209 apparatus under air atmosphere $v = 20 \text{ mL min}^{-1}$ using ceramic crucibles—which were used as a reference material.

3.1.5. X-ray Measurement and Crystal Structure Determination

X-ray diffraction data were collected for ligand L2 on an XtaLAB Synergy Dualflex Pilatus 300K diffractometer (Rigaku Corporation, Tokyo, Japan). Using Olex2 [30], the structure was solved with the SHELXT [31] using Intrinsic Phasing and refined with the SHELXL [32] using Least Squares minimization. All non-hydrogen atoms were refined anisotropically. All hydrogen atoms were found from the Fourier difference map and refined using the "riding" model. The details concerning X-ray diffraction data and structure refinement are given in Table 7.

Table 7. A-ray diffraction data and structure reinfement details for figand L2.

Compound	L2
Empirical formula	C ₁₆ H ₂₀ ClN ₃ S
Formula weight	321.86
Temperature (K)	100(1)
Crystal system	monoclinic
Space group	$P2_1/c$
a (Å)	11.62470(10)
b (Å)	17.19590(10)
<i>c</i> (Å)	8.25110(10)
α (°)	90
β (°)	109.8670(10)
γ (°)	90
Volume (Å ³)	1551.21(3)
Z	4
Calculated density (g/cm ³)	1.378
Absorption coefficient (mm^{-1})	3.399
F(000)	680.0
Crystal size (mm)	0.089 imes 0.054 imes 0.028
Radiation	Cu K α (λ = 1.54184)
θ Range for data collection (°)	8.088 to 157.678
Index ranges	$-14 \leq h \leq 13$, $-21 \leq k \leq 21$, $-10 \leq l \leq 9$
Reflections collected/independent	30763/3280
R _{int}	0.0285
Completeness (%)	100.0
Data/restraints/parameters	3280/0/191
Goodness-of-fit on F^2	1.048
Final <i>R</i> indices $[I > 2\sigma(I)]$	$R_1 = 0.0241, wR_2 = 0.0610$
Final <i>R</i> indexes [all data]	$R_1 = 0.0256, wR_2 = 0.0618$
Largest diff. peak/hole ($e \bullet Å^{-3}$)	0.31/-0.18

3.2. Synthesis of Organic Ligands (L1-L3)

The 1-(4-*n*-propyl)piperazine thioamide (1) was directly obtained by the reaction of the 1-*n*-propylpiperazine dihydrobromide with potassium thiocyanate in an aqueous solution [33]. Synthesis of 1-[2-thiazol-4-yl-(4-substitutedphenyl)]-4-*n*-propylpiperazines, outlined in Scheme 1, was accomplished according to the procedure described previously in [34].



Scheme 1. Synthesis of the 1-[2-thiazol-4-yl-(4-substitutedphenyl)]-4-n-propylpiperazines.

To a stirred solution of the corresponding 2-bromo-1-phenylethanone (1.60 mmol) in *n*-propanol (16 mL), the 1-(4-*n*-propyl)piperazine thioamide (1) (1.60 mmol) was added. The reaction mixture was heated at 90 °C for 5 h. After cooling, the precipitate was filtered and washed with a small amount of cold *n*-propanol and ethyl ether. The hydrobromide salt/product of the 1-[2-thiazol-4-yl-(4-substitutedphenyl)]-4-*n*-propylpiperazine was mixed with 5% sodium hydroxide aqueous solution for 1 h at room temperature, and then the aqueous layer was extracted with ethyl ether (3 × 30 mL). The organic extracts were washed with water (3 × 30 mL), dried over Na₂SO₄, and filtered. The solvent was evaporated to give the corresponding ligand.

L1 (C₁₆H₂₁N₃S; M = 287.42); mp 50–51 °C, yield: 0.245g, 53% white solid. ¹H NMR (600 MHz, CDCl₃) δ 7.85–7.81 (m, 2H, H_{arom}), 7.38–7.34 (m, 2H, H_{arom}), 7.28–7.24 (m, 1H, H_{arom}), 6.76 (s, 1H, H_{thiazole}), 3.61–3.53 (m, 4H, CH_{2piper}), 2.61–2.53 (m, 4H, CH_{2piper}), 2.39–2.33 (m, 2H, NCH₂CH₂CH₃), 1.59–1.49 (m, 2H, NCH₂CH₂CH₃), 0.93 (t, J = 7.4 Hz, 3H, NCH₂CH₂CH₃).¹³C NMR (151 MHz, CDCl₃) δ 171.1, 152.0, 135.2, 128.6 (2C), 127.7, 126.2 (2C), 101.5, 60.7, 52.6 (2C), 48.5 (2C), 20.1, 12.0. TLC (dichloromethane:methanol 9:1), Rf = 0.76;

L2 (C₁₆H₂₀ClN₃S; M = 321.87); mp 102–103 °C, yield: 0.338g, 65% white solid. ¹H NMR (600 MHz, CDCl₃), δ 7.82–7.72 (m, 2H, H_{arom}), 7.39–7.27 (m, 2H, H_{arom}), 6.74 (s, 1H_{thiazole}), 3.63–3.49 (m, 4H, CH_{2piper}), 2.63–2.52 (m, 4H, CH_{2piper}), 2.39–2.31 (m, 2H, NCH₂CH₂CH₂CH₃), 1.61–1.49 (m, 2H, NCH₂CH₂CH₃), 0.93 (t, J = 7.4 Hz, 3H, NCH₂CH₂CH₂CH₃). ¹³C NMR (151 MHz, CDCl₃) δ 171.1, 150.7, 133.6, 133.2, 128.6 (2C), 127.3 (2C), 101.7, 60.6, 52.4 (2C), 48.4 (2C), 20.0, 11.9. TLC (dichloromethane:methanol 9:1), Rf = 0.72;

L3 ($C_{17}H_{23}N_3OS$; M = 317.45); mp 65–66 °C, yield: 0.421g, 83% white solid. ¹H NMR (600 MHz, CDCl₃) δ 7.83–7.70 (m, 2H, H_{arom}), 6.96–6.85 (m, 2H, H_{arom}), 6.62 (s, 1H_{thiazole}), 3.82 (s, 3H,OCH₃), 3.61–3.50 (m, 4H, CH_{2piper}), 2.62–2.50 (m, 4H, CH_{2piper}), 2.37–2.33 (m, 2H, N<u>CH₂CH₂CH₃</u>), 1.59–1.47 (m, 2H, NCH₂<u>CH₂CH₃</u>), 0.93 (t, J = 7.4 Hz, 3H, NCH₂CH₂<u>CH₃</u>). ¹³C NMR (151 MHz, CDCl₃) δ 171.1, 159.3, 151.8, 128.3, 127.4 (2C), 113.9 (2C), 99.7, 60.8, 55.4, 52.6 (2C), 48.5 (2C), 20.1, 12.1. TLC (dichloromethane:methanol 9:1), Rf = 0.79.

3.3. Synthesis of Coordination Compounds

Figure 8 illustrates the general synthesis steps for all complexes.

 $Cu(L1)_2Cl_2$. The complex was synthesized by mixing the organic ligand, L1 (167.9 mg), and anhydrous CuCl₂ (81.2 mg), which were dissolved in *n*-propanol (30 mL). The mixture was rapidly stirred and heated for 1.5 h at 40 °C. The formed solid product was filtered and washed with cold *n*-propanol and left for drying.

Cu(*L*1)₂*Cl*₂ (C₃₂H₄₂S₂N₆CuCl₂; M = 709.29); mp 46.7 °C; yield: 15% brown solid. Anal.calcd. for C 54.18%; H 5.96%; N 11.85%; Cu 8.96%; S 9.04%. Found: C 54.31%; H 5.79%; N 11.91% Cu 9.34%; S 9.77%. FTIR spectra (KBr, cm⁻¹): v(NH) 3447,3360; v(CH) 2966–2607; v(C=N) 1599; 1230; v(C=C) 1520; β(CH) 1442; v(CS) 966, 837.



Figure 8. General scheme of the complexation process.

 $Cu(L2)Cl_2$. The complex was synthesized by mixing the organic ligand, L2 (82.6 mg), and anhydrous CuCl₂ (36.6 mg), which were dissolved in ethanol (30 mL). The mixture was rapidly stirred and heated for 1.5 h at 40 °C. The formed solid product was filtered and washed with cold ethanol and left for drying.

 $\begin{aligned} & Cu(L2)Cl_2 \ (C_{16}H_{20}N_3SCuCl_3; M = 456.33); mp \ 98.5 \ ^\circ C; \ yield: \ 40\% \ brown \ solid. \ Anal. \\ & calcd. \ for \ C \ 42.11\%; \ H \ 4.41\%; \ N \ 9.21\%; \ Cu \ 13.93\%; \ S \ 7.03\%. \ Found: \ C \ 42.99\%; \ H \ 4.19\%; \ N \\ & 8.91\%; \ Cu \ 13.36\%; \ S \ 7.87\%. \ FTIR \ spectra \ (KBr, \ cm^{-1}): \ \nu(NH) \ 3446, 3355; \ \nu(CH) \ 2968-2738; \\ & \nu(C=N) \ 1616, \ 1223; \ \nu(C=C) \ 1527; \ \beta(CH) \ 1276, \ 1229; \ \nu(CS) \ 967, 835; \ \nu(CCl) \ 732. \end{aligned}$

 $Cu(L3)Cl_2$. Complex Cu(L3)Cl₂ was synthesized by mixing the organic ligand, L3 (80 mg), and anhydrous CuCl₂ (39.2 mg), which were previously dissolved in ethanol (30 mL). The mixture was rapidly stirred and heated for 1.5 h at 40 °C. The formed solid product was filtered and washed with cold ethanol and left for drying.

Cu(*L*3)*Cl*₂ (C₁₇H₂₃SN₃OCuCl₂; M = 451.90); mp 56.5 °C; yield: 84% brown solid. Anal.calcd. for C 45.18%; H 5.13%; N 9.13%; Cu 14.06%; S 7.10%. Found: C 44.76%; H 4.99%; N 8.89%; Cu 14.89%; S 6.72%. FTIR spectra (KBr, cm⁻¹): v(NH) 3446, 3356; v(CH) 2965–2750; v(C=N) 1606; v(C=C) 1526; β(CH) 1457; v(C-O) 1250; v(CS) 964, 837.

3.4. Biological Activity Predictions

The activity prediction of three organic ligands was investigated using the toolkit service http://www.way2drug.com/ (accessed on 25 December 2021) [35]. The Way2Drug portal has been developed and supported by the multidisciplinary team of researchers working in bioinformatics, cheminformatics, and computer-aided drug discovery for about thirty years. The current version predicts several thousand different biological activities based on the structural formula of a drug-like organic compound.

3.5. ADME Analysis

The ADME analysis was performed using the Swiss ADME toolkit service. The SwissADME web tool gives free access to a pool of fast, yet robust predictive models for physicochemical properties, pharmacokinetics, drug-likeness, and medicinal chemistry friendliness, among which include in-house proficient methods such as the BOILED-Egg, iLOGP, and Bioavailability Radar [36]. The ProTOX II service was used for the prediction of toxicity for the compounds [37–39].

3.6. Microbiology

The antimicrobial activity, which was expressed as the minimum inhibitory concentration (MIC) of the synthesized compounds, was tested using the broth microdilution method according to the protocol established by the Clinical and Laboratory Standards Institute [40,41]. The stock solution of the compounds was prepared by dissolving them in DMSO. Serial dilution in the range of 1024–2 μ g/mL was made using 96-well plates. Secondly, the bacterial or fungal inoculums were added to the wells with Mueller Hinton

II Broth (for bacteria) or RPMI 1640 (for fungus) and diluted substances. Plates were incubated for 24 h and 48 h at 37 °C (bacteria) and 24 h at 37 °C (fungus). Assays were conducted in triplicate. The MIC was determined at the lowest concentration at which the visible growth of microorganisms was inhibited.

3.7. Cell Viability Analysis by MTT Assay

Breast cancer cells MCF7 and mouse fibroblast cell line L929 were used as a cell model for testing the effect of compounds on the viability of different types of cells. MCF7 and L929 cells were selected as cancerous and normal cell lines, respectively. Cells were cultured in Dulbecco's Modified Eagle Medium (DMEM) medium in 96-well plates with a concentration of 10,000 cells per well and incubated for 24 h in a 5% CO₂ atmosphere at 37 °C. After that, cells were treated with different concentrations—1, 10, 50, 100, and 200 μ M—of compounds for 24 h. The viability of cells treated with the tested compounds was assessed by measuring the ability of live cells to metabolize 3-(4,5-dimethylthiazolo-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) to formazan, using a standard protocol. Briefly, after incubation with compounds, the medium of each well was discarded, cells were washed with PBS, and then 100 μ L of fresh media with 10 μ L MTT solution (5 mg mL⁻¹ in PBS) was added to each well. After 4 h incubation at 37 °C, the produced formazan was solubilized by the addition of DMSO and the absorbance of each well was determined at 570 nm using an ELISA reader. The results were expressed as the mean of four replicates as a percentage of control (taken as 100%).

4. Conclusions

Numerous articles report that thiazole-based derivatives exhibit many notable pharmacological activities, including antiparasitic, anti-inflammatory, and antitumor activity [42–45]. This current study has proven the pharmacological activity of three synthesized 1-[2-thiazol-4-yl-(4-substitutedphenyl)]-4-*n*-propylpiperazines. Moreover, three newlysynthesized corresponding Cu(II) complexes were synthesized as well, and were shown to have promising biological activity. Among all synthesized compounds, two copper complexes, Cu(L1)₂Cl₂ and Cu(L3)Cl₂, presented cytotoxic effects. The complexation process has enhanced the pharmacological activity, as the other compounds without copper and Cu(L2)Cl₂ were not cytotoxic in the used concentration. In conclusion, the tests results from minimal inhibitory concentration assays indicate that the tested complexes have antifungal activity. The preliminary results contained in this paper indicate that the coordination compounds with these ligands are biologically active. Our results can be improved by further research.

Supplementary Materials: The following supporting information can be downloaded at: https://www.mdpi.com/article/10.3390/ijms23179844/s1. Which contains the supplementary crystallographic data for this paper and NMR spectra. CCDC 2193571 contains the supplementary crystallographic data for ligand L2. These data can be obtained free of charge via http://www.ccdc.cam.ac.uk/conts/retrieving.html (accessed on 25 December 2021) (or from the CCDC, 12 Union Road, Cambridge CB2 1EZ, UK; Fax: +44-1223-336033; E-mail: deposit@ccdc.cam.ac.uk).

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