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Synthesis, Molecular Docking Study, and Cytotoxic Activity against MCF Cells of New Thiazole–Thiophene Scaffolds

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Abstract: Investigating novel compounds that may be useful in designing new, less toxic, selective, and potent breast anticancer agents is still the main challenge for medicinal chemists. Thus, in the present work, acetylthiophene was used as a building block to synthesize a novel series of thiazole-bearing thiophene derivatives. The structures of the synthesized compounds were elucidated based on elemental analysis and spectral measurements. The cytotoxic activities of the synthesized compounds were evaluated against MCF-7 tumor cells and compared to a cisplatin reference drug, and against the LLC-Mk2 normal cell line using the MTT assay, and the results revealed promising activities for compounds 4b and 13a. The active compounds were subjected to molecular modeling using MOE 2019, the pharmacokinetics were studied using SwissADME, and a toxicity radar was obtained from the biological screening data. The results obtained from the computational studies supported the results obtained from the anticancer biological studies.

Keywords: thiophenes; thiazoles; multicomponent synthesis; hydrazonoyl halides; anticancer; molecular docking; toxicity radar; ADME



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1. Introduction

Breast cancer affects 14% of all women living globally [1]. It is the most frequently diagnosed neoplasm in female patients. The essential treatments for breast cancer are surgery, radiotherapy, and chemotherapy, individually tailored to the patient. Unfortunately, one of the main problems in the pharmacotherapy of cancers, including breast cancer, is the rapid development of drug resistance. Thus, designing and synthesizing more efficient agents with fewer adverse effects is essential [2]. Investigating novel compounds that may be useful in designing new, less toxic, selective, and potent anticancer agents is still the main challenge for medicinal chemists. Several studies have been carried out using various sulfur heterocycles, including thiophene and thiazole, directed towards different pathologies. It is reported in the literature that compounds containing a thiophene core have attracted considerable interest in drug discovery due to their potential anticancer activity [3–7]. Many anticancer agents are available on the market that contain a thiophene nucleus, as shown in Figure 1, and these exhibit their activity via multiple pathways involved in cancer [8,9].

On the other hand, 1,3-thiazole derivatives have been extensively considered by researchers generating novel lead compounds and in drug development. Thiazole derivatives of thiosemicarbazone are the scaffolds of many natural, synthetic, and semi-synthetic drugs which exhibit numerous remarkable pharmacological activities, including antiparasitic, anti-inflammatory, and antineoplastic activities [10–13]. Furthermore, the thiazole ring is present in several anticancer drugs (Figure 1), including Tiazofurin, Dasatinib, Dabrafenib,

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Bleomycin, Ixabepilone, and Epothilone, making this skeleton an ideal candidate for the development of more potent and safer anticancer drugs [14–16].

Figure 1. Lead compounds among thiazoles and thiophenes with anticancer activity.

Molecular hybridization is a beneficial approach to structural alteration involving the integration of a single species of two or more pharmacophores [11,17–23]. Over the last several years, hybrid drug design has been used as a prime method for developing novel anticancer therapies that can solve many of the pharmacokinetic disadvantages of traditional anticancer drugs. Thus, several studies have indicated that thiazole–thiophene hybrids have important anticancer activity [24].

Based on the above-mentioned promising aspects, and in continuation of our previous work to synthesise anticancer agents [11,24–28] from readily available, cheap, laboratory starting materials with anticipated biological activities, the strategy of this present work involved gathering the two bioactive entities, thiophene and thiazole, into one compact hybrid structure that may lead to enhanced anticancer activity due to the synergistic effect of both rings. Therefore, 2-acetylthiophene seemed to be a suitable starting material to fulfill this objective (Figure 2).

Figure 2. The targeted compounds with anticancer activity.

Apoptosis is essential to normal breast development and homeostasis. Pro- and antiapoptotic signals are tightly regulated in normal breast epithelial cells. Dysregulation of this Molecules **2022**, 27, 4639 3 of 19

balance is required for breast tumorigenesis, and this increases the acquired resistance to treatments such as molecularly targeted therapies, radiation, and chemotherapies. Members of the pro- or anti-apoptotic BCL-2 family of proteins are key regulators of the apoptosis process, and 2W3L is one of the proteins in this family. Therefore, 2W3L is a promising target to improve the killing of breast cancer tumor cells [29,30]. The docking process was carried out by simulating the exchange of the most biologically active compounds, 4a,b, 8a, 11b, and 13a,b, with two types of breast cancer proteins (PDB = 2W3L). Moreover, ADME analysis showed that the compounds have drug-like properties. ProToxII is one of the most common tools for predicting pharmacokinetic drug toxicity [31].

2. Results and Discussion

2.1. Chemistry

Heterocyclic compounds with three components have been widely exploited to prepare substituted heterocyclic compounds [25,32,33]. Thus, combining 2-acetylthiophene (1) with thiocarbohydrazide (2) and α -keto hydrazonoyl chlorides (3a–d) in ethanol with a catalytic amount of triethylamine under refluxing conditions led to the formation of 4-methyl5-(arylazo)-2-[(1-(thiophen-2-yl)ethylidene)hydrazineylidene]thiazol-3(2H)-amines (4a–d). Furthermore, this three-component process worked effectively with different substituent aryl groups on the hydrazonoyl chloride molecule (Scheme 1).

Scheme 1. Synthesis of aminothiazole derivatives **4a–d**.

We previously proposed the mechanistic pathway for this transformation starts with in situ condensations of 2-acetylthiophene (1) and thiocarbohydrazide (2) to give the respective hydrazone 5 [34]. Subsequent heterocyclization of intermediates 6a–d gave the isolable products 4a–d (Scheme 1). Assignment of spectral data (via IR, NMR, MS, and elemental analysis) for the isolated products provided significant indications of their structures. For example, in the IR spectra of compounds 4a–d, the appearance of characteristic bands at 3414–3420 and 3226–3230 cm⁻¹ confirms the presence of the amino group, and in ¹H-NMR, the singlet signal at approximately δ = 5 ppm is assigned to the resonance of amino protons on the thiazole ring [35].

Aspects of this methodology were extended to investigate the three-component reaction of 2-acetylthiophene (1), thiocarbohydrazide (2), and ethyl 2-chloro-2-(2-arylhydrazono) acetate (7a,b). Similarly, this process furnished the respective 3-amino-thiazolidine-4-one derivatives 8a,b, as illustrated in Scheme 2.

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Scheme 2. Synthesis of 3-amino-thiazolidin-4-one derivatives 8a,b.

The structures of the isolated products **8a** and **8b** were established from the stretching vibration signals at 3428, 3340, 3235, and 1680 cm⁻¹ in the IR spectra, which were attributed to NH₂, NH, and C=O groups, respectively. In addition, ¹H-NMR revealed two singlet signals at δ = 5.08–5.11 and 10.76–11.24 ppm (D₂O exchangeable) which were assigned to the NH₂ and NH protons.

Compounds 3a and 7a were expeditiously transformed into 4a and 8a, respectively, via their reactions with N'-[1-(thiophen-2-yl)ethylidene]hydrazinecarbothiohydrazide (5) [34], as depicted in Scheme 3.

EtOH / TEA /
$$\Delta$$

EtOH / TEA / Δ

EtOH / TEA / Δ

EtOH / TEA / Δ

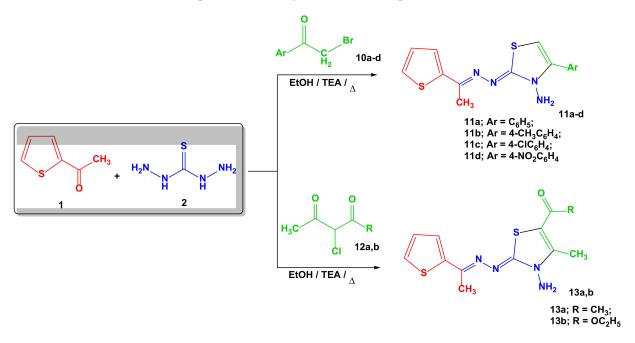
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Scheme 3. Alternative synthesis of compounds 4a and 8a.

To exploit the synthetic approach of a three-component system for the preparation of thiazole-3-amine derivatives, we condensed acetylthiophene (1) and thiocarbohydrazide (2) with either 2-bromo-1-arylethanone (10a–d) or α -chloro-dicarbonyl compounds (12a,b) under the previously employed conditions to obtain 11a–d or 13a,b, respectively (Scheme 4).

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Furthermore, the structures of the isolated products **11a–d** and **13a,b** were elucidated based on their spectral and analytical data (see Experimental Section).



Scheme 4. Synthesis of thiazole derivatives **12a–d** and **14a,b**.

2.2. Cytotoxic Potential

The cytotoxicity of the synthesized thiazoles **4a–d**, **8a,b**, **11a–d**, and **13a,b** was investigated against the human breast cancer MCF-7 and normal LLC-Mk2 cell lines using the MTT assay and cisplatin as the reference drug. The % inhibition was plotted against log concentration, with normalization applied and error bars for the MCF-7 cell line shown, as represented in Figure 3.

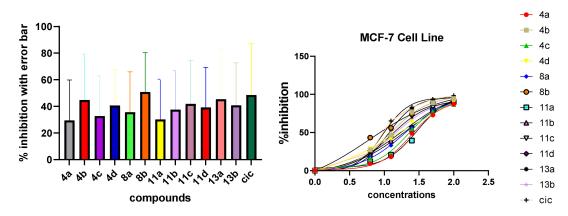


Figure 3. The % inhibition against log concentration, with normalization, and with error bars for MCF-7 cell line.

The results were used to plot a dose–response curve, from which the concentrations of the tested samples required to kill half of the cell population (IC_{50}) were determined. In addition, cytotoxic activities were expressed as the mean IC_{50} calculated from three independent experiments. The results, represented in Table 1, Figures 3 and 4, revealed that most of the tested compounds showed very variable activity compared to the reference drug.

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Table 1. In vitro cytotoxic activity of the newly synthesized thiazoles 4a-d, 8a,b, 11a-d, and 13a	,b
against MCF-7 and LLC-MK2.	

Tested	IC ₅₀ (μM)	CC ₅₀ (µM)	SI Values
Compounds	MCF-7	LLC-MK2	(CC_{50}/IC_{50})
4a	16.3 ± 1.0	183.05 ± 21.31	11.23
4b	10.2 ± 0.8	175.92 ± 18.24	17.25
4c	19.7 ± 1.3	-	-
4d	19.8 ± 0.9	-	-
8a	13.6 ± 0.9	149.46 ± 15.86	10.99
8b	23.7 ± 1.6	-	-
11a	21.0 ± 0.7	-	-
11b	17.9 ± 0.8	231.45 ± 25.03	12.93
11c	38.2 ± 1.4	-	-
11d	54.8 ± 1.6	-	-
13a	11.5 ± 0.7	162.65 ± 19.06	14.14
13b	16.3 ± 1.4	135.22 ± 9.58	8.29
Cisplatin	13.3 ± 0.61	158.75 ± 4.67	11.93

(-): not measured.

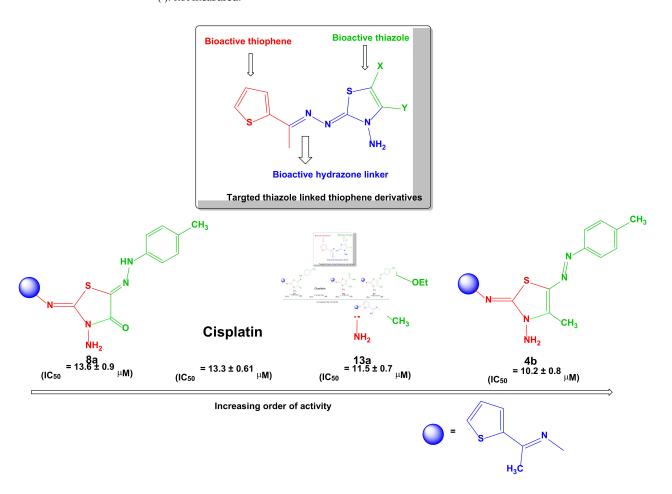


Figure 4. The order of activity toward the MCF-7 cell lines.

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Examination of the SAR leads to the following conclusions:

• The 1,3-thiazole derivatives **4b** and **13a** (IC₅₀ = 10.2 ± 0.7 and 11.5 ± 0.8 μ M, respectively) have promising antitumor activity against the breast carcinoma cell line (MCF-7), and showed greater activities than the cisplatin reference drug (IC₅₀ = 13.3 ± 0.61 μ M);

- The 1,3-thiazole derivatives **11c** and **11d** have poor antitumor activity ($IC_{50} > 38 \mu M$), while the rest of the evaluated thiazoles have moderate activity ($IC_{50} = 13.6-23.7 \mu M$);
- For 1,3-thiazoles 4, 8, and 11: the introduction of an electron-donating group (eg. methyl group) into phenyl group at position 5 in the 1,3-thiazole ring enhances the antitumor activity, while the introduction of an electron-withdrawing group (chlorine) decreases the activity (4b > 4a > 4c > 4d; 8a > 8b; and 11b > 11a > 11c > 11d);
- For the substituent at position 5 of the 1,3-thiazoles: an acetyl group (Ac) gives higher activity than an ester group (CO₂Et). **13a** (IC₅₀ = 11.5 \pm 0.7 μ M) > **13b** (IC₅₀ = 16.3 \pm 1.4 μ M).

The effects of the cisplatin standard drug and the most active compounds, **4a**, **4b**, **8a**, **11b**, **13a**, and **13b**, against LLC-Mk2 (rhesus monkey normal kidney epithelial cells) were also measured, to produce a dose–response curve and to calculate the fifty percent cytotoxic concentration (CC_{50}), as indicated in Table 1. The results showed that all examined compounds are non-toxic, because their CC_{50} toward normal cell lines was higher than 100 uM [36].

The selectivity index (SI) was calculated by dividing CC_{50} by IC_{50} . Our results showed that most of the derivatives presented good selectivity index values, indicating higher potency than the cisplatin anticancer drug. When the test compounds were evaluated for their toxicity against normal cells, they exhibited low toxic effects, indicating the safe use of most of them, but this may require further in vivo and pharmacological studies.

2.3. Molecular Docking Studies

Molecular docking is computational software routinely used for understanding the protein–receptor interaction with complexes. The docking process was carried out by simulating the exchange of the prepared compounds with two types of breast cancer proteins (PDB = 2W3L) [37] for compounds 4a,b, 8a, 11b, and 13a,b (Figure 5). Owing to the anticancer biological study results obtained, only these compounds were subjected to molecular docking because of their higher activity compared to the other synthesized compounds. Cisplatin energy was not calculated as it is difficult to calculate this using the software MOE2019, because it appears in the MOE system as a square planner molecule, not a cis molecule. To solve this issue, we used carboplatin as a reference docking drug, and the new results have been added to the corrected manuscript.

The docking score energies of compounds 4a, 4b, 8a, 11b, 13a, 13b, and CarboPt were (-5.911, -6.011, -6.161, -5.65, -5.436, -5.883, and -4.671 kcal/mol), respectively. These scores are only of moderate activity, not higher, and this can be explained by the following: only compound 4b showed an interaction of the S(8) atom on the ligand to the receptor on the O of Arg 66 (B) by hydrogen donation, while all the other compounds had no measurable interaction, and all of them connected to the dummies by ligand exposure, as shown in Table 2. Having no measurable interactions depends on the exposure of the whole ligand to the protein. The results also indicated that compound 4b was the best one, as its IC_{50} was closest to cisplatin, with good SI values of 46–30. These results are consistent with the data which show that 4b was the only one with measurable interactions of high docking score energy.

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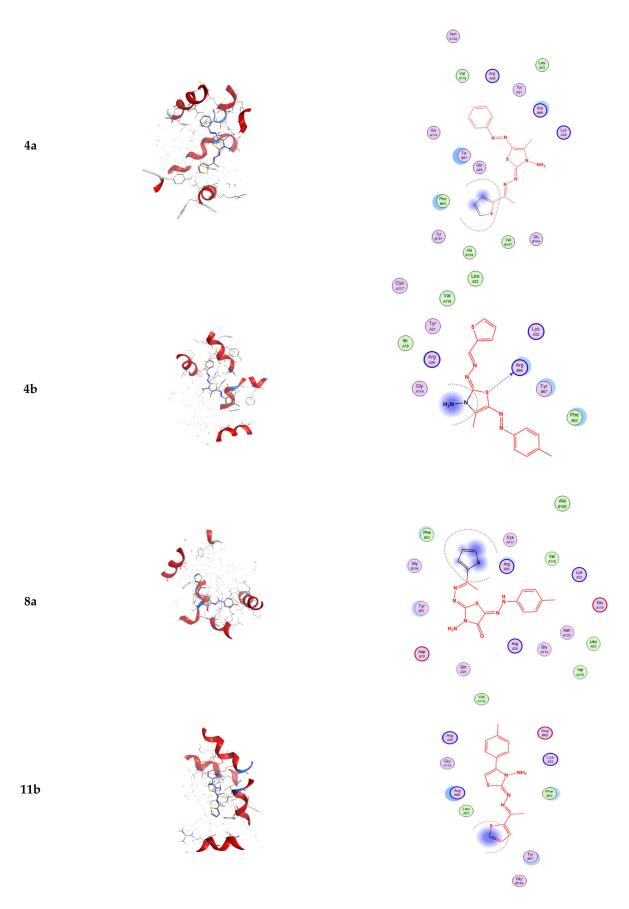


Figure 5. Cont.

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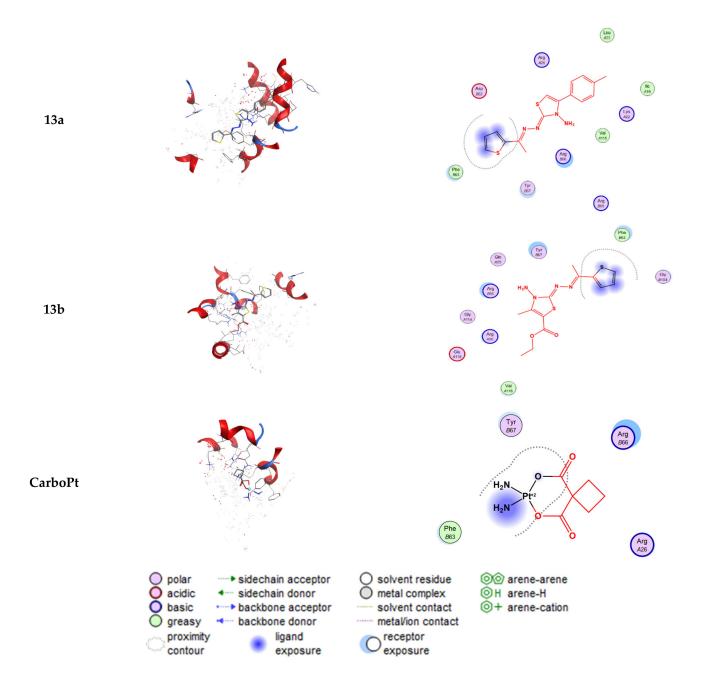


Figure 5. 3D and 2D snapshots showing the hydrophilicity interaction with the (PDB = 2W3L) receptor.

Table 2. Docking interactions of selected compounds with breast cancer cells (PDB = 2W3L).

Compound	Ligand	Receptor	Interaction	Distance	E (kcal/mol)
4a	No measurable interaction				
4b	S (8)	O ARG 66 (B)	H-donor	3.69	-0.5
8a	No measurable interaction				
11b	No measurable interaction				
13a	No measurable interaction				
13b	No measurable interaction				
Carbo-Pt	No measurable interaction				

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The molecular docking studies were also carried out on HAS (Human Serum Albumin) PDB = 1AO6, but only with **4b**, **13a**, and carboplatin (Figure 6). This was to validate the results obtained from the docking studies on PDB=2W3l. The docking score energy of compound **4b** with PDB=1AO6 was -6.3 kcal/mol, while that of compound **13a** was -5.228 kcal/mol. Furthermore, both **4b** and **13a** showed no measurable interactions with 1AO6. The only interaction was via ligand exposure. In addition, the docking score of CarboPt with 1AO6 was -4.78 kcal/mol.

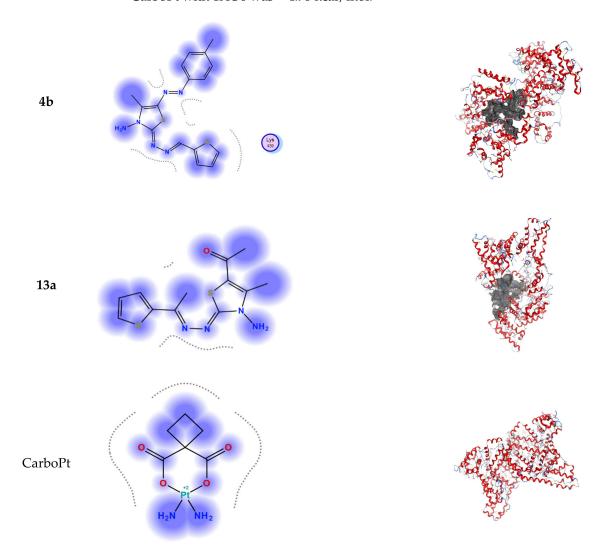


Figure 6. 3D and 2D snapshots showing the hydrophilicity interaction with the (PDB = 1AO6) HAS receptor.

2.4. Toxicity Radar

The ProTox-II data showed that the tested compounds (4a, 4b, 8a, 11b, 13a, and 13b) were predicted to have oral LD₅₀ values ranging from 159 to 3000 mg/kg in a rat model, with (1 s, 4 s)-eucalyptol bearing the highest value, and quercetin holding the lowest one (Figure 7 and Table 3). Therefore, the SI (selectivity index) calculations for these compounds obtained from the biological studies were adequately compatible with the toxicity radar calculations, which validates the results obtained. Through screening the toxicity radar results, we found that compound 4b had a higher predictable LD₅₀, which agrees with the results obtained from the molecular docking and the biological activities. However, the prediction accuracy for all compounds was 12%, so these compounds should be further investigated and screened.

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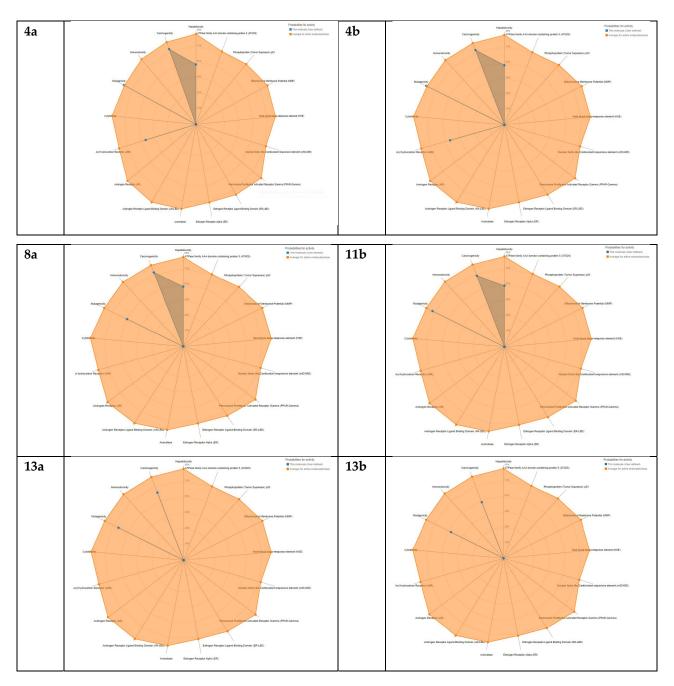


Figure 7. Toxicity radar for compounds 4a, 4b, 8a, 11b, 13a, and 13b.

Table 3. The predicted toxicity for compounds 4a, 4b, 8a, 11b, 13a, and 13b using ProTox-II.

	4a	4b	8a	11b	13a	13b
Predicted LD ₅₀ (mg/kg)	525	3200	1000	1000	300	1000
Predicted toxicity class	4	5	4	4	3	4
Average similarity (%)	29.02	29.63	33.11	38.09	32.55	37.66
Prediction accuracy (%)	12	12	23	23	23	23

2.5. SwissADME Studies

ADME (absorption, distribution, metabolism, and excretion) studies, including druglikeness analysis, are essential in drug discovery, and provide a reasonable decisiveness on whether or not inhibitors should be progressed to a biological system [31]. A potent antagonistic interaction of inhibitors with a receptor protein or enzyme can not guarantee Molecules **2022**, 27, 4639 12 of 19

the ability of an inhibitor to act as a drug; therefore, ADME assessment is essential in drug development. Inhibitors having low ADME properties and high toxicity effects on biological systems are often the dominant reasons for the failure of most medicines in the experimental phase.

Figure 8 shows the output of the ADME studies and the drug-likeness properties (refer to the Supplementary Materials); it was observed that the **4b** and **13a** molecules display one or two violations of Lipinski's rule, and the first violation is the molecular weight rule, with a result of 356.47–425.36 g/mol. The drug-likeness parameters are related to aqueous solubility and intestinal permeability, determining the first step of oral bioavailability [38]. The results also indicated good pharmacokinetic properties, in which compounds **4b** and **13a** have high gastrointestinal absorption.

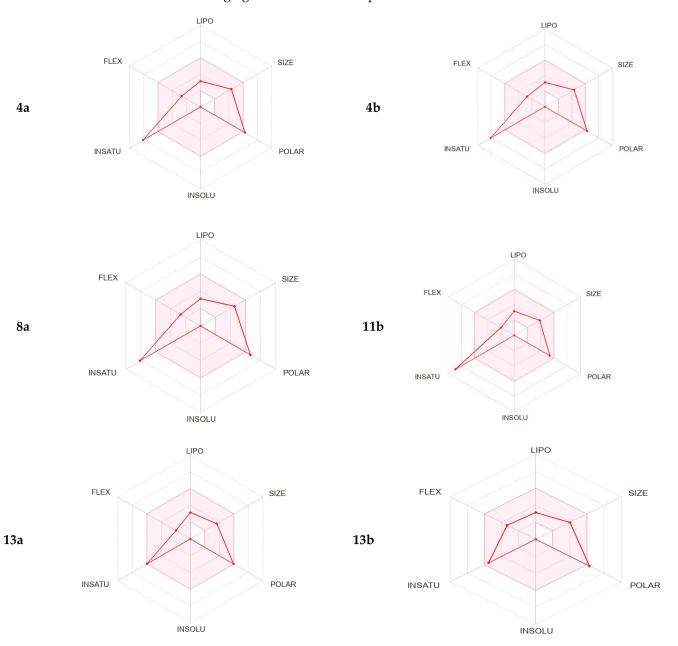


Figure 8. Toxicity profiles of the evaluated compounds using the SwissADME server.

2.6. Pred-hERG

Chemically similar compounds often bind to biologically diverse protein targets, and protein structures do not always recognize identical ligands. Pharmacological and off-

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target relationships between proteins and a ligand set help to improve machine learning confidence by interpolating the output prediction equalized by the compound similarity criteria. This pipeline helps to improve predictions of off-target drug effects, reducing false-negative errors. The Labmole server was used to predict Pred-hERG, and to predict similar compounds with structure–activity relationships (Table 4, Figure 8).

Table 4. Predicted toxicity for compounds 4b and 13a using Pred-hERG software.

Duom outre	Pred-hERG			
Property	4b	13a		
Prediction/Potency	Weak or Moderate	Weak or Moderate		
Confidence (%)	60	70		
Applicability domain (AD)	No (Value = 0.19 and limit = 0.26)	No (Value = 0.23 and limit = 0.26)		

Chemical similarity is one of the most critical concepts in cheminformatics. One commonly used algorithm to calculate these similarity measures is the 2D Tanimoto algorithm employed here (Figures 9–11). The resulting Tanimoto coefficient is fingerprint-based, encoding each molecule to a fingerprint "bit" position (MACCS), with each bit recording the presence ("1") or absence ("0") of a fragment of the molecule. Interpretation of the probability of toxicity for compounds **4b** and **13a** can be explained by the cytotoxicity diagram [31].

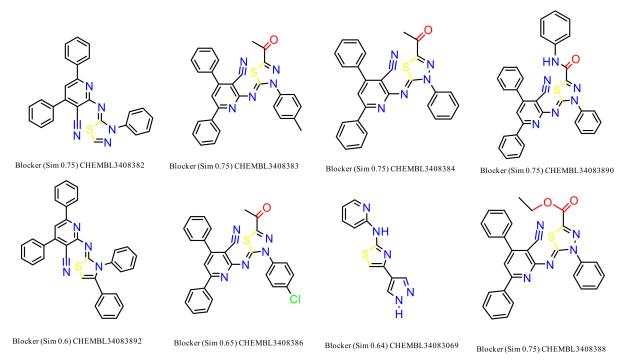


Figure 9. Similar off-target compounds to compound 4b.

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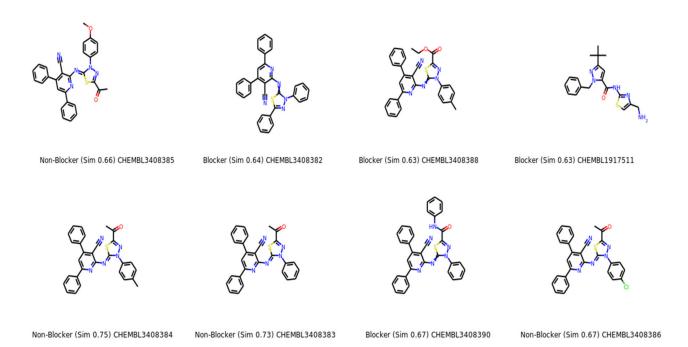


Figure 10. Similar off-target compounds to compound 13a.

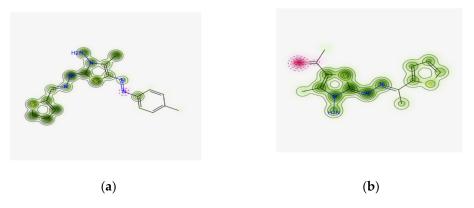


Figure 11. Probability map of HERG for **4b** (**a**) and **13a** (**b**). The more contour lines and the more intense green color means a higher positive contribution of an atom or a fragment to the hERG blockage, while pink coloration means that it contributes to a decrease in the hERG blockage, and gray color means no contribution.

3. Experimental Section

3.1. Chemistry

3.1.1. Experimental Instrumentation

All melting points were determined using electrothermal apparatus, and were left uncorrected. IR spectra were recorded (KBr disc method) using a Shimadzu FT-IR 8201 PC spectrophotometer. ¹H NMR and ¹³C NMR spectra were recorded in DMSO solutions using a BRUKER 400 FT-NMR spectrometer, and chemical shifts were expressed in ppm using TMS as an internal reference. Mass spectra were recorded using a Shimadzu GC-MS QP1000 EX. Elemental analyses were carried out at the Microanalytical Center of Cairo University.

3.1.2. General Procedure for Synthesizing the Thiazole Derivatives **4a–d**, **8a,b**, **11a–d**, and **13a,b**

A mixture of 2-acetylthiophene (1) (0.126 g, 1 mmol) and thiocarbohydrazide (2) (0.106 g, 1 mmol) in ethanol (20 mL) was refluxed with a few drops of hydrochloric acid for one hour. Then, without extraction of the hydrazone product, either the appropriate

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hydrazonoyl chlorides, **3a-d** or **7a,b**, or the α -halocarbonyl compounds, **10a-d** or **12a,b**, (1 mmol), were added with catalytic amounts of triethylamine, and the reaction mixture was refluxed for 4 h (monitored by TLC). Finally, the precipitate formed was isolated by filtration, washed with methanol, dried, and recrystallized from the appropriate solvent to give products **4a-d**, **8a,b**, **11a-d**, or **13a,b**, respectively. The physical properties and spectral data of the isolated products are listed below.

4-Methyl-5-(phenyldiazenyl)-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene] thiazol-3(2H)-amine (4a). Red solid, 74% yield, m.p. 173–175 °C (EtOH); IR (KBr): v 3414, 3229 (NH₂), 1602 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.25 (s, 3H, CH₃-thiazole), 2.46 (s, 3H, CH₃-C=N), 5.08 (s, 2H, Ar-H), 7.07–7.63 (m, 8H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 13.21 (CH₃-thiazole), 15.84 (CH₃-C=N), 110.68, 125.11, 126.14, 127.39, 128.25, 129.27, 134.87, 142.47, 144.41, 145.32, 158.24, 161.04; MS m/z (%): 356 (M⁺, 27). Analysis calculated for C₁₆H₁₆N₆S₂ (356.09): C, 53.91; H, 4.52; N, 23.58; S, 17.99; Found: C, 53.80; H, 4.42; N, 23.39; S, 18.08%.

4-Methyl-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene]-5-((p-tolyldiazenyl) thiazol-3(2H)-amine (4b). Red solid, 76% yield, m.p. 190–192 °C (EtOH); IR (KBr): v 3420, 3230 (NH₂), 1598 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ 2.27 (s, 3H, CH₃-thiazole),), 2.39 (s, 3H, Ar-CH₃), 2.46 (s, 3H, CH₃-C=N), 5.11 (s, 2H, Ar-H), 7.02–7.61 (m, 7H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 12.89 (CH₃-thiazole), 16.14 (CH₃-C=N), 21.28 (Ar-CH₃), 109.98, 125.13, 126.32, 127.19, 128.64, 130.27, 133.87, 140.47, 143.41, 145.32, 158.74, 160.81; MS m/z (%): 370 (M⁺, 39). Analysis calculated for C₁₇H₁₈N₆S₂ (370.10): C, 55.11; H, 4.90; N, 22.68; S, 17.31; Found: C, 55.03; H, 4.77; N, 22.51; S, 17.25%.

5-((4-Chlorophenyl)diazenyl)-4-methyl-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene] thiazol-3(2*H*)-amine (4c). Dark red solid, 75% yield, m.p. 205–207 °C (DMF); IR (KBr): v 3418, 3226 (NH₂), 1600 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.28 (s, 3H, CH₃-thiazole),), 2.46 (s, 3H, CH₃-C=N), 5.03 (s, 2H, Ar-H), 7.08–7.63 (m, 7H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 12.91 (CH₃-thiazole), 16.04 (CH₃-C=N), 111.18, 124.81, 125.94, 126.99, 127.53, 128.85, 132.81, 142.49, 143.41, 145.31, 158.17, 160.94; MS m/z (%): 392 (M⁺+ 2, 12), 390 (M⁺, 32). Analysis calculated for C₁₆H₁₅ClN₆S₂ (390.05): C, 49.16; H, 3.87; N, 21.50; S, 16.40; Found: C, 49.27; H, 3.71; N, 21.44; S, 16.52%.

5-((2,4-Dichlorophenyl)diazenyl)-4-methyl-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene] thiazol-3(2*H*)-amine (4d). Brown solid, 79% yield, m.p. 227–229 °C (DMF); IR (KBr): v 3415, 3228 (NH₂), 1600 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.28 (s, 3H, CH₃-thiazole),), 2.48 (s, 3H, CH₃-C=N), 5.11 (s, 2H, Ar-H), 7.08–7.81 (m, 6H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 13.27 (CH₃-thiazole), 16.02 (CH₃-C=N), 110.54, 124.91, 126.18, 127.89, 128.23, 130.27, 132.58, 134.87, 135.76, 142.43, 144.41, 145.65, 156.24, 162.14; MS m/z (%): 424 (M⁺, 25). Analysis calculated for C₁₆H₁₄Cl₂N₆S₂ (424.01): C, 45.18; H, 3.32; N, 19.76; S, 15.07; Found: C, 45.03; H, 3.25; N, 19.68; S, 15.14%.

3-Amino-2-[(1-(thiophen-2-yl)ethylidene)hydrazineylidene]-5-(2-(p-tolyl)hydrazineylidene) thiazolidin-4-one (8a). Yellow solid, 70% yield, m.p. 155–157 °C (EtOH); IR (KBr): v 3428, 3340, 3235 (NH $_2$ & NH), 1680 (C=O), 1598 (C=N) cm $^{-1}$; 1 H-NMR (DMSO- d_6): δ = 2.37 (s, 3H, Ar-CH $_3$), 2.47 (s, 3H, CH $_3$ -C=N), 5.38 (s, 2H, NH $_2$), 7.08–7.75 (m, 7H, Ar-H), 10.76 (s, 1H, NH); 13 C-NMR (DMSO- d_6): δ = 15.12 (CH $_3$ -C=N), 21.14 (Ar-CH $_3$), 120.11, 122.13, 124.32, 125.19, 127.64, 128.55, 129.27, 140.47, 146.87, 153.41, 159.74 (Ar-Cs), 168.81 (C=O); MS m/z (%): 372 (M $^+$, 62). Analysis calculated for C $_{16}$ H $_{16}$ N $_{6}$ OS $_{2}$ (372.08): C, 51.60; H, 4.33; N, 22.56; S, 17.22; Found: C, 51.45; H, 4.19; N, 22.61; S, 17.35%.

3-Amino-5-(2-(4-chlorophenyl)hydrazineylidene)-2-[(1-(thiophen-2-yl)ethylidene) hydrazineylidene]thiazolidin-4-one (8b). Yellow solid, 74% yield, m.p. 177–179 °C(DMF-EtOH); IR (KBr): v 3428, 3342, 3236 (NH₂ & NH), 1681 (C=O), 1599 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.46 (s, 3H, CH₃-C=N), 5.32 (s, 2H, NH₂), 7.01–7.62 (m, 7H, Ar-H), 11.24 (s, 1H, NH); ¹³C-NMR (DMSO- d_6): δ = 15.12 (<u>CH₃-C=N</u>), 120.08, 121.93, 124.31, 125.21, 127.94, 128.51, 130.17, 141.41, 146.82, 151.87, 159.72 (Ar-Cs), 168.87 (C=O); MS m/z (%): 394 (M⁺+ 2, 6), 392 (M⁺, 16). Analysis calculated for C₁₅H₁₃ClN₆OS₂ (392.03): C, 45.86; H, 3.34; N, 21.39; S, 16.32; Found: C, 45.71; H, 3.45; N, 21.26; S, 16.44%.

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4-Phenyl-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene]thiazol-3(2*H***)-amine (11a). Yellowish-white crystals, 79% yield, m.p. 166–168 °C (EtOH); IR (KBr): v 3424, 3220 (NH₂), 1599 (C=N) cm⁻¹; ¹H-NMR (DMSO-d_6): \delta = 2.37 (s, 3H, CH₃-C=N), 4.77 (s, 2H, NH₂), 7.02 (s, 1H, thiazole-H), 7.08–7.83 (m, 8H, Ar-H); ¹³C-NMR (DMSO-d_6): \delta = 15.12 (<u>CH₃-C=N</u>), 112.08, 124.13, 125.31, 126.21, 127.94, 128.53, 135.17, 139.11, 144.54, 151.87, 159.72, 164.87 (Ar-Cs); MS m/z (%): 314 (M⁺, 51). Analysis calculated for C₁₅H₁₄N₄S₂ (314.07): C, 57.30; H, 4.49; N, 17.82; S, 20.39; Found: C, 57.16; H, 4.57; N, 17.69; S, 20.51%.**

2-[((1-(Thiophen-2-yl)ethylidene)hydrazineylidene]-4-(*p***-tolyl)thiazol-3(2***H***)-amine (11b).** Yellowish-white crystals, 75% yield, m.p. 149–151 °C (EtOH); IR (KBr): v 3430, 3214 (NH₂), 1598 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.31 (s, 3H, Ar-CH₃), 2.37 (s, 3H, CH₃-C=N), 4.81 (s, 2H, NH₂), 7.08 (s, 1H, thiazole-H), 7.23–7.79 (m, 7H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 14.82 (CH₃-C=N), 21.18 (Ar-CH₃), 112.08, 124.04, 125.11, 126.18, 127.24, 128.53, 134.21, 139.11, 143.54, 152.87, 159.72, 163.94 (Ar-Cs); MS m/z (%): 328 (M⁺, 100). Analysis calculated for C₁₆H₁₆N₄S₂ (328.08): C, 58.51; H, 4.91; N, 17.06; S, 19.52; Found: C, 58.63; H, 4.80; N, 17.11; S, 19.63%.

4-(4-Chlorophenyl)-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene]thiazol-3(2H)-amine (11c). Yellow solid, 73% yield, m.p. 180–182 °C (DMF); IR (KBr): v 3428, 3220 (NH₂), 1603 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.41 (s, 3H, CH₃-C=N), 4.86 (s, 2H, NH₂), 7.08 (s, 1H, thiazole-H), 7.11–7.84 (m, 7H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 14.82 (<u>CH₃-C=N</u>), 110.11, 120.13, 125.31, 126.21, 127.94, 128.53, 132.17, 135.11, 145.54, 149.87, 155.72, 164.87 (Ar-Cs); MS m/z (%): 350 (M⁺ + 2, 19), 348 (M⁺, 46). Analysis calculated for C₁₅H₁₃ClN₄S₂ (348.03): C, 51.64; H, 3.76; N, 16.06; S, 18.38; Found: C, 51.50; H, 3.84; N, 15.93; S, 18.44%.

4-(4-Nitrophenyl)-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene]thiazol-3(2H)-amine (11d). Brown solid, 81% yield, m.p. 194–196 °C (DMF); IR (KBr): v 3424, 3226 (NH₂), 1600 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.43 (s, 3H, CH₃-C=N), 4.84 (s, 2H, NH₂), 7.05 (s, 1H, thiazole-H), 7.11–7.84 (m, 7H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 14.59 (<u>CH₃-C=N</u>), 110.11, 123.13, 124.31, 125.21, 126.94, 127.53, 142.17, 145.11, 149.54, 151.87, 155.72, 162.87 (Ar-Cs); MS m/z (%): 359 (M⁺, 73). Analysis calculated for C₁₅H₁₃N₅O₂S₂ (359.05): C, 50.13; H, 3.65; N, 19.49; S, 17.84; Found: C, 50.05; H, 3.51; N, 19.37; S, 17.72%.

1-[3-Amino-4-methyl-2-((1-(thiophen-2-yl)ethylidene)hydrazineylidene]-2,3-dihydrothiazol-5-yl)ethan-1-one (13a). Yellowish-white crystals, 77% yield, m.p. 156–158 °C (EtOH); IR (KBr): v 3416, 3220 (NH₂), 1715 (C=O), 1591 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 2.25 (s, 3H, CH₃-thiazole), 2.41 (s, 3H, CH₃-C=N), 2.49 (s, 3H, COCH₃), 5.12 (s, 2H, NH₂), 7.02–7.62 (m, 3H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 9.05 (CH₃-thiazole), 15.37 (CH₃-C=N), 40.52 (COCH₃), 126.12, 127.49, 128.18, 142.11, 143.99, 145.66, 157.51, 164.47 (Ar-Cs), 181.91 (C=O); MS m/z (%): 294 (M⁺, 49). Analysis calculated for C₁₂H₁₄N₄OS₂ (294.06): C, 48.96; H, 4.79; N, 19.03; S, 21.78; Found: C, 48.79; H, 4.63; N, 19.01; S, 21.68%.

Ethyl 3-amino-4-methyl-2-[((1-(thiophen-2-yl)ethylidene)hydrazineylidene]-2,3-dihydro thiazole-5-carboxylate (13b). Yellowish-white crystals, 71% yield, m.p. 141–143 °C (EtOH); IR (KBr): v 3420, 3222 (NH₂), 1721 (C=O), 1595 (C=N) cm⁻¹; ¹H-NMR (DMSO- d_6): δ = 1.14 (t, 3H, CH₃-CH₂), 2.28 (s, 3H, CH₃-thiazole), 2.46 (s, 3H, CH₃-C=N), 3.02 (q, 2H, CH₂CH₃), 5.11 (s, 2H, NH₂), 7.02–7.62 (m, 3H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 9.11 (CH₃-thiazole), 15.05 (CH₃-CH₂), 15.36 (CH₃-C=N), 56.52 (CH₂CH₃), 126.51, 127.45, 128.04, 143.11, 144.19, 145.68, 157.52, 163.47 (Ar-Cs), 182.61 (C=O); MS m/z (%): 324 (M⁺, 27). Analysis calculated for C₁₃H₁₆N₄O₂S₂ (324.07): C, 48.13; H, 4.97; N, 17.27; S, 19.76; Found: C, 48.06; H, 4.83; N, 17.19; S, 19.61%.

3.1.3. Alternate Synthesis of 4a and 8a

A mixture of N'-[1-(thiophen-2-yl)ethylidene]hydrazinecarbothiohydrazide (5) (0.214 g, 1 mmol) with the appropriate 2-oxo-N'-phenylpropanehydrazonoyl chloride (3a) or ethyl 2-chloro-2-(2-(p-tolyl)hydrazineylidene)acetate (7a) (1 mmol) in ethanol (20 mL) containing a catalytic amount of TEA was refluxed for 4 h (monitored by TLC). The precipitate formed was isolated by filtration, washed with methanol, dried, and recrystallized from EtOH to

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ensure the product was identical in all respects (m.p., mixed mp, and IR spectra) with the products **4a** or **8a**, respectively.

3.2. In Vitro Cytotoxic Activity

The cytotoxic potentials of the newly synthesized compounds was carried out at the Regional Center for Mycology and Biotechnology at Al-Azhar University, Cairo, Egypt. Cells were purchased from the Egyptian Holding Company for Biological Products and Vaccines (VACSERA, Giza, Egypt) and kept in a tissue culture unit. Cells were grown in Roswell Park Memorial Institute (RPMI)-1640 medium supplemented with 10% heatinactivated fetal bovine serum (FBS), 50 units/mL penicillin, and 50 mg/mL streptomycin, and maintained in a humidified atmosphere containing 5% CO₂ [11,25]. Cells were maintained as monolayer cultures using serial subculture. Cell culture reagents were obtained from Lonza (Basel, Switzerland). The anticancer activities of the rest of the compounds were evaluated in MCF-7 (breast cancer) cells. In addition, the sulforhodamine B (SRB) assay method, as described previously in [39,40], was used to determine cytotoxicity. Exponentially growing cells were collected using 0.25% trypsin-EDTA and seeded in 96-well plates at 1000-2000 cells/well in RBMI-1640 supplemented medium. After 24 h, cells were incubated for 72 h with various concentrations of the compounds tested. Following 72 h of incubation, the cells were fixed with 10% trichloroacetic acid for 1 h at 4 °C. Wells were stained for 10 min at room temperature with 0.4% sulforhodamine B (SRBC) dissolved in 1% acetic acid. Plates were air-dried for 24 h, and the dye was solubilized with Tris-HCl for 5 min on a shaker at 1600 rpm. The optical density (OD) of each well was measured spectrophotometrically at 564 nm using an ELISA microplate reader (ChroMate-4300, Palm City, FL, USA). IC₅₀ values were calculated using a Boltzmann sigmoidal concentration response curve using non-linear regression fitting models (Graph Pad, Prism Version 9, GraphPad Software, San Diego, CA, USA).

4. Conclusions

In summary, acetylthiophene was employed as a critical intermediate to synthesize a novel series of thiazole-bearing thiophene derivatives. The assigned structure for all of the newly synthesized compounds was elucidated by elemental and spectral analysis data, and the mechanisms accounting for their formation were discussed. The in vitro growth inhibitory activity of the synthesized compounds against MCF-7 tumor cells was investigated in comparison with cisplatin as a standard drug using the MTT assay, and the results revealed promising activities for compounds **4b** and **13a**. The results obtained from the computational studies, including molecular modeling, pharmacokinetics, and toxicity radar, supported the results obtained from the anticancer biological studies.

Supplementary Materials: The following supporting information can be downloaded at: https://www.mdpi.com/article/10.3390/molecules27144639/s1, Figure S1: The extra ADME results.

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Conflicts of Interest: The authors declare that there are no conflicts of interest regarding the publication of this paper.

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