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Novel quinazolines bearing 1,3,4-thiadiazole-aryl urea derivative as anticancer agents: design, synthesis, molecular docking, DFT and bioactivity evaluations

Sara Masoudinia¹, Marjaneh Samadizadeh^{1*}, Maliheh Safavi², Hamid Reza Bijanzadeh³ and Alireza Foroumadi^{4,5*}

Abstract

A novel series of 1-(5-((6-nitroquinazoline-4-yl)thio)-1,3,4-thiadiazol-2-yl)-3-phenylurea derivatives **8** were designed and synthesized to evaluate their cytotoxic potencies. The structures of these obtained compounds were thoroughly characterized by IR, ¹H, and ¹³C NMR, MASS spectroscopy and elemental analysis methods. Additionally, their in vitro anticancer activities were investigated using the MTT assay against A549 (human lung cancer), MDA-MB231 (human triple-negative breast cancer), and MCF7 (human hormone-dependent breast cancer). Etoposide was used as a reference marketed drug for comparison. Among the compounds tested, compounds 8b and 8c demonstrated acceptable antiproliferative activity, particularly against MCF7 cells. Considering the potential VEGFR-2 inhibitor potency of these compounds, a molecular docking study was performed for the most potent compound, **8c**, to determine its probable interactions. Furthermore, computational investigations, including molecular dynamics, frontier molecular orbital analysis, Fukui reactivity descriptor, electrostatic potential surface, and in silico *ADME* evaluation for all compounds were performed to illustrate the structure–activity relationship (SAR).

Keywords Quinazoline, Anticancer, Molecular docking, In silico ADME, MTT assay

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Introduction

Cancer is a complex and severe human disease characterized by abnormal cell growth that can potentially spread to other parts of the body. It is currently recognized as the second leading cause of death worldwide [1]. Developed countries are experiencing an alarming increase in cancer-related mortality rates, with approximately 10 million deaths recorded in 2020. Predictions suggest that this number will rise to 16.3 million by 2040 [2, 3].

In response to this growing concern, extensive efforts have been made in the field of medicinal chemistry to develop innovative multi-target chemotherapeutics. Various compounds with anticancer properties, such as thiadiazole, quinazolines, and urea derivatives, have been explored [4–12]. Quinazoline, in particular, is a significant aza-heterocycle that not only exhibits strong antiproliferative effects but also demonstrates a wide range of biological activities, including antioxidant, antibacterial, anti-inflammatory, antimutagenic, antifungal, and antiviral properties [13–21].

Furthermore, several commercially available anticancer drugs, including Erlotinib, Lapatinib, Vandetanib, Afatinib, and Gefitinib, contain quinazoline as a key component and are commonly used worldwide.

Thiadiazole is another common 5-membered heterocyclic system containing one sulfur atom and two nitrogen atoms. This heterocyclic core structure is present in numerous commercially available drugs and biologically active natural products. The remarkable aromaticity, non-toxicity, and exceptional in vivo stability of thiadiazole make it highly desirable for use in bioactive compounds [22].

In recent studies, various thiadiazoles, particularly 1,3,4-thiadiazole derivatives, have been extensively investigated for their potential antitumor activity These compounds have shown significant potency, attracting considerable attention as a promising avenue of research for developing potential anti-cancer agents [23–31].

The urea functionality plays a crucial role in drug development and medicinal chemistry due to its ability to form multiple stable hydrogen bonds with various protein and receptor targets. Urea and its derivatives have been utilized to modulate drug potency, selectivity, and improve drug properties in the development of anticancer, antibacterial, anticonvulsive, anti-HIV, and antidiabetic agents [32–34].

Notably, several FDA-approved drugs containing the urea functionality, such as Regorafenib, Sorafenib, and Tivozanib, have been successfully developed and are widely used as effective anticancer drugs in the global market [35–37] (Fig. 1)

Considerable attention has been focused on ongoing research dedicated to discovering innovative and effective anti-cancer compounds that can target multiple pathways involved in the growth and spread of cancer cells. The process of angiogenesis, which encompasses the formation and maintenance of blood vessels, plays a vital role in cell development and replication. In cancer cells, this physiological process takes on a similar role and becomes crucial in the progression and metastasis of tumors. Therefore, utilizing efficient anti-angiogenesis agents may be proposed as a suitable approach to combat cancer [38].

Vascular endothelial growth factors (VEGF) and their receptors (VEGFR) have been proven to play significant regulatory roles in pathological angiogenesis, particularly in cancer. Among the three subtypes of VEGFRs, VEGFR-2 plays a critical role in promoting tumor angiogenesis. The interaction between VEGF and VEGFR-2 results in dimerization and subsequent auto-phosphorylation of tyrosine residues (Tyr1059 and Tyr1054) within the receptor. This activation initiates signaling pathways that contribute to angiogenesis in the tumor microenvironment [39].

Hence, the use of antitumor substances that disrupt tyrosine phosphorylation demonstrates inhibitory effects on angiogenesis [40]. Therefore, numerous compounds bearing valuable scaffolds particularly quinazolines, thiadiazole, and urea functionality showing great VEGFR-2 inhibitory potencies have been developed. For example, the compound SKLB1002 is a type of quinazoline-bearing 1,3,4-thiadiazole that has been identified as a potent inhibitor of VEGFR-2.

It has shown minimal toxicity in studies [41, 42]. Diaryl ureas have traditionally been used to inhibit VEGFR-2 activity, but there have been advancements in modifying their structure to enhance their effectiveness. For instance, researchers have substituted the urea functional group with a thiourea moiety and incorporated additional rings such as 1,3,4-thiadiazole, oxadiazole, and 1,2,3-triazole to improve binding affinity with the receptor [22, 43, 44].

In the present study, the researchers aimed to develop novel quinazoline conjugates with 1,3,4-thiadiazole and diaryl urea, and to investigate their anticancer activities [45–51]. Therefore, an efficient, multi-step, synthetic pathway was executed to produce the desired compounds, which were then assessed for their ability to inhibit cell proliferation in A549 cells (human lung cancer), MDA-MB-231 (human triple-negative breast cancer), and MCF7 (human hormone-dependent breast cancer) in comparison with Etoposide as the reference drug. Moreover, comprehensive computational

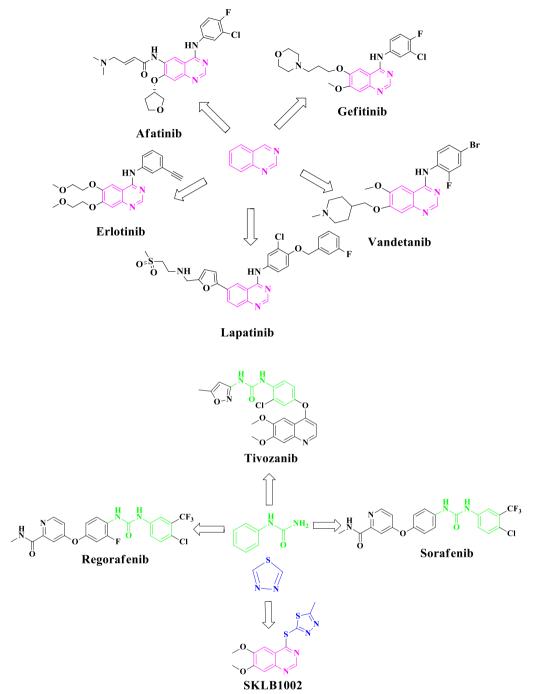


Fig. 1 Structures of FDA-approved EGFR and VEGFR-2 inhibitor drugs featuring quinazoline, thiadiazole, and urea moieties

investigations, including molecular dynamics, molecular docking studies, frontier molecular orbital analysis using DFT calculations, and in silico *ADME* studies, were performed to confirm the biological results (Fig. 2)

To reveal the significant role of the nitro group on this scaffold, it must be noted this functionality is a unique functional group in medicinal chemistry. It has a powerful electron-attracting potency that creates localized electron-deficient locations within molecules and interacts with biological nucleophiles present in living systems, such as amino acids, proteins, nucleic acids, and enzymes. The interaction can happen by nucleophilic addition and electron transfer involving reduction and oxidation, or also simply by molecular complexation, to

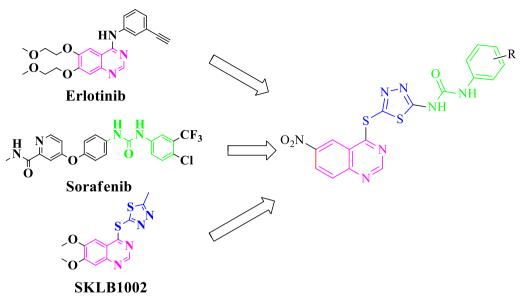


Fig. 2 Design strategy of the target 1-(5-((6-nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)-3-phenylurea derivatives as anticancer agents

induce undesired or desired biological changes [52, 53]. Therefore, countless medicinal chemistry campaigns have been initiated to investigate compounds containing nitro groups. Already, drugs bearing nitro groups have a long history of use as antibiotic, antineoplastic, and antiparasitic agents, as well as fungicides, tranquilizers, insecticides, and herbicides [53–59]. Moreover, some medicinal compounds including PR-104, Tarloxotinib (TH-4000), and Paclitaxel prodrug possess nitro group [60, 61].

Experimental section Materials

The solvents and reagents utilized in this study were obtained from reputable companies such as Merck or Sigma, and no additional purification was required. Melting points were measured using a Kofler hot-plate microscope apparatus, and the values reported were unadjusted. The NMR spectra were obtained using a Bruker FT-500 and Bruker FT-300 MHz spectrometer, with either DMSO-d6 or CDCl₃ serving as the solvent, while TMS was used as the internal standard. The chemical shifts and coupling constants are presented in δ (ppm) and J (Hz), respectively. Additionally, all reactions underwent monitoring via TLC on plastic sheets coated with silica gel 60 F254. The Perkin-Elmer Spectrum Version 10.03.06 was used to obtain IR spectra, specifically using potassium bromide disks. Mass spectra were obtained using the HP Agilent Technologies 5937 instrument with an ionization potential of 70 eV. Elemental analysis for carbon (C), hydrogen (H), and nitrogen (N) was conducted using the elemental analyzer GmbH VarioEL.

General chemistry

Synthesis of quinazoline-4(3H)-one (2)

A mixture of 2-aminobenzoic acid **1** (0.137 g, 1 mmol) and formamide (0.045 g, 14 mmol) was heated with vigorous stirring at 150 °C for 6h. After completion of the reaction according to the TLC analysis, the reaction mixture was cooled down to the ambient temperature. The precipitated product was filtered and washed with sufficient amount of water to remove the excess of formamide and to afford pure compound as a white solid with m.p. 212–214 °C [62].

Synthesis of 6-nitroquinazolin-4(3H)-one (3)

Under the ice bath conditions, quinazoline-4(3H)-one (0.146 g, 1 mmol) **2** was slowly added into an acid mixture (concentrated sulfuric acid and concentrated nitric acid with a ratio of 1:1) (2 mL) within almost 1h. When the addition was complete, the temperature of the system was slowly raised up to 95 °C, and at this temperature, the reaction was heated for 1 h. After completion of the reaction according to the TLC analysis, the reaction mixture was poured into ice water (25 mL), and it was stirred till yellow solid was precipitated. The precipitation was filtered to afford the pure 6-nitroquinazolin-4(3H)-one **3** with m.p. 279–283 °C [63]. ¹H NMR (500 MHz, DMSO-*d*6) δ 11.68 (s, 1H), 8.76 (d, *J*=2.7 Hz, 1H), 8.57–8.52 (m, 1H), 8.48 (s, 1H), 7.85 (d, *J*=9.0 Hz, 1H).

Synthesis of 4-Chloro-6-nitroquinazoline (4)

6-Nitroquinazolin-4(3H)-one **3** (0.209 g, 1 mmol) was added to the solution of thionyl chloride (SOCl₂, 2 mL) in DMF (5 mL). The mixture was heated under reflux

with stirring for almost 4 h. Then, ice MeOH (15 mL) was added slowly, and the obtained mixture was extracted with dichloromethane (DCM). The organic layer was dried using MgSO₄ and then concentrated to give 4-chloro-6-nitroquinazoline **4** as a yellow solid with m.p. 134–135 °C [64]. ¹H NMR (500 MHz, DMSO-*d*6): δ 8.81 (d, *J*=2.7 Hz, 1H), 8.56 (dd, *J*=9.0, 2.7 Hz, 1H), 8.37 (s, 1H), 7.89 (d, *J*=8.9 Hz, 1H).

Synthesis of 5-amino-1,3,4-thiadiazole-2-thiol (6)

In a reaction flask, thiosemicarbazide 5 (1 mmol) and Na_2CO_3 (1 mmol) were combined with absolute ethanol (5 mL) and heated to 60 °C for 30 min. A solution of carbon disulfide (CS₂, 3 mmol) in absolute EtOH (5 mL) was then slowly added drop by drop to the mixture. The resulting mixture was refluxed overnight. Once the reaction was complete, as confirmed by TLC analysis, the solvent was evaporated under reduced pressure. The remaining residue was diluted with water (25 mL), followed by the slow addition of concentrated HCl solution (5 mL). This resulted in the precipitation of compound 6 as a light yellow solid with a yield of 78% and a melting point range of 233–235 [65].

Synthesis

of 1-(5-mercapto-1,3,4-thiadiazol-2-yl)-3-phenylurea derivatives (8a-l)

5-Amino-1,3,4-thiadiazole-2-thiol (0.133 g, 1 mmol) was dissolved in acetonitrile (CH₃CN) (3 mL) and stirred at ambient temperature for 30 min. Next, the aryl isocyanate derivative 7a-l (1.2 mmol) was dissolved in CH₃CN (3 mL) and slowly added to the solution while stirring. The resulting mixture was stirred overnight at room temperature. The solid that formed was filtered and washed with diethyl ether (Et₂O), resulting in pure white solids known as adducts 8a-l. These adducts were then used in the subsequent step without requiring any additional purification steps [66].

Synthesis of 1-(5-((6-nitroquinazolin-4-yl)

thio)-1,3,4-thiadiazol-2-yl)-3-phenylurea Derivatives (8a-l)

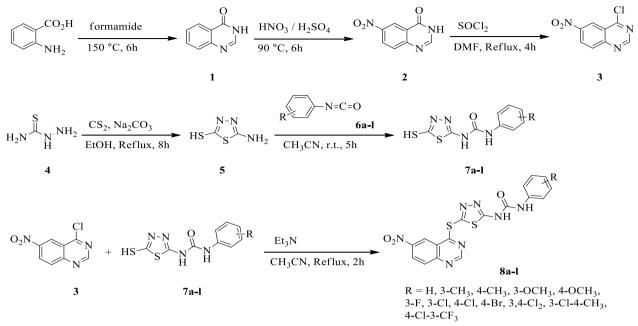
In order to synthesize the desired products 8a-l, a solution of 1-(5-mercapto-1,3,4-thiadiazol-2-yl)-3-phenylurea derivatives 7a-l (1 mmol) in acetonitrile (5 mL) was prepared. To this solution, a mixture of 4-chloro-6-nitro-quinazoline (0.209 g, 1 mmol) and Et_3N (0.101 g, 1 mmol) was added. The resulting mixture was then refluxed for 3 h. Once the reaction was complete as confirmed by TLC analysis, the solvent was evaporated under reduced pressure and the crude product obtained underwent recrystallization from DMF/Water. This process yielded pure desired products 8a-l in high yields.

l-(5-((6-Nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)-3-phenylurea (8a) Yellow solid; yield: 68%; mp: 225–227 °C; IR (KBr) (ν_{max} /cm⁻¹): 3247, 1697 (C=O), 1562, 1311. ¹HNMR (300 MHz, DMSO- d_6): δ 10.80 (s, 1H), 9.18 (s, 1H), 8.98 (s, 1H), 8.73 (d, *J*=9.2 Hz, 1H), 8.23 (d, *J*=9.1 Hz, 1H), 7.74 (d, *J*=8.1 Hz, 2H), 7.31 (dd, *J*=7.9 Hz, 2H), 6.99 (dd, *J*=7.3 Hz, 1H); ¹³C NMR (75 MHz, DMSO- d_6): δ 170.51, 169.14, 155.81, 154.96, 150.67, 145.60, 145.03, 139.75, 130.94, 128.76 (2CH), 128.22, 122.15, 121.41, 120.22, 118.43 (2CH); ESI-MS m/z: 426.0 [M+H]⁺; Anal. Calcd for C₁₇H₁₁N₇O₃S₂: C, 47.99; H, 2.61; N, 23.05. Found: C, 48.64; H, 2.55; N, 23.44.

l-(5-((6-Nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)-3-(m-tolyl)urea (8b) Yellow solid; yield: 78%; mp: 240–24 °C; IR (KBr) (ν_{max} /cm⁻¹): 3370, 1612 (C=O), 1554, 1319; ¹HNMR (300 MHz, DMSO- d_6): δ 10.27 (s, 1H), 9.19 (s, 1H), 9.00 (s, 1H), 8.74 (d, *J*=9.0 Hz, 1H), 8.25 (d, *J*=8.9 Hz, 1H), 7.56 (s, 1H), 7.44 (d, *J*=9.3 Hz, 1H), 7.18 (dd, *J*=7.5 Hz, 1H), 6.81 (d, *J*=7.2 Hz, 1H), 2.28 (s, 3H); ¹³C NMR (75 MHz, DMSO- d_6): δ 170.34, 168.64, 155.74, 154.54, 150.64, 145.55, 145.22, 139.49, 137.86, 130.92, 128.55, 128.16, 122.90, 121.36, 120.14, 118.94, 115.58, 21.29; ESI-MS m/z: 440.0 [M+H]⁺; Anal. Calcd for C₁₈H₁₃N₇O₃S₂: C, 49.19; H, 2.98; N, 22.31. Found: C, 49.43; H, 3.12; N, 23.92.

l-(5-((6-Nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)-3-(p-tolyl)urea (8c) Yellow solid; yield: 75%; mp: 257–259 °C; IR (KBr) (ν_{max} /cm⁻¹): 3390, 1673 (C=O), 1538, 1322. ¹HNMR (300 MHz, DMSO-d₆): δ 10.25 (s, 1H), 9.19 (s, 1H), 9.00 (s, 1H), 8.74 (d, *J*=9.2 Hz, 1H), 8.25 (d, *J*=9.2 Hz, 1H), 7.56 (d, *J*=8.5 Hz, 2H), 7.09 (d, *J*=8.1 Hz, 2H), 2.24 (s, 3H). ¹³C NMR (75 MHz, DMSO-d₆): δ 170.45, 169.17, 155.76, 154.86, 150.64, 145.53, 144.89, 137.21, 130.91, 129.12, 128.13 (2CH), 121.36, 120.15, 118.38, 20.40; ESI-MS m/z: 440.0 [M+H]⁺; Anal. Calcd for C₁₈H₁₃N₇O₃S₂: C, 49.19; H, 2.98; N, 22.3. Found: C, 48.13; H, 3.65; N, 23.42.

1-(3-Methoxyphenyl)-3-(5-((6-nitroquinazolin-4-yl) thio)-1,3,4-thiadiazol-2-yl)urea (8d) Yellow solid; yield: 62%; mp: 228–230 °C; IR (KBr) (ν_{max} /cm⁻¹): 3386, 1716 (C=O), 1550, 1322. ¹HNMR (500 MHz, DMSO- d_6): δ 11.27 (s, 1H), 9.19 (s, 1H), 9.09 (s, 1H), 8.96 (s, 1H), 8.73 (d, J=9.1 Hz, 1H), 8.24 (d, J=9.1 Hz, 1H), 8.24 (dd, J=9.1 Hz, 1H), 7.14 (s, 1H), 6.99 (d, J=8.2 Hz, 1H), 6.64 (d, J=8.4 Hz, 1H), 3.73 (s, 3H), ¹³C NMR (125 MHz, DMSO- d_6): δ 170.41, 168.49, 155.65, 152.56, 143.29, 142.59, 139.19, 133.09, 130.98, 128.09,127.47, 119.37, 117.23, 112.01, 55.21; ESI-MS m/z: 455.0 [M+H]⁺; Anal. Calcd for C₁₈H₁₃N₇O₄S₂: C, 47.47; H, 2.88; N, 21.53. Found: C, 48.23; H, 2.31; N, 22.48.



Scheme 1 Synthetic route toward 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea 8a-I

1-(4-Methoxyphenyl)-3-(5-((6-nitroquinazolin-4-yl) thio)-1,3,4-thiadiazol-2-yl)urea (8e) Yellow solid; yield: 69%; mp: 242–244 °C; IR (KBr) (ν_{max} /cm⁻¹): 3390, 1712 (C=O), 1550, 1326. ¹HNMR (300 MHz, DMSO-*d*₆): δ 11.31 (s, 1H), 9.20 (s, 1H), 8.978 (s, 1H), 8.74 (d, *J*=6.8 Hz, 1H), 8.25 (d, *J*=9.2 Hz, 1H), 7.39 (d, *J*=9.0 Hz, 2H), 6.91 (d, *J*=9.0 Hz, 2H), 3.73 (s, 3H), ¹³C NMR (75 MHz, DMSO-*d*₆): δ 169.82, 164.52, 159.18, 155.73, 155.46, 150.70, 147.57, 130.99, 128.41, 121.44, 120.97 (2CH), 120.27, 114.09 (2CH), 55.20; ESI-MS m/z: 455.0 [M + H]⁺; Anal. Calcd for C₁₈H₁₃N₇O₄S₂: C, 47.47; H, 2.88; N, 21.53. Found: C, 46.52; H, 2.07; N, 22.98.

1-(3-*Fluorophenyl*)-3-(5-((6-*nitroquinazolin*-4-*yl*) *thio*)-1,3,4-*thiadiazol*-2-*yl*)*urea* (8*f*) Yellow solid; yield: 72%; mp: 230–232 °C; IR (KBr) (ν_{max} /cm⁻¹): 3386, 1720 (C=O), 1592, 1322. ¹HNMR (500 MHz, DMSO-*d₆*): δ 11.45 (s, 1H), 9.47 (s, 1H), 9.20 (s, 1H), 8.95 (s, 1H), 8.73 (d, *J*=9.1 Hz, 1H), 8.25 (d, *J*=9.1 Hz, 1H), 7.46 (d, *J*=11.5 Hz, 1H), 7.35 (d, *J*=6.8 Hz, 1H), 7.22 (d, *J*=8.7 Hz, 1H), 6.88 (dd, *J*=8.5 Hz, 1H). ¹³C NMR (125 MHz, DMSO-*d₆*): δ 170.15, 168.02, 159.72, 156.67, 152.40, 147.57, 145.64, 143.11, 133.12, 131.07, 130.91, 128.30, 124.02, 122.99, 121.40, 119.37, 118.62; ESI–MS m/z: 443.0 [M+H]⁺; Anal. Calcd for C₁₇H₁₀FN₇O₃S₂: C, 46.05; H, 2.27; N, 22.11. Found: C, 47.35; H, 2.84; N, 23.59.

1-(3-Chlorophenyl)-3-(5-((6-nitroquinazolin-4-yl) thio)-1,3,4-thiadiazol-2-yl)urea (8g) Yellow solid; yield: 65%; mp: 258–260 °C; IR (KBr) (ν_{max} /cm⁻¹): 3370, 1720

(C=O), 1535, 1322. ¹HNMR (300 MHz, DMSO- d_6): δ 10.97 (s, 1H), 9.18 (s, 1H), 8.96 (s, 1H), 8.71 (d, J=9.2 Hz, 1H), 8.23 (d, J=9.2 Hz, 1H), 7.88 (s, 1H), 7.53 (d, J=8.2 Hz, 1H), 7.32 (dd, J=8.1 Hz, 1H), 7.02 (d, J=5.7 Hz, 1H), ¹³C NMR (75 MHz, DMSO- d_6): δ 170.15, 167.30, 155.77, 154.00, 150.66, 145.65, 140.87, 133.25, 130.96, 130.45, 128.30, 121.99, 120.22, 117.68, 116.78; ESI-MS m/z: 460.0 [M+H]⁺; Anal. Calcd for C₁₇H₁₀ClN₇O₃S₂: C, 44.40; H, 2.19; N, 21.32. Found: C, 45.35; H, 2.62; N, 22.34.

1-(4-*Chlorophenyl*)-3-(5-((6-*nitroquinazolin*-4-*yl*) *thio*)-1,3,4-*thiadiazol*-2-*yl*)*urea* (8*h*) Yellow solid; yield: 67%; mp: 246–248 °C; IR (KBr) (ν_{max} /cm⁻¹): 3390, 1720 (C=O), 1538, 1322. ¹HNMR (300 MHz, DMSO-*d*₆): δ 11.44 (s, 1H), 9.32 (s, 1H), 9.19 (s, 1H), 8.96 (s, 1H), 8.73 (d, *J*=9.2 Hz, 1H), 8.24 (d, *J*=9.2 Hz, 1H), 7.52 (d, *J*=8.9 Hz, 2H), 7.36 (d, *J*=8.9 Hz, 2H), ¹³C NMR (75 MHz, DMSO*d*₆): δ 169.61, 160.56, 155.68, 150.69, 145.74, 137.16, 130.99, 128.79 (2CH), 128.41, 126.91, 121.42, 120.61, 120.25 (2CH); ESI-MS m/z: 460.0 [M+H]⁺; Anal. Calcd for C₁₇H₁₀ClN₇O₃S₂: C, 44.40; H, 2.19; N, 21.32. Found: C, 43.24; H, 2.87; N, 22.43.

1-(4-Bromophenyl)-3-(5-((6-nitroquinazolin-4-yl) thio)-1,3,4-thiadiazol-2-yl)urea(8i) Yellow solid; yield: 60%; mp: 236–238 °C; IR (KBr) (ν_{max} /cm⁻¹): 3386, 1716 (C=O), 1531, 1319. ¹HNMR (300 MHz, DMSO-*d*₆): δ 10.64 (s, 1H), 9.19 (s, 1H), 9.00 (s, 1H), 8.74 (d, *J*=6.8 Hz, 1H), 8.25 (d, *J*=9.2 Hz, 1H), 7.65 (d, *J*=8.9 Hz, 2H), 7.46 (d, *J*=8.9 Hz, 2H), ¹³C NMR (75 MHz, DMSO-*d*₆): δ 170.45, 159.72, 156.67, 155.84, 150.68, 145.70, 138.94, 131.54 (2CH), 130.95, 128.32, 124.27, 121.45, 120.27 (2CH); ESI-MS m/z: 505.9 [M+H]⁺; Anal. Calcd for $\rm C_{17}H_{10}BrN_7O_3S_2$: C, 40.48; H, 2.00; N, 19.44. Found: C, 41.18; H, 2.30; N, 18.14.

1-(3,4-Dichlorophenyl)-3-(5-((6-nitroquinazolin-4-yl) thio)-1,3,4-thiadiazol-2-yl)urea (8j) Yellow solid; yield: 65%; mp: 252–254 °C; IR (KBr) (ν_{max} /cm⁻¹): 3382, 1727 (C=O), 1585, 1322. ¹HNMR (300 MHz, DMSO- d_6): δ 11.54 (s, 1H), 9.44 (s, 1H), 9.19 (s, 1H), 8.91 (d, J=2.5 Hz, 1H), 8.71 (d, J=9.2 Hz, 1H), 8.22 (d, J=9.2 Hz, 1H), 7.83 (s, 1H), 7.52 (d, J=8.8 Hz, 1H), 7.39 (d, J=8.9 Hz, 1H), ¹³C NMR (75 MHz, DMSO- d_6): δ 170.41, 168.49, 156.69,

155.83, 152.56, 143.29, 139.19, 133.09, 130.98, 128.09, 127.47, 119.37, 118.62, 118.00, 117.23; ESI-MS m/z: 493.9 $[M+H]^+$; Anal. Calcd for $C_{17}H_9C_{12}N_7O_3S_2$: C, 41.30; H, 1.84; N, 19.83. Found: C, 40.35; H, 2.01; N, 20.73.

1-(3-Chloro-4-methylphenyl)-3-(5-((6-nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)urea (8k) Pale-yellow solid; yield: 69%; mp: 234–236 °C; IR (KBr) (ν_{max} / cm⁻¹): 3178, 1616 (C=O), 1542, 1311. ¹HNMR (300 MHz, DMSO- d_6): δ 10.49 (s, 1H), 9.19 (s, 1H), 8.98 (s, 1H), 8.74 (d, *J*=6.8 Hz, 1H), 8.25 (d, *J*=9.2 Hz, 1H), 7.84 (s, 1H), 7.43 (d, *J*=6.0 Hz, 1H), 7.25 (d, *J*=8.8 Hz, 1H), 2.25 (s, 3H), ¹³C NMR (75 MHz, DMSO- d_6): δ 170.02, 168.04,

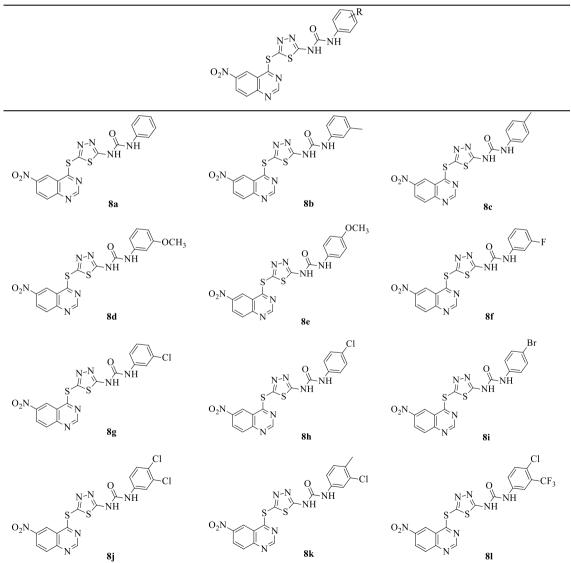


Table 1 Substrate scope for the synthesis of 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea 8a-I

155.64, 154.33, 150.60, 145.64, 145.48, 138.72, 133.12, 131.07, 130.91, 128.49, 128.09, 121.27, 120.04, 118.07, 116.94, 18.83; ESI-MS m/z: 474.0 $[M+H]^+$; Anal. Calcd for C₁₈H₁₂ClN₇O₃S₂: C, 45.62; H, 2.55; N, 20.69. Found: C, 46.70; H, 3.01; N, 21.99.

1-(4-chloro-3-(trifluoromethyl)phenyl)-3-(5-((6-nitroquinazolin-4-yl)thio)-1,3,4-thiadiazol-2-yl)urea (8l) Yellow solid; yield: 58%; mp: 240–245 °C; IR (KBr) (ν_{max} / cm⁻¹): 3363, 1704 (C=O), 1527, 1319. ¹HNMR (300 MHz, DMSO- d_6): δ 11.65 (s, 1H), 9.61 (s, 1H), 9.20 (s, 1H), 8.94 (s, 1H), 8.72 (d, J=9.3 Hz, 1H), 8.24 (d, J=9.1 Hz, 1H), 8.07 (s, 1H), 7.73 (d, J=8.8 Hz, 1H), 7.64 (d, J=8.8 Hz, 1H), ¹³C NMR (75 MHz, DMSO- d_6): δ 170.15, 163.09, 159.51, 155.77, 154.00, 150.66, 143.11, 133.25, 130.96, 128.49, 128.09, 127.62, 126.91, 121.40, 120.23, 117.67; ESI-MS m/z: 527.9 [M+H]⁺; Anal. Calcd for C₁₈H₉ClF₃N₇O₃S₂: C, 40.95; H, 1.72; N, 18.57. Found: C, 41.25; H, 1.89; N, 19.53.

In vitro cytotoxicity assay

The MTT assay was used to test the cytotoxicity of novel 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8** against the A-549 (human lung cancer), MCF-7 (human hormone dependent breast cancer), MDA-MB-231 (human triple negative breast cancer), and HDF (Human Dermal Fibroblasts) cell lines. All of the cell lines were obtained from the Pastor Institute's National Cell Bank in Tehran, Iran, and were cultured in either RPMI-1640 or DMEM supplemented with 10% fetal bovine serum (Gibco, Milano, Italy). The cancerous cells were cultured at 37 °C in a humidified 5% CO₂ incubator after being seeded onto 96-well micro-plates at a

Table 2 In vitro anti-proliferative effects (IC_{50} , μ M) of compounds **8a-I** against A549, MDA-MB231 and MCF7 cell lines

Compound	R	A549	MDA-MB231	MCF7
8a	Н	>50	>50	>50
8b	3-CH ₃	>50	>50	29 ± 0.005
8c	4-CH ₃	>50	>50	25 ± 0.001
8d	3-OCH ₃	> 50	>50	>50
8e	4-OCH3	>50	>50	>50
8f	3-F	>50	>50	>50
8g	3-Cl	>50	>50	39.1 ± 0.001
8h	4-Cl	>50	>50	>50
8i	4-Br	>50	>50	>50
8j	3,4DiCl	>50	>50	>50
8k	3-CI-4-CH ₃	>50	46 ± 0.003	>50
81	4-CI-3-CF ₃	>50	> 50	>50
Etoposide	-	25.8 ± 0.0008	27.9 ± 0.002	29.8 ± 0.0007

Values were the means of three replicates ± standard deviation (SD)

density of 1×10^4 cells/mL. After overnight incubation at 37 °C, compounds were added to the cells at varying final concentrations (1 µg/mL, 5 µg/mL, 20 µg/mL, and 40 µg/mL) after being dissolved in dimethyl sulfoxide (DMSO). Fresh phenol red free RPMI-1640 containing 0.5 mg/mL of MTT was added after incubation. Using a multiwell plate reader (Gen5, Epoch, BioTek) optical density was measured at 492 nm. Both Etoposide and 1% DMSO were employed as positive and negative controls, respectively. Three separate dose–response curves were used to get the IC₅₀ values (the dosage needed to suppress cell growth by 50%).

Results and discussion

Chemistry

The synthetic approach toward the desired novel 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea 8a-l is illustrated in Scheme 1. Heating a mixture of 2-aminobenzoic acid and formamidine under the reflux conditions led to obtain quinazoline-4(3H)-one 1 which subsequently underwent the electrophilic substitution with the mixture of nitric acid and sulfuric acid at the ambient temperature to give 6-nitroquinazolin-4(3H)-one 2 The chlorination of carbonyl group in this compound with thionyl chloride in DMF was carried out the reflux conditions to obtain 4-chloro-6-nitroquinazolin-4(3H)-one 3. On the other hand, the production of 1-(5-mercapto-1,3,4-thiadiazol-2-yl)-3-arylurea 7a-l was achieved through a condensation reaction involving thiosemicarbazide 4 and carbon disulfide. This reaction took place in the presence of sodium carbonate under reflux conditions in absolute EtOH (compound 5). Subsequently, various substituted phenyl isocyanates 6a-l were added to the mixture in CH₃CN to yield the corresponding derivatives 7a-l. Finally, chlorine moiety compound 3 underwent nucleophilic aromatic substitution with synthesized 1-(5-mercapto-1,3,4-thiadiazol-2-yl)-3-arylurea **7a-l** in the presence of Et_3N in CH_3CN under the reflux conditions to afford our desirable 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea derivatives 8a-l. The spectra of the compounds are provided in the supplementary information file.

The generality of the substrate scope was investigated using various substituted phenyl isocyanates **6a-1** bearing electron-donating groups like methyl and methoxy, as well as electron-withdrawing groups like fluorine, chlorine, bromine, and trifluoromethyl. The structures of isolated 6-nitro quinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea derivatives **8a-1** are summarized in Table 1. They are deduced based on their IR, ¹H, and ¹³C NMR, as well as MASS spectroscopy and elemental analysis. Partial assignments of these resonances are given in the Experimental Part.

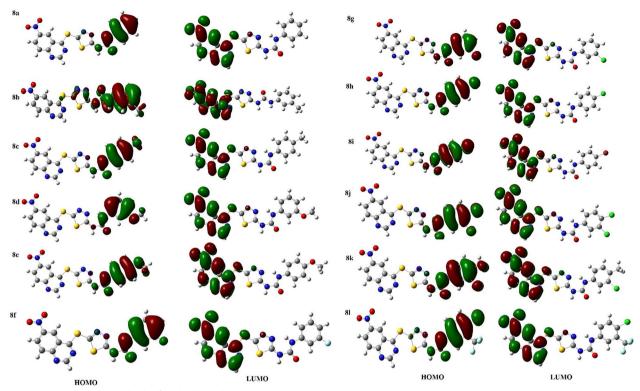


Fig. 3 Frontier molecular orbitals of studied ligands

In Vitro antiproliferative activity

The targeted 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8a-1** were evaluated for their in vitro antitumor activities against three human cancerous cell lines, including A549, MDA-MB231, and MCF7 cell lines by using the MTT colorimetric assay to investigate the role of substituents on the phenyl

Table 3 The quantum reactivity indexes including chemical potential (μ), hardness (η), and electrophilicity index (ω) in eV

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Compound	Eg	μ	η	ω
8a	2.68	- 4.59	1.34	7.86
8b	2.96	-4.57	1.48	7.05
8c	2.50	-4.49	1.25	8.06
8d	2.49	-4.47	1.24	8.05
8e	2.16	-4.31	1.08	8.60
8f	2.81	-4.69	1.40	7.85
8g	2.88	-4.74	1.44	7.80
8h	2.71	-4.65	1.35	8.00
8i	2.67	- 4.63	1.33	8.06
8j	2.86	-4.77	1.43	7.95
8k	2.70	-4.63	1.35	7.93
81	2.97	-4.83	1.48	7.88

moiety originated from arylisocyanates. The cytotoxicity is expressed as the concentration which inhibits 50% of cell viability (IC₅₀). In present study, Etoposide was used as a positive control having IC_{50} values of 25.8 ± 0.0008 µM against A549, 27.9±0.002 µM against MDA-MB231, and $29.8 \pm 0.0007 \mu$ M against MCF7. These compounds were shown better cytotoxicity comparable to Etoposide on MCF7 cells, as some derivatives named 8b, 8c, and 8g exhibited IC₅₀ values of $29.0 \pm 0.005 \ \mu\text{M}$, $25.0 \pm 0.001 \ \mu\text{M}$, $39.1 \pm 0.001 \mu$ M, respectively. Other results were noticeably weaker than standard drug. It seems that the presence of methyl as a mild electron-donating group particularly at C-4 position (compound 8c) improved the cytotoxic potency (Table 2). Furthermore, we assessed the cytotoxic impact of 8b and 8c as most potent cytotoxic compounds on normal HDF. Remarkably, no toxic properties were observed in HDF cells even at a concentration as high as 50 µM.

Computational details

To investigate the electronic features of synthesized 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8a-l**, quantum calculations were performed at the B3LYP/6-31 g (d, p) level of theory. Frequency calculations were performed to confirm the nature of minima

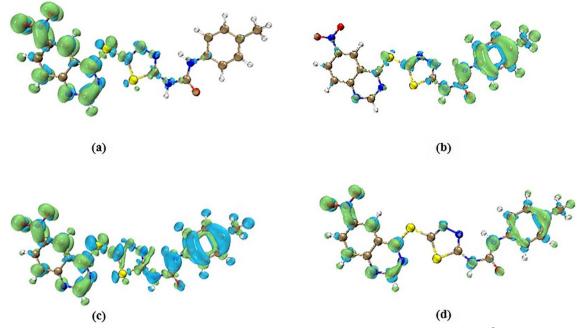


Fig. 4 Fukui functions for **a** nucleophilic attack $f^{+}(r)$ **b** electrophilic attack $f^{-}(r)$ **c** dual descriptor (Δf), and **d** radical attack f^{0} for **8c** ligand at B3LYP/6–31 g (d, p) level. The green and blue regions show positive and negative values of these functions

structures. All the DFT quantum calculations were conducted using GAMESS [67]. The frontier molecular orbital analysis was performed to study the chemical reactivity of ligands **8**. The electrostatic surface potential (ESP) analysis was performed by Multiwfn 3.3.8 code [68]. To reveal the relationship between the structure and activity of compounds, the quantitative ESP analysis was done. To evaluate the plausible interactions between most potent compounds, **8c**, as a potential VEGFR-2 inhibitor, and VEGFR-2 target, the molecular docking calculation were performed using Autodock 4.2.1. The crystallographic structure of 3WZE was selected as a VEGFR-2 target.

Molecular dynamic simulation was performed with GROMACS 2020. Amber force field was used for building the topological files for protein and ligand. To consider the solvation, the TIP3 water model was employed. The box system including protein, ligand, water, and ions was minimized foe 2175 steps. The equilibration process was done in the NVT and NPT ensembles at the pressure of 1 atm and temperature of 300K for 200 ps. The MD simulation was performed for 100 ns. The RMSD, RMSF, and hydrogen bond analysis were performed on ythe trajectory file.

Frontier molecular orbital analysis

The frontier molecular orbitals could be used to elucidate the chemical reactivity of a molecule. The energy gap can be considered as an indicator of the stability and chemical reactivity. Molecules with smaller band gap categorized as more polarizable molecules with higher chemical reactivity. In all 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8a-1**, HOMO is located on the phenyl urea moieties whereas the LUMO is localized on the nitro quinazoline moiety. Therefore, according to different distribution of HOMO and LUMO, one can expect that there is a favorable intramolecular charge transfer for these series of ligands (Fig. 3).

The HOMO and LUMO orbital energies can be used to assess the quantum chemical reactivity indexes including electrophilicity and nucleophilicity indexes, chemical potential, softness and hardness parameters as:

$$\mu = \frac{E_{HOMO} + E_{LUMO}}{2} \quad \eta = \frac{E_{LUMO} - E_{HOMO}}{2} \quad \omega = \frac{\mu^2}{2\eta}$$

The quantum chemical indexes including chemical potential, chemical hardness, and global electrophilicity indexes are calculated for all obtained compounds **8a-1**, and the results are summarized in Table 3. These parameters are calculated based on the ground state optimized geometries. It can be seen that the chemical reactivity of all ligands is negative which can be assign to the stability of these compounds **8**.

Atom	q(N)	q(N+1)	q(N-1)	f⁻	f+	f ⁰	DD
N39	0.2497	0.1986	0.2524	0.0028	0.0510	0.0269	0.0483
O40	-0.1935	-0.2946	-0.1769	0.0166	0.1011	0.0588	0.0844
O41	-0.1941	-0.2842	-0.1864	0.0077	0.0901	0.0489	0.0824
N42	-0.1487	-0.1849	-0.1450	0.0036	0.0362	0.0199	0.0326
N43	-0.1598	-0.2139	-0.1365	0.0233	0.0541	0.0387	0.0308
C23	0.0130	0.0140	0.0360	0.0230	-0.0009	0.0110	-0.0239
C21	0.0820	0.0612	0.0868	0.0048	0.0209	0.0128	0.0160
N4	-0.0861	-0.0946	-0.0528	0.0333	0.0085	0.0209	-0.0248
C1	0.2080	0.2045	0.2268	0.0188	0.0035	0.0111	-0.0153
O20	-0.2963	-0.3128	-0.2550	0.0413	0.0165	0.0289	-0.0249
N2	-0.0819	-0.0842	-0.0224	0.0596	0.0023	0.0309	-0.0573
C11	-0.0462	-0.0502	-0.0046	-0.0415	0.0040	0.0228	-0.0375
C6	0.0447	0.0487	0.1009	0.0562	-0.0040	0.0261	-0.0602
C7	-0.0544	-0.0572	-0.0067	0.0477	0.0028	0.0252	-0.0449
C8	-0.0568	-0.0563	-0.0098	0.0470	-0.0005	0.0232	-0.0475
С9	-0.0437	-0.0495	-0.0019	0.0419	0.0058	0.0238	-0.0361
C13	-0.0024	-0.0101	0.0721	0.0745	0.0077	0.0411	-0.0668
C16	-0.0817	-0.0845	-0.0608	0.0209	0.0027	0.0118	-0.0181
C28	-0.0131	-0.0282	-0.0091	0.0040	0.0151	0.0096	0.0111
C29	-0.0202	-0.0923	-0.0202	0.0001	0.0721	0.0361	0.0720
C30	0.0654	0.0476	0.0718	0.0065	0.0178	0.0121	0.0113
C31	0.0960	0.0314	0.1019	0.0058	0.0647	0.0352	0.0588
C32	0.0334	-0.0084	0.0451	0.0117	0.0417	0.0267	0.0300
C34	-0.0211	-0.0748	-0.0098	0.0113	0.0537	0.0325	0.0424
C36	-0.0159	-0.0420	-0.0033	0.0126	0.0261	0.0193	0.0134
S22	0.0762	0.0596	0.1224	0.0462	0.0166	0.0314	-0.0296
S26	0.0933	0.0477	0.1498	0.0565	0.0456	0.0511	-0.0109
N24	-0.1283	-0.1555	-0.1053	0.0230	0.0272	0.0251	0.0042
N25	-0.1300	-0.1474	-0.1137	0.0163	0.0174	0.0169	0.0011

Table 4 The values of Fukui functions $f^{-}(r)$, $f^{+}(r)$, and $f^{0}(r)$ and dual descriptors ($\Delta f(r)$) for active sites on the potent **8c** ligand

Fukuie reactivity descriptor

A key parameter to describe the reactive area in a molecule is Fukui function, $f(\mathbf{r})$, which is based on the conceptual density functional theory. It can be described as $(r) = \left[\frac{\partial \rho(r)}{\partial N}\right]_{v}$ where N and v can be defined as number of electrons in the system and the external potential due to nuclear charges, respectively. Finding the reactive centers is important to elucidate the interaction between the compound and biological target. For this purpose, the Fukui parameters are calculated for the most potent 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diarylurea, **8c**. In general, the centers with high activity show the more values of Fukui functions. This function can be evaluated for three conditions including nucleophilic attack $f^+(r)$, electrophilic attack $f^-(r)$, and radical attack $f^0(r)$ according to:

$$f^{+}(r) = \rho_{N+1}(r) - \rho_{N}(r)$$
$$f^{-}(r) = \rho_{N}(r) - \rho_{N-1}(r)$$
$$f^{0}(r) = \frac{\rho_{N+1}(r) - \rho_{N-1}(r)}{2}$$

Dual descriptor is another parameter for the evaluation of the nucleophilic and electrophilic centers at the same time. The regions with $\Delta f > 0$ and $\Delta f < 0$ can be described as suitable centers for a nucleophilic attack and electrophilic attack, respectively. Δf can be evaluated as

$$\Delta f(r) = f^+(r) - f^-(r) = \rho_{N+1}(r) - 2\rho_N(r) + \rho_{N-1}(r)$$

The graphical view of these indexes was indicated in Fig. 4.

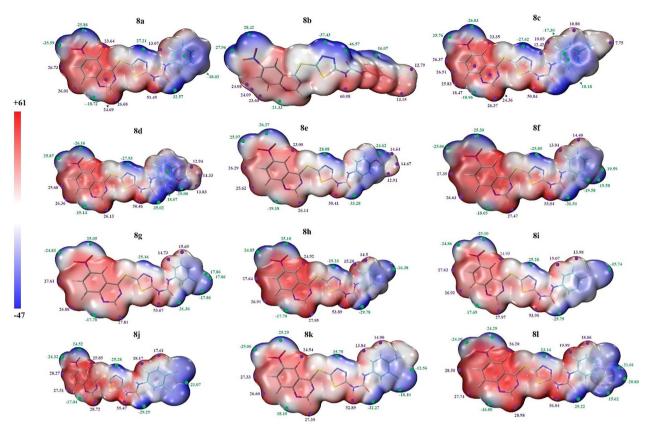


Fig. 5 The quantitative ESP analysis on the molecular vdW surface of studied ligands. The unit is in kcal/mol. The green and magenta spheres represent surface local minima and maxima of ESP values, respectively

Ligand	MW	TPSA	GI	BBB	MlogP	Lipinski	N rotatable bond	N H-bond acceptor	N H-bond donor	C Log P _{o/w}
8a	425.44	192.05	Low	No	1.48	Yes	7	7	2	2.24
8b	439.47	192.05	Low	No	1.71	Yes	7	7	2	2.64
8c	439.47	192.05	Low	No	1.71	Yes	7	7	2	2.64
8d	455.47	201.28	Low	No	1.22	Yes	8	8	2	2.23
8e	455.47	201.28	Low	No	1.22	Yes	8	8	2	2.29
8f	443.43	192.05	Low	No	1.86	Yes	7	8	2	2.60
8g	459.89	192.05	Low	No	1.98	Yes	7	7	2	2.84
8h	459.89	192.05	Low	No	1.98	Yes	7	7	2	2.79
8i	504.34	192.05	Low	No	2.10	Yes	7	7	2	2.88
8j	494.33	192.05	Low	No	2.48	Yes	7	7	2	3.29
8k	473.92	192.05	Low	No	2.21	Yes	7	7	2	3.13
81	527.89	192.05	Low	No	2.82	Yes	8	10	2	3.85

Table 5 Drug likeness properties of ligands 8a-I

Furthermore, to quantify the amount of reactivity in each site of the molecules, the condensed Fukui functions are calculated and the results are reported in Table 4.

It can be seen that regarding the nucleophilic attack, the potential active sites are distributed mainly on the nitro quinazoline part. As to electrophilic attack, the

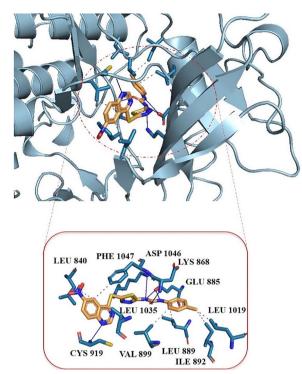


Fig. 6. 3dimensional (3D) binding interactions of compound **8c**. The hydrophobic interactions and hydrogen bonds are indicated as black dotted and blue lines, respectively

plausible active sites are localized on the phenyl urea moiety. It can be seen that the thiadiazole ring is common in both nucleophilic and electrophilic attacks. To clarify and asses the nature of this position, the dual descriptor parameter could be useful. It can be seen that the DD parameter at thiadiazole ring is mainly positive. Therefore, it has potential for nucleophilic attack. For the other fragments in the compound **8c**, the results of dual parameter are in the line with the results of $f^+(r)$ and $f^-(r)$ Fukui functions. For radical attack, the plausible active sites are located mainly on the nitro and phenyl ring.

Electrostatic potential surface

The electrostatic potential iso-surfaces can be considered as a valuable tool to conduct a close link between the distribution of electrons in a molecule and the biological activity. It can be used to predict and interpret the weak non-covalent interactions between the small ligands and the macromolecule target. The ESP analysis elucidate for us the regions with positive and negative electrostatic potential with low and high amount of electron densities, respectively. The map of electron density for two compounds is plotted in Fig. 5.

The regions with low and high electron densities are shown with red and blue color, respectively. The quantitative electrostatic potential analysis was performed to shed light more accurately on the regions with high and low electron densities based on the optimized structures of molecules on the ground state. The positive regions on the hydrogen atoms are due to the lower electronegativities of this element. The electronegative elements with high electronegativities show negative regions. The white color shows region with moderate electronegativities. It is expected that the regions with negative potential conduct strong attraction with positive regions on the target and the positive regions on the ligand conduct attraction interactions with negative regions on the target. It can be seen that different position of methyl group on compounds **8b** and **8c** led to different distribution of electron density that has an important role in the bioactivity of these compounds.

To have a favorable interaction between ligand and bio target, a complementary electron density distribution must be existed. Different substituents on the phenyl ring led to different distribution of electron density and consequently the different bioactivity. In all 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8a-1**, hydrogen atoms with lower electronegativity show positive potential region (red color). The atoms with more electronegativity including O, F, and N show negative potential regions with blue color.

In silico ADME evaluation

The physiochemical and pharmacokinetic properties for all 6-nitroquinazoline conjugated with 1,3,4-thiadiazole and diaryl-urea **8a-1** were computed using Swiss *ADME* online (http://www.swissadme.ch/index.php) toolkit [69]. Through this in silico study under the Lipniski's rule, several determined drug-likeness parameters were compared with the known drugs [70–74]. These evaluated parameters are summarized in Table 5. In terms of drug likeness properties, it can be seen that there is not any violation to Lipinski rules for these compounds. Consensus Log $P_{o/w}$ is predicted in the range of 2.23–3.85. The calculations predict no Blood Brain Barrier (BBB).

Molecular docking studies

To investigate the interactions between **8c** and VEGFR2 target, the molecular docking calculation was performed. There are two types of interactions between SM1 ligand and target VEGFR2. The ligand forms hydrophobic interactions with LEU 840, LEU 889, ILE 892, VAL 899, LEU 1019, LEU 1035, PHE 1047 and hydrogen bonds with LYS 868, GLU 885, CYS 919, and ASP 1046 (Fig. 6).

To assess the stability of the complex including protein and ligand, an MD simulation was performed for 100 ns. It can be seen that, in Fig. 7, the number of

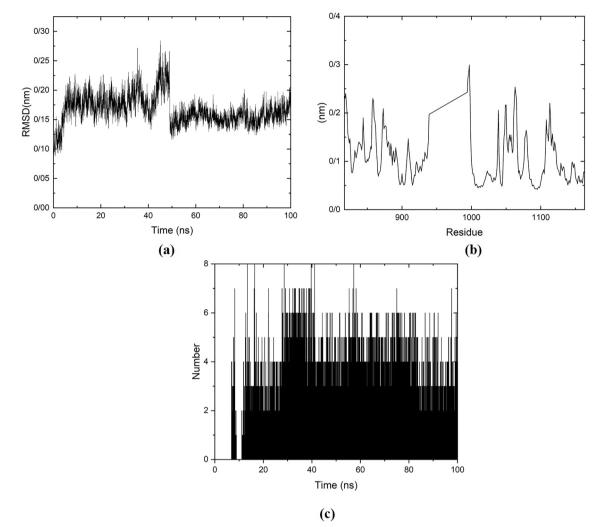


Fig. 7 RMSD plot for the backbone atoms (a), RMSF plot of the complex (b), and the number of hydrogen bonds between the protein target VEGFR2 and compound 8c

hydrogen bonds between protein and ligand vary in the range of 1–4 in most of the time of the simulation. The RMSD plot which is an indicator of stability of the protein, remains stable 0.15 nm after 50 ns. RMSF, which is used to evaluate the mobility of the residues, show low fluctuations and the structure remains stable during simulation time, the peaks are related to the residues out of the active site of protein. Thus, according to the results of RMSD, RMSF, and H-bond interactions, we can see that the structure remains stable during the 100 ns.

Conclusion

A novel series of 1-(5-((6-nitroquinazoline-4-yl) thio)-1,3,4-thiadiazol-2-yl)-3-phenylurea derivatives**8a**–**1**were designed, synthesized, investigated for their potential as an anticancer agent against A549, MDA-MB231,

and MCF7. Among them, compound 8b and 8c bearing methyl group at C-3 and C-4 positions of phenyl ring showed good in vitro anti-proliferative potency with IC_{50} value of 29.0 $\pm 0.005~\mu M$ and 25.0 $\pm 0.001~\mu M$ against MCF7, respectively, which was comparable with Etoposide (IC₅₀=29.8 \pm 0.0007 μ M). The ADME properties were calculated to assess the drug-like features. The quantitative electrostatic potential analysis was done to have a better understanding of the relationship between the electronical structure and activity. The stability of the protein-ligand complex was verified through 100 ns MD simulation. Finally, considering better cytotoxicity of compound 8c, the molecular docking study for this ligand was performed, showing several hydrogen bonds, hydrophobic interactions, and van der Waals forces between it and residues in the active sites of VEGFR-2.

Although it has been anticipated that these compounds possess great anti-proliferative potencies, this goal was not achieved which might be related to the adverse role of the nitro group. Therefore, changing this functionality in an attempt to find further potent anticancer agents could be an interesting goal for future studies.

Supplementary Information

The online version contains supplementary material available at https://doi. org/10.1186/s13065-024-01119-0.

Additional file 1. Additional file 1 of Novel quinazolines bearing 1,3,4-thiadiazole-aryl urea derivative as anticancer agents: Design, Synthesis, Molecular docking, DFT and Bioactivity evaluations

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Author contributions

AF and MSZ designed the study and conducted the experiments. SM synthesized the targeted compounds, wrote the manuscript, and analyzed the characterization data. MS performed the inhibition investigations. HRB prepared the Additional file 1.

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Availability of data and materials

The authors confirm that the data supporting the finding of this study are available within the manuscript.

Declarations

Ethics approval and consent to participate Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare no competing interests.

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References:

- Elzahabi HS, et al. Anticancer evaluation and molecular modeling of multi-targeted kinase inhibitors based pyrido [2, 3-d] pyrimidine scaffold. J Enzyme Inhib Med Chem. 2018;33(1):546–57. https://doi.org/10.1080/ 14756366.2018.1437729.
- Guruswamy DKM, et al. Novel 3-(3, 5-difluoro-4-hydroxyphenyl)-1-(naphthalen-2-yl) prop-2-en-1-one as a potent inhibitor of MAP-kinase in HeLa cell lines and anti-angiogenic activity is ediated by HIF-1α in EAC animal model. Oncotarget. 2020;11(50):4661. https://doi.org/10.18632/ oncotarget.27836.
- Arnold M, et al. Current and future burden of breast cancer: global statistics for 2020 and 2040. The Breast. 2022;66:15–23. https://doi.org/10. 1016/j.breast.2022.08.010.

- Zhao Y, et al. Discovery of arylamide-5-anilinoquinazoline-8-nitro derivatives as VEGFR-2 kinase inhibitors: synthesis, in vitro biological evaluation and molecular docking. Bioorg Med Chem Lett. 2019;29(23): 126711. https://doi.org/10.1016/j.bmcl.2019.126711.
- Fan H, et al. Discovery of Dioxino [2, 3-f] quinazoline derivative VEGFR-2 inhibitors exerting significant antipro-liferative activity in HUVECs and mice. Eur J Med Chem. 2019;175:349–56. https://doi.org/10.1016/j. ejmech.2019.04.015.
- Türkeş C, et al. Synthesis, biological evaluation and in silico studies of novel N-substituted phthalazine sulfonamide compounds as potent carbonic anhydrase and acetylcholinesterase inhibitors. Bioorg Chem. 2019;89: 103004. https://doi.org/10.1016/j.bioorg.2019.103004.
- Aziz MA, et al. Discovery of potent VEGFR-2 inhibitors based on furopyrimidine and thienopyrimidne scaffolds as cancer targeting agents. Sci Rep. 2016;6(1):1–20. https://doi.org/10.1038/srep24460.
- Mahmoud HK, et al. Novel 2-indolinone thiazole hybrids as sunitinib analogues: design, synthesis, and potent VEGFR-2 inhibition with potential anti-renal cancer activity. Eur J Med Chem. 2020;208: 112752. https://doi. org/10.1016/j.ejmech.2020.112752.
- Eldehna WM, et al. Indoline ureas as potential anti-hepatocellular carcinoma agents targeting VEGFR-2: synthesis, in vitro biological evaluation and molecular docking. Eur J Med Chem. 2015;100:89–97. https://doi. org/10.1016/j.ejmech.2015.05.040.
- Wang C, et al. Biphenyl derivatives incorporating urea unit as novel VEGFR-2 inhibitors: design, synthesis and biological evaluation. Bioorg Med Chem. 2014;22(1):277–84. https://doi.org/10.1016/j.bmc.2013.11. 027.
- 11. Shan Y, et al. Discovery of novel anti-angiogenesis agents. Part 11: development of PROTACs based on active molecules with potency of promoting vascular normalization. Eur J Med Chem. 2020;205: 112654. https://doi.org/10.1016/j.ejmech.2020.112654.
- Sun W, et al. Design, synthesis and biological evaluation of pyrimidinebased derivatives as VEGFR-2 tyrosine kinase inhibitors. Bioorg Chem. 2018;78:393–405. https://doi.org/10.1016/j.bioorg.2018.04.005.
- Luo H, et al. Synthesis and biological evaluation of novel 6-chloroquinazolin derivatives as potential antitumor agents. Eur J Med Chem. 2014;84:746–52. https://doi.org/10.1016/j.ejmech.2014.07.053.
- Kung P-P, et al. Structure—activity relationships of novel 2-substituted quinazoline antibacterial agents. J Med Chem. 1999;42(22):4705–13. https://doi.org/10.1021/jm9903500.
- Srivastava SK, et al. Synthesis of quinazolines as tyrosine kinase inhibitors. Anti Cancer Agents Med Chem. 2009;9(3):246–75. https://doi.org/10. 2174/1871520610909030246.
- Kuneš J, et al. Quinazoline derivatives with antitubercular activity. Il Farmaco. 2000;55(11–12):725–9. https://doi.org/10.1016/S0014-827X(00) 00100-2.
- Avery M, Alvim-Gaston M. New type of febrifugine analogues, bearing a quinolizidine moiety, show potent antimalarial activity against plasmodium malaria parasite. Chemtracts-New York. 2000;13(1):57–61. https:// doi.org/10.1021/jm990131e.
- Jatav V, Kashaw S, Mishra P. Synthesis, antibacterial and antifungal activity of some novel 3-[5-(4-substituted phenyl) 1, 3, 4-thiadiazole-2-yl]-2-styryl quinazoline-4 (3H)-ones. Med Chem Res. 2008;17(2):169–81. https://doi. org/10.1002/ardp.200800005.
- Oggu S, et al. Synthesis, cytotoxicity and molecular docking studies of chalcone incorporated 1, 2, 3-Triazol-1, 3, 5-Triazin-quinazoline as anticancer agents. J Mol Struct. 2022;1266: 133412. https://doi.org/10.1016/j. molstruc.2022.133412.
- Maluleka MM, Mokoena TP, Mampa RM. Synthesis, crystal, and Hirschfeld surface, DFT and molecular docking studies of 6-(3-chloro-4-fluorophenyl)-4-ethoxy-2-(4-methoxyphenyl) quinazoline derivative. J Mol Struct. 2022;1255: 132439. https://doi.org/10.1016/j.molstruc.2022. 132439.
- Dilebo KB, et al. Synthesis, in vitro cytotoxic, anti-Mycobacterium tuberculosis and molecular docking studies of 4-pyridylamino-and 4-(ethynylpyridine) quinazolines. J Mol Struct. 2021;1243: 130824. https://doi.org/10. 1016/j.molstruc.2021.130824.
- 22. Aghcheli A, et al. Design, synthesis, and biological evaluation of 1-(5-(benzylthio)-1, 3, 4-thiadiazol-2-yl)-3-phenylurea derivatives as

anticancer agents. Med Chem Res. 2020;29(11):2000–10. https://doi.org/ 10.1007/s00044-020-02616-2.

- Yang X-H, et al. Synthesis, biological evaluation, and molecular docking studies of cinnamic acyl 1, 3, 4-thiadiazole amide derivatives as novel antitubulin agents. Bioorg Med Chem. 2012;20(3):1181–7. https://doi.org/ 10.1016/j.bmc.2011.12.057.
- Hosseinzadeh L, Khorand A, Aliabadi A. Discovery of 2-P henyl-N-(5-(trifluoromethyl)-1, 3, 4-thiadiazol-2-yl) acetamide derivatives as apoptosis inducers via the caspase pathway with potential anticancer activity. Arch Pharm. 2013;346(11):812–8. https://doi.org/10.1002/ardp.20130 0180.
- Aliabadi A, et al. Synthesis and In-vitro cytotoxicity assessment of N-(5-(Benzylthio)-1, 3, 4-thiadiazol-2-yl)-2-(4-(trifluoromethyl) phenyl) acetamide with potential anticancer activity. Iranian J Pharm Res. 2013;12(4):687.
- Polkam N, et al. Synthesis, in vitro anticancer and antimycobacterial evaluation of new 5-(2, 5-dimethoxyphenyl)-1, 3, 4-thiadiazole-2-amino derivatives. Bioorg Med Chem Lett. 2015;25(7):1398–402. https://doi.org/ 10.1016/j.bmcl.2015.02.052.
- Gomha SM, et al. One pot single step synthesis and biological evaluation of some novel bis (1, 3, 4-thiadiazole) derivatives as potential cytotoxic agents. Molecules. 2016;21(11):1532. https://doi.org/10.3390/molecules2 1111532.
- Vudhgiri S, et al. Synthesis and biological evaluation of 5-fattyacylamido-1, 3, 4-thiadiazole-2-thioglycosides. Bioorg Med Chem Lett. 2017;27(15):3370–3. https://doi.org/10.1016/j.bmcl.2017.06.004.
- Janowska S, et al. New 1, 3, 4-thiadiazole derivatives with anticancer activity. Molecules. 2022;27(6):1814. https://doi.org/10.3390/molecules2 7061814.
- Bhat MA, et al. Synthesis of novel 2, 5-bis (substituted thio)-1, 3, 4-thiadiazoles by acid catalyzed intermolecular cyclization reactions of substituted dithiocarbazates as a possible 2019-nCoV main protease inhibitor. J Mol Struct. 2022;1253: 132252. https://doi.org/10.1016/j.molstruc.2021. 132252.
- Choodamani B, et al. Synthesis, molecular docking, and preliminary cytotoxicity study of some novel 2-(naphthalen-1-yl)-methylimidazo [2, 1-b][1, 3, 4] thiadiazoles. J Mol Struct. 2021;1234: 130174. https://doi.org/ 10.1016/j.molstruc.2021.130174.
- Gallou I. Unsymmetrical ureas. Synthetic methodologies and application in drug design. Organic Prep Proc Int. 2007;39(4):355–83. https://doi.org/ 10.1080/00304940709458592.
- Regan J, et al. Pyrazole urea-based inhibitors of p38 MAP kinase: from lead compound to clinical candidate. J Med Chem. 2002;45(14):2994– 3008. https://doi.org/10.1021/jm020057r.
- Faraji A, et al. Design, synthesis and evaluation of novel thienopyrimidinebased agents bearing diaryl urea functionality as potential inhibitors of angiogenesis. Eur J Med Chem. 2021;209: 112942. https://doi.org/10. 1016/j.ejmech.2020.112942.
- 35. Keating GM. Sorafenib: a review in hepatocellular carcinoma. Target Oncol. 2017;12:243–53. https://doi.org/10.1007/s11523-017-0484-7.
- Dhillon S. Regorafenib: a review in metastatic colorectal cancer. Drugs. 2018;78:1133–44. https://doi.org/10.1007/s40265-018-0938-y.
- Caquelin L, et al. Tivozanib in renal cell carcinoma: a systematic review of the evidence and its dissemination in the scientific literature. BMC Cancer. 2022;22(1):381. https://doi.org/10.1186/s12885-022-09475-7.
- Musumeci F, et al. Vascular endothelial growth factor (VEGF) receptors: drugs and new inhibitors. J Med Chem. 2012;55(24):10797–822. https:// doi.org/10.1021/jm301085w.
- Ferrara N, Gerber H-P, LeCouter J. The biology of VEGF and its receptors. Nat Med. 2003;9(6):669–76. https://doi.org/10.1038/nm0603-669.
- Zhang L, et al. Discovery and evaluation of triple inhibitors of VEGFR-2, TIE-2 and EphB4 as anti-angiogenic and anti-cancer agents. Oncotarget. 2017;8(62): 104745. https://doi.org/10.18632/oncotarget.20065.
- Zhang S, et al. SKLB1002, a novel potent inhibitor of VEGF receptor 2 signaling, inhibits angiogenesis and tumor growth in vivoSKLB1002 inhibits angiogenesis and tumor growth. Clin Cancer Res. 2011;17(13):4439–50. https://doi.org/10.1158/1078-0432.CCR-10-3109.
- Zhang QY, et al. SKLB1002, a potent inhibitor of VEGF receptor 2 signaling, inhibits endothelial angiogenic function in vitro and ocular angiogenesis in vivo. Mol Med Rep. 2020;21(6):2571–9. https://doi.org/10.1158/1078-0432.CCR-10-3109.

- Li C, et al. Discovery of novel anti-angiogenesis agents. Part 7: multitarget inhibitors of VEGFR-2, TIE-2 and EphB4. Eur J Med Chem. 2017;141:506– 18. https://doi.org/10.1016/j.ejmech.2020.112942.
- Abdelhaleem EF, et al. Design and synthesis of thienopyrimidine urea derivatives with potential cytotoxic and pro-apoptotic activity against breast cancer cell line MCF-7. Eur J Med Chem. 2018;143:1807–25. https://doi.org/10.1016/j.ejmech.2017.10.075.
- Ayati A, et al. A review on progression of epidermal growth factor receptor (EGFR) inhibitors as an efficient approach in cancer targeted therapy. Bioorg Chem. 2020;99: 103811. https://doi.org/10.1016/j.bioorg.2020. 103811.
- Ayati A, et al. Pyrimidine-based EGFR TK inhibitors in targeted cancer therapy. Eur J Med Chem. 2021;221: 113523. https://doi.org/10.1016/j. ejmech.2021.113523.
- Ayati A, et al. Thiazole in the targeted anticancer drug discovery. Future Med Chem. 2019;11(16):1929–52. https://doi.org/10.4155/fmc-2018-0416.
- Toolabi M, et al. 6-Cinnamoyl-4-arylaminothienopyrimidines as highly potent cytotoxic agents: design, synthesis and structure-activity relationship studies. Eur J Med Chem. 2020;185: 111786. https://doi.org/10. 1016/j.ejmech.2019.111786.
- Tehrani MB, et al. Phthalimide-1, 2, 3-triazole hybrid compounds as tyrosinase inhibitors; synthesis, biological evaluation and molecular docking analysis. J Mol Struct. 2019;1176:86–93. https://doi.org/10.1016/J.MOLST RUC.2018.08.033.
- Peytam F, et al. Design, synthesis, molecular docking, and in vitro α-glucosidase inhibitory activities of novel 3-amino-2, 4-diarylbenzo [4, 5] imidazo [1, 2-a] pyrimidines against yeast and rat α-glucosidase. Sci Rep. 2021;11(1):1–18. https://doi.org/10.1038/s41598-021-91473-z.
- Moghimi S, et al. Synthesis, in-vitro evaluation, molecular docking, and kinetic studies of pyridazine-triazole hybrid system as novel a-glucosidase inhibitors. Bioorg Chem. 2021;109: 104670. https://doi.org/ 10.1016/j.bioorg.2021.104670.
- Ju K-S, Parales RE. Nitroaromatic compounds, from synthesis to biodegradation. Microbiol Mol Biol Rev. 2010;74(2):250–72. https://doi.org/10. 1128/MMBR.00006-10.
- Strauss MJ. The nitroaromatic group in drug. Design pharmacology and toxicology (for Nonpharmacologists). Ind Eng Chem Prod Res Dev. 1979;18(3):158–66. https://doi.org/10.1021/i360071a002.
- Gulley JL, Drake CG. Immunotherapy for prostate cancer: recent advances, lessons learned, and areas for further research. Clin Cancer Res. 2011;17(12):3884–91. https://doi.org/10.1158/1078-0432.CCR-10-2656.
- Program NT. Eleventh report on carcinogens. National Institute of Environmental Health Sciences. National Institutes of Health, 2005. 19826456.
- Nigro ML, Carballo M. Genotoxicity and cell death induced by tinidazole (TNZ). Toxicol Lett. 2008;180(1):46–52. https://doi.org/10.1016/j.toxlet. 2008.05.017.
- 57. Raether W, Hänel H. Nitroheterocyclic drugs with broad spectrum activity. Parasitol Res. 2003;90(1):S19–39. https://doi.org/10.1007/s00436-002-0754-9.
- Whitmore G, Varghese A. The biological properties of reduced nitroheterocyclics and possible underlying biochemical mechanisms. Biochem Pharmacol. 2013;35:97–103.
- 59. Boelsterli UA, et al. Bioactivation and hepatotoxicity of nitroaromatic drugs. Curr Drug Metab. 2006;7(7):715–27. https://doi.org/10.2174/13892 0006778520606.
- Phillips RM. Targeting the hypoxic fraction of tumours using hypoxiaactivated prodrugs. Cancer Chemother Pharmacol. 2016;77(3):441–57. https://doi.org/10.1007/s00280-015-2920-7.
- Guise CP, et al. The bioreductive prodrug PR-104A is activated under aerobic conditions by human aldo-keto reductase 1C3Aerobic activation of PR-104A by AKR1C3. Can Res. 2010;70(4):1573–84. https://doi.org/10. 1158/0008-5472.CAN-09-3237.
- Pitta E, et al. Differential characterization using readily accessible NMR experiments of novel N-and O-alkylated quinolin-4-ol, 1, 5-naphthyridin-4-ol and quinazolin-4-ol derivatives with antimycobacterial activity. Eur J Med Chem. 2017;125:890–901. https://doi.org/10.1016/j.ejmech.2016.10. 014.
- Bourkoula A, et al. Synthesis and characterization of rhenium and technetium-99m tricarbonyl complexes bearing the 4-[3-bromophenyl] quinazoline moiety as a biomarker for EGFR-TK imaging. Eur J Med Chem. 2009;44(10):4021–7. https://doi.org/10.1016/j.ejmech.2009.04.033.

- Hou X, et al. Design, synthesis and bioactivities evaluation of novel quinazoline analogs containing oxazole units. Chin J Chem. 2014;32(6):538–44. https://doi.org/10.1002/cjoc.201400271.
- Talath S, Gadad A. Synthesis, antibacterial and antitubercular activities of some 7-[4-(5-amino-[1, 3, 4] thiadiazole-2-sulfonyl)-piperazin-1-yl] fluoroquinolonic derivatives. Eur J Med Chem. 2006;41(8):918–24. https:// doi.org/10.1016/j.ejmech.2006.03.027.
- Al-Sanea MM, et al. Design, synthesis, and biological investigation of oxadiazolyl, thiadiazolyl, and pyrimidinyl linked antipyrine derivatives as potential non-acidic anti-inflammatory agents. J Enzyme Inhib Med Chem. 2023;38(1):2162511. https://doi.org/10.1080/14756366.2022.21625 11.
- Schmidt MW, et al. General atomic and molecular electronic structure system. J Comput Chem. 1993;14(11):1347–63. https://doi.org/10.1002/ jcc.540141112.
- Lu T, Chen F. Multiwfn: A multifunctional wavefunction analyzer. J Comput Chem. 2012;33(5):580–92. https://doi.org/10.1002/jcc.22885.
- Daina A, Michielin O, Zoete V. SwissADME: a free web tool to evaluate pharmacokinetics, drug-likeness and medicinal chemistry friendliness of small molecules. Sci Rep. 2017;7(1):42717. https://doi.org/10.1038/srep4 2717.
- Lipinski CA. Lead-and drug-like compounds: the rule-of-five revolution. Drug Discov Today Technol. 2004;1(4):337–41. https://doi.org/10.1016/j. ddtec.2004.11.007.
- Sehout I, et al. Synthesis, crystal structure, Hirshfeld surface analysis, biological evaluation, DFT calculations, and in silico ADME analysis of 4-arylidene pyrazolone derivatives as promising antibacterial agents. J Mol Struct. 2021;1229: 129586. https://doi.org/10.1016/j.molstruc.2020. 129586.
- Coşar ED, et al. Anticholinesterase activities of novel indole-based hydrazide-hydrazone derivatives: design, synthesis, biological evaluation, molecular docking study and in silico ADME prediction. J Mol Struct. 2022;1247: 131398. https://doi.org/10.1016/j.molstruc.2021.131398.
- 73. Dincel ED, et al. Design, biological evaluation, molecular docking study and in silico ADME prediction of novel imidazo [2, 1-b] thiazole derivatives as a novel class of α-glucosidase inhibitors. J Mol Struct. 2021;1245: 131260. https://doi.org/10.1016/j.molstruc.2021.131260.
- Alnoman RB, et al. New quinoline-based BODIPYs as EGFR/VEGFR-2 inhibitors: molecular docking, DFT and in vitro cytotoxicity on HeLa cells. J Mol Struct. 2022;1247: 131312. https://doi.org/10.1016/j.molstruc.2021. 131312.

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