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Synthesis and tyrosinase inhibitory activities of novel isopropylquinazolinones

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Abstract

To find new anti-browning and whitening agents in this study, new series of isopropylquinazolinone derivatives were designed and synthesized. All derivatives were evaluated as possible tyrosinase inhibitors and compound **9q** bearing 4-fluorobenzyl moieties at the R position exhibited the best potencies with an IC_{50} value of $34.67 \pm 3.68 \mu M$. The kinetic evaluations of **9q** as the most potent derivatives recorded mix-type inhibition. Compounds **9o** and **9q** also exhibited potent antioxidant capacity with IC_{50} values of 38.81 and 40.73 μM , respectively confirming their antioxidant potential. Molecular docking studies of **9q** as the most potent derivative were executed and it was shown that quinazolinone and acetamide moieties of compound **9q** participated in interaction with critical His residues of the binding site. The obtained results demonstrated that the **9q** can be considered a suitable pharmacophore to develop potent tyrosinase inhibitors.

Keywords Isopropylquinazolinone, Synthesis, Tyrosinase, Enzyme inhibition, Kinetic evaluation, In silico studies

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Introduction

Melanin pigment is produced by the complex melanogenesis process in melanocytes and plays a significant role in determining the color of human skin, eyes, and hair [1]. Melanin also protects humans against the harmful effects of ultraviolet (UV) [2]. On the other hand, excess secretion of melanin from melanocytes leads to dermatologic disorders such as melasma, freckles, post-inflammatory melanoderma vitiligo, and cancer. Also, there are correlations between melanogenesis disorders and neurodegenerative diseases, including Parkinson's, Alzheimer's, and Huntington's diseases [3, 4].

Therefore, more attention has been paid to develop novel agents and reduce the side effects of hyperpigmentation. Tyrosinase (EC.1.14.18.1) is known as the main rate-limiting enzyme that regulates the biosynthesis of melanin production by the conversion of L-tyrosine to L-dopa and then dopaquinone via monophenolase and diphenolase activities [5, 6]. In this context, tyrosinase inhibitors represent key inhibitory agents to overcome the excess production of melanin. A wide variety of standard tyrosinase inhibitors such as kojic acid, tropolone, and hydroquinone are clinically effective. Most of these inhibitors have associated toxic effects and low efficacy problems. Thus, the production of efficient tyrosinase inhibitors is essential for use in the cosmetic, food, medical and agricultural industries [7, 8].

Quinazolinone is a common structural motif with the fused heterocyclic system that is ubiquitous in several pharmaceuticals and natural products. Quinazolin-4(3H)-one exhibited varied ranges of biological properties such as antibacterial, antifungal, anti-inflammatory, anti-convulsant, anticancer, anti-HIV and analgesic activities

[9–11]. In a recent effort to look for the possible application of quinazolin-4(3H)-one as a tyrosinase inhibitor, 2–4-fluorophenyl-quinazolin-4(3H)-one was reported as tyrosinase inhibitor with $IC_{50} = 120 \pm 2 \mu M$ against tyrosinase in diphenolase activity. Compound **A** was found to be a reversible and mixed-type inhibitor in kinetic studies. In silico studies revealed that this compound affected the mass transfer rate by blocking the enzyme catalytic center [12]. Oxoquinazolin-3(4H)-ylfuran-2-carboxamide (compound **B**) was found to be another potent inhibitor with an IC_{50} value of 0.028 to $1.775 \mu M$ [13]. In 2021, thio-quinazolinone (compound **C**) demonstrated good potency against tyrosinase and reduced the melanin content in the B16F10 cell line with limited toxicity [14]. Yaru Huang et al. reported the synthesis of derivative **D** with mix type inhibition pattern. Molecular docking studies showed that quinazolinone ring participated in several H-bound interactions with tyrosinase enzyme [15].

It seems that the quinazolinone scaffold could be categorized as tyrosinase inhibitor and S, or SH groups on the designed pharmacophore, making it possible to interact with residues of the tyrosinase active site. As can be seen in compounds **B** and **C** (Fig. 1) increased bulkiness at the 3-position of quinazolinone ring improved the potencies. Recently the effects of different types of substitutions at the 3 position of quinazolinone ring were evaluated and it was shown that isopropyl moiety (compounds **E** and **F**, Fig. 1) compared to different aliphatic and aromatic moieties significantly improves the tyrosinase inhibition [16].

As result in this study, isopropyl moiety as aliphatic and bulk groups were conjugated into the 3-position of quinazolinone compared to previous studies bearing

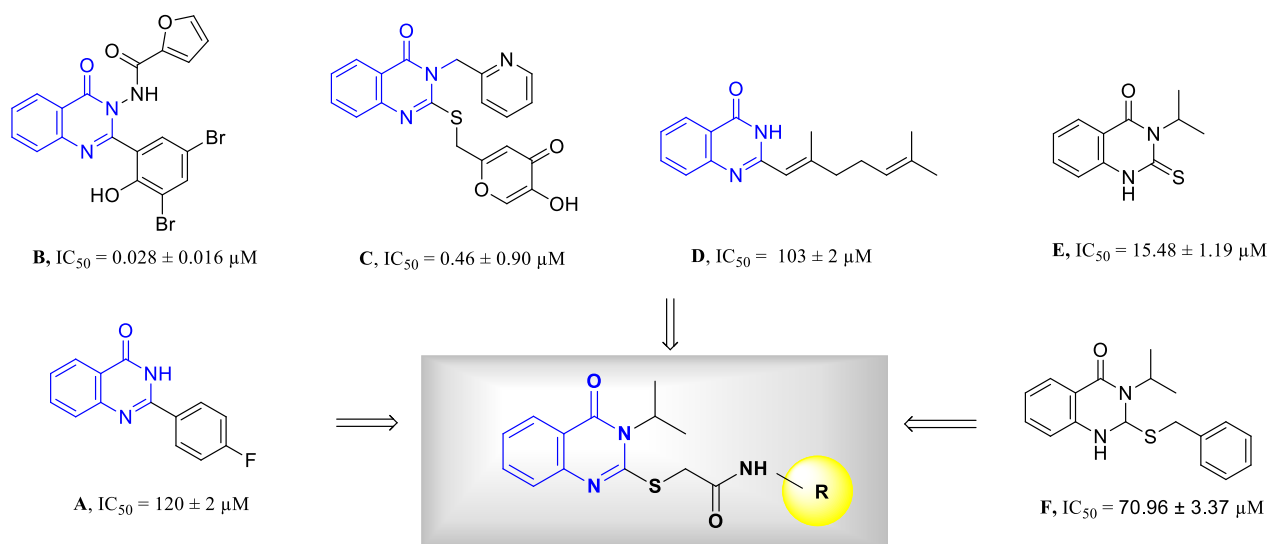
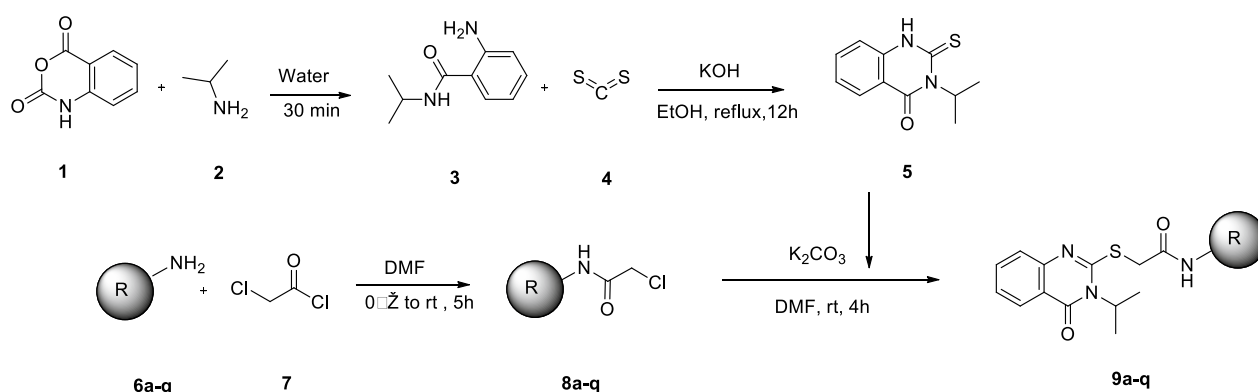


Fig. 1 Chemical structures of some reported tyrosinase inhibitors and newly designed compound



Scheme 1 Synthesis of isopropylquinazolinone derivatives **9a–q**

aromatic substitutions in this position. Furthermore, aryl acetamide moieties might be involved in several hydrophobic and hydrophilic interactions with the binding site via occupying the critical cavities of the enzyme. Derivatization was performed on thioquinazolinone moiety using the different aryl acetamides which could affect the tyrosinase inhibition and antioxidant potential.

Hence, the present studies illustrate the design and synthesis of new isopropylquinazolinones with various substituted analogs on acetamide moiety. After successful synthesis and characterization, all compounds were examined for their tyrosinase activity and antioxidant potential. Moreover, the kinetic and molecular docking studies of the most potent derivatives were performed.

Results and discussion

Chemistry

A mixture of isatoic anhydride (compound **1**) and appropriate amine (compound **2**) was stirred for 30 min at room temperature in water (Scheme 1). The resulting 2-amino-N-isopropylbenzamide (compound **3**), carbon disulfide (compound **4**), and KOH in ethanol were heated under reflux for 12 h to synthesize 3-isopropyl-2-thioxo-2,3-dihydroquinazolin-4(1H)-one (compound **5**).

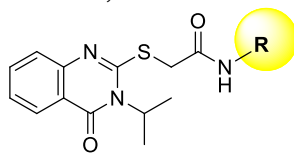
On the other hand, to the stirring solution of amine derivatives (compound **6a–q**) in DMF at 0 °C, chloroacetyl chloride (compound **7**) was added dropwise followed by stirring for 5 h at room temperature. After completion of the reaction, the mixture was poured into ice water to obtain, 2-chloro-N-phenylacetamide derivatives (**8a–q**). Finally, 3-isopropyl-2-thioxo-2,3-dihydroquinazolin-4(1H)-one (compound **5**), were added to the 2-chloro-N-phenylacetamide derivatives (**8a–q**) in DMF in presence of potassium carbonate at room temperature for 4 h. The structure of final products **9a–q** was confirmed using NMR and IR spectroscopy as well as elemental analysis.

Mushroom tyrosinase inhibition and structure–activity relationship

All newly synthesized compounds (**9a–q**) were examined for their inhibitory potentials against the tyrosinase enzyme and the results were presented in Table 1 and the structure–activity relationships (SARs) were constructed. From the experimental data, it appears that all of the synthesized compounds have an inhibitory effect against tyrosinase for the oxidation of L-dopa as compared with kojic acid as the positive control ($IC_{50} = 23.64 \pm 2.56 \mu\text{M}$).

The parent compound **9a** (% inhibition at 50 μM = 21.64) exhibited weak inhibitory activity against tyrosinase in comparison with the standard kojic acid. The halogen-substituted analogs (**9b–e**) displayed improvement in the potencies compared to unsubstituted analogs. Amongst, 2-chlorophenyl (**9b**, % inhibition at 50 μM = 37.97) exhibited the best results as compared to the rest of the halogen-substituted groups. On the other hand, evaluations of *para*-halogen substituted groups did not present significant differences so that **9d** (R = 4-chlorophenyl, % inhibition at 50 μM = 26.86) and **9e** (R = 4-bromo-phenyl, % inhibition at 50 μM = 24.77) exhibited similar inhibition at 50 μM . On the other hand, **9f** with *para*-nitro moiety was recorded as the second most active derivative among the electron-withdrawing analogs which could be due to the hydrophilicity of NO_2 group as well as its strong electron-withdrawing power.

Furthermore, in most cases, the mono electron-donating substituted analogs (**9g–k**) showed improvement in the inhibitory potential compared to the unsubstituted derivative, **9a**. The best potency among electron-donating derivatives was seen in **9j** (R: 4-hydroxyphenyl) and **9g** (R: 2-methylphenyl) with 35.48% and 34.62% inhibition at 50 μM . However, changing the position from *ortho*-methyl to *para*-methyl reduced the activity in **9h** compared to **9g**. Also, the replacement of methyl with methoxy did not

Table 1 Tyrosinase inhibitory activity of compounds **9a–q**


Compounds	R	% Inhibition at 50 μM	IC ₅₀ (μM)
9a	Phenyl	21.64 \pm 3.79	NA ^a
9b	2-Chlorophenyl	37.97 \pm 4.68	NA
9c	3-Chlorophenyl	27.76 \pm 1.19	NA
9d	4-Chlorophenyl	26.86 \pm 4.67	NA
9e	4-Bromophenyl	24.77 \pm 3.12	NA
9f	4-Nitrophenyl	32.68 \pm 5.67	NA
9g	2-Methylphenyl	34.62 \pm 4.67	NA
9h	4-Methylphenyl	26.71 \pm 6.42	NA
9i	4-Methoxyphenyl	25.37 \pm 3.13	NA
9j	4-Hydroxyphenyl	35.48 \pm 4.91	NA
9k	4-Ethylphenyl	20.44 \pm 4.19	NA
9l	2,3-Dimethylphenyl	18.20 \pm 5.52	NA
9m	2,4-Dimethylphenyl	27.31 \pm 3.43	NA
9n	2,6-Dimethylphenyl	16.26 \pm 6.57	NA
9o	Naphthyl	35.22 \pm 3.88	NA
9p	Benzyl	29.08 \pm 5.24	NA
9q	4-Fluorobenzyl	54.47 \pm 3.94	34.67 \pm 3.68
Kojic acid ^b	-		23.64 \pm 2.56

Data presented here are the mean \pm S.E

^a NA means not active

^b Positive control

improve the potency in **9i** with just 25.37% inhibition at 50 μM . The activity pattern is a bit the same in the case of the *para*-ethyl substituted compound (**9k**) which recorded 20.44% inhibition at 50 μM similar to an unsubstituted derivative. In the case of one electron-donating substitution, it was assumed that increased bulkiness or lipophilicity at the *para* position is not favorable and the exception in this trend came back to **9j** bearing *para*-hydroxyphenyl.

Compounds **9l–n** with dimethyl substitution mostly displayed inferior inhibitory potential compared to the unsubstituted derivative. It seems that increased steric hinder deteriorated the potencies although **9m** exhibited 27.31% inhibition at 50 μM . Noteworthy, the replacement of phenyl (**9a**) with a naphthyl ring (**9o**) resulted in improved inhibition.

Assessment of phenyl analogs compared with benzyl derivatives recorded interesting results so that derivative **9p** (R = benzyl with 29.08% inhibition at 50 μM) revealed better potencies as compared to **9a** analogs. Also, **9q** bearing 4-fluorobenzyl moiety was discovered to be the most active derivative of the series.

Overall the precise mechanism of action of the isopropylquinazolinones is unknown. However, the increase of benzyl derivatives' potencies compared to the phenyl compounds might indicate better interaction of benzyl analogs with the hydrophobic pocket of the tyrosinase enzyme and the possibility of better rotation of substitutions in the binding pocket of the enzyme. Also, SARs analysis demonstrated mostly, better inhibitory activities of *ortho*-substituted analogous (**9b** and **9g**) as compared to the *para*-substituted derivatives (**9d** and **9h**).

Enzyme kinetic studies

According to Fig. 2a, the Lineweaver–Burk plot showed that **9q** is a mix-type inhibitor in which the inhibitor may bind to the enzyme whether or not the enzyme has already bound to the substrate. Furthermore, the plot of the slope of the lines versus different concentrations of inhibitor gave an estimate of the inhibition constant (K_I) of 13.60 μM (Fig. 2b). Compound **9q** also recorded the inhibition constant with the enzyme–substrate complex (K_{IS}) of 87.32 μM (Fig. 2c) [17, 18].

Inspecting the antioxidant properties of isopropylquinazolinones

In vitro antioxidant activities of the synthesized compounds were determined quantitatively using DPPH radical scavenging assay. DPPH is a stable free radical at room temperature and accepts an electron or hydrogen radical from an antioxidant to become a stable diamagnetic molecule [19]. The results of antioxidant potential are presented in Table 2. These results delineate that the unsubstituted compounds **9a** and **9p** had no antioxidant capacity while mostly any types of substitutions improved the antioxidant activities. The best potencies came back to **9o** bearing naphthyl moiety with an IC₅₀ value of 38.81 μM followed by **9q** containing 4-fluorobenzyl (IC₅₀ = 40.73 μM). Also, 4-bromophenyl (% inhibition at 100 μM = 41.44) and 4-chloro-phenyl (% inhibition at 100 μM = 42.78) derivatives were other potent antioxidant agents. Similar to previous studies, halogens especially fluorine, increases antioxidant activity [20]. Assessments of electron donating groups showed compound **9n** bearing 2,6-dimethyl-phenyl was the most potent agent among electron donating derivatives with 44.38% inhibition at 100 μM .

Docking study

In the first step, molecular docking validation was performed with the redocking of tropolone as a native ligand inside the tyrosinase. Alignment of the best pose of tropolone in the active site of the enzyme and crystallographic ligand recorded an RMSD value of 0.42 Å (RMSD should be less than 2 Å) which confirms the accuracy of

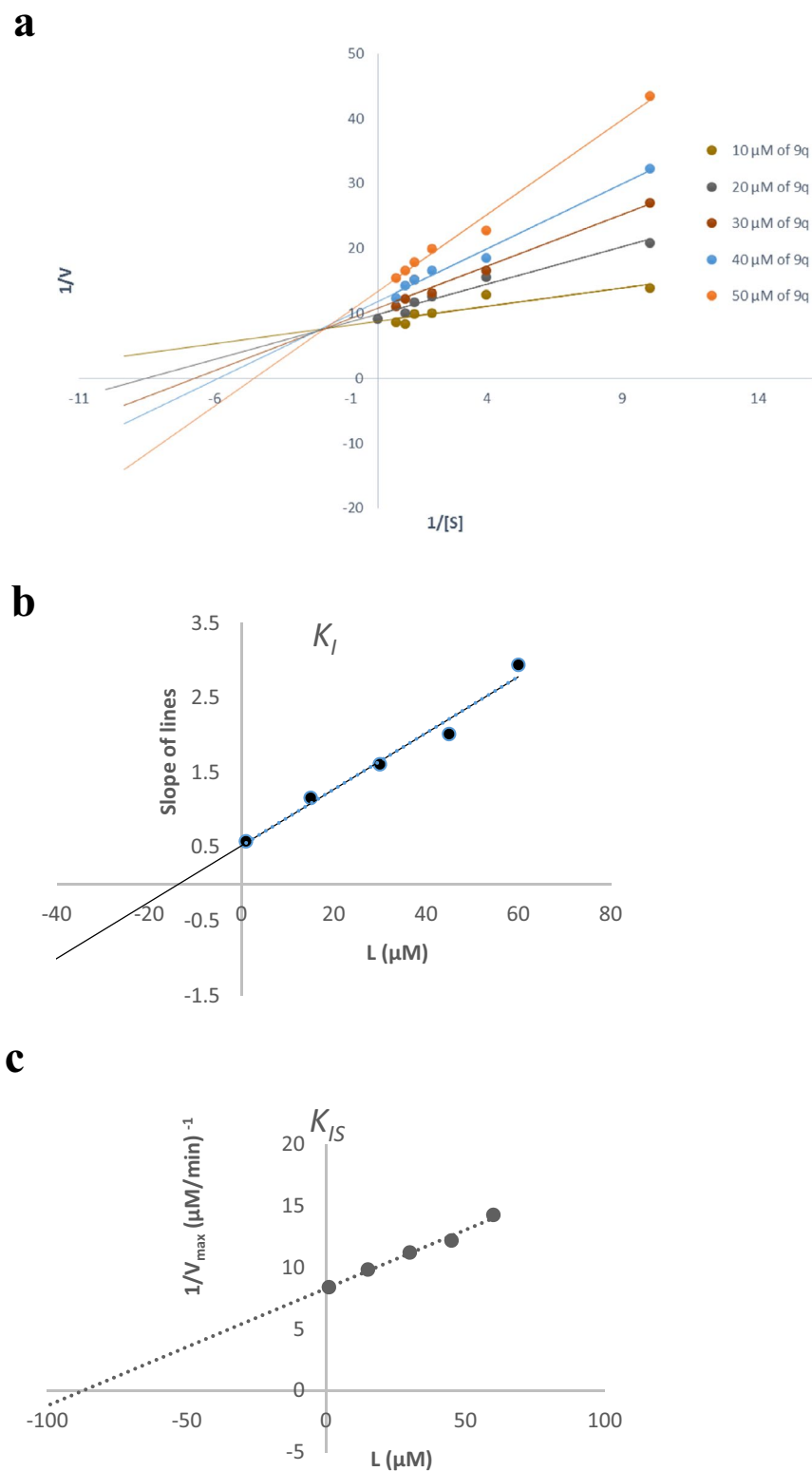


Fig. 2 Kinetics of tyrosinase inhibition by 9q. **a** The Lineweaver–Burk plot in the absence and presence of different concentrations of 9q; **b** The secondary plot of slopes of the lines vs various concentrations of 9q; **c** The secondary plot of $1/V_{\text{max}}$ vs various concentrations of 9q

Table 2 Antioxidant properties of synthesized compounds using DPPH assay

Compound	R	% Inhibition at 100 μ M	IC ₅₀ (μ M) \pm RSD ^a
9a	Phenyl	2.941 \pm 1.76	NA
9b	2-Chlorophenyl	26.47 \pm 2.48	NA
9c	3-Chlorophenyl	18.71 \pm 5.39	NA
9d	4-Chlorophenyl	42.78 \pm 4.86	NA
9e	4-Bromophenyl	41.44 \pm 3.85	NA
9f	4-Nitrophenyl	4.87 \pm 1.65	NA
9g	2-Methylphenyl	20.71 \pm 2.71	NA
9h	4-Methylphenyl	29.41 \pm 1.76	NA
9i	4-Methoxyphenyl	20.05 \pm 3.47	NA
9j	4-Hydroxyphenyl	40.83 \pm 4.53	NA
9k	4-Ethylphenyl	10.16 \pm 4.27	NA
9l	2,3-Dimethylphenyl	5.08 \pm 2.14	NA
9m	2,4-Dimethylphenyl	21.12 \pm 2.99	NA
9n	2,6-Dimethylphenyl	44.38 \pm 5.03	NA
9o	Naphtyl	68.44 \pm 4.84	38.81 \pm 3.06
9p	Benzyl	4.47 \pm 2.35	NA
9q	4-Fluorobenzyl	60.33 \pm 5.53	40.73 \pm 2.77
Quercetin ^b		–	18.56 \pm 2.19

Data presented here are the mean \pm S.E of three independent experiments

^a NA means not active

^b Positive control

docking. Molecular docking results revealed a fit-well binding pattern of compound **9q** with the adaptation of several key interactions with active site residues (Fig. 3).

Numerous significant interactions with catalytic residues were found so that quinazolinone rings interacted through π - π staking interaction with His263 (one of the critical residues of the active site) whereas, the C=O and NH groups of acetamide moieties formed two critical hydrogen bonds with His85 and His244 of target tyrosinase protein. Literature data ensured the importance of these His residues in bonding with tyrosinase inhibitors, which strengthen our docking results.

Conclusions

In summary, in this study, a series of isopropylquinazolinones was developed and evaluated as tyrosinase inhibitors. The SARs was presented by analyzing the impact of varying substitutions on tyrosinase inhibitory potential. The 4-fluorobenzyl-substituted compound **9q** displayed the best tyrosinase inhibition of the series. The kinetic studies of the most active compound, **9q**, showed that this compound had a mixed mode of inhibition. Also, their antioxidant properties were determined using DPPH assay and SARs analysis of antioxidant capacities showed that the *para*-halogen substituted compounds displayed better antioxidant activities. The binding conformation of the most potent derivative was determined by a molecular docking study and revealed that structural features of **9q** as an active compound play a vital role in the binding interactions of this compound with the enzyme.

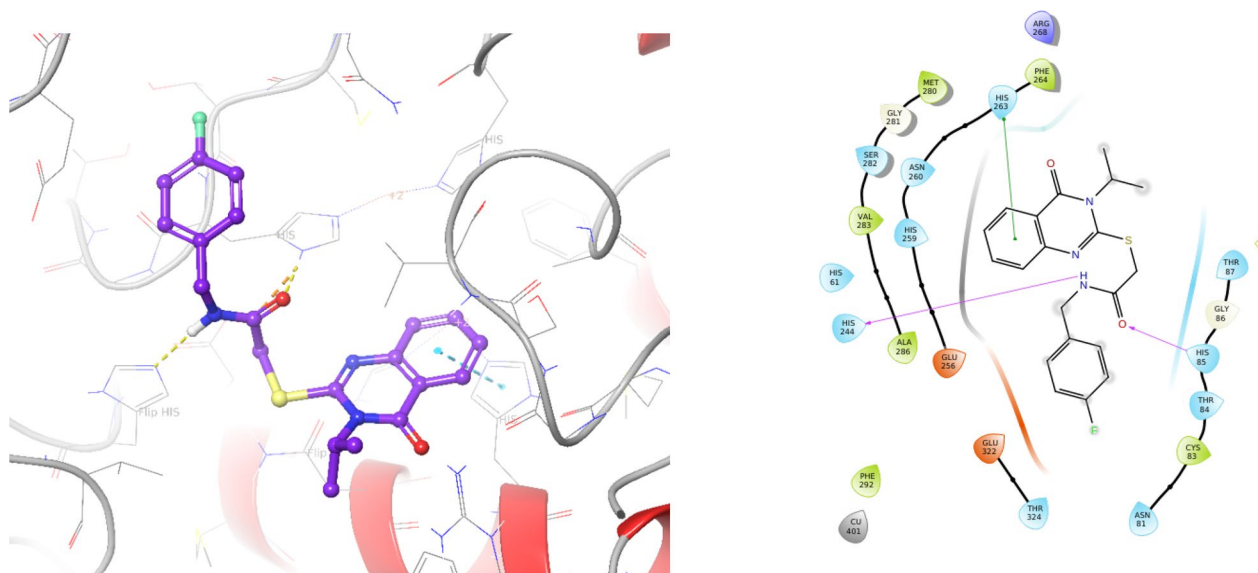


Fig. 3 3D and 2D diagrams of compound **9q** within the binding pocket of tyrosinase

Material and methods

Synthesis

A mixture of isatoic anhydride (compound **1**, 10 mmol) and appropriate amine (compound **2**, 10 mmol) was stirred for 30 min at room temperature in water (20 mL). After completion of the reaction (checked by TLC), the resulting white 2-amino-N-isopropylbenzamide was filtered off and used for the next reaction without further purification. 2-amino-N-isopropylbenzamide (compound **3**, 1 mmol), carbon disulfide (compound **4**, 1.2 mmol), and KOH (1.5 mmol) in ethanol (5 mL) were heated under reflux for 12 h. After completion of the reaction (checked by TLC), the reaction mixture was poured into cold water which was acidified with HCL 6N (pH=5). The precipitates were filtered off, and washed with cold water to obtain the pure 3-isopropyl-2-thioxo-2,3-dihydroquinazolin-4(1H)-one (compound **5**). To a solution of amine derivatives (compound **6a–q**, 1 mmol) in DMF (4 mL) at 0 °C, chloroacetyl chloride (compound **7**), was added. The mixture was stirred at room temperature for 5 h and poured into ice water and then filtered to get the obtained solids (**8a–q**) were then filtered, dried, and recrystallized from ethanol.

A mixture of 3-isopropyl-2-thioxo-2,3-dihydroquinazolin-4(1H)-one (compound **5**, 1 mmol), 2-chloro-N-phenylacetamide derivatives (**8a–q**, 1 mmol) and, potassium carbonate (1.5 mmol) in DMF (5 ml) was stirred at rt for 4 h. Next, the mixture reaction was poured in cold water and the residue was filtrated and purified using recrystallization from ethanol to give 2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-phenylacetamide derivatives (**9a–q**).

Analytical HPLC evaluation of **9b**, **9o**, and **9q** was performed on a Kanuer HPLC system equipped with UV detectors using an RP column (C18, 5 µm, 150×4.6 mm) and solvent acetonitrile and water with flow rate, 1 mL/min; detection, 254 nm; injection volume, 10 µL (Additional file 1: Figs. S1, S2 and S3).

2-((3-Isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-phenylacetamide (9a)

Cream solid; Yield:77%; MP = 190–192 °C; IR (KBr, ν_{\max}) 3252 (NH), 3025 (CH Aromatic), 2970(CH Aliphatic), 1660 (C=O) Cm^{-1} ; ^1H NMR (500 MHz, DMSO- d_6) δ 10.39 (s,1H, NH_{Amide}), 8.01 (d, $J=7.90$ Hz, 1H, H₅), 7.71 (t, $J=7.60$ Hz, 1H, H₇), 7.61 (d, $J=8.10$ Hz, 2H, H₂, H₆), 7.42 (d, $J=8.10$ Hz, 1H, H₈), 7.38 (t, $J=7.50$ Hz, 1H, H₆), 7.29 (t, $J=7.80$ Hz, 2H, H₃, H₅), 7.03 (t, $J=7.30$ Hz, 1H, H₄), 4.75–4.40 (m, 1H, CH_{Isopropyl}), 4.17 (s, 2H, CH₂), 1.59 (d, $J=6.70$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO-d₆): δ 166.2, 161.3, 156.6, 146.6, 139.5, 134.9, 129.2, 126.6, 126.3, 125.8, 123.8, 120.3, 119.5; ESI–MS (C₁₉H₁₉N₃O₂S): calculated m/z 353.12

[M+H]⁺, observed m/z 353.19[M+H]⁺; Anal. Calcd: C₁₉H₁₉N₃O₂S; C, 64.57; H, 5.42; N, 11.89; Found; C, 64.69; H, 5.57; N, 12.02.

N-(2-chlorophenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9b)

Brown solid; Yield:78%; MP = 207–209 °C; IR (KBr, ν_{\max}) 3357 (NH), 3070 (CH Aromatic), 2980 (CH Aliphatic), 1677 (C=O) Cm^{-1} ; ^1H NMR (500 MHz, DMSO- d_6) δ 9.86 (s,1H, NH_{Amide}), 8.03 (d, $J=7.90$ Hz, 1H, H₅), 7.80–7.69 (m, 2H, H₇, H₆), 7.50 (d, $J=8.10$ Hz, 1H, H₈), 7.45 (d, $J=8.00$ Hz, 1H, H₃), 7.39 (t, $J=7.50$ Hz, 1H, H₆), 7.28 (t, $J=7.70$ Hz, 1H, H₅), 7.14 (t, $J=7.60$ Hz, 1H, H₄), 4.83–4.54 (m, 1H, CH_{Isopropyl}), 4.27 (s, 2H, CH₂), 1.59 (d, $J=6.50$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 166.8, 161.3, 156.4, 146.6, 135.2, 134.8, 129.9, 127.9, 126.6, 126.3, 126.2, 126.0, 125.8, 120.3, 37.0, 19.6; ESI–MS (C₁₉H₁₈ClN₃O₂S): calculated m/z 387.08 [M+H]⁺, observed m/z 387.13 [M+H]⁺; Anal. Calcd: C₁₉H₁₈ClN₃O₂S; C, 58.83; H, 4.68; N, 10.83; Found; C, 59.13; H, 4.85; N, 11.01.

N-(3-chlorophenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9c)

Brown solid; Yield:74%; MP = 203–205 °C; IR (KBr, ν_{\max}) 3363 (NH), 3070 (CH Aromatic), 2980 (CH Aliphatic), 1681 (C=O) Cm^{-1} ; ^1H NMR (500 MHz, DMSO- d_6) δ 10.60 (s,1H, NH_{Amide}), 8.01 (d, $J=8.20$ Hz, 1H, H₅), 7.81 (s, 1H, H₂), 7.71 (t, $J=7.60$ Hz, 1H, H₇), 7.48 (d, $J=8.50$ Hz, 1H, H₈), 7.40–7.36 (m, 2H, H₆, H₆), 7.34 (t, $J=8.00$ Hz, 1H, H₅), 7.10 (d, $J=8.50$ Hz, 1H, H₄), 4.74–4.54 (m, 1H, CH_{Isopropyl}), 4.18 (s, 2H, CH₂), 1.59 (d, $J=6.50$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 166.7, 161.3, 156.5, 146.6, 140.9, 134.9, 133.5, 130.9, 126.6, 126.3, 125.8, 123.5, 120.3, 119.0, 117.9, 37.7, 19.6; ESI–MS (C₁₉H₁₈ClN₃O₂S): calculated m/z 387.08 [M+H]⁺, observed m/z 387.17 [M+H]⁺; Anal. Calcd: C₁₉H₁₈ClN₃O₂S; C, 58.83; H, 4.68; N, 10.83; Found; C, 59.03; H, 4.81; N, 10.98.

N-(4-chlorophenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9d)

Brown solid; Yield:79%; MP = 211–213 °C; IR (KBr, ν_{\max}) 3378 (NH), 3065 (CH Aromatic), 2990 (CH Aliphatic), 1703 (C=O) Cm^{-1} ; ^1H NMR (500 MHz, DMSO- d_6) δ 10.54 (s,1H, NH_{Amide}), 8.01 (d, $J=8.00$ Hz, 1H, H₅), 7.70 (t, $J=7.60$ Hz, 1H, H₇), 7.64 (d, $J=8.50$ Hz, 2H, H₂, H₆), 7.41–7.36 (m, 2H, H₈, H₆), 7.35 (d, $J=8.50$ Hz, 2H, H₃, H₅), 4.81–4.54 (m, 1H, CH_{Isopropyl}), 4.17 (s, 2H, CH₂), 1.58 (d, $J=6.60$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 166.4, 161.3, 156.5, 146.6, 138.4, 134.9, 129.1, 127.4, 126.6, 126.3, 125.8, 121.0, 120.3, 37.7, 19.6; ESI–MS (C₁₉H₁₈ClN₃O₂S): calculated m/z 387.08

[M+H]⁺, observed m/z 387.21 [M+H]⁺; Anal. Calcd: C₁₉H₁₈ClN₃O₂S; C, 58.83; H, 4.68; N, 10.83; Found; C, 59.11; H, 4.89; N, 10.98.

N-(4-bromophenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9e)

Brown solid; Yield: 84%; MP = 203–205 °C; IR (KBr, ν_{\max}) 3378 (NH), 3020 (CH Aromatic), 2885 (CH Aliphatic), 1661 (C=O) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 10.54 (s, 1H, NH_{Amide}), 8.01 (d, *J* = 7.90 Hz, 1H, H₅), 7.70 (t, *J* = 7.60 Hz, 1H, H₇), 7.59 (d, *J* = 8.10 Hz, 2H, H₂, H₆), 7.49–7.44 (m, 2H, H₈, H₆), 7.40–7.34 (m, 2H, H₃, H₅), 4.75–4.52 (m, 1H, CH_{Isopropyl}), 4.17 (s, 2H, CH₂), 1.58 (d, *J* = 6.70 Hz, 6H, 2xCH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 166.5, 161.3, 156.5, 146.6, 138.8, 134.9, 132.0, 126.6, 126.3, 125.8, 121.4, 120.3, 115.4, 37.8, 19.6; ESI-MS (C₁₉H₁₈BrN₃O₂S): calculated m/z 431.03 [M+H]⁺, observed m/z 431.09 [M+H]⁺; Anal. Calcd: C₁₉H₁₈BrN₃O₂S; C, 52.78; H, 4.20; N, 9.72; Found; C, 52.91; H, 4.40; N, 9.97.

2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-(4-nitrophenyl)acetamide (9f)

Brown solid; Yield: 81%; MP = 216–218 °C; IR (KBr, ν_{\max}) 3343 (NH), 3040 (CH Aromatic), 2980 (CH Aliphatic), 1674 (C=O), 1560–1355 (NO₂) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 11.03 (s, 1H, NH_{Amide}), 8.22 (d, *J* = 7.50 Hz, 2H, H₃, H₅), 8.00 (d, *J* = 7.90 Hz, 1H, H₅), 7.87 (d, *J* = 8.10 Hz, 2H, H₂, H₆), 7.68 (t, *J* = 7.60 Hz, 1H, H₇), 7.37 (t, *J* = 7.20 Hz, 1H, H₆), 7.37 (d, *J* = 7.20 Hz, 1H, H₈), 4.84–4.43 (m, 1H, CH_{Isopropyl}), 4.22 (s, 2H, CH₂), 1.60 (d, *J* = 6.10 Hz, 6H, 2xCH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 167.5, 161.3, 156.4, 146.5, 145.6, 142.7, 135.0, 126.6, 126.3, 125.7, 125.5, 125.4, 119.2, 37.8, 19.6; ESI-MS (C₁₉H₁₈N₄O₄S): calculated m/z 398.10 [M+H]⁺, observed m/z 398.28 [M+H]⁺; Anal. Calcd: C₁₉H₁₈N₄O₄S; C, 57.27; H, 4.55; N, 14.06; Found; C, 57.43; H, 4.81; N, 14.25.

2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-(o-tolyl)acetamide (9g)

Cream solid; Yield: 76%; MP = 196–198 °C; IR (KBr, ν_{\max}) 3238 (NH), 3035 (CH Aromatic), 2965 (CH Aliphatic), 1682 (C=O) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 9.69 (s, 1H, NH_{Amide}), 8.04 (d, *J* = 7.90 Hz, 1H, H₅), 7.75 (t, *J* = 7.60 Hz, 1H, H₇), 7.50 (d, *J* = 8.10 Hz, 1H, H₈), 7.41 (t, *J* = 7.50 Hz, 1H, H₆), 7.36 (d, *J* = 7.80 Hz, 1H, H₃), 7.18 (d, *J* = 7.30 Hz, 1H, H₆), 7.13 (t, *J* = 7.50 Hz, 1H, H₅), 7.06 (t, *J* = 7.30 Hz, 1H, H₄), 4.81–4.57 (m, 1H, CH_{Isopropyl}), 4.22 (s, 2H, CH₂), 2.18 (s, 3H, CH₃), 1.59 (d, *J* = 6.50 Hz, 6H, 2xCH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 166.2, 161.3, 156.6, 146.6, 136.6, 134.9, 132.3, 130.7, 126.6, 126.4, 126.3, 125.9, 125.8, 125.4, 120.4, 34.0, 19.6,

18.2; ESI-MS (C₂₀H₂₁N₃O₂S): calculated m/z 387.08 [M+H]⁺, observed m/z 387.13 [M+H]⁺; Anal. Calcd: C₂₀H₂₁N₃O₂S; C, 65.37; H, 5.76; N, 11.44; Found; C, 65.52; H, 5.91; N, 11.67.

2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-(p-tolyl)acetamide (9h)

Cream solid; Yield: 76%; MP = 199–201 °C; IR (KBr, ν_{\max}) 3400 (NH), 3020 (CH Aromatic), 2975 (CH Aliphatic), 1701 (C=O) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 10.44 (s, 1H, NH_{Amide}), 8.01 (d, *J* = 7.90 Hz, 1H, H₅), 7.71 (t, *J* = 7.60 Hz, 1H, H₇), 7.50 (d, *J* = 8.30 Hz, 2H, H₂, H₆), 7.42 (d, *J* = 8.10 Hz, 1H, H₈), 7.38 (t, *J* = 7.50 Hz, 1H, H₆), 7.09 (d, *J* = 8.10 Hz, 2H, H₃, H₅), 4.75–4.57 (m, 1H, CH_{Isopropyl}), 4.17 (s, 2H, CH₂), 2.22 (s, 3H, CH₃), 1.59 (d, *J* = 6.60 Hz, 6H, 2xCH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 166.0, 161.3, 156.6, 146.6, 137.0, 134.9, 132.7, 129.5, 126.5, 126.2, 125.9, 120.3, 119.5, 37.7, 20.8, 19.6; ESI-MS (C₂₀H₂₁N₃O₂S): calculated m/z 367.14 [M+H]⁺, observed m/z 367.28 [M+H]⁺; Anal. Calcd: C₂₀H₂₁N₃O₂S; C, 65.37; H, 5.76; N, 11.44; Found; C, 65.51; H, 5.97; N, 11.59.

2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-(4-methoxyphenyl)acetamide (9i)

Cream solid; Yield: 88%; MP = 205–207 °C; IR (KBr, ν_{\max}) 3253 (NH), 3030 (CH Aromatic), 2975 (CH Aliphatic), 1661 (C=O) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 10.25 (s, 1H, NH_{Amide}), 8.02 (d, *J* = 7.90 Hz, 1H, H₅), 7.72 (t, *J* = 7.60 Hz, 1H, H₇), 7.51 (d, *J* = 8.20 Hz, 2H, H₂, H₆), 7.44 (d, *J* = 8.10 Hz, 1H, H₈), 7.39 (t, *J* = 7.60 Hz, 1H, H₆), 6.87 (d, *J* = 8.20 Hz, 2H, H₃, H₅), 4.83–4.51 (m, 1H, CH_{Isopropyl}), 4.14 (s, 2H, CH₂), 3.70 (s, 3H, CH₃), 1.59 (d, *J* = 6.50 Hz, 6H, 2xCH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 165.7, 161.3, 156.6, 155.7, 146.6, 134.9, 132.6, 126.6, 126.3, 125.9, 121.1, 120.3, 114.3, 55.5, 37.7, 19.6; ESI-MS (C₂₀H₂₁N₃O₃S): calculated m/z 383.13 [M+H]⁺, observed m/z 383.29 [M+H]⁺; Anal. Calcd: C₂₀H₂₁N₃O₃S; C, 62.64; H, 5.52; N, 10.96; Found; C, 62.82; H, 5.75; N, 11.13.

N-(4-hydroxyphenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9j)

Cream solid; Yield: 77%; MP = 193–195 °C; IR (KBr, ν_{\max}) 3295 (NH), 3075 (CH Aromatic), 2970 (CH Aliphatic), 1675 (C=O) Cm^{-1} ; ¹H NMR (500 MHz, DMSO-*d*₆) δ 11.98 (s, 1H, OH), 10.25 (s, 1H, NH_{Amide}), 8.03 (d, *J* = 7.70 Hz, 1H, H₅), 7.74 (t, *J* = 7.50 Hz, 1H, H₇), 7.52 (d, *J* = 8.00 Hz, 2H, H₂, H₆), 7.43 (d, *J* = 7.90 Hz, 1H, H₈), 7.39 (t, *J* = 7.50 Hz, 1H, H₆), 6.88 (d, *J* = 8.10 Hz, 2H, H₃, H₅), 4.67 (m, 1H, CH_{Isopropyl}), 4.25 (s, 2H, CH₂), 1.60 (d, *J* = 6.50 Hz, 3H, CH₃), 1.59 (d, *J* = 6.40 Hz, 3H, CH₃) ppm. ¹³C NMR (125 MHz, DMSO-*d*₆): δ 165.5, 161.4, 156.7,

155.8, 147.6, 134.9, 132.6, 127.6, 127.3, 126.8, 122.1, 120.5, 115.4, 38.7, 19.6; ESI-MS ($C_{19}H_{19}N_3O_3S$): calculated m/z 369.11 $[M+H]^+$, observed m/z 370.11 $[M+H]^+$; Anal. Calcd: $C_{19}H_{19}N_3O_3S$; C, 61.77; H, 5.18; N, 11.37; Found; C, 62.05; H, 5.81; N, 11.17.

N-(4-ethylphenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9k)

Cream solid; Yield:81%; MP=195–197 °C; IR (KBr, ν_{max}) 3401 (NH), 3065(CH Aromatic), 2950(CH Aliphatic), 1670 (C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 10.44 (s, 1H, NH_{Amide}), 8.01 (d, $J=7.40$ Hz, 1H, H₅), 7.72 (t, $J=7.60$ Hz, 1H, H₇), 7.52 (d, $J=8.30$ Hz, 2H, H₂, H₆), 7.43 (d, $J=8.20$ Hz, 1H, H₈), 7.39 (t, $J=7.60$ Hz, 1H, H₆), 7.12 (d, $J=8.10$ Hz, 2H, H₃, H₅), 4.76–4.59 (m, 1H, CH_{Isopropyl}), 4.17 (s, 2H, CH₂), 2.52 (q, 2H, CH₂_{Ethyl}), 1.58 (d, $J=6.60$ Hz, 6H, 2xCH₃), 1.13 (t, 3H, CH₃_{Ethyl}) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 166.0, 161.3, 156.6, 146.6, 139.1, 137.2, 134.9, 128.3, 126.5, 126.3, 125.9, 120.3, 119.6, 37.7, 28.0, 19.6, 16.1. ESI-MS ($C_{21}H_{23}N_3O_2S$): calculated m/z 381.15 $[M+H]^+$, observed m/z 381.27 $[M+H]^+$; Anal. Calcd: $C_{21}H_{23}N_3O_2S$; C, 66.12; H, 6.08; N, 11.01; Found C, 66.37; H, 6.24; N, 11.35.

N-(2,3-dimethylphenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9l)

Brown solid; Yield:69%; MP=217–219 °C; IR (KBr, ν_{max}) 3371 (NH), 3040 (CH Aromatic), 2985 (CH Aliphatic), 1669 (C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 9.86 (s, 1H, NH_{Amide}), 8.03 (d, $J=7.90$ Hz, 1H, H₅), 7.75 (t, $J=7.20$ Hz, 1H, H₇), 7.51 (d, $J=8.20$ Hz, 1H, H₈), 7.41 (t, $J=7.50$ Hz, 1H, H₆), 7.09 (d, $J=7.50$ Hz, 1H, H₆), 7.03–6.95 (m, 2H, H₄, H₅), 4.75–4.61 (m, 1H, CH_{Isopropyl}), 4.21 (s, 2H, CH₂), 2.20 (s, 3H, CH₃), 2.04 (s, 3H, CH₃), 1.59 (d, $J=6.60$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 166.2, 161.4, 156.6, 146.7, 137.4, 136.4, 134.9, 131.8, 127.5, 126.6, 126.3, 126.0, 125.6, 123.9, 120.3, 37.0, 20.5, 19.6, 14.5; ESI-MS ($C_{21}H_{23}N_3O_2S$): calculated m/z 381.15 $[M+H]^+$, observed m/z 381.24 $[M+H]^+$; Anal. Calcd: $C_{21}H_{23}N_3O_2S$; C, 66.12; H, 6.08; N, 11.01; Found; C, 66.33; H, 6.27; N, 11.21.

N-(2,4-dimethylphenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9m)

Brown solid; Yield: 69%; MP=222–225 °C; IR (KBr, ν_{max}) 3376 (NH), 3045 (CH Aromatic), 2995 (CH Aliphatic), 1674 (C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 9.57 (s, 1H, NH_{Amide}), 8.04 (d, $J=7.90$ Hz, 1H, H₅), 7.77 (t, $J=7.60$ Hz, 1H, H₇), 7.53 (d, $J=8.80$ Hz, 1H, H₆), 7.42 (t, $J=7.50$ Hz, 1H, H₆), 7.09–6.98 (m, 3H, H₈, H₃, H₅), 4.81–4.60 (m, 1H, CH_{Isopropyl}), 4.23 (s, 2H, CH₂), 3.32 (s, 3H, CH₃), 2.09 (s, 3H, CH₃), 1.60 (d, $J=6.60$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 165.7, 161.4,

156.5, 146.7, 135.7, 135.3, 134.9, 128.0, 127.0, 126.6, 126.3, 126.0, 34.4, 19.6, 18.5; ESI-MS ($C_{21}H_{23}N_3O_2S$): calculated m/z 381.15 $[M+H]^+$, observed m/z 381.31 $[M+H]^+$; Anal. Calcd: $C_{21}H_{23}N_3O_2S$; C, 66.12; H, 6.08; N, 11.01; Found; C, 66.28; H, 6.34; N, 11.17.

N-(2,6-dimethylphenyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9n)

Cream solid; Yield:68%; MP=215–217 °C; IR (KBr, ν_{max}) 3235 (NH), 3030 (CH Aromatic), 2970 (CH Aliphatic), 1684 (C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 9.57 (s, 1H, NH_{Amide}), 8.04 (d, $J=8.00$ Hz, 1H, H₅), 7.77 (t, $J=7.70$ Hz, 1H, H₇), 7.53 (d, $J=8.20$ Hz, 1H, H₈), 7.42 (t, $J=7.50$ Hz, 1H, H₆), 7.05–7.00 (m, 3H, H₄, H₃, H₅), 4.79–4.64 (m, 1H, CH_{Isopropyl}), 4.23 (s, 2H, CH₂), 2.09 (s, 6H, 2xCH₃), 1.60 (d, $J=6.60$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 165.7, 161.4, 156.5, 146.7, 135.7, 135.3, 134.9, 128.0, 127.0, 126.6, 126.3, 126.0, 120.4, 36.4, 19.6, 18.5; ESI-MS ($C_{21}H_{23}N_3O_2S$): calculated m/z 381.15 $[M+H]^+$, observed m/z 381.24 $[M+H]^+$; Anal. Calcd: $C_{21}H_{23}N_3O_2S$; C, 66.12; H, 6.08; N, 11.01; Found; C, 66.23; H, 6.29; N, 11.26.

2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)-N-(naphthalen-2-yl)acetamide (9o)

Cream solid; Yield:65%; MP=217–219 °C; IR (KBr, ν_{max}) 3291 (NH), 3065 (CH Aromatic), 2980 (CH Aliphatic), 1675 (C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 10.36 (s, 1H, NH_{Amide}), 8.17 (d, $J=7.8$ Hz, 1H, H₈), 8.05 (d, $J=7.80$ Hz, 1H, H₅), 7.86 (t, $J=7.60$ Hz, 1H, H₇), 7.84–7.80 (m, 1H, H₈), 7.75 (t, $J=7.50$ Hz, 1H, H₆), 7.64 (d, $J=7.80$ Hz, 1H, H₇), 7.61–7.56 (m, 2H, H₃, H₆), 7.46 (t, $J=7.50$ Hz, 1H, H₄), 7.13 (t, $J=7.60$ Hz, 1H, H₅), 4.25–4.12 (m, 1H, CH_{Isopropyl}), 4.01 (s, 2H, CH₂), 0.80 (d, $J=6.70$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO- d_6): δ 165.9, 164.2, 161.2, 156.8, 147.7, 135.8, 135.5, 134.5, 132.2, 132.0, 131.2, 128.8, 127.0, 126.4, 126.3, 121.2, 121.2, 115.9, 115.7, 37.4, 13.7; ESI-MS ($C_{23}H_{21}N_3O_2S$): calculated m/z 403.14 $[M+H]^+$, observed m/z 403.27 $[M+H]^+$; Anal. Calcd: $C_{23}H_{21}N_3O_2S$; C, 68.46; H, 5.25; N, 10.41; Found; C, 68.65; H, 5.42; N, 10.60.

N-benzyl-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9p)

Cream solid; Yield:81%; MP=194–196 °C; IR (KBr, ν_{max}) 3264 (NH), 3025(CH Aromatic), 2960 (CH Aliphatic), 1662(C=O) Cm^{-1} ; 1H NMR (500 MHz, DMSO- d_6) δ 8.73 (t, $J=6.20$ Hz, 1H, NH_{Amide}), 8.03 (d, $J=8.10$ Hz, 1H, H₅), 7.73 (t, $J=7.60$ Hz, 1H, H₇), 7.45–7.38 (m, 2H, H₈, H₆), 7.24–7.18 (m, 5H, H_{Benzyl}), 4.77–4.56 (m, 1H, CH_{Isopropyl}), 4.32 (d, $J=5.90$ Hz, 2H, CH₂_{Benzyl}), 4.05 (s, 2H, CH₂), 1.58 (d, $J=6.50$ Hz, 6H, 2xCH₃) ppm. ^{13}C

NMR (125 MHz, DMSO-*d*₆): δ 167.2, 161.4, 156.5, 146.6, 139.6, 134.8, 128.6, 127.5, 127.5, 127.1, 126.5, 126.2, 126.1, 120.3, 43.0, 36.5, 19.6; ESI-MS ($C_{20}H_{21}N_3O_2S$): calculated m/z 367.14 $[M+H]^+$, observed m/z 367.26 $[M+H]^+$; Anal. Calcd: $C_{20}H_{21}N_3O_2S$; C, 65.37; H, 5.76; N, 11.44; Found; C, 65.49; H, 5.98; N, 11.63.

N-(4-fluorobenzyl)-2-((3-isopropyl-4-oxo-3,4-dihydroquinazolin-2-yl)thio)acetamide (9q)

Cream solid; Yield:76%; MP=224–226 °C; IR (KBr, ν_{max}) 3314 (NH), 3050(CH Aromatic), 2960(CHAliphatic), 1671(C=O) cm^{-1} ; 1H NMR (500 MHz, DMSO-*d*₆) δ 8.83 (t, $J=6.10$ Hz, 1H, NH_{Amide}), 8.02 (d, $J=8.00$ Hz, 1H, H₅), 7.73 (t, $J=7.60$ Hz, 1H, H₇), 7.41 (t, $J=7.50$ Hz, 1H, H₆), 7.36 (d, $J=8.10$ Hz, 1H, H₈), 7.29–7.23 (m, 2H, H₂, H₆), 7.04–6.97 (m, 2H, H₃, H₅), 4.72–4.58 (m, 1H, CH_{Isopropyl}), 4.28 (d, $J=5.90$ Hz, 2H, CH_{2Benzyl}), 4.03 (s, 2H, CH₂), 1.56 (d, $J=6.50$ Hz, 6H, 2xCH₃) ppm. ^{13}C NMR (125 MHz, DMSO-*d*₆): δ 167.2, 162.5, 161.4, 160.5, 156.5, 146.6, 135.9, 135.8, 134.8, 129.5, 129.5, 126.5, 126.2, 126.0, 120.3, 115.3, 115.2, 42.2, 36.5, 19.6; ESI-MS ($C_{20}H_{20}FN_3O_2S$): calculated m/z 385.13 $[M+H]^+$, observed m/z 385.26 $[M+H]^+$; Anal. Calcd: $C_{20}H_{20}FN_3O_2S$; C, 62.32; H, 5.23; N, 10.90; Found; C, 62.54; H, 5.37; N, 11.07.

Screening of tyrosinase inhibitory activity

Tyrosinase inhibitory activity was determined based on the previously described procedure with a slight modification [3, 21]. In brief, the test reaction mixture comprised 10 μ L of each derivative, and 10 μ L of mushroom tyrosinase (500 units; Sigma-Aldrich, St Louis, MO, USA) in 160 potassium phosphate buffer (pH=6.8). The reaction mixture was incubated at 37 °C for 20 min and then 20 μ L of L-Dopa (0.5 mmol/L) was added to each well, and the absorption was measured at 475 nm. The absorbance of the same mixture without tyrosinase was used as the control. Kojic acid was used as a positive control. The optical density of the inhibition in the control was considered to represent 100%. The data are expressed as mean percentages and the results were repeated in triplicate.

Enzyme kinetic studies

The mode of inhibition of the most active compound **9q** identified with the lowest IC_{50} , was investigated against tyrosinase activity with different concentrations of L-Dopa (0.1.5 mM) as substrate in the absence and presence of **9q** at different concentrations (10, 20, 30, 40 and 50 μ M). A Lineweaver–Burk plot was generated to identify the type of inhibition and the Michaelis–Menten constant (K_m) value was determined from the plot between the reciprocal of the substrate concentration ($1/[S]$) and reciprocal of enzyme rate ($1/V$) over various inhibitor concentrations.

Antioxidant activities

The DPPH antioxidant potencies of all derivatives were performed according to previously published articles [22].

Molecular docking

The molecular docking studies were performed using the Maestro Molecular Modeling platform (version 10.5) by Schrödinger, LLC. The X-ray crystal structure of the receptor (PDB ID: 2Y9X) was extracted from the PDB database. The protein is then prepared using a protein preparation wizard, the co-crystallized ligands and all water molecules were removed [23], the missing side chains and loops were filled using the prime tool, and PROPKA assigned H-bonds at pH: 7.4. To prepare the ligands, the 2D structures of the ligands were drawn in ChemDraw and converted into SDF files and subjected to ligprep module [24]. Ligands were prepared by OPLS_2005 force field using EPIK. The grid box was generated for each binding site using entries with a box size of 20 Å, the derivative was docked on binding sites using induced-fit docking, reporting 10 poses per ligand to form the final complex.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s13065-023-00978-3>.

Additional file 1: Figure S1. HPLC results of 9a. **Figure S2.** HPLC results of 9n. **Figure S3.** HPLC results of 9p. **Figure S4.** 1H NMR spectrum of compound 9a. **Figure S5.** ^{13}C NMR spectrum of compound 9a. **Figure S6.** 1H NMR spectrum of compound 9b. **Figure S7.** ^{13}C NMR spectrum of compound 9b. **Figure S8.** 1H NMR spectrum of compound 9c. **Figure S9.** ^{13}C NMR spectrum of compound 9c. **Figure S10.** 1H NMR spectrum of compound 9d. **Figure S11.** ^{13}C NMR spectrum of compound 9d. **Figure S12.** 1H NMR spectrum of compound 9e. **Figure S13.** ^{13}C NMR spectrum of compound 9e. **Figure S14.** 1H NMR spectrum of compound 9f. **Figure S15.** ^{13}C NMR spectrum of compound 9f. **Figure S16.** 1H NMR spectrum of compound 9g. **Figure S17.** ^{13}C NMR spectrum of compound 9g. **Figure S18.** 1H NMR spectrum of compound 9h. **Figure S19.** ^{13}C NMR spectrum of compound 9h. **Figure S20.** 1H NMR spectrum of compound 9i. **Figure S21.** ^{13}C NMR spectrum of compound 9i. **Figure S22.** 1H NMR spectrum of compound 9j. **Figure S23.** ^{13}C NMR spectrum of compound 9j. **Figure S24.** 1H NMR spectrum of compound 9k. **Figure S25.** ^{13}C NMR spectrum of compound 9k. **Figure S26.** 1H NMR spectrum of compound 9l. **Figure S27.** ^{13}C NMR spectrum of compound 9l. **Figure S28.** 1H NMR spectrum of compound 9m. **Figure S29.** ^{13}C NMR spectrum of compound 9m. **Figure S30.** 1H NMR spectrum of compound 9n. **Figure S31.** ^{13}C NMR spectrum of compound 9n. **Figure S32.** 1H NMR spectrum of compound 9o. **Figure S33.** ^{13}C NMR spectrum of compound 9o. **Figure S34.** 1H NMR spectrum of compound 9p. **Figure S35.** ^{13}C NMR spectrum of compound 9p. **Figure S36.** 1H NMR spectrum of compound 9q. **Figure S37.** ^{13}C NMR spectrum of compound 9q.

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Author contributions

A.H., M.N., N.D., S.E.S.E., F.J.Z., M.A.D and M.M synthesized compounds and contributed to the design and characterization of compounds. K.M, M.H. E.S

and B.L. performed the biological assay. N.F.H. C.I and A.I. performed in silico study and contributed to the preparation of the manuscript. A.I. supervised the biological tests. M.M. supervised all phases of the study. All authors read and approved the article.

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Availability of data and materials

The datasets generated and/or analyzed during the current study are available in the Worldwide Protein Data Bank (wwPDB) repository; (<https://www.rcsb.org/structure/2y9x>) with PDB <https://doi.org/10.2210/pdb2Y9X/pdb>.

Declarations

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

Not applicable.

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