



Article Green Route Synthesis and Molecular Docking of Azines Using Cellulose Sulfuric Acid under Microwave Irradiation

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Abstract: A novel catalytic approach for the synthesis of symmetric and asymmetric azines has been developed. The environmentally benign protocol was achieved via condensation of 1-[(2-thienyl)ethylidene]hydrazine (1) with different aromatic aldehydes **2a**–**h** and acetyl heterocyclic compounds (**4**, **6**, **8**, **10**, and **12**) in the presence of cellulose sulfuric acid (CSA) as the green catalyst. These procedures offer an interesting method for the large-scale industrial manufacture of azines due to their high percentage yield, mild reaction conditions, broad substrate range, and utilization of an economical and environmentally acceptable catalyst. Additionally, the molecular docking of the products to the monoamine oxidase (MAO-A) target protein was achieved to highlight the possible binding interaction with the amino acid residues Arg51, Glu43, Gly22, Gly49, Gly443, Ala272, Ile335, and Tyr407 at the point of binding. The binding interaction energy was discovered to be (– 6.48 kcal/mol) for the protein MAO-A (PDB ID: 2Z5X). The most effective azine derivatives **7** and **13** revealed some major conserved interactions between the MAO-A protein's binding site amino acid residues and the PDB co-crystal ligand 2Z5X. Moreover, azine derivatives **3a** and **3f** showed the lowest binding activity with the target MAO-A.

Keywords: Schiff base; condensation reaction; azines; molecular docking; MAO-A target protein

1. Introduction

Monoamine oxidase (MAO) is overexpressed in dysthymia, which leads to neurological system disorders including Parkinson's and Alzheimer's disease and inhibits the amounts of monoaminergic transmitters, and as a result, the nerve system's ability to communicate. MAO inhibitors can keep MAO levels within cells and in communication [1]. A significant enzyme called MAO, which is found on the membrane enclosing the mitochondria in neuronal cells, contains flavins. The C-terminal transmembrane polypeptide loop provides the connection [2], and it is inserted into the membrane via ubiquitin, with ATP acting as the source of energy [3], controlling monoaminergic homeostasis and perhaps neurotransmission in neuronal, glial, and other cells. The deficiency of certain neurotransmitters in the brain, such as γ -aminobutyric acid (GABA), norepinephrine, dopamine, and serotonin, leads to depression and other mental problems. A key finding in the monoamine theory of depression was the discovery in the 1950s of the antidepressant effects of MAO inhibitors (MAOIs). However, earlier MAOIs that were used in care medicine were terminated because of unfavorable side effects, including hepatotoxicity, orthostatic hypotension, and the reported "cheese effect" that had hypertensive crises as its feature. The firstgeneration antidepressants being used were MAOIs, for decades, in the treatment for those



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Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). who suffered from unusual depression [4], a tremendous range of depression, bipolar, anxiety resistant to therapy depression [5], specific phobias, post-traumatic stress disorder (PTSD), and migraines that did not respond to other therapies [6]. The research of MAOIs was sparked by the unexpected result of antidepressant effects in patients taking iproniazid, a hydrazine-based antitubercular drug chemically related to isoniazid. This discovery, coupled with evidence that iproniazid was a potent MAOI [7], sparked the development of further MAOIs, such as phenelzine. Among the family of MAOs is monoamine oxidase A, which has the ability to accelerate the deamination of serotonin, noradrenaline, and adrenaline. Dopamine, tryptamine, and tyramine can also be metabolized by monoamine oxidases (MAO-A) [9], and moclobemide (MAO-A) [10]. The MAO-A inhibitor is suggested for the treatment of anxiety and depression in the brain [11–13]. An entirely new family of chemicals has emerged as a result of recent efforts to find reversible and selective MAOIs.

Sustainable green catalysts with environmental and economical advantageousness are considered to be of interest in the development of green chemistry. In synthetic organic chemistry, the utilization of green techniques is supportive to reduce the usage of hazardous toxic chemicals [14], minimize the reaction time, and increase selectivity [15]. That simplifies the purification and isolation of the isolated products rather than conventional methods [16]. Cellulose sulfuric acid (CSA) is an attractive sustainable candidate in green technology because it is a biodegradable material and has a renewable source. It has emerged as a biopolymeric solid support acid catalyst for the synthesis of pyridotriazolopy-rimidines [17], dihydropyrimidinones [18], pyrazoles [19], and α -aminonitriles [20]. On the other hand, aldazines and ketazines with the conjugated hydrazone system (C=N–N=C) have been widespread and employed in pharmacological research and the industrial field. Both symmetric and asymmetric azines have revealed biological activities as antimicrobial [21], antibacterial [22], and antitumor [23] agents. In addition, azines were used as building blocks for the construction of nonlinear optical materials [24] and covalent organic frameworks (COFs) which served as chemo-sensing detectors [25].

Based on the aforementioned precedents and in continuation of our efforts in green chemistry research [26–35], we have presently employed cellulose sulfuric acid as a catalyst for the synthesis of symmetric and asymmetric azines in this context. Additionally, it was also shown that molecular modelling may be used to estimate the binding affinity and determine the interactions of the tested compounds at the MAO-A (PDB: 2Z5X) binding site where the synthesized azines were established.

2. Results and Discussion

Recently, environmentally benign protocols for the condensation of carbonyl compounds and amines were achieved in the presence of meglumine [36], L-tyrosine [37], and citric acid [38] as green catalysts. These reports encouraged us to synthesize symmetric and asymmetric azines by using CSA as a biodegradable and reusable catalyst. Thus, the condensation of 1-[(2-thienyl)ethylidene]hydrazine (1) with aromatic aldehydes **2a–h** in ethanolic solution, containing 0.05 g of CSA, under microwave irradiation at 120 °C for 20–30 min, gave the corresponding azines **3a–h** (Scheme 1). The required products were separated in good to excellent yields regardless of the kind of substituents (electronwithdrawing or electron-donating group). The reaction was slightly sensitive to the steric hindrance of 2-substituted benzaldehydes **2f–2h** which decreased the reactivity and yield percentage of products **3f–3h** (Table 1).



Scheme 1. Synthesis of unsymmetric azines 3a–h.

Table 1. Yield percentage of azine derivatives 3a-h.

| Compd. No. | R ¹ | R ² | Yield (%) |
|------------|------------------|------------------|-----------|
| 3a | Н | Н | 95 |
| 3b | Н | OCH ₃ | 92 |
| 3c | Н | Cl | 92 |
| 3d | Н | Br | 91 |
| 3e | Н | NO ₂ | 90 |
| 3f | OH | Н | 86 |
| 3g | OCH ₃ | ОН | 84 |
| 3h | Cl | Cl | 83 |

The structural elucidation of azines **3a–h** was characterized by elemental and spectroscopic data. In IR spectra, the stretching absorption of the (C=N) group was observed at about 1599–1904 cm⁻¹. In the ¹H-NMR spectra of compounds, the Schiff base proton (N=CH) was determined at about δ = 8.54–8.96 ppm and the aromatic protons were assigned at the expected region.

Notably, the catalytic activity of cellulose sulfuric acid was applicable for the preparation of symmetric and asymmetric ketazines. Thus, different acetyl heterocyclic compounds (4, 6, 8, 10, and 12) were subjected to react with 1-[(2-thienyl)ethylidene] hydrazine (1) under the employed conditions to give the respective *bis*-hydrazones (5, 7, 9, 11, and 13) in good yields (Scheme 2).



Scheme 2. Synthesis of ketazine derivatives 5, 7, 9, 11, and 13.

3. Molecular Docking

A version of the Molecular Operating Environment (MOE) software 2015.10 was used to conduct the molecular docking and validated through the re-docking of the native ligand high-resolution melt (HRM) within the active site of MAO-A, giving docked scores of -6.48 kcal/mol with a root mean square deviation (RMSD) value of 0.84 A° between the docked configurations and the co-crystallized ligand. HRM demonstrated the important interactions with the residues at the active site of MAO-A (Figure 1).



Figure 1. 2D-binding of HRM (a reference ligand) with MAO-A.

Therefore, using the native co-crystallized ligand HRM700 as a docking partner into the appropriate MAO-A binding sites, the verification of the MAO-A docking accuracy was examined (PDB: 2Z5X). After being re-docked into the binding site, the co-crystallized ligand HRM700 interacted hydrophobically with Gln215, an essential amino acid for MAO-A. Within an RMSD of 0.84 Å, this auto-dock success rate was extraordinarily good (Figure 1).

Hydrazones, as a moiety open chain with aryl or bioactive azoles, were reported as MAO-inhibiting agents [39,40]. The synthesized azines all showed excellent inhibitory potency and selectivity towards MAO-A in general. Molecular docking studies showed that the synthesized azines (3a–h, 5, 7, 9, 11, and 13) have binding energy values between –8.09 and –6.36 kcal/mol. According to the molecular docking analysis, docking scores for compounds 7 and 13 were the highest (–7.32 kcal/mol and –8.09, respectively) (Table 2).

3.1. SAR Analysis for a Series of 1-(arylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3a-h)

The substituted phenyl ring was preferred over an unsubstituted one. Thus, compound (**3a**) (-6.36 kcal/mol) displayed the least decrease in the activity for the investigated series. The *para* substitution of the electron-donating methoxy group (**3b**) (-7.16 kcal/mol) or electron-withdrawing moiety such as the NO₂ group (**3e**) (-7.15 kcal/mol) was found to have a significant impact on the inhibition of the receptor. The activity indicated that the alkoxy (-OCH₃)-substituted compound (**3b**) (-7.16 kcal/mol) was more active when compared to the halogen (Cl and Br)-substituted ones (**3c** and **3d**) (-6.94 and -6.83 kcal/mol, respectively). The small size electron-withdrawing moiety of Cl was less than that of Br; thus, compound (**3c**) showed more activity than bulk compound (**3d**) (-6.94 and -6.83 kcal/mol, respectively). The presence of the hydroxy group at the *ortho* position in (**3f**) (-6.65 kcal/mol) derivative was not favorable for binding, and thus showed poor inhibition activity. The monosubstituted one's steric groups (**3h**) (-6.78 kcal/mol).

| Compd. No. | RMSD (A°) | Energy Score (S) (kcal/mol) |
|------------|--------------------|--------------------------------|
| 3a | 1.50 | -6.36 |
| 3b | 1.64 | -7.16 |
| 3c | 1.41 | -6.94 |
| 3d | 1.62 | -6.83 |
| 3e | 1.11 | -7.15 |
| 3f | 0.97 | -6.65 |
| | 1.10 | -6.94 |
| 3h | 0.68 | -6.78 |
| 5 | 1.39 | -7.18 |
| 7 | 1.86 | -7.32 |
| 9 | 1.72 | -7.28 |
| 11 | 0.99 | -7.19 |
| 13 | 1.27 | -8.09 |
| HRM | 0.84 | -6.48 |

Table 2. RMSD (A°) and energy scores (kcal/mol) for azine derivatives (3a–h, 5, 7, 9, 11, and 13) compared to the native ligand (HRM).

3.2. SAR Analysis for a Series of Symmetric and Asymmetric Azines (5, 7, 9, 11, and 13)

It was found that compound 7 (-7.32 kcal/mol), with different hetero atoms such as sulfur and nitrogen in the thiazole ring, had more inhibitory activity than compound 5 (-7.18 kcal/mol) possessing only the sulfur atom in the thiophen ring. From another structural viewpoint, the azine derivative 9 with indole moiety (-7.28 kcal/mol) exhibited improved inhibitory properties over compound 11 with isatin moiety (-7.19 kcal/mol)[41,42]. The most potent azine derivative was compound 13 because the synthetic coumarin derivative has been characterized as MAOIs [43,44].

Numerous hydrogen bonds and hydrophobic interactions between the azine derivatives and the receptor's active residues were the cause of the binding affinity as shown in Figure 2. As demonstrated, compounds 3a, 3c, 3f and 3b, 3d, 3e, 5, and 13 provided the π - π interaction between the phenyl and thiophene ring with the essential amino acid Try407, respectively. Additionally, in compound **3e**, the carbon atom of the phenyl ring interacted as an *H*-bond donor with the Cys323. Moreover, compound **3g** showed the π -*H* interaction between the thiophene ring with the essential amino acid Try407. The 3D model of the binding mode of compound 7 showed one H-bond was between the nitrogen of the amino group afforded the *H*-bond donor and the use of the side chain of Gly443, and additionally the π - π interaction between the thiazole ring and the essential amino acid Try407. As demonstrated, compound 9 provided two π - π interactions between the indole ring and Try407. The 3D model of the binding mode of compound **11** showed three π - interactions, one of which was between the thiophene ring and permitted the π - π to utilize the side chain of Try407, and the other two π -H interactions were between indolin-2-one moiety and the side chain of Gln215 and Phe208. Moreover, compound **13** showed the *H*-bond acceptor between the oxygen atom of coumarin moiety and the side chain of Phe208. The results showed that the examined ligands were positioned and oriented similarly inside the predicted binding site of HRM (Figure 2).



Figure 2. Cont.



Figure 2. Cont.



Figure 2. Cont.



Figure 2. 3D- and 2D-ligand interactions within the binding site of 2Z5X for azine derivatives (3a–h, 5, 7, 9, 11, and 13).

4. Experimental Section

The ¹H-NMR and ¹³C-NMR spectra were measured using a Varian Mercury VXR-300 spectrometer (300 MHz and 75 MHz for ¹H-NMR and ¹³C-NMR, respectively). The melting temperatures of the newly formed azine derivatives were measured using an electrothermal Gallenkamp device. The mass spectra of the azine derivatives were measured using a GCMS 5988-A HP spectrometer at 70 eV of ionizing voltage. IR spectra were measured using a Pye-Unicam SP300 device in KBr discs. The CEM Discover Labmate microwave device was used for the microwave tests (Discover, SP, NC, USA, 300 W). At the Microanalytical Center at Cairo University in Giza, Egypt, elemental studies were performed.

4.1. General Reactions of 1-[(2-thienyl)ethylidene]hydrazine (1) with Aldehydes (2a-h) or Ketones (4, 6, 8, 10, and 12)

The 1-[(2-Thienyl)ethylidene]hydrazine (1) (0.14 g, 1 mmol) was allowed to react with arylaldehydes (2a–h) or acetylthiophene (4), or 2-amino-4-methyl-5-acetylthiazole (6) or 3-acetylindole (8), or isatin (10) or 3-acetylcoumarine (12) (1 mmol of each) in EtOH (10 mL) containing 0.05 g of CSA. The reaction mixture was irradiated by MW at 120 °C in a closed Teflon vessel until completeness (20–30 min). CSA was carefully removed by filtration of the hot solution. The excess solvent was removed, and the reaction mixture was triturated with methanol. The separated product was then filtered and finally recrystallized from EtOH to give the products 3a-h or 5 or 7 or 9 or 11 or 13, respectively.

1-(Benzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3a)

Pale yellow crystals, mp 117–120 °C IR (KBr) v = 2917 (CH), 1602 (C=N) cm⁻¹; MS, m/z (%) 228 (M⁺, 30); ¹H-NMR (DMSO- d_6) $\delta = 2.48$ (s, 3H, CH₃), 7.42–7.86 (m, 8H, Ar-H), 8.68 (s, 1H, CH=N); ¹³C-NMR (DMSO- d_6): $\delta = 15.3$ (CH₃), 123.4, 125.8, 127.5, 128.9, 129.7, 130.2, 134.1, 143.8, 158.4, 162.5 (Ar-C and C=N). Anal. calcd for C₁₃H₁₂N₂S (228.07): C, 68.39; H, 5.03; N, 12.27; S, 14.04. Found: C, 68.44; H, 4.92; N, 12.41; S, 14.18%.

1-(4-Methoxybenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3b)

Yellow crystals, mp 159–161 °C IR (KBr) v = 2919 (CH), 1601 (C=N) cm⁻¹; MS, m/z (%) 258 (M⁺, 43); ¹H-NMR (DMSO- d_6) $\delta = 2.49$ (s, 3H, CH₃), 3.78 (s, 3H, OCH₃), 7.01–7.78 (m, 7H, Ar-H), 8.59 (s, 1H, CH=N); ¹³C-NMR (DMSO- d_6): $\delta = 15.3$ (CH₃), 56.1 (OCH₃), 114.9, 127.1, 128.2, 129.7, 130.1, 130.5, 144.2, 157.5, 160.9, 162.2 (Ar-C and C=N). Anal. calcd for C₁₄H₁₄N₂OS (258.08): C, 65.09; H, 5.46; N, 10.84; S, 12.41. Found: C, 64.94; H, 5.22; N, 10.71; S, 12.58%.

1-(4-Chlorobenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3c)

Pale yellow crystals, mp 207–209 °C IR (KBr) v = 2921 (CH), 1599 (C=N) cm⁻¹; MS, m/z (%) 262 (M⁺, 18); ¹H-NMR (300 MHz, DMSO- d_6) $\delta = 2.47$ (s, 3H, CH₃), 7.11–7.87 (m, 7H, Ar-H), 8.68 (s, 1H, CH=N); ¹³C-NMR (75 MHz, DMSO- d_6): $\delta = 15.7$ (CH₃), 124.4, 125.6, 127.4, 128.8, 129.5, 131.1, 134.5, 144.2, 158.8, 162.7 (Ar-C and C=N). Anal. calcd for C₁₃H₁₁ClN₂S (262.03): C, 59.43; H, 4.22; N, 10.66; S, 12.20. Found: C, 59.54; H, 4.12; N, 10.40; S, 12.38%.

1-(4-Bromobenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3d)

Pale yellow crystals, mp 228–230 °C IR (KBr) v = 2918 (CH), 1600 (C=N) cm⁻¹; MS, m/z (%) 305 (M⁺, 12); ¹H-NMR (DMSO- d_6) $\delta = 2.47$ (s, 3H, CH₃), 7.11–7.87 (m, 7H, Ar-H), 8.67 (s, 1H, CH=N); ¹³C-NMR (DMSO- d_6): $\delta = 15.8$ (CH₃), 124.8, 125.7, 127.9, 128.5, 130.1, 132.1, 134.2, 145.2, 158.7, 162.9 (Ar-C and C=N). Anal. calcd for C₁₃H₁₁BrN₂S (305.98): C, 50.83; H, 3.61; N, 9.12; S, 10.44. Found: C, 50.69; H, 3.52; N, 9.24; S, 10.32%.

1-(4-Nitrobenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3e)

Brown crystals, mp 285–287 °C IR (KBr) v = 2917 (CH), 1601 (C=N) cm⁻¹; MS, m/z (%) 273 (M⁺, 32); ¹H-NMR (DMSO- d_6) $\delta = 2.47$ (s, 3H, CH₃), 7.52–8.34 (m, 7H, Ar-H), 8.68 (s, 1H, CH=N); ¹³C-NMR (DMSO- d_6): $\delta = 15.7$ (CH₃), 124.4, 125.8, 126.5, 127.9, 129.3, 133.2, 138.1, 144.8, 157.8, 162.6 (Ar-C and C=N). Anal. calcd for C₁₃H₁₁N₃O₂S (273.06): C, 57.13; H, 4.06; N, 15.37; S, 11.73. Found: C, 57.34; H, 3.92; N, 15.41; S, 11.88%.

1-(2-Hydroxybenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3f)

Yellow crystals, mp 173–175 °C IR (KBr) v = 3420 (OH), 2920 (CH), 1604 (C=N) cm⁻¹; MS, m/z (%) 244 (M⁺, 28); ¹H-NMR (DMSO- d_6) $\delta = 2.36$ (s, 3H, CH₃), 6.93–7.66 (m, 7H, Ar-H), 8.96 (s, 1H, CH=N), 11.12 (s, 1H, OH); ¹³C-NMR (DMSO- d_6): $\delta = 15.9$ (CH₃), 118.9, 123.1, 124.2, 125.8, 126.7, 128.1, 130.5, 132.2, 133.8, 156.5, 160.1, 163.5 (Ar-C and C=N). Anal. calcd for C₁₃H₁₂N₂OS (244.07): C, 63.91; H, 4.95; N, 11.47; S, 13.12. Found: C, 63.74; H, 5.12; N, 11.61; S, 13.28%.

1-(2-Hydroxy-4-methoxybenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3g)

Yellow crystals, mp 144–146 °C IR (KBr) v = 3425 (OH), 2918 (CH), 1602 (C=N) cm⁻¹; MS, m/z (%) 274 (M⁺, 46); ¹H-NMR (DMSO- d_6) $\delta = 2.46$ (s, 3H, CH₃), 3.78 (s, 3H, OCH₃), 7.40–8.54 (m, 6H, Ar-H), 8.66 (s, 1H, CH=N), 11.32 (s, 1H, OH); ¹³C-NMR (DMSO- d_6): $\delta = 15.8$ (CH₃), 56.8 (OCH₃), 108.9, 112.1, 116.7, 124.2, 125.8, 126.7, 128.1, 130.5, 133.8, 155.8, 160.1, 164.5 (Ar-C and C=N). Anal. calcd for C₁₄H₁₄N₂O₂S (274.08): C, 61.29; H, 5.14; N, 10.21; S, 11.69. Found: C, 61.44; H, 5.19; N, 10.41; S, 11.48%.

1-(2,4-Dichlorobenzylidene)-2-[1-(2-thienyl)ethylidene]hydrazine (3h)

Yellow crystals, mp 108–110 °C IR (KBr) v = 2921 (CH), 1599 (C=N) cm⁻¹; MS, m/z (%) 295 (M⁺, 16); ¹H-NMR (DMSO- d_6) $\delta = 2.48$ (s, 3H, CH₃), 7.10–7.62 (m, 6H, Ar-H), 8.54 (s, 1H, CH=N); ¹³C-NMR (DMSO- d_6): $\delta = 15.4$ (CH₃), 124.4, 125.4, 126.2, 127.5, 128.1, 128.7, 129.5, 130.1, 131.5, 134.2, 158.8, 162.7 (Ar-C and C=N). Anal. calcd for C₁₃H₁₀Cl₂N₂S (295.99): C, 52.54; H, 3.39; N, 9.43; S, 10.79. Found: C, 52.44; H, 3.22; N, 9.52; S, 10.88%.

1,2-Bis[1-(2-thienyl)ethylidene]hydrazine (5)

Orange powder, mp 97–99 °C, lit mp = 95–96 °C [45]; IR (KBr) v = 2918 (CH), 1596 (C=N) cm⁻¹; MS, m/z (%) 248 (M⁺, 6); ¹H-NMR (DMSO- d_6) δ = 2.36 (s, 6H, 2CH₃), 7.09–7.61 (m, 6H, Ar-H); ¹³C-NMR (DMSO- d_6): δ = 15.2 (CH₃), 125.5, 126.9, 128.7, 129.2, 162.5 (Ar-C and C=N). Anal. calcd for C₁₂H₁₂N₂S₂ (248.04): C, 58.03; H, 4.87; N, 11.28; S, 25.82. Found: C, 58.14; H, 4.92; N, 11.42; S, 25.68%.

4-Methyl-5-{1-[((1-(2-thienyl)ethylidene)hydrazono]ethyl}thiazol-2-amine (7)

Yellow crystals, mp 95–97 °C IR (KBr) v = 3342, 3218 (NH₂), 2923 (CH), 1601 (C=N) cm⁻¹; MS, m/z (%) 278 (M⁺, 30); ¹H-NMR (DMSO- d_6) $\delta = 2.35$ (s, 3H, CH₃),

2.36 (s, 3H, CH₃), 2.47 (s, 3H, CH₃), 7.11–7.73 (m, 5H, Ar-H + NH₂); ¹³C-NMR (DMSO-*d*₆): δ = 14.9 (CH₃), 15.3 (CH₃), 16.9 (CH₃), 124.8, 125.5, 126.9, 127.7, 128.2, 129.6, 131.2, 160.4, 162.5 (Ar-C and C=N). Anal. calcd for C₁₂H₁₄N₄S₂ (278.07): C, 51.77; H, 5.07; N, 20.13; S, 23.03. Found: C, 51.64; H, 4.99; N, 20.31; S, 23.18%.

3-{1-[((1-(2-Thienyl)ethylidene)hydrazono]ethyl}-1H-indole (9)

Yellow crystals, mp 284–286 °C IR (KBr) v = 3214 (NH), 2920 (CH), 1600 (C=N) cm⁻¹; MS, m/z (%) 281 (M⁺, 24); ¹H-NMR (DMSO- d_6) $\delta = 2.42$ (s, 3H, CH₃), 2.48 (s, 3H, CH₃), 7.12–8.48 (m, 8H, Ar-H), 11.39 (s, 1H, NH); ¹³C-NMR (DMSO- d_6): $\delta = 15.4$ (CH₃), 16.1 (CH₃), 112.1, 116.6, 120.8, 122.6, 123.6, 125.6, 126.4, 127.8, 128.4, 129.3, 130.4, 137.7, 158.3, 160.5 (Ar-C and C=N). Anal. calcd for C₁₆H₁₅N₃S (281.10): C, 68.30; H, 5.37; N, 14.93; S, 11.39. Found: C, 68.14; H, 5.19; N, 14.81; S, 11.19%.

3-{[(1-(2-Thienyl)ethylidene]hydrazono}indolin-2-one (11)

Orange crystals, mp 196–198 °C IR (KBr) v = 3210 (NH), 2922 (CH), 1660 (C=O), 1599 (C=N) cm⁻¹; MS, m/z (%) 269 (M⁺, 51); ¹H-NMR (DMSO- d_6) $\delta = 2.47$ (s, 3H, CH₃), 6.93–7.33 (m, 7H, Ar-H), 10.63 (s, 1H, NH); ¹³C-NMR (DMSO- d_6): $\delta = 15.8$ (CH₃), 120.1, 125.9, 126.8, 127.6, 128.6, 129.6, 130.4, 131.8, 132.4, 137.7, 139.4, 141.3, 160.5 (Ar-C and C=N). Anal. calcd for C₁₄H₁₁N₃OS (269.06): C, 62.44; H, 4.12; N, 15.60; S, 11.90. Found: C, 62.64; H, 4.19; N, 15.81; S, 12.09%.

3-{1-[1-((2-Thienyl)ethylidene)hydrazono]ethyl}-2H-chromen-2-one (13)

Orange crystals, mp 221–223 °C IR (KBr) v = 2917 (CH), 1703 (C=O), 1604 (C=N) cm⁻¹; MS, m/z (%) 310 (M⁺, 25); ¹H-NMR (DMSO- d_6) $\delta = 2.42$ (s, 3H, CH₃), 2.47 (s, 3H, CH₃), 7.44–8.79 (m, 8H, Ar-H); ¹³C-NMR (DMSO- d_6): $\delta = 15.6$ (CH₃), 16.4 (CH₃), 116.1, 118.6, 119.8, 123.8, 124.6, 125.6, 126.6, 127.4, 128.8, 129.4, 130.3, 132.4, 147.7, 158.3, 161.5 (Ar-C and C=N). Anal. calcd for C₁₇H₁₄N₂O₂S (310.08): C, 55.79; H, 4.55; N, 9.03; S, 10.33. Found: C, 55.84; H, 4.69; N, 9.18; S, 10.19%.

4.2. Molecular Docking

Chemdraw 12.0 was used to sketch out the chemical structures of the most active compounds. To further refine the findings, London dG force and force field energy were employed. Using Merck molecular force field 94 (MMFF 94), all minimizations were carried out until an RMSD gradient of 0.1 kcal·mol⁻¹Å⁻¹ was reached. The MOE software's scoring function and dock function (S, kcal/mol) were used to assess the ligand's binding affinity. The protein data bank was used to download the X-ray crystal structure of the enzyme in PDB format (PDB ID: 2Z5X, resolution: 2.2 Å) [46]. The enzyme was prepared for docking studies: (i) The waters were eliminated from the protein. (ii) The structure was then supplied with hydrogen atoms in their typical geometries, then the broken bonds were reconnected and the potential fixed [47]. (iii) MOE Alpha Site Finder was utilized for the large site search in the enzyme structure, and dummy atoms were generated from the alpha spheres that were produced [48]. (iv) The interaction of the ligand with the amino acids of the active site was analyzed. The active ligands' largest negative value yielded the highest docking score. All docking procedures and scoring were recorded according to established protocols [49-51]. Overall docking results were compared to the crystal structure of the bound natural ligand-protein complex (HRM700). The London dG score tool and Triangle Matcher placement method were used for docking. After docking, the 2D and 3D interactions with residues of amino acids were visualized.

5. Conclusions

Novel series of symmetric and asymmetric azines have been synthesized via the condensation of 1-[(2-thienyl)ethylidene]hydrazine with different aromatic aldehydes and acetyl heterocyclic compounds in the presence of CSA as green catalysts with environmental and economical advantageousness. The attractive features of this protocol are the mild reaction conditions, high percentage yield, wide substrate scope, and use of an inex-

pensive and environmentally friendly catalyst, all of which make it an attractive strategy for the large-scale industrial preparation of azines. Additionally, the binding interaction energies between azines and the monoamine oxidase (MAO-A) target protein were determined using molecular docking techniques. The ligand in discussion exhibits a binding interaction energy of (-6.48 kcal/mol) for protein 2Z5X. The visual representation of the intermolecular interactions between the designated ligand and protein is investigated.

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